Essays on Dynamic Marketing Intercommunications

by

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To my parents - Seung Hyun Lee and Young Nam Byun

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ABSTRACT

The dissertation examines two distinct problems related to "marketing communication dynamics". The main goal of this line of research is to help firms provide individually tailored marketing contents to their customers. In these two essays, I develop statistical models to first understand customers' responses, and then explore methods to optimize firms' reactions accordingly. Essay 1 examines "scale attraction effects" in a charitable donation context, introducing novel constructs ("compliance degree", "pulling amount", "accumulated pulling amount") to describe attraction effects for multi-point appeals scales. The proposed model jointly accounts for donation incidence and amount using a Tobit 2 formulation, and allows heterogeneity in seasonality and pulling effects. Results suggest substantial scale attraction effects that vary across donors, stronger "pulling down" than "pulling up", and heterogeneous seasonal donation patterns. A significantly negative error correlation between donation incidence and donation amount underscores the importance of accounting for selectivity effects. The effects of individually tailoring appeals scales is demonstrated through simulation.

Essay 2 investigates mate-seeking users' decision rules in an online dating context. I develop an empirical two-stage mate choice model that can accommodate compensatory and non-compensatory decision rules in each of two stages: browsing and writing. A mixture of logits model with changepoints allows for distinct decision rules across stages and heterogeneity in rule use across site users. Most importantly, it allows us to identify and compare attribute-level decision rules ("deal-breakers" and "deal-makers") over the two stages. Results suggest the existence of heterogeneity in decision rules across (1) genders, (2) stages, and (3) site users. Additionally, it suggests the existence of potential

deal-breakers/makers across both discrete and continuous attributes.

CHAPTER I

INTRODUCTION

This dissertation develops models of *dynamic intercommunication*: between firms and their customers and among customers themselves. In these essays, I am interested in not only measuring customers' responses to firms' marketing communication actions (stimuli), but also how firms can optimize such stimuli to individual customers in a dynamic manner. This stream of research is particularly intriguing to me, as small changes in marketing communications - such as scaling (1^{st} essay), and matching algorithm (2^{nd} essay) can exert great impact on customers' behaviors by simply "nudging" them appropriately. Such nudging can help firms to maximize the efficiency of marketing communications, as well help customers to find the products/services that best fit their needs. That is, without investments in costly new campaigns or infrastructure, firms can (algorithmically, optimally) tune their personalized marketing intercommunications, over time.

The first, "Modeling Scale Attraction Effects: An Application to Charitable Donations and Optimal Laddering" examines *scale attraction effects* when approaching individual potential donors. Panel data from a unique 3.5 year quasi-experiment enables a joint account of both whether a donation is made (incidence) and, if so, its size (amount). The model incorporates heterogeneity across donors in scale attraction effects and donation patterns (e.g., seasonality), and allows tests of distinct operationalizations of internal and external reference price theories. Results suggest that scale points do exert substantial attraction ef-

fects; that these vary markedly across donors; that donors are more easily persuaded to give less than more; and that seasonal donation patterns are pronounced. A significantly negative error correlation between (latent) donation propensity and (observed) donation amount highlights the importance of accounting for selectivity effects. We illustrate the framework with a speculative application to "laddering": how much charities should increase amounts subsequently requested of individual donors, based on their donation histories.

The second essay, "Modeling mate choice behavior: A two-stage mate choice model with potentially non-compensatory decision rules" allows us to empirically evaluate alternative decision rules used by individuals who interacted via an online dating service provider in the US. The proposed model captures the intrinsically multistage behavior involved in many online transactions, but in particular dating, where one decides which profiles to browse and then (stage 1), conditional on having browsed, whom to write to (stage 2), if anyone. Here, we account for these two distinct activities by modeling the binary decisions (of browsing and writing). The model can accommodate compensatory and noncompensatory decision rules in each stage; it allows decision rules to differ across stages; different attributes can be modeled as having distinctly different utility 'shapes'; and heterogeneity in rule use across site users provides interpretable profiles of different types of mate-seeking behavior. Finally, and most importantly, we directly model the utility functions of attributes to identify and compare attribute-level decision rules ("deal-breakers" and "deal-makers") over the two distinct stages. Based on the model and direct extensions to other data settings, firms can offer more accurate, targeted search among potential dyads, in online dating, social networks, and even among various users of online shopping sites.

CHAPTER II

Modeling Scale Attraction Effects: An Application to Charitable Donations and Optimal Laddering

2.1 Abstract

Charities seeking donations usually employ an "appeals scale," a set of specific suggested amounts presented directly to potential donors. Choosing them well is crucial: if charities select overly high scale points, they risk their being ignored or even alienating donors and receiving nothing, while overly low scale points may encourage donors to give less than they'd have otherwise. Despite their widespread use, little is known about the degree to which the points on such scales affect both whether a donation is made and, if so, its size. Using panel data from a 3.5 year quasi-experiment, we develop a joint model accounting for both donation incidence and amount. The model incorporates heterogeneity across donors in both scale attraction effects and in donation patterns (e.g., seasonality), and allows tests of distinct operationalizations of internal and external reference price theories. Results suggest that scale points do exert substantial attraction effects; that these vary markedly across donors; that donors are more easily persuaded to give less than more; and that there are strong seasonal donation patterns in giving. A significantly negative correlation between error terms in (latent) donation propensity and (observed) donation amount highlights the importance of accounting for selectivity effects. We illustrate the framework

with a speculative application to "laddering": how much charities should increase amounts subsequently requested of individual donors, based on their donation histories.

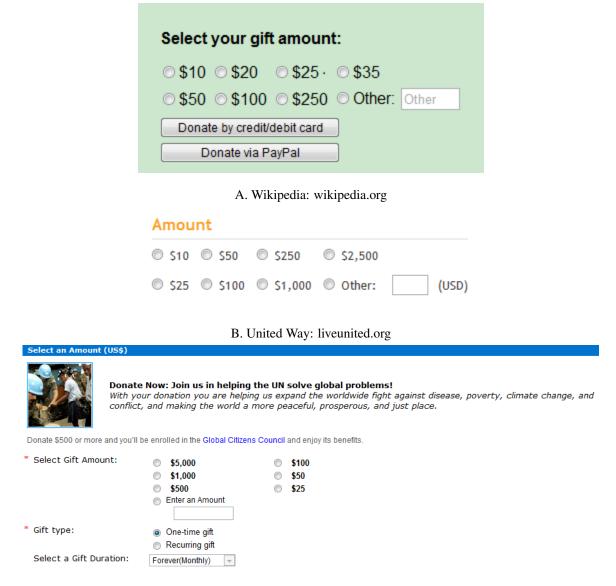
2.2 Introduction

Solicitations for charitable donations are a part of everyday life, with requests being made at physical locations (stores, workplaces), through the mail, various media (radio, television), and increasingly via the Internet (e-mail, websites, social networks). The Center on Philanthropy at Indiana University (2010) recently listed over 1.2 *million* charitable organizations in the United States alone, as of 2009. The total amount of giving in the US has increased rapidly over the past decade, with \$303B in 2009, an 80% inflation-adjusted increase over 20 years earlier. These donations account for 2.1% of U.S. GDP, a quantity ahead of all but 3 corporations in revenues and all in profits.

Private citizens have been generous to charities, even during the recent economic downturn, with 65% of US households donating per annum, \$1,940 on average (including nondonors). Household-level donation, \$227.4B in 2009, accounts for 75% of the U.S. total, followed by foundations (\$38.4B; 13%), bequests (\$23.8B; 8%), and corporations (\$14.1B; 4%).

Potential donors are typically presented with an "appeals scale", a list of suggested amounts or scale points selected by fundraisers. Figure 2.1 presents three such scales, used for recent funding drives by Wikipedia, the United Way, and the U. N. Foundation. Each features the most common sort of appeals scale: a series of specific donation amounts, along with an "open" category. The appeals scale serves several functions, but its main role is to provide concrete anchors to help donors select an appropriate quantity; donors can, of course, also choose to give nothing, or some amount not listed on the scale, including amounts outside the range of listed values.

¹2009 is the last year for which comprehensive statistics are presently available, so it is adopted consistently for comparison purposes.



C. United Nations Foundation: unfoundation.org

Figure 2.1: Examples of Appeals Scales used by Charities

Holding aside questions involving the design of an entire *scale*, an immediate practical concern for fundraisers is simply about how much to ask for: too little, and a donor may be more likely to give, but to give less; too much, and a donor may fail to be influenced by the request, or simply not donate at all. Charities wish to maximize donations, and so must attempt to tailor their requests to avoid asking for inappropriate or suboptimal donation amounts.

Despite their ubiquity in charitable requests and fundraising, there is neither theory nor a body of empirical findings on whether and to what degree such requests, and the scales comprising them, affect individual donor behavior. As a result, fundraisers have little rigorous guidance in assessing and optimizing their appeal requests, instead falling back on prior experience, coupled with summary metrics arising from trial and error (which, as we shall see, can produce misleading or even null results). Part of the problem in providing such guidance is the need for household-level, longitudinal data on both charitable requests and outcomes - "whether" and "how much" - which charities typically possess, along with a (suitably heterogeneous) statistical model for scale attraction effects, which they typically do not.

Some of these issues have been addressed in prior literature, for example, Desmet & Feinberg (2003) and De Bruyn & Prokopec (2011), each of whom examined scale effects statistically via recourse to both internal (donors' latent, planned amounts) and external (how much one is asked for) reference points (Mayhew & Winer, 1992; Mazumdar & Papatla, 2000). Although both detected scale-based effects, neither was able to incorporate heterogeneity (the basis of individually-tailored appeals), seasonal variation in giving (which is pronounced in our empirical application), nor simultaneously account for whether and how much to give, which can lead to selection biases (Van Diepen et al., 2009; Wachtel & Otter, 2011). In this paper, we resolve these and several other issues via a novel model that measures *individual-level* scale attraction effects. The model, which builds upon a classic Type 2 Tobit formulation, is calibrated on donation history panel data from a French charity.

The remainder of the paper is organized as follows. We first provide a concise overview of prior literature on scale attraction, donation behavior, reference effects, and related areas. We then describe our empirical application, develop the model, and present both empirical results and model comparisons. An illustrative simulation exercise examining the effect of tailored appeals scales is followed by potential limitations and associated future research.

2.3 Literature Review

The contextual effects of scaling on responses have been intensively examined in social psychology over the past two decades. Schwarz (1999)'s comprehensive review suggests that features of research instruments - question wording, format, and scaling, among others - can substantially affect respondents' self-reported behaviors and attitudes. In particular, response scales presented to respondents are far more than a simple "measurement device," but can work as reference frames that directly influence respondents' judgments (Schwarz et al., 1991).

Researchers working in the area of social norms have found them to systematically influence human behaviors. Individuals seek out social norms to better understand or more effectively react to social situations they encounter, especially under high uncertainty (Cialdini & Goldstein, 2004). Fisher & Ackerman (1998) support this "normative" perspective in their studies on volunteerism, and several studies have examined the effects of social information on donation behavior specifically. It has long been observed that manipulating such information (i.e., what *other* donors gave) can strongly affect donation behaviors (Reingen, 1982); Shang et al. (2008) and Shang & Croson (2009) found exactly this in a field test for a national radio fundraising campaign. When other donors' behavior is not disclosed during a donation appeal (which is typical), respondents are more uncertain in deciding a donation amount, so a given set of response alternatives - an appeals scale - can provide contextually normative information via the location (i.e., distribution) of its scale points (Schwarz et al., 1991).

Many studies have addressed charitable donations directly, and examined the role of request size on donation behavior (amount and compliance) in laboratory and field data (Doob & McLaughlin, 1989; Fraser et al., 1988; Schibrowsky & Peltier, 1995; Weyant & Smith, 1987). Although contexts and methods vary across them, these studies largely confirm scale manipulation effects, yet differ as to whether they affect donation likelihood, donation amount, or both (see De Bruyn & Prokopec, 2011, for review). These differences

may have originated from variations in compliance techniques, solicitation methods, and the suggested donation amounts. One particularly compelling potential source for inconsistencies across prior studies is lack of an account of *internal* referents. In the words of De Bruyn & Prokopec (2011), "... most fundraising research to date has overlooked the crucial role of a donor's internal reference point in moderating the impact of appeals scales on behavior." In marketing specifically, reference price theory has been a cornerstone of consumer behavior research, and supported empirically in dozens of studies (Kalyanaram & Winer, 1995, provide an extensive review).

We make especial use of one of the key findings from this literature: that two distinct kinds of reference prices play a role in choice decisions. One is *internal* reference prices, consumer-specific, memory-based amalgams of actual, recent (and "fair") prices; the other is external reference prices, present at the time of purchase. It is well-known that both internal and external reference points play a role in consumer purchase decisions (Mayhew & Winer, 1992); in donation contexts, the former is characterized by what the donor typically gives and/or plans to give, the latter what the donor is asked to. Specifically, the internal referent is an unobservable construct that must be inferred from other observable information (e.g., past donation behavior), while external referents are those presented at the time of the request via the appeals scale. Prior work in donations was unable to employ both referents, since individual-specific donation histories were lacking. Thus, researchers were unable to avail of potential donors' internal referents when designing scales for experiments. This may have led to inconsistent scale manipulation results as reported by Weyant & Smith (1987) vs. those of Doob & McLaughlin (1989). Weyant & Smith (1987) found no significant difference in the average donation amount between the "smaller request" and "larger request" conditions, only in donation rate. Assimilation-contrast theory (Sherif et al., 1958) suggests that stimuli are evaluated with regard to a point of reference based on previous experience, and so depend on a "latitude of acceptance"; Doob & McLaughlin (1989) suggest that the listed amounts in the "larger request" condition fell outside this

latitude of acceptance, and so had little effect on donors. When more plausible amounts (i.e., lower) were substituted in the larger request condition, they found a significant difference in the average donation *amount*, but none in donation *rate*. In short, taking account of appropriate *internal* referents literally reversed the pattern of substantive results.

Another potential source of inconsistencies involves not accounting for heterogeneity in internal referents. Most previous studies could avail only of aggregate data (e.g., control / experimental group, or segment level; e.g., Desmet & Feinberg, 2003) to assess the mean scale manipulation effect across conditions. Because donor-specific internal referents were unavailable, group-level may dilute the effect of scale manipulation. In this regard, De Bruyn & Prokopec (2011) were unique in having obtained each donor's last donation before the field experiment, used it a proxy for a donor's internal referent. Despite this advance, the "one shot", before-after nature of their data precludes incorporating parametric, "unobserved" heterogeneity, which likewise plagues all prior studies relying on crosssectional data. In a similar vein, no previous study of which we are aware reflects seasonal variation in donation patterns: donors are more likely to give, and/or give more, at certain times of year, such as Christmas in the U.S.; results may therefore be sensitive to when data are collected, especially so for field experiments. For these and other reasons, a panel of individual donors provides by far the best platform to detect and measure scale effects. Panel data further enables us to examine donors' internal referents evolve over time, as well as provide a fully heterogeneous account of scale attraction effects. This information is critical in designing optimal, dynamic appeals for each donor separately.

Lastly, none of the studies that employed scale manipulation provided a unified account of both donation incidence and donation amount. Models should not simply presume that *whether* to donate and *how much* to donate are behaviorally or econometrically unrelated. Doing so could introduce well-known measurement errors (Heckman, 1979). An especially appealing modeling framework is afforded by a Type 2 Tobit model, which comprises two components: one accounts for selection ("did they donate?"), the other the conditional

output of interest ("if so, how much?"). In marketing, Type 2 Tobit models have been deployed to analyze disparate consumer decisions making processes (e.g., Donkers et al., 2006; Van Diepen et al., 2009; Ying et al., 2006; Zhao et al., 2009), with the degree of selectivity between incidence and amount represented by a correlation parameter. Most relevant to our research, although not involving scale manipulation specifically, Donkers et al. (2006) and Van Diepen et al. (2009) used such a model in donation contexts, but with somewhat different results: Donkers et al. (2006) found a small negative correlation, while Van Diepen et al. (2009) found a very large positive correlation. We return to this point later when discussing our own results.

2.4 Data Description

Our data were provided by a French charity that conducted a large-scale field experiment as part of a national fundraising campaign. The charity holds three fund-raising drives a year, at Easter, June, and Christmas. Data were collected for three and a half years, from Easter 2000 to Easter 2003, for 10 donation appeals in total. The database contains household-level records for the appeals scale presented to donors, whether a donation was made and, if so, the donation amount. Donation appeals were made by door-to-door canvassing to "regular" donors; the charity judged regularity based on each donor's frequency (the number of donations during past two years) and recency (the number of periods since last donation). Subjects were partitioned into two groups ("levels" 1 and 2) according to their average donation amounts over the two years prior to the start of the experiment. Household-averaged donations in the level 1 and 2 groups fall within 100 FF-199 FF and 200 FF-399 FF, respectively.²

²The currency unit in the data is French Francs (FF), trading during the collection window at approximately 7 to the US dollar.

Table 2.1: Appeals Scales used in the Field Experiment

		Standard Scale						
	100 FF	150 FF	250 FF	500 FF	1000 FF	Other		
Prior Donation level		Test Scales						
1	120 FF					Other		
2	120 FF	200 FF	350 FF	500 FF	750 FF	Other		

The charity sought to better understand the role of appeals scales in donation behavior, so manipulated it by randomly assigning respondents to receive either a "standard" or a "test" scale. The standard scale had previously been used for all subjects prior to the experiment, and thereby helps establish a baseline. Scales all consisted of five suggested amounts (e.g., 100, 150, 250, 500, 1000 FF for the standard scale), as well as an "Other" category, which allowed donations below or above all five scale points, or between any adjacent pair. The test appeals scales manipulated these five suggested amounts; these all appear in Table 2.1.

The charity thereby implemented a 2×2 design: (prior donation) "level 1" or "level 2" \times random assignment of either a "standard" or "test" appeals scale. It is important to note that the charity was collecting real donations, and therefore did not have the luxury of 'optimally' designing the scale for the purposes of the experiment, such as orthogonalizing, including extreme values, and the like. Thus, the points comprising the "test" scale for the level 2 (higher prior) donation group were higher than those used in the test scale for the level 1 group. This 'endogeneity' is a data limitation over which we had no control, and our model will take care to treat scales as a collection of anchor points, in part to mitigate this concern.

Table 2.2: Average Donation Amounts and Frequencies

Prior Donation	Sagles	Average Dona	Yield Rate	
Level	Scales	per Household	per Occasion	Heia Kale
1	Standard	430.2	136.5	31.73%
1	Test	434.3	137.3	31.61%
2	Standard	844.7	286.2	33.88%
2	Test	839.5	283.8	33.81%

Two hundred households from each of the four groups were randomly selected for analysis. Table 2.2 presents descriptive statistics for each, average donation amount (per household and per occasion), and yield rate. Level 1 and 2 differ substantially in per-household and in per-occasion average donation amounts; this is unsurprising, as the baseline donation amount was used by the charity to partition donors into different levels. However, yield rates are remarkably similar across the four groups, with all between 32% and 34%. Moreover, each of the descriptive statistics - yield rate and both per-occasion and per-household amount - fails to differ across the standard and test scales, within a donation level (1 or 2). One might therefore conclude that there were *no* effects attributable to the use of the test scale. As our analysis will show, such a conclusion based on aggregated metrics is not only premature, but highly misleading.

Table 2.3 suggests a clear (aggregate) seasonal pattern in both yield rate and average donation amount: people give more, and more often, at Easter than during June or Christmas. The difference in yield rates is striking - approximately $\frac{3}{4}$ of respondents donate at Easter (an important holiday in France), while under $\frac{1}{4}$ do at the other times of year - and these proportions are nearly identical in the level 1 and 2 donation groups (the latter, by construction, has higher donation *amounts* across the board). Holding aside any aggregate patterns, there is nonetheless sizable variation in *household*-level donation profiles. Table 2.4 presents donation histories for five households from the level 1/standard scale group, for illustrative

purposes; considerable heterogeneity in timing (and some in amounts) is apparent. For example, households #3, 66, and 118 seem to be a "100FF in Easter, only", a "not in June", and a "never at Christmas" giver, respectively. By contrast, household #148 has no obvious seasonal or amount pattern. It is these variations in donation patterns - both incidence, and amount - that we will model, in order to estimate the degree of "pull" of the appeals scale, which itself will vary across donors.

Table 2.3: Yield Rate and Average Amount of Observed Donations across Seasons

	Level 1			Level 2		
	Easter	June	Christmas	Easter	June	Christmas
Yield Rate	72.8%	18.6%	19.6%	75.3%	17.6%	20.9%
Average Donation	140.1	126.7	129.7	265.0	221.7	215.3

2.5 Model Development

2.5.1 Internal and External Reference Points

The model hinges on two assumptions, as discussed previously: that, for a particular request, each donor has some (latent) quantity, which serves as an *internal* referent (r^I) ; and that the request itself provides a set of alternatives, in the form of the appeals scale, that serve as *external* referents (r^E) . If an appeals scale contains multiple points, we denote the k^{th} as $r^{E,k}$.

The internal referent admits different operationalizations; because it is unobserved, it must be inferred based on data and the model. The reference pricing literature offers several contenders; among the most common are last price paid (Krishnamurthi et al., 1992; Mayhew & Winer, 1992) and a (perhaps weighted) average of past prices (Kalyanaram & Little, 1994; Lattin & Bucklin, 1989; Mazumdar & Papatla, 2000; Rajendran & Tellis, 1994), and

we will empirically compare them. We include two additional specifications that can account for seasonal donation variations; so, the four (donor-specific) internal reference point models estimated are: the average of all prior observed donation amounts (IR-1); the last observed donation amount (IR-2); the average observed donation amount at the same time of year (IR-3); and the last observed donation amount at the same time of year (IR-4).³

That the external reference points are observable might make them appear simple to model. This might be so were there only a single requested amount. But, in practice, there are usually many, and so it is unclear how they exert their "joint pull": perhaps only the extremes are noticed; or only those nearest the internal reference have any influence; or some summary measure of all points (like the average or median); or something else entirely. We therefore empirically examine five such formulations, where influence is exerted: by all scale points (ER-1); by the two scale points closest (above and below) the internal referent (ER-2); by the largest and the smallest scale point (ER-3); by the median (i.e., middle) of the scale points (ER-4); or by the mean of all scale points, which itself is typically *not* a point on the scale (ER-5). We consider such a wide range of possibilities because there is no prior theory to suggest how a *group* of referents exert collective influence. In fact, we view this as among the most intriguing open questions that our data and model can help address. Note that, when multiple points are presumed to exert influence (as in ER-1, ER-2, and ER-3), we must also specify the *weight* associated with each point; we address this in detail subsequently.

2.5.2 Modeling Scale Attraction Effects

In the absence of any appeals scale - for example, if a potential donor is simply asked how much s/he would like to give - whether and how much is donated would be influenced by the internal referent, not any external ones. However, when presented with (the external

³Instead of exponential or geometric time discounting, we used simple averaging, i.e., equal weights. Given the small number of observations per donor (2.99, on average), the difference between the formulations is minor.

referents of) the appeals scale, observed behavior may be affected by both the internal and external referents. One way to visualize this is that the internal referent is "pulled" by the external ones, and that these separate pulls (if indeed more than one external referent is "noticed", as in ER-1-3) can cumulate in their effects. A simple metric for how influential a scale point is its "compliance degree," which we describe next.

2.5.2.1 Compliance Degree

We define CD^k , "compliance degree" of the k^{th} external reference point as the proportional increase (or decrease) in donation amount from a donor's *internal* reference point (r^I) to an *external* one $(r^{E,k})$. More formally (with DA = Donation Amount received):

$$CD^k = \frac{DA - r^I}{r^{E,k} - r^I} \tag{2.1}$$

For example, if a donor is "planning" to give (i.e., has an internal referent of) \$100, but is asked for \$101, he will be very likely to comply, in which case both the numerator and denominator are \$1 and $CD^1 = 100\%$ (the superscript "1" indicates there was just one external reference point). However, if the same donor is asked to give \$200 more (i.e., \$300), the donor is less likely to fully comply; if the resulting donation is instead \$140, $CD^1 = (\$140 - \$100)/(\$300 - \$100) = .2$, or 20%. In simple language, the donor "came up 20%" from a \$100 baseline. An analogous calculation pertains to external referents below the internal one.

It is convenient to define the *distance*, d^k , between the k^{th} external and the internal referent as an incremental/decremental ratio.

$$distance(d^k) = \frac{\left\| r^{E,k} - r^I \right\|}{r^I}$$
 (2.2)

This allows both compliance degree as well and the pulling amount (described later) to be expressed as a dimensionless quantity for each donor. This in turn helps to unify the model;

for example, it can treat the response of a donor planning to give \$10, but asked to donate \$20, similarly to that of one planning to donate \$100, but asked for \$200.

We will model both upward and downward "compliance degree curves", which satisfy three properties:

- 1) $CD^k \approx 1$ for $d^k \approx 0$: "Maximal compliance occurs near donors' internal referents."
- 2) CD^k decreases monotonically in d^k : "Compliance is worse for requests further from the internal referent."
 - 3) $CD^k \ge 0$: "Compliance can't be worse than zero."

Properties 1 and 2 suggest donation is highly responsive to asking for amounts close to what was 'planned' (the internal referent), but increasingly less so for distant amounts. This is consistent with "latitude of acceptance" in Assimilation-Contrast Theory (Sherif et al., 1958), which has found prior support in a donation context (Doob & McLaughlin, 1989). Property 3 simply suggests that requests can be ignored, but do not literally repel donors from a scale point.

There are many ways to specify compliance degree curves satisfying these three properties, including using fully parametric (e.g., polynomial), semi-parametric, or non-parametric formulations. We select a translated gamma kernel function, for two reasons. First, it provides a parsimonious, yet flexible, functional form that naturally satisfies properties 1-3; this parsimony is important for a *heterogeneous* account to be identified, given the small number of responses per donor during the data window. Second, the gamma kernel enables the pulling *amount* curves (described later) to follow a unimodal, yet flexibly-shaped, distribution, which in turn facilitates eventual optimization. Thus, we arrive at an especially simple form:

$$CD^{k} = \frac{\exp\left(-\frac{d^{k}+1}{\theta}\right)}{\exp\left(-\frac{1}{\theta}\right)} = \exp\left(-\frac{d^{k}}{\theta}\right); \qquad \theta = \begin{cases} \exp(\beta^{U}), & r^{E,k} \ge r^{I} \\ \exp(\beta^{D}), & r^{E,k} < r^{I} \end{cases}$$
(2.3)

where $\theta > 0$ is the gamma kernel scale parameter.

The compliance degree curve follows from a gamma kernel with "shape parameter" 1 and "scale parameter" θ . ⁴ This is then both translated and normalized - first horizontally translated by -1 so that it crosses the y-axis, then normalized to have a value of 1 at the origin - after which it follows a translated gamma kernel, anchored at (0,1) with curvature determined by the scale parameter. Note that there are actually two different compliance degree curves, depending on the relative location of the internal and the external referents. When $r^{E,k} \geq r^I$, we have an "upward" compliance degree curve, and otherwise a "downward" one.

Since the scale parameter (θ) must be positive, we specify β^U or $\beta^D = \ln(\theta)$, where β^U and β^D are the "upward" and "downward" parameters in (2.3). Figure 2.2 depicts both curves, which can have a variety of shapes, for different values of β^U and β^D . However, $\beta^U = \beta^D$ does not imply identical upward and downward curves, because the domain of the downward curve is bounded by 100%, since one cannot give less than zero (i.e., a 100% decrement).

⁴Fixing the shape parameter at 1 yields a non-negative, monotonically decreasing, convex curve (with regard to the origin), satisfying properties 1-3. Numerous simulations showed recovery of *two* parameters (both scale and shape) was very poor, suggesting weak identification in data generated to resemble ours.

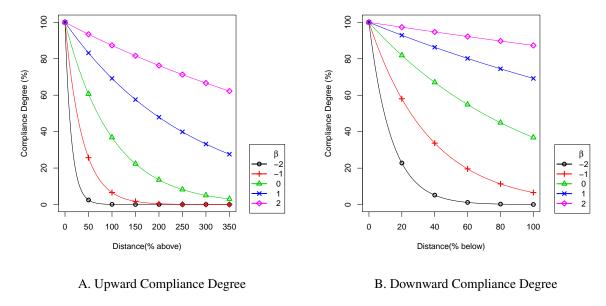


Figure 2.2: Compliance Degree Curves

2.5.2.2 Pulling Amount

The pulling amount (PA^k) represents the size of effect exerted by a scale point, a simple matter of multiplying compliance by the (Euclidean) distance between the internal (r^I) and the kth external reference point $(r^{E,k})$:

$$PA^k = CD^k \times ||r^{E,k} - r^I|| \tag{2.4}$$

The pulling amount suggests a trade-off between asking for too little and asking for too much: If a charity asks too little - that is, just a bit more than the internal referent - compliance (CD^k) may be high, but the potential surplus $(\|r^{E,k}-r^I\|)$ is small. On the other hand, if a charity asks too much, the compliance degree may be low, while the surplus is large. In light of this trade-off (where the extremes are literally zero), optimizing donation drives requires considering both elements, that is, asking for a judiciously chosen amount from each donor.

Each of the two compliance degree curves therefore gives rise to a "pulling amount"

curve: $r^{E,k} \geq r^I$ corresponds to "upward" pulling, $r^{E,k} < r^I$ to "downward". The simple nature of (2.4) implies that these curves also follow a gamma kernel, with shape parameter 2 and scale parameters $\exp(\beta^U)$ and $\exp(\beta^D)$. As depicted in Figure 2.3, these curves can have many shapes: the upward pulling curve has domain $[0,\infty)$, is unimodal (and thus has a unique maximum), with zero at the origin and asymptoting to zero for large d (for any β^U). The domain of the downward pulling amount curve is [0,1]; it is unimodal (with unique maximum) if $\beta^D < 0$, and is monotonically increasing otherwise (with maximum at 1). These internal maxima allow us to derive a closed-form expression for optimal, donor-specific scale points, discussed in the section on the effect of individually tailored appeals scales.

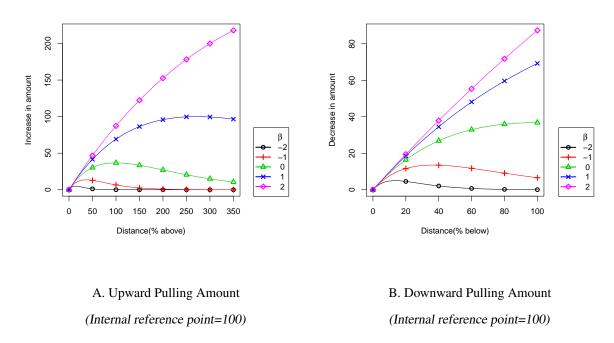


Figure 2.3: Pulling Amount Curves

2.5.2.3 Accumulating Scale Attraction Effects

Because real appeals scales almost always comprise multiple amounts, their effects need to be somehow combined. Figure 2.4 illustrates the "accumulated pulling amount" accruing from multiple external referents; to match our empirical application, five external referents

are depicted, with two distinct upward and downward curves on either side of the graph. Here, scale points 1, 2, and 3 are greater than the internal referent (set by convention to d = 0), so each induces an upward pull on donation amount, tending to increase it. By contrast, scale points 4 and 5 are lower than the internal referent, tending to pull the observed donation downward.

The discussion thus far concerns the pulling amount for individual scale points, not how to combine them. Just as we considered a number of specifications for the effects of the internal and external referents, we will do so for this combination. Before detailing these, we highlight one simplifying assumption: that the effect of a particular scale point can be modeled separately from the existence or the location of the others. This is dictated by a data limitation: the charity did not change scales (over the course of the experiment, nor within each of the four donation groups), so that identifying *interactions* between scale points is not possible. Even were this not the case, such interactions would greatly weaken gamma kernel parameter (β^U and β^D) identification, owing to the small number of observations per donor (and again to the lack of within-donor scale variation during the experiment).

While at first blush such independence assumptions may appear unrealistic, they are mitigated by the weighted-averaging schemes explored for the "accumulated pulling amount", or APA. We examine three: i) the sum; ii) the mean; and iii) the weighted mean of the pulling amounts. Each is described as follows, along with potential caveats. In general:

$$APA = \sum_{k=1}^{K} w^{k} \times I^{k} \times PA^{k}; \qquad I^{k} = \begin{cases} 1, & r^{E,k} \ge r^{I} \\ -1, & r^{E,k} < r^{I} \end{cases}$$
 (2.5)

$$Sum: w^k = 1;$$
 $Mean: w^k = \frac{1}{K};$ $Weighted\ Mean: w^k = \frac{PA^k}{\sum_{k=1}^K PA^k}$ (2.6)

Sum. Simple summation appears to be the most direct way to accumulate the separate pulling amounts. However, this specification has two inherent problems. First, the

predicted donation amount can lie above the largest, or below the smallest, scale point. Although this is not impossible, our data contains very few instances in which the donation exceeds the largest scale point. Second, the effect of including *additional* scale points can be overstated (something that, in our data, will not be testable, since the charity fixed this at 5). For example, given an internal referent of 50, the APA of an appeals scale with four points of 9, 11, 99, and 101 is about twice as large as for one with two scale points of 10 and 100, which seems decidedly unrealistic.

Mean. The mean specification retains additivity and resolves the two problems with the sum, but is not without problems of its own, owing to equal-weighting. For example, if a donor is asked to give \$2000 when the planned amount is \$100, the effect of such a large scale point on APA might be small or negligible. However, equal weighting forces a large scale point like \$2000 to have tremendous effect on the APA by substantially lowering the accumulated pulling amount. A straightforward fix involves the use of a weighted mean, as follows.

Weighted Mean. This makes use of weights, w^k , which one might imagine were estimable. Two data limitations prevent this, (once again) the lack of within-donor scale variation, and that only three different appeals scales (one standard and two test) appeared in the experiment. For this reason, and because we include heterogeneity β^U and β^D , even homogeneous w^k proved impossible to estimate.⁵ Therefore, the weight is set in proportion to the size of pulling amount, based both on conceptual appeal and trial of multiple alternate schemes (which we do not report here). The key point is that the weighted mean allows a scale point with a larger pulling amount to contribute more to the total pull, unlike for either of the previous two specifications.⁶

⁵De Bruyn & Prokopec (2011) tried to estimate each scale point's weight, which they term "absolute attraction weight". However, they could estimate only the weight of the smallest of four scale points, while fixing those of the other three to 1. They attribute this identification problem to the inherent correlation across suggested donation amounts on the appeals scale (i.e., suggested amounts increase monotonically and are highly correlated).

⁶In fact, we found model with weighed mean specification (explained next) fits better than that with equal weight mean specification, keeping all other model components the same: The RMSE and MAD of the former are 0.257 and 0.188 respectively; for the latter, are 0.271 and 0.195.

2.5.3 General Model(*Type 2 Tobit*)

We begin by outlining the general model structure. The model has been set up to allow a "dimensionless" account of pulling effects, so that heterogeneity can be specified across the log-scale for donation amount. As discussed earlier, we use a Type 2 Tobit model (Amemiya, 1985) to jointly account for donation incidence and amount, as follows:

$$y^{s*} = X^{s}\beta^{s} + \epsilon^{s}$$

$$y^{a*} = \ln(r^{I} + APA) + X^{a}\beta^{a} + \epsilon^{a}, where:$$

$$y^{s} = 1, \text{ if } y^{s*} \ge 0; \text{ 0 otherwise}$$

$$y^{a} = y^{a*}, \text{ if } y^{s} = 1; \text{ unobserved otherwise}$$

$$(\epsilon^{s}, \epsilon^{a}) \sim BVN(0, \Sigma_{\epsilon}); \quad \Sigma_{\epsilon} = \begin{bmatrix} 1 & \rho\sigma \\ \rho\sigma & \sigma^{2} \end{bmatrix}$$

The subscripts i and t (for donor and time) are suppressed, and X^s and X^a are covariates in the selection (s) and amount (a) equations, respectively, which we detail below.

In the amount equation, let y^{a*} denote the log of the latent donation amount, which is observed only when a donation is made, that is, when y^s is 1, which occurs when the latent variable $y^{s*} \geq 0$. The error terms of the selection and amount equations (ϵ^s and ϵ^a) follow a bivariate normal distribution; the variance of ϵ^s is fixed to 1 for identification. It is important to note that we model the logarithm of donation amount, for several reasons: first, it allows ϵ^a to be plausibly homoscedastic; second, it allows all effects in the amount equation to enter multiplicatively; and third, it allows for coefficient heterogeneity to act on a dimensionless quantity, which we address in detail shortly.

The amount equation (for y^{a*}) contains two deterministic components. The first is the sum of a donor's internal referent (r^I) and the accumulated pulling amount (APA). The second is all factors $(X^a\beta^a)$ that affect the donation, other than those stemming from

the appeals scale. Scale-based effects do not appear directly in the selection equation, because in our data all scales used were set in "reasonable" ranges for every donor (recall that these were real donors, and the charity was understandably reluctant to alienate them with unrealistically high requests, or lose funds with low ones). Hypothetically, were all or many of the suggested amounts exceedingly large, it is possible that the donor would become annoyed and give nothing. Therefore, although we cannot preclude this possibility for all data settings, for ours the appeals scale can exercise influence on donation incidence only indirectly, via the correlation, ρ . [We did estimate a model allowing for scale effects in selection; the APA coefficient in selection was ns.]

Table 2.4: Examples of Donation Histories for Several Randomly Selected Households

id#		2000		2001			2002			2003
	Easter	June	Xmas	Easter	June	Xmas	Easter	June	Xmas	Easter
3	100	0	0	100	0	0	100	0	0	100
20	0	150	0	0	150	0	150	0	0	250
66	200	0	150	200	0	150	150	0	150	250
118	100	100	0	100	100	0	100	0	0	150
148	0	90	0	100	0	100	150	150	100	150

2.5.4 Explanatory variables and Heterogeneity

2.5.4.1 Explanatory variables

Selection Equation

The selection equation contains three types of explanatory variable (X^s) , which we detail subsequently: seasonal indicators, (log of) prior donation, and "level" fixed effects. Table 2.3 reveals strong aggregate seasonal variation in donation likelihood, by far highest at Easter; Table 2.4 suggests household-level variation as well. Previous studies, which were mostly one-shot, could not account for such seasonal variations, which are critical in our data. Three dummies - Easter (X_{it}^E) , June (X_{it}^J) , and Christmas (X_{it}^C) - represent when the donation request occurred.

The log of (1+ amount the donor gave on the last request), donated X_{it}^{lag} , is included to examine carryover effects; as discussed previously, logs help retain error homoscedasticity, among other benefits. The donation amount itself is not censored, but truncated, at 0; so, X_{it}^{lag} is 0 when we observe no donation taking place. The directly prior donation (0 or otherwise) may affect the decision to donate for several reasons: these are regular donors, so are likely aware of their donation patterns, and the interval between solicitations is relatively short. ⁷

Although Table 2.3 shows only modest differences in yield rate between the "larger" (level 2) and "smaller" (level 1) donation groups, we still include an appropriate dummy (X_i^{level}) among the selection covariates, to control for potential differences in baseline donation likelihood *after* accounting for seasonal patterns. Coefficients for the three seasonal dummies, the log-donation lag, and the level dummy, are denoted β^E , β_i^J , β_i^C , β^{lag} , and $\beta^{level,s}$, respectively.

In the experiment, donors were randomly assigned to receive either a standard or a test appeals scale, so no dummies were entered for this difference (in either selection, or amount). Doing so failed to improve fit, in any case, so we do not discuss these again.

Table 2.5: Mean and SE of observed amounts (FF) for each donor

	Observed Don	ation Amounts
Prior Donation level	Mean	SE
1	137.71	22.60
2	259.05	47.43

⁷We estimated a similar model by replacing the log-donation lag with an indicator variable for *whether* one in the previous period, finding poorer in-sample fit. This may be because the continuous variable (log-donation) carries additional information, compared with a simple indicator variable.

Amount Equation

Based on examination of the data and unimproved fit of models including them, seasonal dummies are not included in the amount equation; the somewhat higher amounts indicated at Easter in Table 2.3, for example, will be well-explained by other covariates, like lags in setting "internal" referents (such as in IR-3 and IR-4). Table 2.4 shows far greater household variation in when to give, not how much; and both Figure 2.5 and Table 2.5 suggest that household-level seasonal variation in amount is very small for most donors. Similarly, we do not include a lag for prior donation amount. This may seem paradoxical, but recall that there is little *within-donor* variation in observed donation amount, suggesting that that people do not say, in effect, "I give more than usual last time, so will give less this time," or vice versa. ⁸ Lastly, although donation amount is mainly predicted by a donor's internal referent and scale effects, a level dummy (X_i^{level}) is included to account for the difference in baseline donation amount between the two groups, denoted $\beta^{level,a}$.

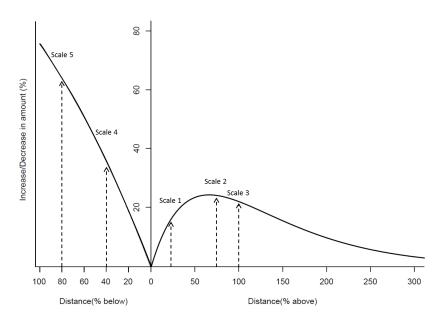


Figure 2.4: Pulling amounts owing to multiple scale (external reference) points

⁸To verify this choice, we estimated a series of models with $\ln(r^I + APA + \beta * PriorDonation)$ instead of the analogous term in the amount equation. For all reference point models (IR1-4), AIC, BIC, and in-sample fit does not show any improvement.

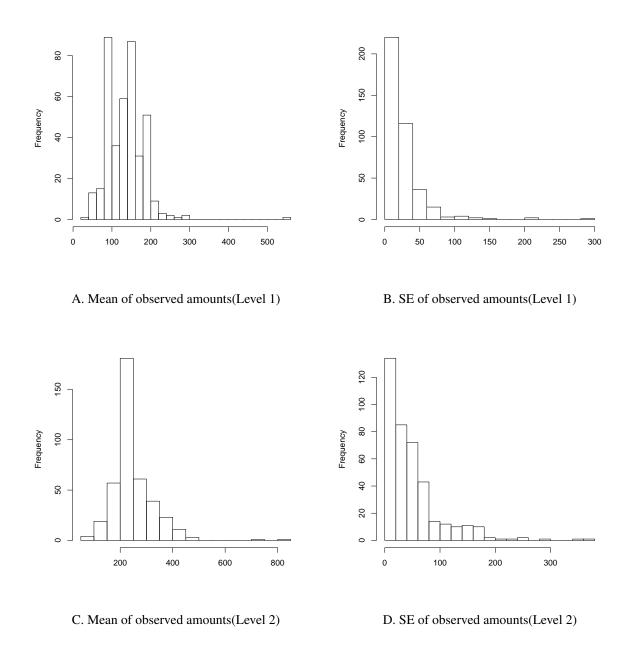


Figure 2.5: Distribution of Mean and SE of observed amounts for each donor

2.5.4.2 Heterogeneity

It is critical, in a model for household-level behavior, to incorporate "unobserved" heterogeneity, which we do in several ways. Given the large household-level seasonal donation variation, we model heterogeneity in the seasonal dummies for the June and Christmas

coefficients (β_i^J) and β_i^C . Our empirical results suggested that household-level seasonal donation patterns were well reflected in heterogeneity for β_i^J and β_i^C , owing perhaps to much larger variation in giving in June and at Christmas.

Importantly, since our model is primarily meant to capture scale attraction effects, the two gamma kernel parameters (β_i^U and β_i^D) in the amount equation are heterogeneous. Imposing heterogeneity on the gamma parameters - especially "upwards", β_i^U - is crucial for formulating tailored appeals scales, which require identifying the request amount with maximum effect in "pulling" up a donor's internal referent. If β_i^U were homogeneous, each donor's optimum would be the *same* percentage above his/her internal referent. This might still provide a helpful guideline for fundraisers, but presumes all donors are equally 'elastic' in being cajoled upwards. Our results, in fact, will strongly weigh against this presumption. We similarly account for heterogeneity in the "downward" parameter, β_i^D , though it will play a lesser role in optimization.

Our formulation therefore specifies four heterogeneous parameters, to be recovered from the relatively short data window of 7 occasions; the 42.7% aggregate yield rate suggests that about 3 of these 7 requests resulted in donations, on average. Although it may appear ambitious to account for 4 household-level parameters based on relatively little data, simulations showed good recovery for all four heterogeneous parameters, and excellent recovery of the others.

2.6 Estimation

The full model (see Appendix A) is estimated using Markov chain Monte Carlo methods. Data augmentation (Tanner & Wong, 1987) converts the model to a Bayesian Hierarchical Seemingly Unrelated Regression. We obtain posterior draws via Metropolis-within-Gibbs

 $^{^9}$ Extensive simulations for data matching ours in marginal (summary) statistics failed to recover the true parameters - mean vector and covariance matrix for β_i^E , β_i^J , β_i^C - when the Easter, June and Christmas coefficients were all heterogeneous. Restricting the most common donation period (Easter, with a 74.1% yield rate) to be homogeneous led to nearly perfect parameter recovery.

algorithms: Gibbs sampling (Geman & Geman, 1984) if the full conditional of a parameter block is of known form, and Metropolis-Hastings, with a random walk proposal (Chib & Greenberg, 1995), otherwise. We set diffuse priors for all parameters of interest; detailed procedures appear in Appendix B. All estimates are based on 100,000 draws. We discard the first 50,000 draws for burn-in, and use the last 50,000 (thinned to every tenth) to calculate posterior densities. Gelman-Rubin scale reduction factors, using 5 chains with different stating points, are below 1.1 for almost all parameters, suggesting good convergence (Brooks & Gelman, 1998).

Table 2.6: Parameter Estimates for Full Model

	Coefficient	mean	SE	95% HDR
	correlation (ρ)	-0.387	0.049	(-0.479, -0.288)
SN	sd of log amount (σ)	0.296	0.005	(0.286, 0.307)
eneoi	Easter dummy (β^E)	0.782	0.035	(0.714, 0.852)
Homogeneous	level dummy in selection $(\beta^{level,s})$	0.073	0.039	(0.002, 0.156)
He	level dummy in amount $(\beta^{level,a})$	0.273	0.019	(0.237, 0.311)
	log amount lag in selection (β^{lag})	-0.131	0.009	(-0.149, -0.113)
	June dummy (β_i^J)	-0.554	0.058	(-0.668, -0.441)
	Christmas dummy (β_i^c)	-1.019	0.070	(-1.163, -0.889)
	"gamma up" (β_i^U)	-0.418	0.070	(-0.563, -0.297)
	"gamma down" (β_i^D)	1.278	0.224	(0.858, 1.731)
	sd(June)	0.406	0.071	(0.277, 0.554)
sn	sd(Christmas)	0.831	0.088	(0.667, 1.010)
Heterogeneous	sd(gamma up)	0.478	0.040	(0.404, 0.565)
terog	sd(gamma down)	0.932	0.154	(0.672, 1.260)
He	corr(June, Christmas)	0.742	0.108	(0.506, 0.906)
	corr(June, gamma up)	-0.004	0.050	(-0.101, 0.094)
	corr(June, gamma down)	0.003	0.050	(-0.096, 0.103)
	corr(Christmas, gamma up)	0.002	0.049	(-0.097, 0.096)
	corr(Christmas, gamma down)	0.006	0.050	(-0.093, 0.102)
	corr(gamma up, gamma down)	0.380	0.145	(0.071, 0.626)

2.7 Results

For brevity, we only present full estimation results for the model with IR-1 (average of all observed donation amounts) and ER-1 (all scale points), as these provided the best fit compared with all possible combinations of the other internal and external references point

formulations (IR 2-4 and ER 2-5). Table 2.6 summarizes posterior means and standard errors for all parameters, and detailed model comparison statistics appear in the following section.

2.7.1 Error Correlation in Selection and Amount equations

The mean of the marginal posterior for the correlation (ρ) between the selection and amount equation errors is negative (-0.387), and the 95% highest density region does not include zero. This suggests that unmeasured factors influencing selection are correlated with those influencing amount, and operate in opposite directions. The size of the correlation is moderate: neither close to 0 nor to 1. This differs from findings in previous research using related model formulations; for example, Donkers et al. (2006) found the correlation to be negligible and negative (-0.033), while Van Diepen et al. (2009) found it to be very large and positive (0.958). A very small correlation fails to help correct for potential selection biases, and could reflect large, independent sources of error in each equation. Conversely, a large correlation might suggest important variables omitted in both equations.

It is difficult to generalize such results, since our model accounts for scale attraction effects, while prior ones do not. We did, however, find significant, moderate, negative values of ρ across a very wide range of candidate models, indicating that selectivity needs to be accounted for in our data. One interpretation of this finding, which is apparently robust, is that, knowing one has donated, the *conditional* expectation of the donation is smaller. Thus, models that account for "whether" and "how much" *separately* may overestimate total expected yield.

2.7.2 Selection: Seasonality

Comparing the Easter coefficient (0.782) to the means of the (heterogeneous) June and Christmas coefficients (-0.554, -1.019 respectively) accords with the aggregate benchmark, that giving is much more likely for Easter than June or at Christmas, on average (a finding

that should not be extrapolated beyond French donors to a nationwide "general purpose" charity.). There is a substantial seasonal heterogeneity: the SD of individual-level parameters for June and Christmas are 0.406 and 0.831, respectively. The high (positive) correlation between these individual-level parameters (0.742) largely reflects the fact the yield rates in June and Christmas are both low (18.1%, 20.3%) and a high proportion of donors (65.6%) gave at neither time.

2.7.3 Level Dummies and Lagged Log-Amount

The level dummy is only marginally significant (mean 0.073, SE 0.039) in selection, but significantly positive in amount (mean 0.273, SE 0.019). So, as aggregate statistics suggest, level 2 donors give more than those in level 1, but with large difference in yield rates. The coefficient of the log-donation lag in selection is significantly negative (-0.131), indicating that a larger donation *amount* last time leads to being less likely to give *at all* this time.

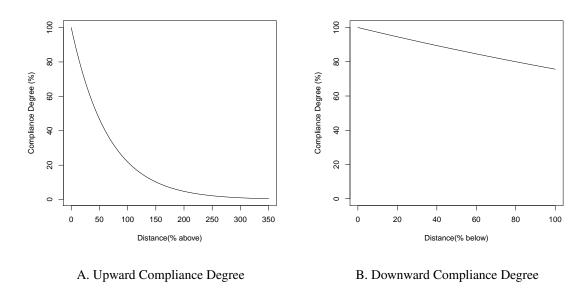


Figure 2.6: Upward and downward compliance curves at gamma posterior mean

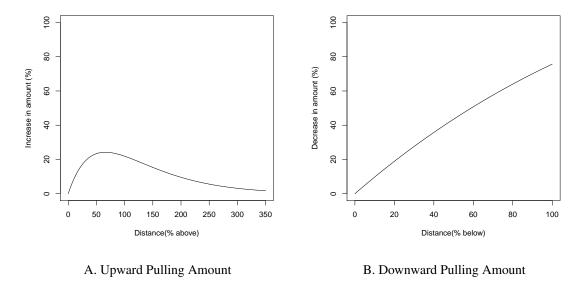


Figure 2.7: Upward and downward pulling amount curves at gamma posterior mean

2.7.4 "Pulling Effects": Gamma Kernel Parameters in Donation Amount

The values of β_i^U and β_i^D determine each donor's degree of compliance ("pull") to the scale points above and below the internal referent. Because the domains of the two compliance curves differ, we should not compare β_i^U directly to β_i^D . Instead (and ignoring for the moment the considerable variation in these across donors), Figure 2.6 shows both compliance curves at the posterior means of β_i^U and β_i^D . The downward compliance curve is far less 'pitched' than the upward. This makes intuitive sense: asking for much more than one is willing to give will eventually result in almost zero compliance, unlike asking for much less.

The compliance curves are, by construction, monotonic. By contrast, the pulling *amount* curves need not be. These are depicted, at the posterior means for β_i^U and β_i^D , in Figure 2.7. The upward pulling amount curve is inverted-U (i.e., unimodal), indicating a single "best request" value, to which we return later. By contrast, the downward curve decreases monotonically, suggesting that donors tend to give less as the suggested amount decreases (with lower bound 0).

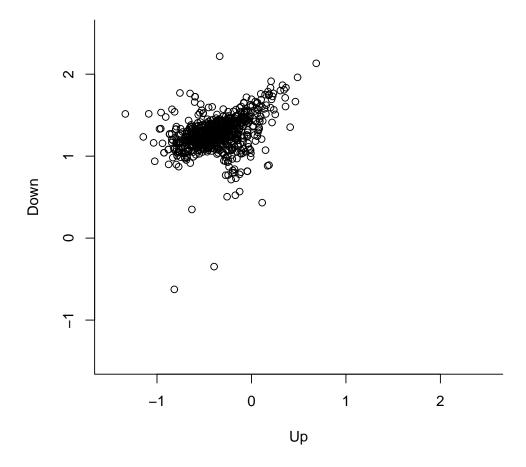


Figure 2.8: Gamma parameters (up and down) for each donor

Figure 2.8 in some sense encapsulates our main results: the upward and downward pulling parameters (β_i^U and β_i^D) for each donor. There is clearly a good deal of heterogeneity, indicating differing degrees of susceptibility to the appeals scale, *despite only modest differences in prior donation behavior*. The upward pulling parameter (β_i^U) displays larger variation (SD 0.93) than the downward (SD 0.48). This might be expected: most everyone can go along with being asked for less, but people react to being asked for more very differently.

By allowing a bivariate density for (β_i^U, β_i^D) , the model helps assess overall scale com-

pliance. Specifically, we find a substantial correlation (0.380) in these values, suggesting that donors who are "upward compliant" tend to be "downward compliant" as well. There is no reason to expect these should be correlated at all, let alone positively, and we believe this finding to be the first of its kind. This bivariate density for (β_i^U, β_i^D) leads immediately to the joint distribution of *maximal pulling amounts*, those scale points that lead to the greatest overall effects; we do not call these "optimal", since a large *downward* pull is to be avoided.

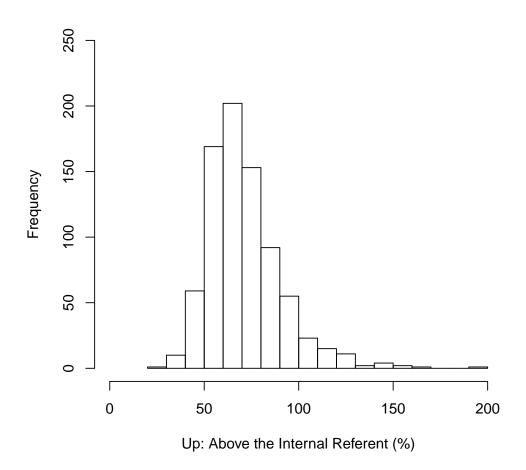


Figure 2.9: Scale point with maximum upward pull

Heterogeneity in (β_i^U, β_i^D) leads to substantial variation in maximally effective potential scale point locations, depicted for the "upward" pull in Figure 2.9 (we omit the analogous

"downward" distribution, as for most respondents these are zero). The model suggests that the scale point with maximal upward pull, which varies across donors, ranges from 27.0% to 198.5%, with a mean of 71.7%, above one's internal referent, which seems reasonable. This non-trivial variation has an important implication: that it may be possible to substantially increase donations by *personalizing* an appeals request, based on each donor's history. We discuss this possibility later, along with associated calculations.

Figure 2.10 in some sense integrates the key elements of the model, and presents its main substantive findings in the context of the original data, specifically: How much can a maximally-effective appeal (either up or down) pull from one's reference donation? It depicts, across donors, the maximal percentage increase and decrease (see Appendix E for derivation). This also allows a direct comparison of the "strength" of upward and downward scale attraction effects, heterogeneously, which was not sensible using (β_i^U, β_i^D) , given their different domains of operation. The maximum percentage increase ranges from 9.9% to 73.0% (mean = 26.3%; SD = 7.2%); maximum percentage decrease ranges from 20.5% to 89.9% (mean = 74.3%; SD = 4.8%). The correlation in these values is 0.298 (echoing the 0.380 value for β_i^U and β_i^D). Figure 2.10 suggests that the maximum percentage decrease is greater than the analogous increase for most donors: 81.6% of the donors lie above the diagonal (dotted) line. This is reminiscent of the asymmetric effects in Desmet & Feinberg (2003), whose lack of individual-level data precluded any distributions across donors, and De Bruyn & Prokopec (2011), who only had one-shot (i.e., "before" and "after") data unsuited to modeling heterogeneity or carryover effects.

¹⁰Discussions with a large university's fundraising team suggested that the success of "laddering" dropped nearly to zero when appeals hit 200% above a donor's typical or last donation amount.

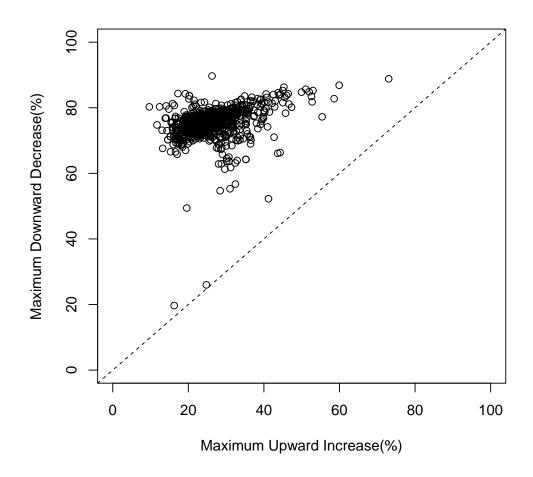


Figure 2.10: Maximum pulling up and down amounts for each donor

2.8 Model comparison

The data give clear indication of the existence of scale attraction effects. But one might reasonably question whether these were strongly dependent on the particular form of the model, four of its elements in particular: 1) internal reference point specification; 2) external reference point specification; 3) the importance of including correlation (Type 2 Tobit), seasonality, and scale effects; and 4) incorporating response heterogeneity. We examine each of these in some detail, to assess relative "contribution" to overall model fit.

With respect to internal reference formulation, we compare four, as described in the model development section, each donor-specific: the average of all prior donation amounts (IR-1); the last amount (IR-2); the average of all amounts at the same time of year (IR-3); and the last amount at the same time of year (IR-4).¹¹ We similarly examine the five external reference formulations explained earlier: all scale points (ER-1); the two scale points closest (above and below) the internal referent (ER-2); the largest and the smallest point in an appeals scale (ER-3); the median (i.e., middle) of all scale points (ER-4); and the mean of all scale points (ER-5).

We call the model with all the aforementioned components - internal and external referents; error correlation; seasonality; heterogeneity - the "full model". Alternative models include those lacking: error correlation ("no correlation"), scale effects ("no scale effect"), and both ("simple regression"). We similarly examine the effects of homogenous seasonality, homogenous scale effects, and both of these.

¹¹For IR-3, IR-4, if we don't observe donation at a certain time of year in the initialization period (first full year, or three data points), we initialize using the mean of the all observed amounts in each group.

Table 2.7: In-sample fit of observed donation amounts

1. Heterogeneous seasonality and scale effects

	The state of the s															
	Full model				No correlation				No scale effect				Simple regression			
	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4
RMSE	0.264	0.287	0.294	0.290	0.276	0.296	0.298	0.301	0.334	0.351	0.320	0.344	0.319	0.331	0.335	0.320
MAD	0.194	0.207	0.215	0.213	0.204	0.215	0.218	0.223	0.229	0.227	0.233	0.228	0.233	0.240	0.238	0.232
LL	-2745	-2950	-2910	-2915	-2762	-2923	-2917	-2935	-3201	-3345	-3093	-3270				
AIC	5532	5943	5862	5873	5563	5885	5874	5910	6427	6715	6210	6563				
BIC	5671	6082	6001	6012	5696	6018	6006	6043	6506	6794	6290	6643				

2. Heterogeneous seasonality and Homogeneous scale effects

	Full model				No correlation				No scale effect				Simple regression			
	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4
RMSE	0.302	0.312	0.320	0.326	0.308	0.320	0.320	0.330	0.334	0.351	0.320	0.344	0.319	0.331	0.335	0.320
MAD	0.224	0.227	0.233	0.240	0.231	0.235	0.233	0.242	0.229	0.227	0.233	0.228	0.233	0.240	0.238	0.232
LL	-3067	-3174	-3102	-3200	-3002	-3090	-3093	-3163	-3201	-3345	-3093	-3270				
AIC	6163	6377	6232	6427	6031	6205	6212	6351	6427	6715	6210	6563				
BIC	6255	6470	6325	6520	6117	6291	6298	6438	6506	6794	6290	6643				

3. Homogeneous seasonality and Heterogeneous scale effects

		Full	model		No correlation				No sca	le effect		Simple regression				
	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4
RMSE	0.266	0.286	0.296	0.291	0.276	0.296	0.298	0.301	0.334	0.351	0.320	0.345	0.319	0.331	0.335	0.320
MAD	0.196	0.208	0.217	0.214	0.204	0.215	0.218	0.223	0.229	0.228	0.233	0.228	0.233	0.240	0.238	0.232
LL	-3062	-3242	-3314	-3270	-3169	-3330	-3348	-3366	-3608	-3718	-3508	-3690				
AIC	6152	6512	6656	6568	6364	6686	6721	6758	7234	7454	7035	7399				
BIC	6245	6605	6749	6661	6450	6773	6807	6844	7294	7514	7094	7459				

4. Homogeneous seasonality and scale effects

BIC

	5															
	Full model				No correlation				No scale effect				Simple regression			
	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4	IR-1	IR-2	IR-3	IR-4
RMSE	0.304	0.313	0.320	0.328	0.308	0.320	0.320	0.330	0.334	0.351	0.320	0.345	0.319	0.331	0.335	0.320
MAD	0.226	0.228	0.233	0.241	0.231	0.235	0.233	0.241	0.229	0.228	0.233	0.228	0.233	0.240	0.238	0.232
LL	-3380	-3462	-3508	-3551	-3428	-3515	-3518	-3589	-3608	-3718	-3508	-3690				
AIC	6781	6945	7037	7125	6875	7049	7056	7197	7234	7454	7035	7399				

Table 2.8: In-sample fit of observed donation amounts (Full model with IR-1)

	RMSE	MAD	LL	AIC	BIC
ER-1: All five scale points	0.264	0.194	-2745	5532	5671
ER-2: Two closest scale points from the internal referent	0.280	0.195	-2871	5784	5923
ER-3: Largest and smallest scale points	0.293	0.208	-2899	5840	5980
ER-4: Middle scale point	0.285	0.209	-2932	5906	6045
ER-5: Mean of all five scale points	0.312	0.217	-3078	6198	6337

For model comparisons, owing to short donation histories (which preclude 'squandering' an entire year for prediction purposes), we compare fit in-sample. "Fit" is assessed via mean absolute deviation (MAD) and root mean square error (RMSE) for donation amount predictions, which appear in Tables 2.7 and 2.8. Table 2.7 shows that the proposed model (the "full" model with IR-1, ER-1 and both seasonal and scale effect heterogeneity) provides a better fit than all alternatives; moreover, including error correlation and scale effects improves fit regardless of internal reference formulation (IR1-5) and the inclusion of heterogeneity. Table 2.7 also allows us to judge relative contribution to overall model fit: scale effects easily best both correlation and seasonality. For example, in subtable 2.7-1, which incorporates heterogeneity in both seasonality and scale effects, failing to account for scaling effects inflates RMSE and MAD approximately 30%; the corresponding figure for removing correlation alone is $\approx 5\%$; and for dropping heterogeneity entirely, $\approx 20\%$. Even in the final subtable (2.7-4), for homogeneous seasonality and scale effects, discarding scale attraction inflates RMSE and MAD by approximately 10%; dropping correlation, about 1%. These comparisons suggest that scale attraction effects appear to explain more variation in giving than those typically modeled in prior donation research combined, although it will require additional data applications and distinct settings to verify whether this holds generally.

In terms of internal reference point specification, IR-1, the average of all prior donation

amounts appeared to dominate across the board. The degree of dominance was not trivial, sometimes hovering near 10%. To our knowledge, such a test of 'internal' referents is unique in donation contexts, and we know of no prior theory that would have anticipated it. We found as well that allowing for heterogeneous scale effects (subtables 1 and 3) provides superior fits compared with analogous homogeneous scale effects models (subtables 2 and 4). However, allowing heterogeneity in seasonality parameters offered rather small increases in fit. So, although heterogeneity is itself important, overall, it is much more so in terms of scale effects.

Based on the results of Table 2.7, we restrict our attention to the "full" model with IR-1, and Table 2.8 summarizes fits of five distinct *external* reference specifications (ER 1-5) for this model. ER-1, with all five scale points included, clearly dominates. We hesitate to term this a general finding, as the charity did deliberately choose all scale points to be "reasonable", since they were understandably more focused on revenue than testing reference point theories. Regardless, the "full" model with IR-1 and ER-1 was verified, using all discussed metrics, to provide the best fit to the data among the $2 \times 2 \times 2 \times 2 \times 4 \times 5$ (scale effects?; scale effect heterogeneity?; seasonality heterogeneity?; error correlation?; IR1-4; ER1-5) design.

To guard against concluding in favor of a potentially overparameterized model based on in-sample fit, we report log-likelihood (LL), AIC, and BIC.¹² LL is based on the Tobit 2 likelihood, calculated at parameters' posterior means. The proposed model has the largest LL and smallest AIC and BIC, consistent to the in-sample fit test result.

¹²We do not report two common Bayesian model comparison measures, the Bayes factor and DIC, for the following reasons. The marginal likelihood (on which the Bayes factor is based) is well known to be sensitive to prior specification and calculation method. We in fact computed marginal likelihoods under several different diffuse priors and common calculation methods, finding it highly variable across them. DIC is also not appropriate here, as it has multiple plausible definitions in missing data models (Celeux et al., 2006). Parameters for AIC and BIC are based on Level-I for homogeneous parameters and Level-II for heterogeneous ones. AIC and BIC are included for comparison purposes only, not as definitive bases for Bayesian model selection.

2.9 Illustrative Application: Effect of individually tailored appeals scales

We conclude with a preliminary examination of setting a superior appeal. While a laudable goal might be to devise optimal donor-specific appeals scales or even a single optimal scale applicable to the entire donor pool, the nature of the available data do not permit this, for a number of reasons. Foremost among these was that the scales used by the charity changed neither during the course of the experiment nor across the four donation groups ([level 1, $2] \times [\text{test, standard}]$). Although the *existence* (and nature) of scale attraction effects was strongly verified, this lack of within-donor variation made identifying *interactions* between scale points impossible, since they were perfectly confounded with the experimental conditions themselves. Such interactions are necessary to avoid placing *all* points in a k-point scale at the same "optimal" spot, an obvious absurdity never seen in real appeals scales. Detailed simulations (available from the authors) verified these claims, which we view simply as a data limitation stemming from constraints put on the charity provider.

We instead focus on choosing a single optimum request - often referred to as an attempt to "ladder" an individual donor - and the effect of tailoring these to each donor, as opposed to using a common best ladder (increase percentage over the internal referent) for each, the ubiquitous practice among real fundraisers, even those with donation histories at their disposal. Finding a single optimal request may appear paradoxical, but we must remember that the charities that use appeals scales do *not* vary them across individuals - the very issue that provoked the modeling effort here - and moreover would never have reason to ask for *less* than an internal referent, given its small pulling effect.¹³

¹³Although neither our data nor model allows us to rectify this issue, it's unclear whether using *multiple* scale points is useful in charitable appeals, let alone how many. An appeals scale with multiple points might be beneficial when the charity cannot estimate donors' internal referents, as when soliciting from first-time donors. If there is wide variation in internal referents, a suitably-spaced appeals scale may help donors find a reasonable anchoring point. Regardless, whether it is beneficial to present multiple scale points to regular donors is an empirical question awaiting suitable data, in which the *number* of scale points is systematically varied.

Given these data limitations, we conducted a simulation study involving a single request amount, and a three-period (i.e., one full year) look-ahead. We immediately discovered that seasonal patterns, which were exceedingly strong in our data, vastly swamped any benefit of potential strategies of "let's ask for less now, so they'll give somewhat less, and use that carryover to make them more likely to give again next time." It simply did not matter. Thus, the "three-period look-ahead" optimization, which involved discrete dynamic programming, gave results identical to the three "myopic one-period look-ahead" optimizations, on which we report. Our sole focus, as stated earlier, is quantifying the results of the appeal itself when it is generated heterogeneously, based on donation history, as opposed to a "one ladder fits all" strategy.

To run the simulation, the optimal suggested amount and the expected value of donation amount are calculated, based on β_U . The optimal external referent, r^{E*} , can be calculated by solving a first-order condition for the upward pulling amount, PA (see Appendix C for derivation).

$$r^{E*} = (\exp(\beta_U) + 1)r^I$$
 (2.8)

The expected value of donation amount can also be calculated analytically. We use the generic symbols ν^s and ν^a for the deterministic parts in the selection and donation amount equations in (2.7). Because y^{a*} is a logged quantity, it needs to be exponentiated to calculate the expected donation amount, which is as follows (see Appendix D for derivation):

$$\mathbb{E}[\exp(y^a)] = \exp\left(\nu^a + \frac{\sigma^2}{2}\right) \times \Phi(\nu^s + \sigma\rho)$$
 (2.9)

Three different types of appeals were used in the simulation: i) a "pseudo" appeal, with no suggested amount; ii) a *group*-level optimized suggested amount; and iii) an *individually* customized suggested amount. In the first case, where no external referent is provided, we presume donors hew to their internal referents; this thus serves as a convenient benchmark.

The group-level request is the single common value that would optimize overall donations, given the heterogeneous distribution of β_i^U . Individual-level amounts are calculated from each donor's β_i^U separately. Simply put, the first scenario does not account for scale effects at all; the second type does, but presumes a common "best request" for everyone (as per current practice in the fundraising industry); and the third accounts for individual-level scale effects.

In each scenario, we compare the expected donation amount of 1,000 donors in each of two hypothetical groups (level 1 and 2). These hypothetical donors are generated via the homogeneous parameters (e.g., group dummies) and draws from the joint multivariate normal density obtained from the field experiment (Table 2.6). One challenge in conducting a simulation study is that we cannot "observe" internal reference points and lag-amounts for hypothetical donors. We thus use the group-wise mean of all observed amounts in the real data as a proxy for the group-wise internal reference points for hypothetical donors. As a proxy for the last donation amount (the lag amount in each period), we use the group-wise mean of all observed amounts for Easter, in June, and at Christmas. Then, for each period, the proportion of hypothetical donors giving in each period was set to reflect the response rate in the real data.

Table 2.9: Expected value of average donation amount in simulation

		Lev	el I		Level 2					
	Easter	June	Xmas	Level 1 Total	Easter	June	Xmas	Level 2 Total		
Individually Tailored Request	127.6	28.6	32.2	188.4	321.3	70.0	81.1	472.4		
Group-wise Request	124.8	28.0	31.4	184.2	314.7	68.5	79.4	462.6		
No suggested amount	100.4	22.5	25.3	148.3	252.8	55.0	63.8	371.5		

Table 2.9 shows the effect of an individually tailored appeals scale compared to a groupwise appeals scale. Improvements (from the group-wise to the individual-level appeals scale) are larger in the level 2 group (than in level 1), because their baseline donation amounts are higher. Likewise, Easter entails the largest increase due to its much higher response rate. The full-year expected donations, for an average level 1 and level 2 donor combined, are 519.8FF, 646.8FF, and 660.7FF, for the "no appeal", "group appeal", and "individual appeal" cases, respectively. The latter two both show marked improvement over "no appeal", 24.4% and 27.1%, respectively, offering a "scale-free" (if we may use that term) assessment of the power of making a donation request. By contrast, making an *individually-tailored* request offers a more modest improvement of 2.2% over a group-wise request. While this may seem comparatively small, it is certainly not so for charities. Moreover, charities are *already* making common group-wise requests (albeit, probably not close to optimal ones), so the question is how to leverage the individual donation histories they already have to boost total yield more. Here, we have done so by simultaneously modeling the effects of such a request on both yield rate and donation amount, as well as the intercorrelation of their unobserved influences (errors), and donor-level response heterogeneity.

Our simulation study should be taken as an illustrative exercise awaiting more detailed field data from a properly orthogonalized, custom design (although one wonders which charity will risk potential losses from such an experiment). Most notably, the nature of our data made it impossible to optimize an entire *scale*. However, the results clearly suggest both the importance of making an appropriate request, and of incorporating heterogeneity. We believe the platform developed here would allow for optimal "laddering," which is the dominant practice in real fundraising, for individual donors

2.10 Conclusion

Charities have long relied on appeals scales as cornerstones of their donation requests, though with little theory or measurement to guide them. In that light, the model developed here offers a heterogeneous, joint account of donation incidence and amount, addressing

potential selectivity in modeling these separately. Importantly, different specifications for internal and external reference point theories can be assessed via model comparison.

Results suggest that variation across donors in scale attraction effects and seasonal donation patterns can be substantial. Such a finding depends critically on the availability of donation *histories*, explaining its absence from prior studies. A moderate, significantly negative, correlation between donation incidence and amount indicates the selectivity-based pitfalls of separately modeling incidence and amount. In terms of internal and external referents, we found that the mean of the previous donation amounts (internal referents) and including all points in an appeals scale (external referents) offered the best fit with our data. Importantly, the model allowed for a preliminary exploration of *laddering*, the common practice of asking for successively greater donation amounts.

Our study has some notable limitations, stemming mainly from the data. The charity that designed and carried out the study was not interested in optimal experimental practice, but in gaining some degree of insight within the context of a live donation drive. So, there is a potential issue with appeals scale amounts roughly tracked prior donation level in each segment. Second, because of the lack of group- and time-wise variation in appeals scale, the weight of each scale point - let alone any potential *interactions* among them - simply cannot be estimated; nor could the optimal *number* of points be ascertained. Third, because the appeals scales in the experiment contained only 'reasonable' amounts, effects of extreme scale points, such as ignoring them or even of alienating donors, could not be measured. Despite these data limitations, the model showed clear and strong evidence for scale attraction, in both upward and downward directions, and that the degree of attraction varied greatly across donors.

Some of the data limitations suggest clear directions for future experimental and field research. First and foremost would be some scheme for orthogonalizing appeals scale amounts across various donor groups, and even the number of points on the scale, so that a truly "optimal" scale for each donor can be devised. Key to doing this is accounting for

interactions among scale points, which is necessary to avoid "bunching up" in optimization. Future research might also identifying subtleties of weighting: do some consumers ignore endpoints, while others anchor on them? Experiments could similarly include extreme scale points, to see whether they are ignored entirely, lead respondents not to donate at all, or something more subtle. Any such data could be analyzed through variants of the basic framework employed here, and would help validate cross-study norms about scale point attraction effects, as well as fashion individually-tailored, multi-period laddering plans.

2.11 Appendix

A. Full Model Specification

As discussed above, we can write the entire model as follows (i = donor; t = time):

$$\begin{split} & y_{it}^{s*} = \beta^E X_{it}^E + \beta_i^J X_{it}^J + \beta_i^C X_{it}^C + \beta^{lag} X_{it}^{lag} + \beta^{level,s} X_i^{level} + \epsilon_{it}^s \\ & y_{it}^{a*} = \ln(r_{it}^I + APA_{it}) + \beta^{level,a} X_i^{level} + \epsilon_{it}^a, \ where: \\ & y_{it}^s = 1, \ \text{if} \ y_{it}^{s*} \geq 0; \ 0 \ \text{otherwise} \\ & y_{it}^a = y^{a*}, \ \text{if} \ y_{it}^s = 1; \ \text{unobserved otherwise} \\ & APA_{it} = \sum_{k=1}^K w_{it}^k \times I_{it}^k \times PA_{it}^k; \quad w_{it}^k = \frac{PA_{it}^k}{\sum_{k=1}^K PA_{it}^k}, \quad I_{it}^k = \begin{cases} 1, & r_i^{E,k} \geq r_{it}^I \\ -1, & r_i^{E,k} \leq r_{it}^I \end{cases} \\ & PA_{it}^k = CD_{it}^k \times \left\| r_i^{E,k} - r_{it}^I \right\| \\ & CD_{it}^k = \exp\left(-\frac{d_{it}^k}{\theta_i}\right); \quad \theta_i = \begin{cases} \exp(\beta_i^U), & r_i^{E,k} \geq r_{it}^I \\ \exp(\beta_i^D), & r_i^{E,k} < r_{it}^I \end{cases}, \quad d_{it}^k = \frac{\left\| r_i^{E,k} - r_{it}^I \right\|}{r_{it}^I} \\ & (\epsilon_{it}^s, \epsilon_{it}^a) \sim BVN(0, \Sigma_\epsilon); \quad \Sigma_\epsilon = \begin{bmatrix} 1 & \rho\sigma \\ \rho\sigma & \sigma^2 \end{bmatrix} \\ & \beta_i \sim MVN(\Delta, \Sigma_\beta), \ where \ \beta_i = (\beta_i^J, \beta_i^C, \beta_i^U, \beta_i^D) \end{cases} \end{split}$$

Note that the internal reference point, r_{it}^I , for donor i can change over the course of the experiment, and is subscripted accordingly. However, since the appeals scale for donor i does not change over time, the k^{th} external reference point for a donor i, $r_i^{E,k}$, lacks a t subscript. Again, the variance of ϵ^s is fixed to 1 for identification. Finally, the vector of heterogeneous parameters (β_i) follows a multivariate normal distribution with mean μ_β and full-rank covariance matrix Σ_β .

B. MCMC Algorithm and Priors

Here we present the prior distributions and sampling algorithm used in estimation. Because the requirement that setting error variance of the binary probit model (for donation incidence) be set to one ruins useful conjugacy properties, we instead make random draws from the unidentified space, as suggested by Edwards & Allenby (2003), and report post-processed estimates. Below, we specify Σ_{ϵ} in the unidentified space as

$$\Sigma_{\epsilon} = \left[egin{array}{ccc} \sigma_s^2 &
ho\sigma_s\sigma_a \
ho\sigma_s\sigma_a & \sigma_a^2 \end{array}
ight]$$

1. Data Augmented Likelihood

$$\prod_{i=1}^{n} \prod_{t=1}^{T} [(y_{it}^{s*}, y_{it}^{a*}) | \beta_h, \beta_i, \Sigma_{\epsilon}] \times \prod_{i=1}^{n} [\beta_i | \mu_\beta, \Sigma_\beta]$$

where $\beta_h = (\beta^E, \beta^{lag}, \beta^{level,s}, \beta^{level,a})$ is a vector of homogeneous parameters and $\beta_i = (\beta_i^J, \beta_i^C, \beta_i^U, \beta_i^D)$ is a vector of heterogeneous parameters.

2. Prior Distribution

We use proper but diffuse priors.

(1)
$$\beta_h \sim MVN(M, V)$$
, where $M = 0$, $V = 10^4 I$

(2)
$$\Sigma_{\epsilon} \sim IW(\nu_{\Sigma_{\epsilon}}, V_{\Sigma_{\epsilon}}), \text{ where } \nu_{\Sigma_{\epsilon}} = 5, V_{\Sigma_{\epsilon}} = 5I$$

(3)
$$\Delta \sim MVN(\bar{\Delta}, A)$$
, where $\bar{\Delta} = \mathbf{0}, A = 10^4 I$

(4)
$$\Sigma_{\beta} \sim IW(\nu_{\Sigma_{\beta}}, V_{\Sigma_{\beta}}), \text{ where } \nu_{\Sigma_{\beta}} = 7, V_{\Sigma_{\beta}} = 7I$$

3. Posterior Distribution

$$\prod_{i=1}^{n} \prod_{t=1}^{T} [(y_{it}^{s*}, y_{it}^{a*}) | \beta_h, \beta_i, \Sigma_{\epsilon}] \times \prod_{i=1}^{n} [\beta_i | \mu_{\beta}, \Sigma_{\beta}] \times [\beta_h | M, V] \times [\Sigma_{\epsilon} | \nu_{\Sigma_{\epsilon}}, V_{\Sigma_{\epsilon}}] \times [\Delta | \bar{\Delta}, A] \times [\Sigma_{\beta} | \nu_{\Sigma_{\beta}}, V_{\Sigma_{\beta}}]$$

4. Sampling Algorithm

Step 1. Draw y_{it}^{s*} and y_{it}^{a*} (Data augmentation step)

$$[(y_{it}^{s*}, y_{it}^{a*})|y_{it}^{s}, y_{it}^{a}, \beta_h, \beta_i, \Sigma_{\epsilon}]$$

1. If $y_{it}^s = 1$ then y_{it}^a is observed. We set $y_{it}^{a*} = y_{it}^a$ and draw y_{it}^{s*} from the truncated normal distribution below:

$$TN(\beta^{E}X_{it}^{E} + \beta_{i}^{J}X_{it}^{J} + \beta_{i}^{C}X_{it}^{C} + \beta^{lag}X_{it}^{lag} + \beta^{level,s}X_{i}^{level} + \frac{\rho\sigma_{s}}{\sigma_{a}}[y_{it}^{a} - (\ln(r_{it}^{I} + APA_{it}) + \beta^{level,a}X^{level})], (1 - \rho^{2})\sigma_{s}^{2}), \ y_{it}^{s*} \ge 0$$

2. If $y_{it}^s = 0$ then y_{it}^a is not observed. We draw $(y_{it}^{s*}, y_{it}^{a*})$ by following steps

$$\text{a. Draw } y_{it}^{s*} \text{ from } TN(\beta^E X_{it}^E + \beta_i^J X_{it}^J + \beta_i^C X_{it}^C + \beta^{lag} X_{it}^{lag} + \beta^{level,s} X_i^{level}, \sigma_s^2), \ \ y_{it}^{s*} < 0$$

b. Draw y_{it}^{a*} conditional on y_{it}^{s*} from normal distribution below:

$$N(\ln(r_{it}^{I} + APA_{it}) + \beta^{level,a}X_{i}^{level} + \frac{\rho\sigma_{a}}{\sigma_{s}}[y_{it}^{s*} - (\beta^{E}X_{it}^{E} + \beta_{i}^{J}X_{it}^{J} + \beta_{i}^{C}X_{it}^{C} + \beta^{lag}X_{it}^{lag} + \beta^{level,s}X_{i}^{level})], (1 - \rho^{2})\sigma_{a}^{2})$$

Step 2. Draw β_i

$$[\beta_i|\beta_h, \Sigma_{\epsilon}, \Delta, \Sigma_{\beta}] \propto \prod_{t=1}^{T} [(y_{it}^{s*}, y_{it}^{a*})|\beta_h, \beta_i, \Sigma_{\epsilon}] \times [\beta_i|\Delta, \Sigma_{\beta}]$$

The full conditional distribution is also of unknown form. Therefore, we use a Metropolis-Hastings algorithm with a normal random walk proposal to make draws.

Step 3. Draw β_h

$$[\beta_h|\{\beta_i\}, \Sigma_{\epsilon}] \propto \prod_{i=1}^n \prod_{t=1}^T [(y_{it}^{s*}, y_{it}^{a*})|\beta_h, \beta_i, \Sigma_{\epsilon}] \times [\beta_h|M, V]$$

Again, we use a Metropolis-Hastings algorithm with a normal random walk proposal to make draws.

Step 4. Draw Σ_{ϵ}

$$[\Sigma_{\epsilon}|\beta_{h}, \{\beta_{i}\}] \propto \prod_{i=1}^{n} \prod_{t=1}^{T} [(y_{it}^{s*}, y_{it}^{a*})|\beta_{h}, \beta_{i}, \Sigma_{\epsilon}] \times [\Sigma_{\epsilon}|\nu_{\Sigma_{\epsilon}}, V_{\Sigma_{\epsilon}}]$$

$$\propto \prod_{i=1}^{n} \prod_{t=1}^{T} BVN \left(\begin{pmatrix} y_{it}^{s*} \\ y_{it}^{a*} \end{pmatrix} | \begin{pmatrix} u_{it}^{s} \\ u_{it}^{a} \end{pmatrix}, \Sigma_{\epsilon} \right) \times IW(\nu_{\Sigma_{\epsilon}}, V_{\Sigma_{\epsilon}})$$

$$[\Sigma_{\epsilon}|\beta_{h}, \{\beta_{i}\}] \sim IW(\tilde{\nu}_{\Sigma_{\epsilon}}, \tilde{V}_{\Sigma_{\epsilon}}),$$

$$\tilde{\nu}_{\Sigma_{\epsilon}} = \nu_{\Sigma_{\epsilon}} + nT,$$

$$\tilde{V}_{\Sigma_{\epsilon}} = V_{\Sigma_{\epsilon}} + \sum_{i=1}^{n} \sum_{t=1}^{T} \begin{pmatrix} y_{it}^{s*} - u_{it}^{s} \\ y_{it}^{a*} - u_{it}^{a} \end{pmatrix} \times \begin{pmatrix} y_{it}^{s*} - u_{it}^{s} \\ y_{it}^{a*} - u_{it}^{a} \end{pmatrix}$$

where

$$u_{it}^s = \beta^E X_{it}^E + \beta_i^J X_{it}^J + \beta_i^C X_{it}^C + \beta^{lag} X_{it}^{lag} + \beta^{level,s} X_i^{level}$$
$$u_{it}^a = \ln(r_{it}^I + APA_{it}) + \beta^{level,a} X_i^{level}$$

Step 5. Draw Δ

$$[\Delta|\{\beta_i\}, \Sigma_{\beta}] \propto \prod_{i=1}^{n} [\beta_i|\Delta, \Sigma_{\beta}] \times [\Delta|\bar{\Delta}, A] \propto MVN_{nk}(B^*|[Z \otimes I_k]\Delta^*, I_n \otimes \Sigma_{\beta})$$
$$\times MVN_{nk}(\Delta^*|\bar{\Delta}, A)$$

where β_i is a vector of length k,

$$B = \begin{bmatrix} \beta_1^T \\ \beta_2^T \\ \vdots \\ \beta_n^T \end{bmatrix}, \quad B^*_{nk \times 1} = vec(B^T), \quad Z = \begin{bmatrix} \iota_1^T \\ \iota_2^T \\ \vdots \\ \iota_n^T \end{bmatrix}, \quad \Delta^* = vec(\Delta^T)$$

$$[\Delta^*|\{\beta_i\}, \Sigma_{\beta}] \sim MVN_{nk}(\Delta^*|\tilde{\Delta}, \tilde{A})$$

$$\text{ where } \tilde{\Delta} = \tilde{A}([Z \otimes \Sigma_{\beta}^{-1}]B^*_{nk \times 1} + A^{-1}\bar{\Delta}), \quad \tilde{A} = [(Z^TZ) \otimes \Sigma_{\beta}^{-1} + A^{-1}]^{-1}$$

Step 6. Draw Σ_{β}

$$[\Sigma_{\beta}|\{\beta_i\},\Delta] \propto \prod_{i=1}^{n} [\beta_i|\Delta,\Sigma_{\beta}] \times [\Sigma_{\beta}|\nu_{\Sigma_{\beta}},V_{\Sigma_{\beta}}] \propto MVN_{nk}(B|Z\Delta,I_n,\Sigma_{\beta}) \times IW(\nu_{\Sigma_{\beta}},V_{\Sigma_{\beta}})$$

$$[\Sigma_{\beta}|\{\beta_i\},\Delta] \sim IW(\tilde{\nu}_{\Sigma_{\beta}},\tilde{V}_{\Sigma_{\beta}})$$

where
$$\tilde{\nu}_{\Sigma_{\beta}} = \nu_{\Sigma_{\beta}} + n$$
, $\tilde{V}_{\Sigma_{\beta}} = V_{\Sigma_{\beta}} + (B - Z\Delta)^T (B - Z\Delta)$

C. Single Optimal Appeal

The single optimal appeal is that (scale) point (r^{E*}) that maximizes the log of the latent donation amount (y^{a*}) . Clearly, r^E should be greater than r^I and maximizes the upward pulling amount (PA), so that r^{E*} solves a first order condition (PA) with respect to r^E). Subscripts for donors and times (i,t) are suppressed for simplicity.

$$y^{a*} = \ln(r^{I} + PA) + X^{a}\beta^{a} + \epsilon^{a}$$

$$PA = \exp\left(-\frac{d}{\theta}\right)(r^{E} - r^{I}), \quad d = \frac{r^{E}}{r^{I}} - 1, \quad \theta = \exp(\beta_{U})$$

$$FOC: \quad \frac{\partial PA}{\partial r^{E}} = 0$$

$$r^{E*} = (\theta + 1)r^{I} = (\exp(\beta_{U}) + 1)r^{I}$$

D. Expected Donation Amount

Following the Type 2 Tobit model specification, let y^{s*} and y^{a*} be the latent dependent variables in the selection and the amount equations, and y^s and y^a be their observed counterparts in (2.9). We set ν^s , ν^a to be deterministic parts of each equation and ϵ^s , $\sigma\epsilon^a$ to be stochastic parts.

Now we have the system of equations:

$$y^{s*} = \nu^s + \epsilon^s$$
$$y^{a*} = \nu^a + \sigma \epsilon^a$$

Because the errors are correlated, we can re-write the system as (writing e^s as e):

$$y^{s*} = \nu^s + \epsilon$$

$$y^{a*} = \nu^a + \sigma(\rho\epsilon + \bar{\rho}z)$$

where z is a standard normal draw uncorrelated with ϵ , and $\bar{\rho} = \sqrt{1 - \rho^2}$. We want to calculate $\mathbb{E}[\exp(y^a)]$, the expected donation amount, and so merely integrate over the two uncorrelated errors, z and ϵ :

$$\mathbb{E}[\exp(y^{a})] = \int_{\epsilon=-\nu^{s}}^{\epsilon=\infty} \int_{z=-\infty}^{z=\infty} \exp[\nu^{a} + \sigma(\rho\epsilon + \bar{\rho}z)]\phi(z)\phi(\epsilon)dzd\epsilon$$

$$= \exp(\nu^{a}) \times \int_{\epsilon=-\nu^{s}}^{\epsilon=\infty} \exp(\sigma\rho\epsilon)\phi(\epsilon)d\epsilon \times \int_{z=-\infty}^{z=\infty} \exp(\sigma\bar{\rho}z)\phi(z)dz$$

The second term can be calculated by completing the square:

$$\int_{\epsilon=-\nu^{s}}^{\epsilon=\infty} \exp(\sigma\rho\epsilon)\phi(\epsilon)d\epsilon = \int_{\epsilon=-\nu^{s}}^{\epsilon=\infty} \exp(\sigma\rho\epsilon)\frac{1}{\sqrt{2\pi}}\exp\left(-\frac{\epsilon^{2}}{2}\right)d\epsilon$$

$$= \int_{\epsilon=-\nu^{s}}^{\epsilon=\infty} \frac{1}{\sqrt{2\pi}}\exp\left(-\frac{(\epsilon-\sigma\rho)^{2}}{2}\right)d\epsilon \times \exp\left(\frac{(\sigma\rho)^{2}}{2}\right) \qquad Let \ u = \epsilon - \sigma\rho$$

$$= \int_{u=-\nu^{s}-\sigma\rho}^{u=\infty} \frac{1}{\sqrt{2\pi}}\exp\left(-\frac{u^{2}}{2}\right)du \times \exp\left(\frac{(\sigma\rho)^{2}}{2}\right) = \Phi(\nu^{2} + \sigma\rho) \times \exp\left(\frac{(\sigma\rho)^{2}}{2}\right)$$

The third term can also be calculated by completing the square:

$$\int_{z=-\infty}^{z=\infty} \exp(\sigma\bar{\rho}z)\phi(z)dz = \int_{z=-\infty}^{z=\infty} \exp(\sigma\bar{\rho}z)\frac{1}{\sqrt{2\pi}}\exp\left(-\frac{z^2}{2}\right)dz$$
$$= \int_{z=-\infty}^{z=\infty} \frac{1}{\sqrt{2\pi}}\exp\left(-\frac{(z-\sigma\bar{\rho})^2}{2}\right)dz \times \exp\left(\frac{(\sigma\bar{\rho})^2}{2}\right) = \exp\left(\frac{(\sigma\bar{\rho})^2}{2}\right)$$

Putting this all together, the expected value of donation amount is:

$$\mathbb{E}[\exp(y^{a})] = \exp(\nu^{a}) \times \Phi(\nu^{2} + \sigma\rho) \times \exp\left(\frac{(\sigma\rho)^{2}}{2}\right) \times \exp\left(\frac{(\sigma\bar{\rho})^{2}}{2}\right)$$

$$= \exp(\nu^{a}) \times \Phi(\nu^{2} + \sigma\rho) \times \exp\left(\frac{\sigma^{2}\rho^{2}}{2}\right) \times \exp\left(\frac{\sigma^{2}(1-\rho)^{2}}{2}\right)$$

$$\therefore \bar{\rho} = \sqrt{1-\rho^{2}}$$

$$= \exp\left(\nu^{a} + \frac{\sigma^{2}}{2}\right) \times \Phi(\nu^{2} + \sigma\rho)$$

E. Distance ratio and the incremental/decremental amount, calculated at the scale point with the maximum pulling amount

1.Upward pulling amount

When $r^E \geq r^I$, the upward pulling amount PA_U follows:

$$PA_U = \exp\left(-\frac{d_U}{\theta}\right)(r^E - r^I), \quad d_U = \frac{r^E}{r^I} - 1, \quad \theta_U = \exp(\beta_U)$$

The scale point with maximum upward pulling amount (r^{E*}) can be calculated by solving first order condition of PA_U with respect to r^E .

$$r^{E*} = (\theta_U + 1)r^I = (\exp(\beta_U) + 1)r^I$$

At the scale point of r^{E*} , the incremental ratio in distance (d_U^*) is determined to be $\exp(\beta_U)$ and the maximum incremental ratio in the amount to be as follows:

$$\frac{\exp\left(-\frac{d_U^*}{\exp(\beta_U)}\right)(r^{E*} - r^I)}{r^I} = \exp(-1)\exp(\beta_U)$$

2. Downward pulling amount

When $0 \le r^E \le r^I$, the downward pulling amount PA_D follows:

$$PA_D = \exp\left(-\frac{d_D}{\theta}\right)(r^I - r^E), \quad d_D = 1 - \frac{r^E}{r^I}, \quad \theta_D = \exp(\beta_d)$$

The scale point with maximum downward pulling amount (r^{E*}) can be calculated by solving first order condition of PA_D with respect to r^E . We should note that there is a corner solution if $\beta_D > 0$.

$$r^{E*} = \begin{cases} (1 - \theta_D)r^I = (1 - \exp(\beta_D))r^I, & \beta_D \le 0\\ 0, & \beta_D > 0 \end{cases}$$

At the scale point of r^{E*} , the decremental ratio in distance (d_D^*) is determined to be

$$\begin{cases} \exp(\beta_D), & \beta_D \le 0 \\ 1, & \beta_D > 0 \end{cases}$$

And the maximum decremental ratio in the amount is determined to be as follows:

$$\begin{cases} \frac{\exp\left(-\frac{d_U^*}{\exp(\beta_U)}\right)(r^I - r^{E*})}{r^I} = \exp(-1) - \exp(\beta_U), & \beta_D \le 0\\ \frac{\exp\left(-\frac{d_U^*}{\exp(\beta_U)}\right)(r^I - r^{E*})}{r^I} = \exp\left(-\frac{1}{\exp(\beta_D)}\right), & \beta_D > 0 \end{cases}$$

CHAPTER III

Modeling Mate Choice Behavior: A Two-Stage Mate Choice Model with Potentially Non-Compensatory Decision Rules

3.1 Abstract

In this research, we develop a two-stage mate choice model that allows us to empirically evaluate the evidence in favor of alternative decision rules using data from an online dating service provider in the US. The proposed model captures the intrinsically multistage behavior involved in many online transactions, but in particular dating, where one decides which profiles to browse and then, conditional on having browsed, whom to write to, if anyone. Here, we account for these two distinct activities by modeling the binary decisions (of browsing and writing). The model can accommodate compensatory and non-compensatory decision rules in each stage; it allows decision rules to differ across stages; different attributes can be modeled as having distinctly different utility 'shapes'; and heterogeneity in rule use across site users provides interpretable profiles of different types of mate-seeking behavior. Finally, and most importantly, we directly model the utility functions of attributes to identify and compare attribute-level decision rules ("deal-breakers" and "deal-makers") over two stages.

3.2 Introduction

There is currently great interest among marketers, policymakers, and government funding agencies in models of human decision-making; the hope is that better understanding of how people make decisions may allow us to develop more effective policies aimed at meeting the needs of a diverse consumer pool, as well as potentially changing behavior (e.g., being more charitable, using fewer resources, etc.) In consumer behavior and social science research, the standard approach is to decompose variation in some individual outcome of interest into portions attributable to different "explanatory" covariates. Causal relationships are typically attributed to variables rather than to human actors, whose behavior is contingent and often difficult to measure. In social science research in particular, the statistical model used in the analysis is rarely a plausible model of the underlying behavior or decision-making process that gave rise to the social phenomenon under investigation; as a result, these analyses are often divorced from the actions or activities that led to a particular outcome. To understand consumer needs or fashion effective interventions, we need plausible models of how individuals navigate their social and physical environment.

Researchers in the field of marketing modeling have, for the past 30 or more years, worked out sophisticated models of decision-making, but these have rarely been applied to social contexts, for example, how people choose among potential partners, neighborhoods, jobs, or entrees in a cafeteria. Among the many reasons for this are cultural differences in emphasis (e.g., drivers of individual-level behavior in marketing vs. "big picture" understanding of entire systems in political science and sociology), data availability (detailed databases in marketing vs. more macro-level variables in social science), and degree of emphasis on methodology per se. Given that both fields concern themselves with understanding human behavior, there is fertile territory at their intersection: extending individual-level choice models from marketing which tend to involve one decision-maker picking from a known set of options under measured conditions to a variety of thorny, open problems in sociology and cognate disciplines.

Specifically, the overarching goal of this study is to extend statistical techniques and theoretical models from marketing and choice theory to studies of sociologically relevant decision-making, with a particular emphasis on what is arguably the most important decision an individual can make: mate choice. In contrast to existing work in social science, we develop a statistical model of choice behavior that allows for an actor with partial information, uncertainty about the most desirable outcome, and difficulty comparing more than a handful of alternatives based on a small number of attributes (see Shocker et al., 1991, for review). Importantly, the proposed multistage choice model allows for the possibility of compensatory and non-compensatory screening rules (as discussed later), and different attributes may be evaluated using different rules.

As mentioned earlier, the substantive application is mate choice, a subject with deep roots in social science, but only recently amenable to individual-level modeling. Here, our data stem from an online dating site, an increasingly common way to meet potential dating or marriage partners. Between 1995 and 2005, there was a rapid increase in the number of opposite-gender couples who met their partners online (Rosenfeld & Thomas, 2012). A study commissioned by Match.com in 2010 reported that 1 in 6 couples married within the past three years met their partner on an online dating site, and 1 in 5 people have dated someone they met on an online dating site (Chadwick, 2010). The increased availability of behavioral data from online dating web sites has led to a number of studies to use patterns of early stage interaction online that is, who browses, contacts, or responds to whom to estimate models of mate choice (Hitsch et al., 2010; Lin & Lundquist, 2013). However, in all cases, the key focus is on "preferences", as reected in the relative magnitude and signicance of coefficients in a statistical model. The statistical technique used in these studies assumes a single-stage decision process, in which mate-seekers with unlimited time and computational resources consider every single potential mate in their metropolitan area using a compensatory model. Thus, despite the behavioral nature of the online dating data, past empirical analyses do not in any way represent the underlying activities that give rise

to choice outcomes. Researchers instead impose the same and statistical techniques that have been used for decades to analyze survey data, but which are arguably implausible for online dating and a variety of related data settings.

Behavioral data provide new opportunities to develop theoretical models of individual choices, but this requires an expanded repertoire of statistical models. Compared with traditional sources of information about how individuals search among potential mates (e.g., observed matches or preferences revealed in surveys), online dating activities provide far richer information understanding mate choice, for two reasons: (1) researchers have access to individual-level, longitudinal interaction histories for each participant, including a standardized (or at least stable) array of demographic covariate information for browsed and written-to profiles, which helps statistically nail down accurate mate preferences for users; and (2) the online world facilitates observation of all intermediary steps undertaken by each user while on the site, such as login history, overall time spent, detailed browsing history, profiles viewed and rejected, etc. Although demographic information was always potentially, if somewhat spottily, available using traditional sources, it is this truly novel information on intermediary behaviors that can shed light on both how matching happens and on users decision process, a level of analysis simply not possible solely using records final matches alone.

3.3 Model development

The main objective in model development is to understand how two-sided matching takes place when there are many, interacting searchers/decision-makers, each with his or her own preferences, which express themselves as utilities. These functions (i.e., utilities based on attributes), as discussed in full later, should not be presumed to have constant marginal disutility; that is, "a year is a year" for age or that each inch in height matters as much as any other. Although the model is tested on dating site data, the overarching objective is

broader, and concerns how to capture multistage, highly multiattribute utility functions in an estimable and relatively parsimonious manner, when we *cannot* simply presume strict linearity (or linear-additivity) in the components of utility across their entire ranges. We next discuss some extant hypothesis regarding mate choice behavior that can be fruitfully addressed by the sort of modeling framework discussed here.

3.3.1 Hypotheses on mate choice behavior

There are three main hypotheses in the sociological literature for how men and women evaluate potential mates. The matching hypothesis assumes that people prefer to marry someone of similar social status (DiMaggio & Mohr, 1985; Kerckhoff & Davis, 1962), while the competition hypothesis assumes that people prefer to marry someone of high status (Elder Jr, 1969). The matching hypothesis is typically applied to attributes that imply some sort of shared values or culture, such as consumption behavior or education, while the competition hypothesis is typically applied to attributes that provide resources that can be shared within the relationship, such as income. The exchange hypothesis postulates that people may either purposefully seek mates who differ from them on key attributes, or are willing to accept someone with lower status on some dimension provided this is offset by a high status on another dimension. For example, under a gendered division of labor a man may exchange his economic status for a woman's domestic skills or physical attractiveness (Becker, 1993). Historically the exchange hypothesis has been most frequently applied to potential tradeoffs that may occur between education and race (Fu, 2001; Merton, 1941; Rosenfeld, 2005), and age/attractiveness and economic assets (Buss & Barnes, 1986; Mc-Clintock, 2011).

Up until quite recently, there was little data to study mate choice. The dominant strategy that scholars used was to examine coefficients from loglinear models estimated from a

¹It is likely that these hypotheses may both be valid but apply to different characteristics; for example, people may prefer to seek someone similar based on cultural background and lifestyle issues but optimize on income (Kalmijn, 1994).

cross-classied table of couples' attributes.² This information is typically obtained from data on marriages, but may also come from cohabiting partners or other relationships. The focal parameters provide information on what kinds of matching patterns persist after accounting for the pairings that would be expected on the basis of random sorting given the joint distribution of men and women's attributes. The problem is that these statistical models cannot disentangle individuals preferences or desires from the structural constraints imposed by the marriage market (Logan, 1996; Logan et al., 2008).³ Also, by specifying the unit of analysis to be the match itself, this approach ignores the two-sided nature of the marriage market. In other words, data on successful matches cannot distinguish between men and women's preferences for mates. It is likely that men and women have different preferences for attributes of partners (England & McClintock, 2009), and also men and women may have different tolerance for remaining single in the face of an unacceptable match.

However, over the past few years the increased availability of behavioral data from online dating web sites has led to a number of studies to use patterns of early stage mate choicethat is, who browses, contacts, or responds to whomto estimate models of mate preferences. For example, using a German online dating website to examine patterns of educational assortative mating, Skopek and colleagues (Skopek et al., 2011) nd that both men and women match on education, and preference for a partner of similar education is most pronounced among the highly educated. A couple of studies have used American online dating data to investigate men and women's preferences for mates. Consistent with the work by Skopek and colleagues, they nd that men and women have strong preferences for a partner who shares their education, but document substantial differences in how men

²There have been some recent debates about the validity of simple versus complex methods of studying marriage patterns, especially in terms of assessing the evidence of exchange (Gullickson & Fu, 2010; Kalmijn, 2010; Rosenfeld, 2010). Zeng, Schwartz, and Xie (2012) argue that loglinear models are more effective than simpler approaches because they control for the differential distribution of attributes among men and women as well as baseline tendencies for homogamy and hypergamy.

³Some proponents of loglinear models claim that the models separate out the effects of population composition from the effects of allocation processes such as individual occupational mobility or mate choice behavior (Hauser, 1978; Hout, 1983). However, Logan (1996) demonstrates via analytical and simulation methods that in two-sided contexts such as labor or marriage markets, log-linear models cannot disentangle the relative contributions of population distribution and behavior to observed outcomes.

and women assess partners with more or less education than themselves (Hitsch et al., 2010). With regards to race, these studies suggest that both matching and competition play important roles in mate choice decisions (Lin & Lundquist, 2013).

However, all past studies of mate choice using online dating data assume an implicit single-stage, compensatory model of choice in which fully informed, rational actors with unlimited computation abilities assess all potential mates based on the relative weights assigned to different attributes. However, especially on online dating sites where users are confronted with potentially hundreds of potential partners, mate seekers are likely to rely on multistage and potentially non-compensatory screening rules. For example, men who want to start a biological family of their own might not even consider a woman past childbearing age, regardless of her other attributes. The proposed multistage model with attribute-level decision rules may provide a different perspective on existing hypotheses and also suggest new hypotheses on mate choice behavior. With a multistage model with decision rules, we can assess whether existing hypotheses are supported by the data, and the boundary conditions where each hypothesis holds. In addition, we can find the new relationship among attributes and identify new hypotheses that could not be observed due to screening rules people use.

3.3.2 Decision rules

Since the pioneering work of Swait (1984), a large body of research has addressed the concepts of "choice sets" or "consideration sets" (Gilbride & Allenby, 2004; Hauser & Wernerfelt, 1990; Shocker et al., 1991). The basic idea is that consumers typically go through a two stage-decision process, where only the subset of choice alternatives that pass through the 1st stage decision rules are re-evaluated, typically using a distinct set of rules, to arrive at a final choice or choices (2nd stage). A variety of screening rules can be used in the 1st stage, including linear compensatory, or non-compensatory rules such as conjunctive, disjunctive, or even complex interactions thereof (Hauser et al., 2010).

Among the challenges of this literature is that the choice set is latent that is, unobserved by the researcher and so disentangling preferences for what enters the consideration set from what is eventually selected from it can be statistically challenging, with nontrivial data requirements (e.g., many observations per decision-maker).

While there is a large literature in marketing on estimating compensatory and noncompensatory decision rules, the vast majority of those studies assume that the same type of decision rule applies to all attributes. For example, the underlying values for an attribute (say, price) may be used directly by the decision maker; or they may be subject to some highly non-linear transform; or even 'thresholded' so that nothing beyond a particular value has any marginal effect, like prices so high one would never even think of paying them. Two past studies have allowed for different decision rules to apply to different attributes. Swait (2001) uses piecewise linear functions to estimate response functions that allow decision-makers to penalize, but not eliminate, alternatives that fail to meet a conjunctive cutoff. However, he uses self-reported cutoffs from stated preference data to identify the cutpoints. Our goal is to estimate the response function, including any noncompensatory cutoffs, using observational data. The study most similar in spirit to our approach, Elrod et al. (2004), uses a non-rectangular hyperbolic function to specify decision rules that allow for an explicit parameterization of compensatory versus non-compensatory functions in observational data. Their model has the advantage that it provides a clean and precise statistical test for whether a decision-rule is conjunctive, disjunctive, or compensatory. However, this elegant approach comes at the cost of enacting some fairly restrictive simplifying assumptions. First, they assume that all compensatory decision rules are linear response functions; non-linear is conflated, if not equated, with 'non-compensatory'. But this seems implausible in the case of mate choice: For example, some mate seekers may find potential partners who are divorced to be disproportionately unattractive, while trading this off against other things. In addition, the non-rectangular hyperbola only allows for monotonic response functions, but there is reason to believe that a number of our attributes

have a non-monotonic response function (e.g., women tend to have a zone of acceptability near their own age, finding much younger or much older potential mates a less good fit).

Finally, Elrod et al. (2004)'s cutoff points are derived from the maximum and minimum observed values, without any statistical error. This assumption only makes sense when there are strict rules to follow, as are sometimes imposed in college admission. By contrast, the model to be developed in the sequel will allow the identification of such "near non-compensatory" rules. This distinction is important. Were a "deal-breaker" truly inviolable, it would be a simple and tautological matter to pull them from observed data. For example, if a particular site user wrote only to people above a certain age, we might declare that that being below that age is a "deal-breaker" However, this would be premature, as determining this would depend on examining the *pool* of potential recipients. It would also ignore important statistical background information: if that respondent wrote to 100 other users, 99 of whom were over fifty, and one of whom was twenty-five, the model should not merely spit out that a "deal-breaker" age was anything under the much lower figure. That is, there needs to be an *error model*, and some notion of being able to statistically test various regions for differing response propensities. In other words: a model-based approach.

While past theories of mate choice have not considered what attributes are most widely used in non-compensatory screening rules, we propose a few preliminary hypotheses. First, we expect that women will impose screening rules based on the height of potential partners. Anecdotal evidence from profile text suggests that women prefer a partner who is "not shorter than [she is] in heels." This implies that some women impose a deal-breaker for potential partners less than 2-3 inches taller than they are. Also, we suspect that men (especially men who wish to have children) may impose a deal-breaker for women over 40, or thereabouts. In addition to height and weight deal-breakers, we also test for the possibility of deal-breakers associated with potential partners' education, race, and parental status. We do not anticipate finding any "deal makers" (attributes that, taken entirely on their own, can serve as sufficient reason for a person being chosen), at least within the

general population; this does not preclude the possibility of an occasional "gold digger" for whom financial resources trump all other deficiencies. In the next section, we show how one can use information on mate seekers' activities on the site (browsing and writing to potential mates) to identify mate choice decision rules. To that end, we now define "generic" utility functions for each portion of the two-stage process, browsing and writing.

3.3.3 Utility functions

A key element in our development is the specification of the utility functions for the two explicit stages of the model: browsing and writing. Ideally, one would seek to impose a fully nonparametric account: a model that can have arbitrary complexity in utility shape, achieved via a modeling framework whose degree of parameterization is not set in advance. Such a framework has been developed before (e.g., Kim et al., 2007), but has strong data requirements (e.g., many observations per respondent), high computational costs, and has not been adapted to multistage, highly multiattribute decisions. The key benefit provided by the nonparametric approach is that utility "shapes" can have multiple regions, with a different degree of trade-off between an attribute and (dis)utility in each. Among the major findings of Kim et al. (2007) were: (1) that piecewise *linear* functions (specifically, splines) sufficed to capture utility as well as allowing segments of the function to be quadratic or higher; and (2) that, for the vast majority of participants in multiple data sets, the modal number of interior knots was 2. Specifically, in a conjoint application with 6 attributes, all 6 were best captured by 2 interior knots (see their Fig. 1); and in three scanner data sets, the modal number of knots in each was 2 or fewer (see their page 349, column 2). Although one must exhibit caution in exporting the findings from their study to a novel context, we propose that a flexible yet relatively parsimonious account of utility should allow it to be piecewise linear, so long at it allows for no fewer than two interior knots. We adopt this convention here, emphasizing that, although the usual linear utility specification requires two parameters (slope and intercept), a two-knotted piecewise linear utility spline requires

six, since each additional utility segment grafted onto a base (linear) function requires two new parameters: one for the location of the knot, and another for the slope change coincident with that knot. Note that this does *not* affect the intrinsically discrete nature of how purely categorical variables are typically handled, which we adopt here for both comparability and maximal flexibility.

In line with the preceding discussion, the utility function for browsing is decomposed into three portions: an intercept; a two-knotted piecewise linear spline for continuous (or ordinal) attributes (e.g., age group); and a conjoint-like representation for intrinsically categorical attributes (e.g., ethnic group); as follows:

$$V_{ij}^{B} = \beta_{0i}^{B} + \sum_{k=1}^{K} [\beta_{1ik}^{B} x_{jk}^{B} + \beta_{2ik}^{B} (x_{jk}^{B} - \delta_{1ik}^{B})_{+} + \beta_{3ik}^{B} (x_{jk}^{B} - \delta_{2ik}^{B})_{+}] + \sum_{l=1}^{L} [\gamma_{il}^{B} x_{jl}^{B}] \quad (3.1)$$

where
$$(y)_+ = \begin{cases} y, & if \ y \ge 0 \\ 0, & if \ y < 0 \end{cases}$$
 and $\delta^B_{1ik} \le \delta^B_{2ik}$

 V^B_{ij} stands for the systematic part of the utility for user i of browsing potential mate j. It is specified as a linear additive model with three components: 1) β^B_{0i} (an intercept term); 2) the sum of the utilities of K continuous attributes; and 3) the sum of the utilities of L discrete attributes. As mentioned previously, the utility functions of K continuous attributes follow a continuous piecewise linear function with (up to) two knots (δ^B_{1ik} and δ^B_{2ik}). This formulation is flexible enough to accommodate linear/non-linear compensatory rules as well as non-compensatory decision rules, such as conjunctive and disjunctive rules when any of the slopes approaches $+\infty$ and $-\infty$ respectively.

It is helpful to visualize these sorts of functions in order to understand what their specification "buys us" substantively. Figure 3.1a depicts linear compensatory rule, Figure 3.1b a non-linear but compensatory one. Figure 3.1c is a conjunctive rule where being outside of the range $(\delta_{1ik}$ and $\delta_{2ik})$ acts as a deal-breaker, and Figure 3.1d is a disjunctive rule where being greater than δ_{2ik} acts as a deal-maker. The utility functions of L discrete attributes are specified by dummy variables. The model can also accommodate non-compensatory decision rules for the categorical attributes as various elements of the parameter vector γ_{il}^B approach $\pm \infty$. Figure 3.2a and 3.2b show a deal-breaker and deal-maker for categorical response variables.

The theoretical and empirical challenge is to distinguish deal-breakers or deal-makers from nonlinear compensatory responses. For a continuous attribute k, if any of the pairwise difference(s) among β_{1ik}^B , β_{2ik}^B , and β_{3ik}^B is ∞ , it represents non-compensatory rule, as in Figure 3.1c and 3.1d. In reality, imposing a difference of ∞ is somewhere between meaningless and too harsh: practically speaking, if the difference is large enough to render all other attributes and their differences irrelevant, a nonlinear compensatory rule can function as deal-breaker or deal-maker. For example, a difference of 10 on the logit scale represents a difference in odds (and thereby, roughly speaking, probability) on the order of 20,000; that is, a difference in (say, browsing) utility of -10 makes it 20,000 times less likely that that person will be written to, which by any reasonable standards represents a deal-breaker. Similar logic can be applied to the L categorical attributes. The pairwise difference in dummy variable γ^B_{il} s determines whether the attribute l functions as deal-breaker or dealmaker. [For categorical attributes, the differences need to be compared to an average, not merely to adjacent ones, since "adjacent" does not mean anything for purely categorical variables, e.g., ethnicity.] How big these differences should be is an important empirical question, one that we shall examine in the context of model estimates for our particular data setting.⁴

We turn next to the utility function for writing, which follows a similar general format:

$$V_{ij}^{W} = \beta_{0i}^{W} + \sum_{m=1}^{M} [\beta_{1im}^{W} x_{jm}^{W} + \beta_{2im}^{W} (x_{jm}^{W} - \delta_{1im}^{W})_{+} + \beta_{3im}^{W} (x_{jm}^{W} - \delta_{2im}^{W})_{+}] + \sum_{n=1}^{N} [\gamma_{in}^{W} x_{jn}^{W}]$$
(3.2)

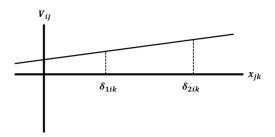
⁴That is, the researcher needs to set some pre-established (significance) level.

where
$$(y)_+ = \begin{cases} y, & \text{if } y \ge 0 \\ 0, & \text{if } y < 0 \end{cases}$$
 and $\delta^W_{1im} \le \delta^W_{2im}$

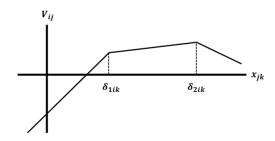
 V_{ij}^W stands for the systematic part of the utility for user i of writing to user j. Although it follows the same specification as V_{ij}^B , the number of continuous and discrete attributes can of course be different from those in the browsing stage. This reflects both the empirical fact, common to all dating sites, that the information available in browsing stage is typically supplemented by additional variables in the writing stage, and also that even information available in browsing might be enhanced after one clicks to reveal a full profile (for example, on some dating sites not the one we use here clicking on a profile often reveals additional photos, which may serve to alter one's prior evaluation of someone's appearance).

Note that the proposed utility functions reflect interaction effects between the attributes of user i and those of potential mate j. Thus, all the attributes of potential mate j (x_j^B and x_j^W) are specified to be relative to user i's attributes.

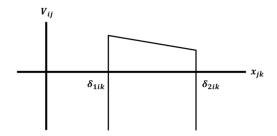
a) Linear Compensatory Rule



b) Non-linear Compensatory Rule



c) Non-compensatory Rule: Conjunctive Rule



d) Non-compensatory Rule: Disjunctive Rule

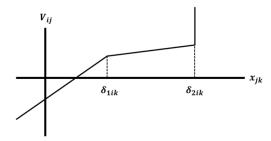


Figure 3.1: Decision rules for continuous (ordinal) attributes

- a) Non-compensatory Rule: Conjunctive Rule
 - $\begin{array}{c|c} v_{ij} \\ \hline & & \\ \hline & & \\ \hline & & \\ \hline & & \\ x_{jl1} & & \\ \hline & \\ \hline & & \\ \hline & \\$

b) Non-compensatory Rule: Disjunctive Rule

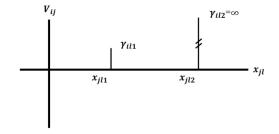


Figure 3.2: Decision rules for categorical attributes

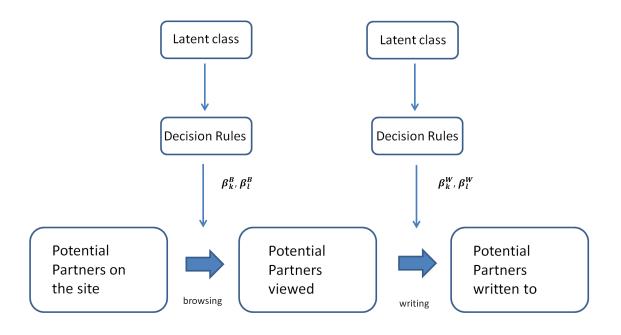


Figure 3.3: Diagram of Mate Choice Process

3.3.4 Two-stage model of mate choice

Both online and offline, mate choice is explicitly a multi-stage process. Online, site users must rst search for potential mates by specifying exclusion criteria based on one or more attributes, and then "browse" potential mates by looking through a list of search results and clicking on attractive proles. Important features of mate choice behavior will be revealed at each stage. For example, a decision to restrict one's search to only members of the same race is quite different from allowing race to be just one of multiple factors determining mate attractiveness at later stages of the selection process. Similarly, offline mate searches do not consider every single person in a given region. Social networks, as well as social venues where people come into contact such as bars, workplaces, and neighborhoods mediate information about available mates. These social environments restrict who is available as a potential mate.

Figure 3.3 provides an overview of the choice process hypothesized in the statistical

model; associated coefficients will be discussed in further detail below. The choice process consists of two stages: the search for available potential mates to consider (the decision to "browse" a particular profile), and the decision to write to a potential mate, given that his or her profile was viewed. The potential choice set includes all men or women on the dating website within the user's metro area at a given time. From these, each user views the profiles for a (typically much smaller) subset, which form the consideration set. From among the browsed profiles, the user may decide to write to one or more potential mates. At each stage, choice is governed by one or more possible decision rules. For example, users mayas implied in past researchadopt a "compensatory" approach in which they compute a weighted sum of *all* potential mates' attributes available at this stage, and browse all those profiles that fall above a user-specific acceptability threshold. Alternatively, users may impose screening rules in which they consider only those profiles that meet some threshold of acceptability on one or more attributes. For example, they may only look for mates within a narrow geographic radius, or with a given level of education or income.⁵

Thus, compared to single stage discrete choice models used in social research (e.g., Bruch & Mare, 2006, 2012; Zeng & Xie, 2008), two stage models better represent the underlying process that people are believed to use in selecting from more than a handful of alternatives. In general, decision rules trade off on effort and accuracy (Johnson & Payne, 1985). In the first stage, when the goal is to reduce the number of potential alternatives to a manageable size, the decision process takes more effort due to the large number of possibilities that must be evaluated. However, the benefit of identifying the most desirable alternatives is small because there is opportunity in the second stage to more fully evaluate possibilities. In the second stage, the number of alternatives to be evaluated is much smaller (thereby reducing costs in terms of cognitive effort) and the benefit of being more accurate

⁵In truth, online dating is a three-stage search process: users first specify a set of search criteria (e.g., restrictions based on geography, age, and education). These search criteria produce a set of potential matches. The user can then "browse" potential matches by clicking on stub versions of their profiles. Finally, after browsing, the user can decide to write a message. In our current data, we do not observe the search criteria or search results; we only observe who was browsed. Thus we collapse the searching and browsing into a single stage.

is greater. As a result, we expect that simpler decision rules are used in the first stage, and more comprehensive roles are used in the second stage. Note that we allow for separate decision rules at each stage, but link the two stages together using latent classes. This allows us to group different response patterns in browsing and writing together. For example, one strategy may be to restrict ones search only to a narrow age range in the browsing stage, but among all profiles who meet the age criteria be indifferent to potential mates' age in the writing stage.

We model each site user's behavior as a sequence of browsing and writing decisions.⁶ In the first stage, the probability that the i^{th} mate seeker will consider (browse) the j^{th} option can be written as a binary logit model:

$$p_{ij}^{B} = \frac{\exp(V_{ij}^{B})}{1 + \exp(V_{ij}^{B})}$$
(3.3)

where V_{ij}^B is the systematic component of utility derived from browsing profile j, described in further detail in the next section. In the second stage, writing behavior (conditional on browsing) is similarly specified as a binary logit model. The probability that user i writes to user j is therefore:

$$p_{ij}^{W}|browsing = \frac{\exp(V_{ij}^{W})}{1 + \exp(V_{ij}^{W})}$$
(3.4)

where V_{ij}^W is the systematic component of utility derived from writing to the j^{th} potential mate. Note that it is not necessary that all salient attributes of potential partners be involved in both the browsing and writing stages of the model. Variables determining the composition of the consideration set and the final choice outcome may overlap partially, completely, or not at all. This reflects the empirical fact, common to all dating sites, that the information available in the browsing stage is typically supplemented by additional variables in the

⁶In this model, we treat the users' entire career on the site as one continuous flow of behavior; we do not distinguish among individual sessions or distinguish between choices made early in one's online dating career and choices made later on. In subsequent work, we plan to focus explicitly on the "learning" that takes place as individuals discover which potential partners are actually plausible options for them online.

writing stage. It also reflects a more general fact about staged decisions making: people may have less information at early stages than at later stages. For example, in housing choice, a decision to view an apartment may be based on a subset of salient attributes: price, location, and number of bedrooms, which are supplemented by an in-person visit.

Because there are large variations in number of profiles browsed and messages sent across users, results from a homogeneous model would differentially reflect the activities of the heaviest site users, and so we consider it especially critical to allow for so-called "unobserved" (i.e., parametric) heterogeneity across users. To avoid presuming that users "clump" near some central tendency on each attribute, we use discrete heterogeneity (Kamakura & Russell, 1989), which also benefits in identifying and interpreting distinct sorts of mate-seeking behaviors. Post-hoc analysis of each group can provide us with new insight of mate preference heterogeneity which cannot be described by a homogeneous model.

3.4 Estimation

3.4.1 ECM algorithm

For estimation, we use the Expectation/Conditional Maximization (ECM) algorithm by Meng & Rubin (1993). The ECM algorithm is a variation of Expectation/Maximization (EM) algorithm (Dempster et al., 1977), which has been commonly used for calculating MLE for finite mixture models (McLachlan & Peel, 2004). In a mixture of regressions with changepoints, the ECM algorithm is typically used, since a single maximization step (M-step) cannot readily handle changepoints and other regression parameters simultaneously. In this case, the M-step is divided into two conditional Maximization steps (CM-steps).

Most relevant to our work, Young (2012) explored mixtures of regressions with changepoints models and developed an ECM algorithm for maximum likelihood estimation. The model includes multiple predictors and changepoints in a mixture of regressions and, importantly, the number of changepoints can vary across components of the mixture. To

explain the ECM algorithm briefly, there are three essential steps: 1) E-step, 2) CM-step 1, and 3) CM-step 2. The E-step is simply the E-step in the usual EM algorithm (McLachlan & Peel, 2004). For the following CM steps, a whole parameter vector is partitioned into two blocks: changepoints, and all other parameters, comprised of regression parameters and weights for the mixing components. In CM-step 1, changepoints are estimated via conditional maximization, by fixing all the other parameters (as given). Maximization is accomplished by using a first order Taylor expansion around the changepoints (Muggeo, 2003); this approach is implemented in the R package "segmented" (Muggeo, 2008). Given changepoints estimated by CM-step 1, CM-step 2 follows the M-step of an EM algorithm for a classic mixture of regressions model (DeSarbo & Cron, 1988, e.g.,). Finally, the ECM algorithm continues until some predefined stopping criterion is met, ordinarily that the degree of improvement in the model log-likelihood falls below some given threshold value for a fixed number of consecutive iterations (See Young, 2012, for details). The described ECM algorithm for the mixtures of regressions with changepoints model is implemented in the "segregmixEM" function of the **R** package "mixtools" (Benaglia et al., 2009). However, the segregmixEM function allows for neither data with multiple observations per individual nor for binary responses. Therefore, we developed a new function that extends segregmixEM in two aspects critical for our application setting: 1) binary responses (with a logit formulation), and 2) accommodating repeated measures per individual.

3.4.2 Mixture of logits model with changepoints

In a "binary response with repeated measurements" context, y_{is} (0 or 1) is the s^{th} observation ($s=1,...,S_i$) for individual i (i=1,...,n), and x_{is} is the corresponding vector of predictors, including an intercept. Following Young (2012)'s notation, m is the number of components and λ_l are the mixing proportions for the component, where $\sum_{l=1}^m \lambda_l = 1$ and $\lambda_l > 0$. β_l is the vector of regression coefficients and γ_l is a vector of changepoints for the l^{th} component. The new ECM algorithm follows the same steps as those in the mixtures of

regressions with changepoints model. However, both the likelihood function and posterior probabilities of component inclusion have been modified accordingly, as follows (See Appendix A for details).

1) Likelihood function In a binary response model (logit), the m-component mixture of logits model with changepoints density is

$$f(y_{is}; x_{is}, \beta, \lambda, \gamma) = \sum_{l=1}^{m} \lambda_l \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}}, \text{ where } \log(\frac{\theta_{isl}}{1 - \theta_{isl}}) = x_{is} (\gamma_l)^T \beta_l$$
 (3.5)

The observed data log likelihood with repeated measurements is

$$ll_o = \sum_{i=1}^n \sum_{s=1}^{S_i} \log(\sum_{l=1}^m \lambda_l \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}})$$
(3.6)

2) Posterior probabilities of component inclusion The posterior probability that individual i is included in component l is

$$p_{il} = \frac{\lambda_l \prod_{s=1}^{S_i} \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}}}{\sum_{h=1}^{m} \lambda_h \prod_{s=1}^{S_i} \theta_{ish}^{y_{is}} (1 - \theta_{ish})^{1 - y_{is}}}$$
(3.7)

These two major modifications are implemented in the *R* function "*replogitregmixEM*" (See Appendix B).

3.4.3 Parameter recovery

To test the *replogitregmixEM* function, simulation studies were conducted. As a first step, instead of generating multiple data sets and checking parameter recovery, we show that "true"(i.e., known) parameters are recovered with multiple distinct starting points. We present the simulation results of 1) a two component mixture of logits model with two

changepoints; and 2) a three component mixture of logits model with two changepoints. In both studies, we generate 200 individuals. Half of the individuals have 20 observations and the other half have 10 observations. Therefore, the total number of observations is 3,000. For clarity, we use one covariate, randomly drawn from U(0,50). The parameters of interest are mixing proportions, an intercept and three slope coefficients, and two changepoints for each component. The true parameters used in the data generation process for study 1 and 2 are represented in Table 3.1 and 3.2 respectively.

Table 3.1: Simulation results for a two component mixture of logits model

		_		1		2	:	3
	Class	True parameters	Starting points	Estimates	Starting points	Estimates	Starting points	Estimates
	1	0.4	0.4	0.3742	0.5	0.3744	0.3	0.3744
mixing proportion	2	0.6	0.6	0.6258	0.5	0.6256	0.7	0.6256
1/:	1	1	1	0.8479	0.5	0.8471	0	0.8457
beta 1(intercept)	2	-1	-1	-0.7706	-2	-0.7714	0	-0.7695
hoto 2/slone 1)	1	-0.3	-0.3	-0.2973	0.1	-0.2970	0	-0.2966
beta 2(slope 1)	2	0.3	0.3	0.2838	0.1	0.2840	0	0.2834
hoto 2/slana 2)	1	0.3	0.3	0.3318	-0.1	0.3318	0	0.3317
beta 3(slope 2)	2	-0.3	-0.3	-0.2968	0.5	-0.2970	0	-0.2965
hota (/slana 2)	1	0.3	0.3	0.2500	0.1	0.2496	0	0.2491
beta 4(slope 3)	2	-0.5	-0.5	-0.4399	0.1	-0.4392	0	-0.4391
CD1/shanganain+1)	1	15	15	15.3637	40	15.3873	10	15.4149
CP1(changepoint1)	2	10	10	10.5185	15	10.5164	20	10.5385
CD2/shanganain+2\	1	35	35	36.0987	40	36.1235	15	36.1141
CP2(changepoint2)	2	40	40	39.8091	35	39.7966	25	39.7993

Table 3.2: Simulation results for a three component mixture of logits model

		T	:	1		2	3	
	Class	True parameters	Starting points	Estimates	Starting points	Estimates	Starting points	Estimates
	1	0.2	0.2	0.1672	0.4	0.1672	0.5	0.1671
mixing proportion	2	0.2	0.2	0.2076	0.3	0.2077	0.3	0.2081
	3	0.6	0.6	0.6252	0.3	0.6252	0.2	0.6248
	1	1	1	2.3858	0.5	2.3811	0	2.3836
beta 1(intercept)	2	-1	-1	-0.3650	-2	-0.3652	0	-0.3623
	3	1	1	0.8869	2	0.8952	0	0.9045
	1	-0.3	-0.3	-0.6642	-0.1	-0.6622	0	-0.6633
beta 2(slope 1)	2	0.3	0.3	0.2469	0.1	0.2469	0	0.2462
	3	-0.2	-0.2	-0.3342	0	-0.3415	0	-0.3502
	1	0.3	0.3	0.5884	0.1	0.5864	0	0.5875
beta 3(slope 2)	2	-0.3	-0.3	-0.2696	-0.1	-0.2695	0	-0.2686
	3	0.2	0.2	0.3436	0.1	0.3503	0	0.3585
	1	0.3	0.3	0.3762	0.2	0.3757	0	0.3761
beta 4(slope 3)	2	-0.5	-0.5	-0.3339	-0.2	-0.3338	0	-0.3335
	3	0.2	0.2	0.1973	-0.1	0.1978	0	0.1985
	1	15	15	5.9396	10	5.9529	10	5.9462
CP1(changepoint1)	2	10	10	10.7608	15	10.7562	15	10.7667
	3	5	5	2.9666	10	2.9181	10	2.8641
	1	35	35	33.9166	40	33.8889	20	33.9084
CP2(changepoint2)	2	40	40	39.8656	30	39.8583	25	39.8538
	3	15	15	14.9505	35	14.9265	30	14.9192

Table 3.1 presents the simulation results of a two component mixture of logits model with two changepoints. It shows that every single estimate across the three simulations converges to a single value close to the true parameter regardless of which of the three distinct starting points at which the algorithm was initialized. The Table 3.2 presents the simulation results of a three component mixture of logits model with two changepoints. Again, every single estimate across the three simulations converges to a single value and most of estimates appear to be relatively close to the true parameters. However, since there is one additional latent class while the number of individuals stays the same, some true parameters are not recovered as well as in the two-component case.

3.5 Empirical results

3.5.1 Data description

Online dating is increasingly becoming a common way to meet one's spouse. Between 1995 and 2005, there was a rapid increase in the number of opposite-gender couples who met their partners online (Rosenfeld & Thomas, 2012). A study commissioned by Match.com in 2010 reported that 1 in 6 couples married within the past three years met their partner on an online dating site, and 1 in 5 people have dated someone they met on an online dating site (Chadwick, 2010). After conditioning on Internet use, Sautter and colleagues found that approximately one third of all people who had been single at some point over the previous ten years had used internet dating websites (Sautter et al., 2010). Some individuals are more likely to use online dating sites than others. Rosenfeld & Thomas (2012) report that the people most likely to use online dating sites are those operating in a thin market, for example, gays, lesbians, and middle-aged heterosexuals. These are the populations who stand most to benefit from the market efficiencies in online dating. The expansion in online dating correlates with an increase in Americans Internet use. According to Current Population Survey data, 55 percent of all households had Internet access; this is more than triple the proportion of the population with Internet access in 1997 (Day et al., 2005). However, some social groups are more likely to have Internet at home than others. Being white, highly educated, high income, and/or having a school-aged child in the household are all positive predictors of Internet use.

This study estimates mate preferences from observed activity on a popular online dating website. These data were originally analyzed by Hitsch et al. (2010), using a single-stage, homogeneous, compensatory response logit model. The sample includes all users active in the San Diego and Boston metro areas over a six-month period. We restrict our sample to users who are 1) seeking opposite gender partners; 2) single, divorced, or "hopeful"; 3) "looking for long-term relationship", "just looking", "making friends", or claimed that a

"friend put me up on this"; and 4) within the ages of 18-65. We also eliminate any user who failed to browse any profiles, since these constitute non-users. This provides 10,271 users in total. There are roughly equal numbers of men (52.49%) and women (47.51%) and about equal numbers of users from Boston (48.19%) and San Diego (51.81%).

The site generates two types of data: 1) user registration information (i.e., profiles), and 2) activities observed on the site. The user registration data contain a variety of user-based attributes, including age, education, income, height, weight, self-rated attractiveness, and an attractiveness rating for extant photos. Our preliminary analysis focuses on two continuous attributes of potential mates: height and age, as well as three discrete attributes of race/ethnicity, having children (or not) and education level. Because we suspect that mate seekers evaluate many characteristics of potential mates relative to their own value, most of these variables are entered into the model either as interactions (to be specified below) or as deviations from ones own value or level.

Table 3.3: user registration information (continuous attributes)

Male

	mean	min	Q1	median	Q3	max
Height	70.8	59	69.5	71.5	73.5	85
Age	37.3	19	28	38	43	63

Female

	mean	min	Q1	median	Q3	max
Height	65.2	59	63.5	65.5	67.5	85
Age	38.3	19	33	38	43	63

Table 3.3 lists the continuous attributes in the user registration data. We see that, on average, men on the site are five inches taller than women on the site and both male and female site users are, again on average, in their late thirties. As mentioned previously, to identify deal-breakers/makers, we focus on *differences* in age and height between users.⁷

⁷We assume that what matters to make seekers is the distance between a potential partner's attribute

Thus, in the analysis, we define a new variable for each attribute:

age_diff = user's age - potential mate's age

height_diff = user's height - potential mate's height

Table 3.4: user registration information (discrete attributes)

Male

Race	White	Black	Hispanic		Asian	Other	
Race	84.96%	3.15%	4.97%		2.63%	4.29%	
17: 1	Yes			No			
Kids	37.61%			62.39%			
	Lower		College		Advanced		
Education Level	15.56%		55.85%		28.58%		

Female

Paco	White	Black	Hispanic		Asian	Other	
Race	84.98%	2.62%	5.35%		3.16%	3.89%	
IX: d -	Yes			No			
Kids	42.62%			57.38%			
	Lov	ver	College		Advanced		
Education Level	12.62%		57.62%		29.72%		

Table 3.4 summarized the categorical attributes taken from user registration data. More than 80 percent of users are white, and less than 5 percent are black. Since whites make up a very high proportion of all users, in the analysis, we combine all the other races into non-white category and examine interactions between and within white (W) and non-white (NW) users. About 38 percent of female users and about 42 percent of male users have children. We examine interactions between users with kids (K) and users without kids (NK) to see whether men and woman only consider potential partners who share their

value and their own attribute value, rather than the absolute value of potential partners' attribute values. For example, a woman does not care how short a man is, as long as he is at least two inches taller than her. In actuality, people probably evaluate potential mates' attributes both in absolute and relative terms.

family status. Finally, users on the site span the full range of education, ranging from less than high school to a doctoral degree. We partitioned 13 categories of education level into three categories: 1) Lower (L), which means less than college degree; 2) College (C), which means college degree; 3) Advanced (A), which means post-college degree. Users currently enrolled in school (high school, college, graduate school) are dropped because the degree of social friction and income might be different from the other users. In the analysis, we use 8 dummy variables for the interaction effects to examine how education level affects mate choice. While these data are not strongly representative of the Boston and San Diego adult populations, they are roughly representative of Internet users in those areas (Hitsch et al., 2010, p137).

The activity data contain time stamped records of all users' browsing and writing history. Table 3.5 and 3.6 show activity data for male and female users, respectively. Each table is divided into "active" (browsing, sending a message) and "passive" (being browsed, receiving a message) activities on the site. These tables show that male users do far more browsing and writing than female users. Male users browsed and sent messages at more than double the volume of female users (active). Consequently, female users were browsed and received messages far more than male users (passive).

3.5.2 The Process of Finding a Mate Online

When they join the dating service, users must fill out a profile providing answers to a number of survey questions as well as several short answer essays. These include measures of a range of demographic attributes (income, marital status, whether they have children, education, race, religion, and age) as well as measures of cultural interests, whether they attend church frequently, spending habits and expectations for a first date. Users are also asked to indicate whether they would be willing to travel to meet a mate, and whether race and religion are important for them in evaluating potential mates. Many users also include one or more photos in their profile.

Once they have completed their profile, users can search for, browse, and write to potential partners. Users typically begin by searching for mates based on a specified age range and geographic region. This query returns a list of "short profiles" containing information on potential partners' age, user name, a brief description, and a photo if available. Users can then decide to "browse" potential mates by clicking on their short profile to access the complete profile containing the full set of profile attributes as well as essay questions, larger versions of the main photo, and additional photos if available. Based on the full profile, users can then decide to write to a potential mate. The data provided a complete moment-by-moment description of users activities, including which profiles he or she browsed, whether or not the photos were viewed, and whether the user sent a first contact message.

Table 3.5: Activity data (male users)

		Total browsing hits			566	,374		
		Male users who browsed			5,233 (83.1%)		
	Active Browsing	Browsing hits per user	mean	min	Q1	median	Q3	max
Active		browsing filts per user	108.23	1	14	45	135	1,396
Active		Total messages sent			57,	338		
	Male users who wrote		3,184 (50.6%)					
	Writing	Messages sent per user	mean	min	Q1	median	Q3	max
		Wiessages sent per user	18.01	1	2	4	14	581
		Female users who got browsed	5,851 (100%)					
	Getting browsed	Browsing hits per user	mean	min	Q1	median	Q3	max
Passive		Browsing into per user	96.80	1	14	32	134	979
Pussive	Receiving a	Female users who received messages	4,693 (80.2%)					
	message	Messages received per	mean	min	Q1	median	Q3	max
		user	12.22	1	2	6	15	144

Table 3.6: Activity data (female users)

		Total browsing hits			257	,242		
	Browsing	Female users who browsed			4765 (81.4%)		
		Browsing hits per user	mean	min	Q1	median	Q3	max
Active		browsing files per user	53.99	1	9	27	67	1,294
7101740		Total messages sent			15,	078		
\\/riting	Female users who wrote		2,391 (40.9%)					
	Writing	Messages sent per user	mean	min	Q1	median	Q3	max
		Wiessages sent per user	6.31	1	1	3	6	120
		Male users who got browsed	6,297 (100%)					
	Getting browsed	Browsing hits per user	mean	min	Q1	median	Q3	max
Passive		browsing files per user	40.85	1	4	12	50	805
Pussive	Receiving a	Male users who received messages	2,985 (47.4%)					
	message	Messages received per	mean	min	Q1	median	Q3	max
		user	5.05	1	1	3	6	65

3.5.3 Estimation results

In this section, we report preliminary model estimation results. We estimate four separate models: one for each of the two stages (browsing and writing) crossed by the two genders (male and female). Each model follows mixture of logits model with changepoints. As described in the model development section, each continuous attribute accommodates two changepoints; that is, it is a piecewise linear spline with (up to) three regions. We specify two latent classes for each model.

To ease preliminary computational burden, we use a random sample of the data for estimation, with the remainder available for holdout testing. We used the activity data from users whose total number of browsed profiles is between 50 and 200. This way, we avoid skewing the activities of typical users with the "long right tail" of light users and the disproportionate influence of a small number of very heavy users, some of whom browsed thousands of profiles. For similar computational and holdout-based reasons, we randomly choose 25% of users from both the male and female respondent pool, yielding 93 male users with 13,899 browsing decision occasions and 7,066 writing decision occasions; and

106 female users with 20,560 browsing decision occasions and 10,576 writing decision occasions.

3.5.3.1 Deal-breaker/maker

Among the main purposes of this project is to find choice (decision) rules characterized by coefficients with "large" absolute values. For discrete attributes, if the coefficient of one of dummy variable is extremely small or large, it implies a potential deal-breaker or -maker, although, once again, we expect the latter to be relatively rare in the general population. For continuous attributes, slopes with "large" absolute values imply deal-breakers or -makers. We informally define a deal-breaker or a deal-maker if the absolute value of a slope is greater than 3 for continuous attributes, or if the difference in dummy coefficients (within one attribute) is greater than 3 for discrete attributes. This corresponds to an increase/decrease in odds ratio of about 20 (by increasing one unit of a certain attribute for continuous attributes, once attributes have been suitably standardized).

3.5.3.2 1st stage - Browsing behavior

Table 3.7 lists the browsing-stage parameter estimates. "Mixing proportion" stands for the proportion of users in each latent class. Among our model covariates, the only information that is visible when viewing a short profile (and thus can inform the decision to browse) is a potential mates age. For this reason, age is the only covariate that allows us to explore directly whether decision rules differ across the two stages. All the other information is only visible once a user decides to view a given profile. There are two changepoints (CP 1 and CP 2) and "slope 1" means the slope of regression line below CP 1. The slope between CP 1 and CP 2 is sum of slope 1 and slope 2, and the slope above CP 2 is the sum of all three slope coefficients; thus, the listed slopes correspond to *differences* between adjacent portions of the piecewise-linear utility function. The table suggests that all three slope parameters are nonsignificant for class 2 users (both male and female). Regarding class 1

users, the absolute values of all the slope coefficients for both men and women are below 3, which suggests that age does not seem to be acting as a deal-breaker in the decision to browse. Figure 3.4 shows the age difference regression lines with two changepoints for male and female users in class 1. It appears, for this two-segment solution, that there are no truly extreme slopes. This suggests that people do not place inviolable or even especially severe restrictions on a potential mate's age when they decide to browse a profile, at odds with our original hypothesis (although the interpretation of this result is partly dependent on the underlying scaling of the age variables). Regardless, age difference plays a distinct role in the writing stage, which will be described later in the section.

Table 3.7: Parameter estimates for Browsing behavior

		Ma	ale	Fen	nale
		Class 1	Class 2	Class 1	Class 2
		Estimate (SE)	Estimate (SE)	Estimate (SE)	Estimate (SE)
Mixing Proportion		0.476 (0.101)	0.524 (0.101)	0.424 (0.093)	0.576 (0.093)
Interc	ept	0.558 (0.754)	-0.014 (0.843)	4.764 (0.903)	0.441 (0.332)
	slope 1	0.159 (0.082) *	0.035 (0.035)	0.403 (0.081) ***	0.064 (0.021)
	slope 2	0.232 (0.264)	-0.163 (0.160)	-0.548 (0.234) ***	-0.328 (0.121)
Age_diff	slope 3	-0.677 (0.313) **	0.134 (0.129)	-0.381 (0.089) ***	0.140 (0.104)
	CP 1	-6.822 (2.138)	10.000 (0.004)	-5.000 (0.081)	5.000 (0.013)
	CP 2	1.652 (0.557)	23.858 (3.252)	-0.109 (0.252)	12.825 (1.916)

*** p < .01 ** p < .05 * p < .1

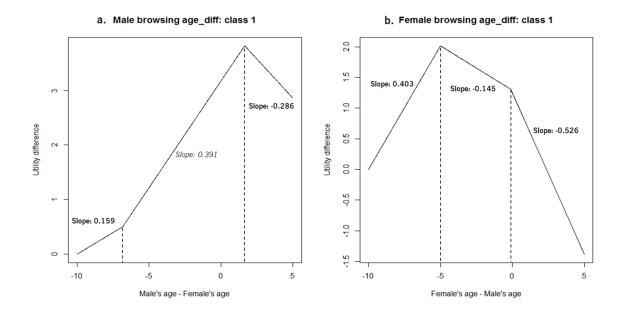


Figure 3.4: Browsing - Age difference for Male and Female users

3.5.3.3 2nd stage - Writing behavior

Table 3.8 lists parameter estimates for the writing stage for both male and female users. As was done for browsing behavior, we specify two latent classes for each model. The table shows that the majority of male users (88.9%) fall into class 2, while a majority of female users (93.8%) do so in class 1. In the writing stage, in addition to a user's age, information on a user's education level, ethnicity, whether living with kids, and height, is available. We use three discrete attributes (Education, Race, and Kids) and two continuous attributes (Age and Height) in the analysis of writing behavior. Since we are interested in how a users attributes interact with a potential mates attributes, we specify covariates to be relative within user-mate pairs. We describe the estimation results of each attribute in turn.

Table 3.8: Parameter estimates for Writing behavior

		Ma	ale	Fen	nale
		Class 1	Class 2	Class 1	Class 2
		Estimate (SE)	Estimate (SE)	Estimate (SE)	Estimate (SE)
Mixing Pro	portion	0.111 (0.015)	0.889 (0.015)	0.938 (0.007)	0.062 (0.007)
Interce	ept	-1.413 (0.947)	49.213 (14.104)	72.440(19.328)	42.604(10.024)
	L-C	2.433 (1.355)	-0.329 (0.463)	-43.630 (9.944) ***	19.747 (6.188) ***
	L-A	2.157 (1.913)	-0.260 (0.667)	-60.852 (9.634) ***	20.496 (5.718) ***
	C-L	1.711 (1.22)	-1.129(0.477)***	-1.898(1.681)	10.412(3.443)***
Education	C-C	2.183 (0.985) **	-0.788 (0.409) **	-0.667 (0.615)	11.808(3.472)***
Education	C-A	2.929 (1.181) ***	-1.311(0.450)***	-0.287 (0.605)	12.092(3.714)***
	A-L	2.029 (1.262)	-0.777 (0.516)	-0.812 (0.783)	-44.314 (6.481) ***
	A-C	5.089 (1.739) ***	-1.496 (0.479) ***	0.290 (0.573)	0.330 (1.926)
	A-A	5.315 (1.387) ***	-1.649 (0.528) ***	0.701 (0.576)	-30.754 (2.917) ***
	W-NW	-0.558 (0.638)	0.036 (0.204)	-98.443 (15.504) ***	-4.168(1.023)***
Race	NW-W	0.089 (0.639)	-19.549 (2.256) ***	0.063 (0.330)	-8.126(2.517)***
	NW-NW	1.081 (1.55)	-0.126 (0.606)	0.196 (0.921)	4.671 (2.226) **
	K-K	0.343 (0.702)	0.339 (0.202) *	-0.363 (0.208) *	7.573 (1.935) ***
Kids	K-NK	-0.501 (0.802)	0.069 (0.240)	-0.265 (0.197)	0.721 (0.913)
	NK-K	-0.919 (0.483) *	-0.459 (0.249) *	-0.215 (0.198)	1.244 (0.699) **
	slope 1	0.339 (0.081) ***	6.583 (1.471) ***	4.472 (0.864) ***	3.564 (0.920) ***
	slope 2	-0.586(0.141)***	-6.579(1.473)***	-4.470(0.862)***	-3.241(0.899)***
Age_diff	slope 3	-8.253 (1.000) ***	-8.968 (5.834)	-15.047 (1.632) ***	-23.162 (3.730) ***
	CP 1	3.649 (0.994)	-9.910 (0.065)	-9.784 (0.095)	-15.481 (0.706)
	CP 2	17.658 (1.000)	14.985 (0.055)	4.954 (0.021)	4.782 (0.040)
	slope 1	9.595 (3.229) ***	13.041(1.146)***	2.060 (1.857)	0.046 (0.124)
	slope 2	-9.937 (3.239) ***	-12.920 (1.142) ***	-2.078 (1.859)	-0.379 (0.211)
Height_diff	slope 3	-5.010(2.377)***	-8.921(1.151)***	-0.721(0.211)***	-21.692 (4.761) ***
	CP 1	-0.018 (0.023)	1.046 (0.154)	-15.010 (0.041)	-5.986 (0.093)
	CP 2	8.501 (0.004)	12.596 (0.044)	-2.386 (0.390)	3.115 (0.681)

*** p < .01 ** p < .05 * p < .1

Education level. As described earlier, there are three education levels: L (Lower), C (College), and A (Advanced). Therefore, the interaction of the education level between a user and a potential mate (User's education Mate's education) can be addressed by 8 parameters, with a baseline (i.e., set for identification) utility of L-L. Table 3.9 and 3.10 represent education parameter estimates for male and female users respectively. The left sub-table in Table 3.9 pertains to male users in class 1. It suggests that, for male users with advanced education (A), having lower education (L) can function as a deal-breaker: differences in the coefficients (A-L and A-C / A-L and A-A) are large (i.e., significantly greater than 3). For male users in class 2 (right sub-table), males users with college education (C) prefer females with college education (C). This largely supports the matching

hypothesis, although the effect itself is not large. In spite of the small differences in the coefficients (and therefore not standing out as deal-breakers or -makers), the table suggest that male users with advanced education (A) in class 2 prefer females with lower education (L), a result opposite to the writing pattern of the male users in class 1. Table 3.10 shows parameter estimates of female users. The left sub-table suggests that, for the female users with lower education (L) in class 1, males with higher education level (C or A) can act as a deal-breaker, supporting the matching hypothesis. However, as can be seen in the right sub-table, males with higher education level (C or A) can serve as a deal-maker for females with lower education (L) in class 2, which supports the competition hypothesis. In addition, having a college degree (C) can function as a deal-maker for females with lower education (L) in class 2.

Table 3.9: Writing - Education level for Male users

Class 1 (11.1%)		Female				
Class 1	(11.170)	L C		Α		
	L	0	2.433	2.157		
Male	С	1.711	2.183(**)	2.929		
	Α	2.029	5.089(***)	5.315(***)		

Class 2 (88.9%)			Female				
Class 2	(00.5%)	L	С	Α			
	L	0	-0.329	-0.26			
Male	С	-1.129(***)	-0.788(**)	-1.311(***)			
	Α	-0.777	-1.496(***)	-1.649(***)			

*** p < .01 ** p < .05 * p < .1

Table 3.10: Writing - Education level for Female users

Class 1 (93.8%)		Male		
		L	С	Α
Female	L	0	-43.63(***)	-60.852(***)
	С	-1.898	-0.667	-0.287
	Α	-0.812	0.29	0.701

Class 2 (6.2%)		Male		
		L	С	Α
Female	L	0	19.747(***)	20.496(***)
	С	10.412(***)	11.808(***)	12.092(***)
	Α	-44.314(***)	0.33	-30.754(***)

*** p < .01 ** p < .05 * p < .1

Race. We use two racial categories for the analysis: White (W) and Non White (NW). The baseline is a match between Whites (W-W). Table 3.8 lists the parameter estimates for race interactions. An intriguing finding regarding race is that there is significant difference across genders. For example, for NW males in class 2 (88.9%), being a W female can

work as a deal-breaker, while W males are far more open to NW females. Contrarily, being a NW male can function as a deal-breaker for W females in class 1 (93.8%), while NW females are substantially more open to W males. This pattern of results suggests that both matching and competition can play important roles in mate choice decisions with regard to race (Lin & Lundquist, 2013).

Kids. Living with kids (K) and not living with kids (NK) are the two categories used, with an identification baseline of NK-NK. Table 3.8 shows the parameter estimates for the Kids interactions. In general, the matching hypothesis holds across genders and latent classes. Specifically, the female users with kids (NK) in class 2 (6.2%) prefer male users with kids (K), to an extent that this can work as something of a deal-maker. In addition, females without kids (NK) in class 2 prefer males with kids (K), which does not happen for male users in either class.

Age differences. As described previously, we use the difference between a site user and potential mates as a covariate for continuous attributes. In the case of age, the new covariate of age difference (age_diff) for a male user is defined as a (male) user's age - a (female) mate's age. Analogously, for a female user, this becomes (female) user's age - a (male) mate's age.

Figure 3.5 represents the effect of age difference, in writing, for male users. Figure 3.5a shows the utility for writing of male users in class 1, which accounts for 11.1% of male users. The slopes of the first two regression lines are relatively flat, and the utility reaches its highest value at the first changepoint (when a male user is older than a female user by approximately 4 years). However, the utility decreases rapidly at the second changepoint (17.658), which suggests that, for males, *being approximately 18 years or older a potential female mate* is a deal-breaker. Figure 3.5b depicts the utility for writing among male users in class 2, which accounts for 88.9% of male users. The regression lines' slopes convey

that utility increases rapidly (slope 1: -9.910) at the first changepoint (CP 1: 0.011) and stays relatively flat until the second changepoint (CP 2: 14.985). At this point, the utility decreases rapidly (slope 3: -8.964). This suggests that *being approximately 10 years or older than female users* can function as a deal-breaker in writing for male users. However, it is premature to make a firm conclusion about deal-breakers on the other side, since the third slope (slope 3) is not statistically significant.

Figure 3.6 represents the effect of age differences in writing for female users. Figure 3.6a shows the utility for writing of female users in class 1, which accounts for 93.8% of female users. All three slope parameters are significant, and there are large changes in slope at the two changepoints, suggesting that there are deal-breakers in age for female users in this class: either being approximately 10 years or younger than male users, or being approximately 5 years or older. That the second condition matters more than the first can be seen in the steeper slope in the right-hand side of the utility function. The utility function for female users in class 2 is illustrated in Figure 3.6b. The pattern is the same as the utility function for class 1 users and there are two slightly different age deal-breakers: either being approximately 15 years or younger than male users, or being approximately 5 years or older.

Compared to the result of age differences in browsing behavior, which does not reveal any evident deal-breaker/makers, this result suggests that the decision rules used to evaluate a given attribute can differ across stages of the choice process. In the specific case of age, people tend to impose stricter restrictions in the writing stage than the browsing stage. This makes intuitive sense: writing is a 'higher cost' activity than browsing, in terms of time, cognitive investment, and potential for regret/rejection; one would therefore expect cutoffs for writing to be more severe than for analogous attributes in the browsing phase.



Figure 3.5: Writing - Age difference for Male users



Figure 3.6: Writing - Age difference for Female users

Height difference. Similar to age, height difference (height_diff) for a male user is defined as a (male) user's height - a (female) mate's height. Analogously, for a female user, it is a (female) user's height - a (male) mate's height.

Figure 3.7 represents the effect of height difference in writing for male users. Figure 3.7a shows the utility for writing of male users in class 1, which accounts for 11.1% of

male users. All three slope parameters are significant, and there are large changes in slope at the two changepoints (-0.018 and 8.501). This suggests that, for males, *either being shorter than female users*, *or being 8.5 inches or taller*, can function as deal-breakers (in writing, for male users). The first condition is apparently more important than the second, as can be seen in the steeper slope in the left-hand side of the utility function. Male users in class 2 (Figure 3.7b) show the same pattern as those in class 1, albeit with slightly different changepoints (1.046 and 12.596). Specifically, for males, *either being approximately 1 inch or shorter than female users, or being approximately a foot or taller*, can serve as deal-breakers.

Figure 3.8 depicts the effect of height difference in writing for female users. For both classes, the first two slopes are statistically nonsignificant, and the slope is not steep (less than 3), while the third slope is significant. Specifically, for female users in class 2 (which accounts for 6.2% of female users), *being approximately 3 inches or taller than males* works as a deal-breaker. We note that this is a relatively small class, and that most women set their "deal-breaker" height for males very close to their own, as the class 1 results suggest.

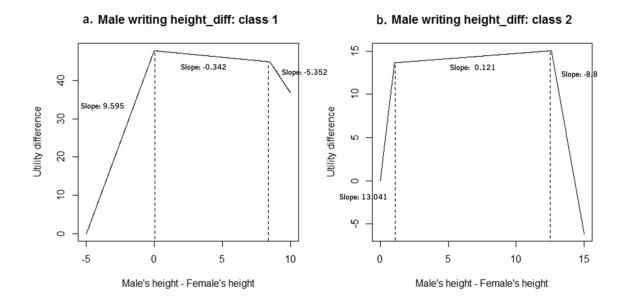


Figure 3.7: Writing - Height difference for Male users

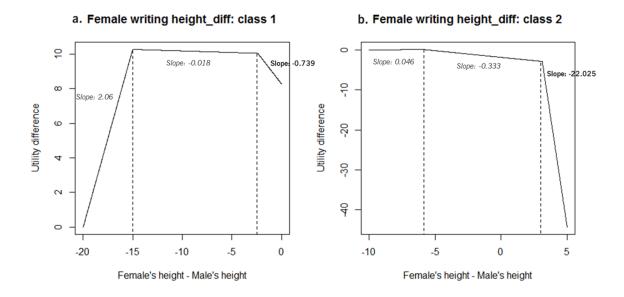


Figure 3.8: Writing - Height difference for Female users

To summarize, these results translate directly into underlying decision rules keyed to each attribute. Different patterns in the dummy coefficients (Table 3.7-3.10) and the shape of utility functions (Figure 3.4-3.8) suggest the existence of heterogeneity in decision rules across 1) gender, 2) stage, and 3) site users given gender and stage (via latent classes). Also, we found potential deal-breakers/makers across both discrete and continuous attributes. Users with distinct types of decision rules might be identified through the model by the extraction of more latent classes, or via a less restrictive (individual-level) analysis using Bayesian nonparametrics, e.g., a Dirichlet process prior. We believe this to be a fruitful avenue for future research in this general application domain.

3.6 Conclusion

In this research, we develop a mate choice model that allows us to identify decision rules at different stages (browsing and writing) of the mate choice process. The proposed model can accommodate compensatory and non-compensatory decision rules, and it allows decision rules to differ across stages. In addition, heterogeneity in decision rules across users within

a stage can be captured by a finite mixture model formulation (among other possibilities). Empirical results suggest that decision rule can vary across stages, as we found for example in the "age difference" variable. Also, the mixture model with two latent classes suggests there is indeed heterogeneity in decision rules across users. We await more thorough results involving increasingly detailed accounts of parametric heterogeneity, but even these admittedly preliminary latent class estimates decisively rule out homogeneity for these data.

The contribution of this research can be summarized along three distinct lines. First, theoretically, we suggest an alternative framework for studying mate preferences and mate choice that does not assume an implicit single-stage and compensatory decision rule. Second, methodologically, this research extends the ECM algorithm for mixtures of regressions with changepoints model (Young, 2012) to accommodate repeated measurements and binary outcomes. Substantively, the empirical result suggests the existence of heterogeneity in choice rules across stages and across users. In terms of specific findings, education level, ethnicity, and differences in both age and height can indeed function as "deal breakers" for certain group of users.

There are some clear limitations that mainly originate with the data. First, there is no information about search criteria used. This can be problematic because search criteria affect users' behavior in the subsequent stage (e.g., either further browsing or writing) by restricting profiles shown to the user. On the other hand, information on search criteria would help identify a user's true preference and decision rules. Second, the data contain activities over a relatively short time window. For example, they cannot account for seasonality and the ability to empirically check such maxims as that, in the spring, "a young man's fancy turns to love". More importantly, it is difficult to capture dynamics among users, among the most important issues in two-sided matching contexts. We expect that people learn about their own desirability through the responses of other users and they adjust their behavior accordingly. Through such a learning process, a users decision rules could change, and a user might well behave strategically, altering his or her profile in order to reveal only the

most advantageous information relative to the pool of potential mates and rivals.

Choice in a two-sided matching context is clearly an under-researched area, one that opens up great opportunities for future investigation. Among these, we consider the development of a dynamic consideration set formation model (through which we can address changes in decision rules over time, and examine strategic behavior among users) to be the most pressing and intriguing topic in the area, and hope to turn our attention to that and related issues in short order.

3.7 Appendix

A. ECM algorithm for the mixture of logits model with changepoints

In a "binary response with repeated measurements" context, y_{is} (0 or 1) is the s^{th} observation ($s=1,...,S_i$) for individual i (i=1,...,n); x_{is} is the corresponding vector of predictors; m is the number of components; and λ_l are the mixing proportions for the component, where $\sum_{l=1}^{m} \lambda_l$ and $\lambda_l > 0$. First, we define terms for the mixture of logits model with changepoints, based on Young (2012)'s notation.

1. Logit model with changepoint

$$\Pr(y_{is} = 1 | x_{is}) = \frac{\exp(u_{is})}{1 + \exp(u_{is})}$$

$$u_{is} = \beta_0 + \beta_1 x_{is} + \beta_2 (x_{is} - \gamma)_+$$

where γ is a changepoint and $(x_{is} - \gamma)_+ = (x_{is} - \gamma)I[x_{is} > \gamma]$

2. Augmented predictor vector

Let c_{jl} be a known number of changepoints for predictor j, j=1,...,p and $\gamma_{jc_{jl}l}$ be a corresponding changepoint in the l^{th} component.

$$x_{isj}(\gamma_{jl}) = (x_{is}, (x_{is} - \gamma_{j1l})_+, ..., (x_{is} - \gamma_{jc_{jl}l})_+)^T$$

where $\gamma_{jl} = (\gamma_{j1l}, ... \gamma_{jc_{jl}l})^T$ is a vector of changepoints for predictor j in the l^{th} component.

3. Vector of predictor j for observation i

$$oldsymbol{x_{ij}(\gamma_{jl})} = (oldsymbol{x_{i1j}(\gamma_{jl})}^T,...,oldsymbol{x_{iS_{ij}}(\gamma_{jl})}^T)^T$$

4. Vector of all p augmented predictors for observation i

$$\boldsymbol{x_i(\gamma_l)} = (1, \boldsymbol{x_{i1}(\gamma_{1l})}^T, ..., \boldsymbol{x_{ip}(\gamma_{pl})}^T)^T$$

where $\gamma_l = (\gamma_{1l}^T, ..., \gamma_{pl}^T)^T$ is a vector of changepoints for the l^{th} component and $c_{.l} = \sum_{j=1}^p c_{jl}$.

5. Augmented design matrix for component l

$$oldsymbol{X}(oldsymbol{\gamma_l}) = (oldsymbol{x_1}(oldsymbol{\gamma_l}),...,oldsymbol{x_n}(oldsymbol{\gamma_l}))$$

with the corresponding regression coefficients vector

$$\beta_{l} = (\beta_{00l}, \beta_{10l}, ..., \beta_{0c_1l}, ..., \beta_{p0l}, ..., \beta_{pc_pl})^{T}$$

where β_{jk^*l} is the k^{*th} regression coefficient for j^{th} augmented predictor in component l.

6. Parameter vector

$$\psi = (\beta_1^T, ..., \beta_m^T, \gamma_1^T, ..., \gamma_m^T, \lambda_1, ..., \lambda_{m-1})^T$$

7. m-component mixture of logit model with changepoints density

$$f(y_{is}; x_{is}, \boldsymbol{\psi}) = \sum_{l=1}^{m} \lambda_l \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}}, \text{ where } \log(\frac{\theta_{isl}}{1 - \theta_{isl}}) = x_{is} (\boldsymbol{\gamma_l})^T \boldsymbol{\beta_l}$$

8. Observed data log likelihood

$$ll_o(\boldsymbol{\psi}) = \sum_{i=1}^n \sum_{s=1}^{S_i} \log(\sum_{l=1}^m \lambda_l \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}})$$

9. Complete data log likelihood

$$ll_c(\boldsymbol{\psi}) = \sum_{i=1}^{n} \sum_{s=1}^{S_i} \sum_{l=1}^{m} Z_{il} log[\lambda_l \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}}]$$

where
$$Z_{ij} = \begin{cases} 1, & \text{if observation } i \text{ belongs to component } l \\ 0, & \text{otherwise} \end{cases}$$

ECM Algorithm

1. E-step

At iteration t, t = 0, 1, ..., calculate the expected complete data log likelihood

$$Q(\boldsymbol{\psi}; \boldsymbol{\psi^{(t)}}) = \sum_{i=1}^{n} \sum_{s=1}^{S_i} \sum_{l=1}^{m} p_{il}^{(t)} log[\lambda_l \theta_{isl}^{y_{is}} (1 - \theta_{isl})^{1 - y_{is}}]$$

where the posterior probability that individual i is included in component l is

$$p_{il}^{(t)} = \frac{\lambda_l^{(t)} \prod_{s=1}^{S_i} \theta_{isl}^{(t)}^{y_{is}} (1 - \theta_{isl}^{(t)})^{1 - y_{is}}}{\sum_{h=1}^{m} \lambda_h^{(t)} \prod_{s=1}^{S_i} \theta_{ish}^{(t)}^{y_{is}} (1 - \theta_{ish}^{(t)})^{1 - y_{is}}}$$

2. CM-step

For CM, partition the parameter vector $\boldsymbol{\psi}$ into $(\boldsymbol{\psi_1}^T, \boldsymbol{\psi_2}^T)$,

$$\psi_1 = ({\gamma_1}^T, ..., {\gamma_m}^T)^T, \ \psi_2 = ({\beta_1}^T, ..., {\beta_m}^T, \lambda_1, ..., \lambda_{m-1})^T$$

2.1 CM-step 1

$$\boldsymbol{\psi_1}^{(t+1)} = \underset{\boldsymbol{\psi_1}}{argmax} Q(\boldsymbol{\psi}; \boldsymbol{\psi}^{(t)})$$

with ψ_2 fixed at $\psi_2^{(t)}$. Maximization is accomplished by using a first order Taylor expansion around the changepoints (Muggeo 2003); this approach is implemented in the R package "segmented" (Muggeo 2008).

2.2 CM-step 2

$$\boldsymbol{\psi_2}^{(t+1)} = \underset{\boldsymbol{\psi_2}}{argmax} Q(\boldsymbol{\psi}; \boldsymbol{\psi}^{(t)})$$

with ψ_1 fixed at $\psi_1^{(t)}$. Then, CM-step 2 is similar to the M-step of an EM algorithm for a classic mixture of regressions model (e.g., DeSarbo and Cron 1988).

$$\lambda_l^{(t+1)} = \frac{1}{n} \sum_{i=1}^n p_{il}^{(t)}$$

$$oldsymbol{eta_l}^{(t+1)} = oldsymbol{P}(oldsymbol{\gamma_l}^{(t+1)})^{-1}oldsymbol{X}(oldsymbol{\gamma_l}^{(t+1)})oldsymbol{W_l}^{(t)}oldsymbol{y}$$

where

$$\boldsymbol{P}(\boldsymbol{\gamma_l}^{(t+1)}) = \boldsymbol{X}(\boldsymbol{\gamma_l}^{(t+1)})^T \boldsymbol{W_l}^{(t)} \boldsymbol{X}(\boldsymbol{\gamma_l}^{(t+1)})$$

$$\boldsymbol{W_l}^{(t)} = diag(p_{1l}^{(t)} \otimes \iota_1, ..., p_{nl}^{(t)} \otimes \iota_n)$$

$$\iota_i = \left[\begin{array}{c} 1\\ \vdots\\ 1 \end{array} \right]_{S_i \times 1}$$

The ECM algorithm continues until some predefined stopping criterion is met.

$$ll_o(\boldsymbol{\psi}^{(t+1)}) - ll_o(\boldsymbol{\psi}^{(t)}) < \epsilon$$

B. R code for replogitsegregmixEM function

```
replogitsegregmixEM=function (y, x, id, lambda = NULL, beta = NULL, k = 2, seg.Z, psi, psi
        .locs = NULL, delta = NULL, epsilon = 1e-5, maxit = 1000000, verb = FALSE, max.
        restarts=100000)
 2
 3
     logit <- function(x) 1/(1 + \exp(-x))
 4
 5
     if (sum(x[,1]==1)==nrow(x)) x=x[,-1]
 6
     x=data.frame(x)
 7
     col.names.x <- colnames(x)</pre>
 8
     xnam <- colnames(x)</pre>
 9
     fmla <- as.formula(paste("y ~ ", paste(xnam, collapse= "+")))</pre>
10
11 if(!is.null(psi.locs)){
12 psi.counts=apply(psi,1,sum)
13
14 for (i in 1:k) {
15
     if(psi.counts[i]>0){
16
       TEMP <- (is.list(psi.locs[[i]]))&(length(psi.locs[[i]])==sum(psi[i,]>0))
17
18
       TEMP <- is.null(psi.locs[[i]])</pre>
19
20
     if(TEMP==FALSE) stop(paste("You must specify a correct changepoint structure!",
21
        "\n"))
22 }
23 }
24
25
     if (!is.null(delta)) {
26
        cat("Estimation performed assuming the changepoints are known.", "\n")
27
       if (is.null(psi.locs)) {
28
          stop(paste("You must specify the changepoints for this setting!",
29
            "\n"))
30
       }
31
      }
32
33
      if ((length(seg.Z) != k) | class(seg.Z) != "list") {
```

```
34
       stop(paste("You must specify a list of length k for the segmented relationships!",
35
          "\n"))
36
37
     if (!identical(all.equal(dim(psi),c(k,ncol(x))),TRUE)) {
38
       stop(paste("You must specify a matrix with the correct dimension for psi!",
39
          "\n"))
40
41
     if (((length(psi.locs) != k) | class(psi.locs) != "list") & !is.null(psi.locs)) {
42
       stop(paste("You must specify a list of length k for the number of changepoints per
           predictor in each component!",
43
         "\n"))
44
45
     tot.cp <- apply(psi,1,sum)</pre>
46
47
     tmp.ind=1
48
     tmp <- try(suppressWarnings(logitsegregmix.init(y=y, x=x, lambda = lambda, beta = beta,</pre>
          k = k, seg.Z=seg.Z, psi=psi, psi.locs = psi.locs)),silent=TRUE)
49
50
     if(class(tmp) == "try-error") {
51
       cat("Generating new initial values.", "\n")
52
       while(tmp.ind<=100){
53
         tmp <- try(suppressWarnings(logitsegregmix.init(y=y, x=x, lambda = NULL, beta = NULL
              , k = k, seg.Z=seg.Z, psi=psi, psi.locs = NULL)),silent=TRUE)
54
         tmp.ind <- tmp.ind+1</pre>
55
         if(tmp.ind==101) stop(paste("Had to reinitialize algorithm too many times.
              Reconsider specified model.", "\n"))
56
         if(class(tmp)!="try-error") tmp.ind=200
57
       }
58
     }
59
60
     x.old=x
61
     x = cbind(1, x)
62
     data.x=cbind(y,x)
63
     lambda <- tmp$lambda
64
     beta <- tmp$beta
65
     k <- tmp$k
66
     psi.locs <- tmp$psi.locs</pre>
67
     sing < - 0
68
     perms=perm(k,k)
69
     perm.ind=nrow(perms)
70
     if (is.null(delta)) delta <- lapply(1:k, function(i) NULL)</pre>
```

```
71
      n <- length(y)
 72
       diff <- 1
 73
       iter <- 0
 74
      z = matrix(nrow = n, ncol = k)
 75
      n_id <-max(id)
 76
       z2 = matrix(nrow = n_id, ncol = k)
 77
       restarts <- 0
 78
 79
       X.aug \leftarrow lapply(1:k, function(i) cbind(1,aug.x(x[,-1],unlist(psi.locs[[i]]),psi[i,],
           delta=delta[[i]])))
 80
      X.aug.old <- X.aug
 81
       psi.locs.old <- psi.locs</pre>
 82
       xbeta <- lapply(1:k, function(i) X.aug[[i]] %*% matrix(beta[[i]],ncol=1))</pre>
 83
       xbeta2<- sapply(1:k, function(i) xbeta[[i]])</pre>
 84
       comp <- dbinom(y, size = 1, prob = logit(xbeta2))*matrix(lambda,n,k,byrow=T)</pre>
 85
       obsloglik <- sum(log(apply(comp, 1, sum)))</pre>
 86
      ll <- obsloglik
       baseint<-10
 87
 88
 89
       while (diff > epsilon && iter < maxit) {
 90
 91
       sing<-0
 92
      null.beta<-0
 93
      lambda.old <- lambda
 94
       beta.old <- beta
 95
 96
             log.fy.temp<-matrix(double(n*k),n,k)</pre>
 97
               for (i in 1:n) {
 98
                 for (j in 1:k) {
 99
                  log.fy.temp[i,j] < -log(logit(xbeta2[i,j])^y[i] * (1-logit(xbeta2[i,j]))^(1-y[i])
100
                  if (\log.fy.temp[i,j] ==-Inf) \{\log.fy.temp[i,j] < --1e+301\}
101
                }
102
               }
103
104
             log.fy<-data.frame(log.fy.temp,id)</pre>
105
             log.prod_fy_id_temp<-ddply(log.fy,"id", numcolwise(sum))[,-1]</pre>
106
             log.prod_fy_id<-data.matrix(log.prod_fy_id_temp)</pre>
107
108
               for (i in 1:n_id) {
109
                 for (j in 1:k) {
```

```
110
                 z.denom<-0
111
                 for (h in 1:k) {
112
                 \verb|z.denom<-z.denom+exp(log(lambda[h])+log.prod_fy_id[i,h]-log(lambda[j])-log.|
                     prod_fy_id[i,j])
113
114
                 z2[i, j] = 1/(z.denom)
115
116
117
118
               lambda.new <- apply(z2, 2, mean)</pre>
119
               for (i in 1:k) {
120
               z[,i] < -rep(z2[,i], rle(id) $length)
121
122
123
               z2.old < -z2
124
               z.old <- z
125
126 if (sum(lambda.new < 1e-05) > 0 \mid | is.na(sum(lambda.new))) {
127
         sing <- 1
128
       }
129
130
      else {
131
        glm.out <- vector("list",k)</pre>
132
        psi.temp=psi.locs
133
        psi.ind=lapply(1:k,function(i) which(psi[i,]!=0))
134
135
         for(i in 1:k){
136
137
           ww<-z[,i]
138
139
           if(is.null(seg.Z[[i]]) | (sum(1-sapply(delta,is.null))>0)){
140
             temp.seg <- glm(fmla,data=data.x,weights=ww,family = quasibinomial(),control=list(</pre>
                 maxit=500))
141
142
           else temp.seg <- try(suppressWarnings(segmented(glm(fmla,data=data.x,weights=ww,</pre>
               family = quasibinomial(),control=list(maxit=500)),seg.Z=seg.Z[[i]],psi=psi.temp
               [[i]],control=seg.control(display=FALSE))),silent=TRUE)
143
144
           if(class(temp.seg)[1] == "try-error"){
145
             sq = 1
146
             temp.names = names(psi.locs.old[[i]])
```

```
147
             cat("1. Error","\n")
148
149
             while (sq < 20) {
150
               psi.temp2 <- vector("list",length(psi.temp[[i]]))</pre>
151
152
               for(ii in 1:length(psi.temp[[i]])){
153
                 x.range <- range(data.x[, which(names(data.x) == temp.names[ii])])</pre>
154
                 psi.temp2[[ii]] \leftarrow psi.temp[[i]][[ii]]+sample(c(-1,1),length(psi.temp[[i])[[ii]])
                      ]]),replace=TRUE)*runif(length(psi.temp[[i]][[ii]]),0,diff(x.range)/
                     baseint)
155
                 if((any(psi.temp2[[ii]]<=x.range[1]))|(any(psi.temp2[[ii]])>=x.range[2]))){
156
                 psi.temp2[[ii]]=psi.temp[[i]][[ii]]
157
158
159
                 psi.temp2[[ii]]=sort(psi.temp2[[ii]])
160
161
162
               names(psi.temp2)=temp.names
163
               temp.seg <- try(suppressWarnings(segmented(glm(fmla,data=data.x,weights=ww,</pre>
                    family = quasibinomial(),control=list(maxit=500)),seg.Z=seg.Z[[i]],psi=psi.
                   temp2,control=seg.control(display=FALSE))),silent=TRUE)
164
               if(class(temp.seg)[1] == "try-error") {
165
                 sq = sq+1
166
               } else {
167
                 sq=40
168
169
170
171
             if(sq==20){
172
               temp.seg <- try(suppressWarnings(segmented(glm(fmla,data=data.x,weights=ww,</pre>
                    family = quasibinomial(),control=list(maxit=500)),seq.Z=seq.Z[[i]],psi=psi.
                   temp[[i]],control=seg.control(display=FALSE))),silent=TRUE)
                                                                                       \#\# lm=>glm
173
               baseint<-baseint*1.1
174
175
176
           glm.out[[i]]=temp.seg
177
178
179
         lambda <- lambda.new</pre>
180
181
         if(sum(sapply(glm.out,class) == "try-error") > 0) {
```

```
182
           newobsloglik=-Inf
183
184
         else
185
186
           if (sum(1-sapply(delta,is.null))>0) {
187
           psi.new <- psi.locs.old
188
           }
189
190
           else {
191
              psi.new <- psi.locs
192
              for(i in 1:k){
193
                if(class(glm.out[[i]])[1] == "segmented"){
194
                  temp.names=names(psi.locs[[i]])
195
                  temp.cumsum=cumsum(sapply(psi.locs[[i]],length))
196
                  TC.ind = length(temp.cumsum)
197
                  seg.temp = glm.out[[i]]$psi[,2]
198
                  psi.new[[i]] = lapply(1:length(psi.locs[[i]]), function(j) as.numeric(glm.out
                       [[i]]$psi[,2]))
199
                  psi.new[[i]] = vector("list", TC.ind)
200
                  psi.new[[i]][[1]] = sort(seg.temp[1:temp.cumsum[1]])
201
                  if(TC.ind>1) for(j in 2:TC.ind) psi.new[[i]][[j]] = sort(seg.temp[(temp.cumsum
                       [j-1]+1):temp.cumsum[j]])
202
                  names (psi.new[[i]]) = temp.names
203
                }
204
205
206
207
           X.aug.new <- lapply(1:k, function(i) cbind(1,aug.x(x[,-1],unlist(psi.new[[i]]),psi[i</pre>
                ,],delta[[i]])))
208
           glm.out2=lapply(1:perm.ind, function(j) lapply(1:k, function(i) glm(y~X.aug.new[[i
                ]][,-1], weights=z[,perms[j,i]],family = quasibinomial(),control=list(maxit=500))
                ))
209
           beta.new <- lapply(1:perm.ind, function(j) lapply(glm.out2[[j]],coef))</pre>
210
            null.perms <- sapply(1:perm.ind,function(i) all(!is.na(lapply(beta.new,unlist)[[i]])</pre>
                ))
211
            null.beta=0
212
213
           if(sum(null.perms)>0){
214
              xbeta.new <- lapply(1:perm.ind, function(j) lapply(1:k, function(i) X.aug.new[[i]]</pre>
                    %*% matrix(beta.new[[j]][[i]],ncol=1)))
215
              \texttt{comp} \gets \texttt{lapply}(\texttt{1:perm.ind}, \; \texttt{function(j)} \; \; \texttt{lapply}(\texttt{1:k}, \; \texttt{function(i)} \; \; \texttt{dbinom(y}, \; \texttt{size} = \texttt{1}, \\
```

```
prob = logit(xbeta.new[[j]][[i]]))*lambda.new[i]))
216
             comp <- lapply(1:perm.ind, function(j) sapply(comp[[j]], cbind))</pre>
217
             compsum <- lapply(1:perm.ind, function(j) apply(comp[[j]], 1, sum))</pre>
218
             newobsloglik <- sapply(1:perm.ind, function(j) sum(log(compsum[[j]])))</pre>
219
             newobsloglik[c(1-null.perms)] = -Inf
220
             IND <- which.max(newobsloglik)</pre>
221
222
             z = z[,perms[IND,]]
223
             z2 = z2[,perms[IND,]]
224
             lambda.new <- apply(z2, 2, mean)</pre>
225
             lambda <- lambda.new</pre>
226
             beta <- beta.new[[IND]]</pre>
227
             xbeta <- xbeta.new[[IND]]</pre>
228
             xbeta2<-sapply(1:k, function(i) xbeta[[i]])</pre>
229
             X.aug <- X.aug.old
230
             psi.locs <- psi.new
231
             newobsloglik <- newobsloglik[IND]</pre>
232
233
           else{
234
             newobsloglik=Inf
235
             null.beta=1
236
237
238
239
         if(((newobsloglik-obsloglik)<(-epsilon) & !is.na(newobsloglik))|null.beta==1){</pre>
240
         glm.out1=lapply(1:perm.ind, function(j) lapply(1:k, function(i) glm(y~X.aug.old[[i
             ]][,-1],weights=z[,perms[j,i]],family = quasibinomial(),control=list(maxit=500))))
241
         beta.new <- lapply(1:perm.ind, function(j) lapply(glm.out1[[j]],coef))</pre>
242
         null.perms <- sapply(1:perm.ind,function(i) all(!is.na(lapply(beta.new,unlist)[[i]])))</pre>
243
244
         if(sum(null.perms)>0){
245
           xbeta.new <- lapply(1:perm.ind, function(j) lapply(1:k, function(i) X.aug.old[[i]] %</pre>
                *% matrix(beta.new[[j]][[i]],ncol=1)))
246
           comp <- lapply(1:perm.ind, function(j) lapply(1:k, function(i) dbinom(y, size = 1,</pre>
                prob = logit(xbeta.new[[j]][[i]]))*lambda.new[i]))
247
           comp <- lapply(1:perm.ind, function(j) sapply(comp[[j]], cbind))</pre>
248
           compsum <- lapply(1:perm.ind, function(j) apply(comp[[j]], 1, sum))</pre>
249
           newobsloglik <- sapply(1:perm.ind, function(j) sum(log(compsum[[j]])))</pre>
250
           newobsloglik[c(1-null.perms)]=-Inf
251
           IND <- which.max(newobsloglik)</pre>
252
```

```
253
           z = z[,perms[IND,]]
254
           z2 = z2[,perms[IND,]]
255
           lambda.new <- apply(z2, 2, mean)</pre>
256
           lambda <- lambda.new</pre>
257
258
           beta <- beta.new[[IND]]</pre>
259
           xbeta <- xbeta.new[[IND]]</pre>
260
           xbeta2<- sapply(1:k, function(i) xbeta[[i]])</pre>
261
           X.aug <- X.aug.old
262
           psi.locs <- psi.locs.old
263
           newobsloglik <- newobsloglik[IND]</pre>
264
         } else{
265
           newobsloglik=Inf
266
           sing=2
267
268
       }
269
270
       if((newobsloglik-obsloglik)<(-epsilon) & !is.na(newobsloglik) & abs(newobsloglik) != Inf
           ) {
271
272
         z2 < -z2.old
273
         z \leftarrow z.old
274
275
         lambda <- apply(z2, 2, mean)</pre>
276
         X.aug.1 \leftarrow lapply(1:k, function(i) cbind(1,aug.x(x[,-1],unlist(psi.locs.old[[i]]),psi[
              i,],delta[[i]])))
277
         glm.out.old < -lapply(1:k, function(i) <math>glm(y^X.aug.1[[i]][,-1], weights=z[,i], family = 0
              quasibinomial(),control=list(maxit=500)))
278
         beta <- lapply(glm.out.old,coef)</pre>
279
         xbeta <- lapply(1:k, function(i) X.aug.1[[i]] %*% matrix(beta[[i]],ncol=1))</pre>
280
         xbeta2<- sapply(1:k, function(i) xbeta[[i]])</pre>
281
         psi.locs <- psi.locs.old
282
         \verb|comp| <- lapply(1:k, function(i) dbinom(y, size = 1, prob = logit(xbeta[[i]]))*lambda[i]| \\
             ])
283
         comp <- sapply(comp, cbind)</pre>
284
         compsum <- apply(comp, 1, sum)</pre>
285
         newobsloglik <- sum(log(compsum))</pre>
286
287
         if(is.na(newobsloglik)){
288
           sing <- 3
289
         } else{
```

```
290
          if((newobsloglik-obsloglik)<(-epsilon)) sing <- 4</pre>
291
        }
292
       }
293
       }
294
295
      if (sing > 0 \mid | is.na(newobsloglik) \mid | abs(newobsloglik) == Inf) {
296
         cat("Need new starting values due to singularity...", "\n")
297
298
        restarts <- restarts + 1
299
        if (restarts > max.restarts) stop("Too many tries!")
300
       tmp.ind=1
301
        while(tmp.ind==1){
302
           if(sum(1-sapply(delta,is.null))>0) psi.temp=psi.locs
303
          tmp <- try(suppressWarnings(logitsegregmix.init(y=y, x=x.old, lambda = NULL, beta =</pre>
               NULL, k = k, seq.Z=seq.Z, psi=psi, psi.locs = NULL)), silent=TRUE)
304
        if(class(tmp)!="try-error") tmp.ind=2
305
306
      lambda <- tmp$lambda
307
308
      beta <- tmp$beta
309
      k <- tmp$k
310
      psi.locs <- tmp$psi.locs</pre>
311
      n <- length(y)
312
      diff <- 1
313
      iter <- 0
314
      X.aug \leftarrow lapply(1:k, function(i) cbind(1,aug.x(x[,-1],unlist(psi.locs[[i]]),psi[i,],
           delta[[i]])))
315
      xbeta <- lapply(1:k, function(i) X.aug[[i]] %*% matrix(beta[[i]],ncol=1))</pre>
316
      xbeta2<- sapply(1:k, function(i) xbeta[[i]])</pre>
317
      comp <- dbinom(y, size = 1, prob = logit(xbeta2))*matrix(lambda,n,k,byrow=T)</pre>
318
      obsloglik <- sum(log(apply(comp, 1, sum)))</pre>
319
      ll <- obsloglik
320
321
       else {
322
       diff <- newobsloglik - obsloglik
323
       obs<-obsloglik
324
       obsloglik <- newobsloglik
325
       ll <- c(ll, obsloglik)</pre>
326
       X.aug.old <- X.aug
327
        psi.locs.old <- psi.locs
328
         iter <- iter + 1
```

```
329
330
                              if (verb) {
331
                                      cat("iteration =", iter, "newobsloglik =", newobsloglik, "obsloglik =", obs, "diff =
                                                      ", diff, "log-likelihood =", obsloglik, "\n")
332
                                     }
333
                        }
334
                         }
335
336
                       if (iter == maxit) {
337
                              warning("Maximum number of iterations reached.", call. = FALSE)
338
339
                       if (iter == 1) {
340
                              cat("Converged in 1 iteration. Consider rerunning with different starting values or
                                               smaller stopping criterion.", "\n")
341
342
                        cat("number of iterations=", iter, "\n")
343
344
                        names(delta) <- c(paste("comp", ".", 1:k, sep = ""))</pre>
345
                        names(seg.Z) <- c(paste("comp", ".", 1:k, sep = ""))</pre>
346
                        names(psi.locs) <- c(paste("comp", ".", 1:k, sep = ""))</pre>
347
                        names(beta) <- c(paste("comp", ".", 1:k, sep = ""))</pre>
348
                        for(i in 1:k){
349
                           names(beta[[i]])[1]="(Intercept)"
350
                            names (beta[[i]]) [2:ncol(x)]=colnames (x) [-1]
351
                              if(!is.null(psi.locs[[i]])){
352
                                    for(j in 1:ncol(psi)){
353
                                             if(psi[i,j]>0 & j==1){
354
                                                    names (beta[[i]]) [(ncol(x)+1): (ncol(x)+cumsum(psi[i,])[j])] = c(paste(colnames(x)[j])) = c(paste(c
                                                                    +1], ".", 1:psi[i,j], sep = ""))
355
                                             } else if (psi[i,j]>0) names (beta[[i]])[(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[j-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i,])[i-1]+1):(ncol(x)+cumsum(psi[i
                                                             cumsum(psi[i,])[j])]=c(paste(colnames(x)[j+1], ".", 1:psi[i,j], sep = ""))
356
357
358
359
                        colnames(z) <- c(paste("comp", ".", 1:k, sep = ""))</pre>
360
                               a = list(x = x, y = y, lambda = lambda, beta = beta, seg.Z = seg.Z, psi.locs = psi.
                                              locs, delta = delta, loglik = obsloglik, posterior = z, all.loglik = 11,
361
                            restarts = restarts, ft = "replogitsegregmixEM")
362
                       class(a) = "mixEM"
363
364 }
```

./ replogits egregmix EM.r

CHAPTER IV

CONCLUSION

The dissertation examines two distinct problems related to "marketing communication dynamics". The main goal of this line of research is to help firms provide individually tailored marketing contents to their customers. In these two essays, I develop statistical models to first understand customers' responses, and then explore methods to optimize firms' reactions accordingly. Essay 1 examines "scale attraction effects" in a charitable donation context, introducing novel constructs ("compliance degree", "pulling amount", "accumulated pulling amount") to describe attraction effects for multi-point appeals scales. The proposed model jointly accounts for donation incidence and amount using a Tobit 2 formulation, and allows heterogeneity in seasonality and pulling effects. Results suggest substantial scale attraction effects that vary across donors, stronger "pulling down" than "pulling up", and heterogeneous seasonal donation patterns. A significantly negative error correlation between donation incidence and donation amount underscores the importance of accounting for selectivity effects. The effects of individually tailoring appeals scales is demonstrated through simulation. Essay 2 investigates mate-seeking users' decision rules in an online dating context. I develop an empirical two-stage mate choice model that can accommodate compensatory and non-compensatory decision rules in each of two stages: browsing and writing. A mixture of logits model with changepoints allows for distinct decision rules across stages and heterogeneity in rule use across site users. Most importantly, it allows us to identify and compare attribute-level decision rules ("deal-breakers" and "deal-makers") over the two stages. Results suggest the existence of heterogeneity in decision rules across (1) genders, (2) stages, and (3) site users. Additionally, it suggests the existence of potential deal-breakers/makers across both discrete and continuous attributes.

Each essay opens up several opportunities for future research. Relevant to Essay 1, since charities find it difficult to attract first-time donors, it is important that they receive ongoing and relatively large donations from existing donors. Therefore, a key issue is that of "optimal laddering": how much charities should increase amounts subsequently requested of individual donors, based on their donation histories. This entails not only how much to ask each time, but also how often to make requests (and thereby the interval between solicitations) becomes a focal issue in maximize donation amounts without repelling existing donors. A substantive but clearly under-researched topic related to Essay 2 is fashioning a "dynamic dyadic choice model" in a two-sided matching context. In two-sided matching, unlike in a traditional choice context (where customers can choose items "unilaterally"), items can - and eventually must, for a match to be made also choose customers back. Therefore decision makers might behave strategically by anticipating other decision makers' utilities. One might expect that decision makers' update their decision rules according to one anothers responses, which might be revealing in regard to strategic behavior. Identifying such behavior, as well as changes in decision rules over time, would add substantially to the current state of understanding about choice and preference dynamics in a broad class of network-based contexts.

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