# Bayesian Estimation of Isotopic Age Differences ${ }^{1}$ 

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#### Abstract

Isotopic dating is subject to uncertainties arising from counting statistics and experimental errors. These uncertainties are additive when an isotopic age difference is calculated. If large, they can lead to no significant age difference by "classical" statistics. In many cases, relative ages are known because of stratigraphic order or other clues. Such information can be used to establish a Bayes estimate of age difference which will include prior knowledge of age order. Age measurement errors are assumed to be log-normal and a noninformative but constrained bivariate prior for two true ages in known order is adopted. True-age ratio is distributed as a truncated log-normal variate. Its expected value gives an age-ratio estimate, and its variance provides credible intervals. Bayesian estimates of ages are different and in correct order even if measured ages are identical or reversed in order. For example, age measurements on two samples might both yield 100 ka with coefficients of variation of 0.2 . Bayesian estimates are 22.7 ka for age difference with a $75 \%$ credible interval of $[4.4,43$. 7] ka .


KEY WORDS: Bayes estimate, dating, credible interval, stalagmite, log-normal.

## INTRODUCTION

Many methods exist for dating rocks and sediments, among which those using radioactive isotopes are prominent. All methods are subject to errors arising from procedural and measurement imprecisions. Classic statistical methods for estimating age differences and their confidence intervals suffice when errors are small and age differences are large. However, if errors are large and ages are similar, classic methods lead to accepting the null hypothesis of zero age difference.

In some cases, age order is known. An important example, which was a primary motivation for this work, is dating of stalagmites by ${ }^{230} \mathrm{Th}^{234} \mathrm{U}$ disequilibrium measurements (e.g., Latham et al., 1986). Stratigraphically, lower samples are known a priori to be older than upper, regardless of date order found from isotope measurements. Bayesian statistical methods permit impos-

[^0]ing such information and estimating properly ordered age differences from even reversed-order data.

Bayesian methods and interpretations used here were adopted from Guttman et al. (1982).

## DISTRIBUTION OF DATING ERRORS

A measured isotopic age $x_{i}$ is assumed to be a sample from a log-normal distribution, with error $\log$-variance $\sigma_{i}^{2}$. Thus

$$
\begin{equation*}
\ln x_{i} \sim N\left(\ln a_{i}, \sigma_{i}^{2}\right) \tag{1}
\end{equation*}
$$

where $a_{1}$ is true age. Standard deviation $\sigma_{i}$ may be estimated from dating statistics as the coefficient of variation of measured age.

Assuming log-normality assures that measured ages, and confidence limits, are non-negative. This usually will be true when variance arises primarily from one major error source such as counting statistics.

Two measured ages $x_{1}$ and $x_{2}$ are assumed independent (dating procedures have no covariance) with a joint probability density function (p.d.f.)

$$
\begin{equation*}
p\left(x_{1}, x_{2} \mid a_{1}, a_{2}\right)=p\left(x_{1} \mid a_{1}\right) p\left(x_{2} \mid a_{2}\right) \tag{2}
\end{equation*}
$$

where true ages are defined $a_{2}>a_{1}>0$. Bayes' theorem gives a posterior p.d.f. for $\left(a_{1}, a_{2}\right)$ as

$$
\begin{equation*}
p\left(a_{1}, a_{2} \mid x_{1}, x_{2}\right)=\frac{p\left(x_{1}, x_{2} \mid a_{1}, a_{2}\right) p\left(a_{1}, a_{2}\right)}{p\left(x_{1}, x_{2}\right)} \tag{3}
\end{equation*}
$$

From the p.d.f. for a log-normal distribution (Eq. 1)

$$
\begin{equation*}
p\left(x_{1}, x_{2} \mid a_{1}, a_{2}\right) \propto \frac{1}{x_{1} x_{2}} \exp \left\{-\frac{1}{2}\left[\left(\frac{1}{\sigma_{1}} \ln \frac{x_{1}}{a_{1}}\right)^{2}+\left(\frac{1}{\sigma_{2}} \ln \frac{x_{2}}{a_{2}}\right)^{2}\right]\right\} \tag{4}
\end{equation*}
$$

## BAYESIAN PRIOR

A noninformative but constrained prior p.d.f. is chosen to be

$$
\begin{align*}
p\left(\ln a_{1}, \ln a_{2}\right) & =k \quad \text { or }  \tag{5}\\
p\left(a_{1}, a_{2}\right) & =\frac{k}{a_{1} a_{2}} \tag{6}
\end{align*}
$$

for $a_{2}>a_{1}>0$, and zero otherwise. This forces non-negativity and the known age order but little more (it has infinite variance). Also, $x_{1}$ and $x_{2}$, and hence
$p\left(x_{1}, x_{2}\right)$, are constants for a given pair of age measurements. From Eqs. 3, 4, and 6

$$
\begin{equation*}
p\left(a_{1}, a_{2} \mid x_{1}, x_{2}\right) \propto \frac{1}{a_{1} a_{2}} \exp \left\{-\frac{1}{2}\left[\left(\frac{1}{\sigma_{1}} \ln \frac{x_{1}}{a_{1}}\right)^{2}+\left(\frac{1}{\sigma_{2}} \ln \frac{x_{2}}{a_{2}}\right)^{2}\right]\right\} \tag{7}
\end{equation*}
$$

## CONSTRAINED BAYES ESTIMATES

Conveniently, $a_{1}$ and $a_{2}$ may be estimated by means of transformed variables

$$
\begin{align*}
& u=\frac{1}{2^{1 / 2}} \ln \left(\frac{a_{2}}{a_{1}}\right)  \tag{8a}\\
& v=\frac{1}{2^{1 / 2}} \ln \left(a_{1} a_{2}\right) \tag{8b}
\end{align*}
$$

whence Eq. 7 becomes

$$
\begin{align*}
p\left(u, v \mid x_{1}, x_{2}\right) \propto & \exp \left\{-\frac{1}{2}\left[\left(\frac{v-u-2^{1 / 2} \ln x_{1}}{2^{1 / 2} \sigma_{1}}\right)^{2}\right.\right.  \tag{9}\\
& \left.\left.+\left(\frac{v+u-2^{1 / 2} \ln x_{2}}{2^{1 / 2} \sigma_{2}}\right)^{2}\right]\right\}
\end{align*}
$$

from which (by integration and normalization)
$p(u)=\exp \left[-\frac{\left[u-\frac{1}{2^{1 / 2}} \ln \left(\frac{x_{2}}{x_{1}}\right)\right]^{2}}{\left(\sigma_{1}^{2}+\sigma_{2}^{2}\right)}\right] /\left[\pi\left(\sigma_{1}^{2}+\sigma_{2}^{2}\right)\right]^{1 / 2} \Phi(\alpha) \quad u>0$
a truncated normal distribution, where

$$
\begin{equation*}
\alpha=\frac{\ln \left(x_{2} / x_{1}\right)}{\left(\sigma_{1}^{2}+\sigma_{2}^{2}\right)^{1 / 2}} \tag{11}
\end{equation*}
$$

and $\Phi(\alpha)$ is the cumulative unit normal distribution function. Expected values for $u$ and $v,\langle u\rangle$ and $\langle v\rangle$, from Eq. 10, are

$$
\begin{equation*}
\langle u\rangle=\frac{1}{2^{1 / 2}} \ln \left(\frac{x_{2}}{x_{1}}\right)+\frac{\left(\sigma_{1}^{2}+\sigma_{2}^{2}\right)^{1 / 2} \exp \left(-\frac{1}{2} \alpha^{2}\right)}{2 \pi^{1 / 2} \Phi(\alpha)} \tag{12a}
\end{equation*}
$$

$$
\begin{equation*}
\langle v\rangle=\frac{1}{2^{1 / 2}} \ln \left(x_{1} x_{2}\right)+\frac{\left(\sigma_{2}^{2}-\sigma_{1}^{2}\right) \exp \left(-\frac{1}{2} \alpha^{2}\right)}{2\left[\pi\left(\sigma_{1}^{2}+\sigma_{2}^{2}\right)\right]^{1 / 2} \Phi(\alpha)} \tag{12b}
\end{equation*}
$$

Bayes age estimates $\hat{a}_{1}$ and $\hat{a}_{2}$ are obtained from Eq. 8 as

$$
\begin{align*}
& \hat{a}_{1}=\exp \left(\frac{\langle v\rangle-\langle u\rangle}{2^{1 / 2}}\right)  \tag{13a}\\
& \hat{a}_{2}=\exp \left(\frac{\langle v\rangle+\langle u\rangle}{2^{1 / 2}}\right) \tag{13b}
\end{align*}
$$

Credible intervals for age ratios are estimated from

$$
\begin{equation*}
\operatorname{Pr}(\underline{u}<\hat{u}<\bar{u})=1-\beta \tag{14}
\end{equation*}
$$

where $\underline{u}$ and $\bar{u}$, lower ( $1-\beta / 2$ ) and upper $(\beta / 2)$ percentage points for Eq. 10 , are given by

$$
\begin{align*}
\Phi(z) & =1-(1-\gamma) \Phi(\alpha) \quad \text { where }  \tag{15}\\
z & =\frac{\langle u\rangle-\left(1 / 2^{1 / 2}\right) \ln \left(x_{2} / x_{1}\right)}{\left[\frac{1}{2}\left(\sigma_{1}^{2}+\sigma_{2}^{2}\right)\right]^{1 / 2}} \tag{16}
\end{align*}
$$

and $\gamma=\beta / 2$ and $1-\beta / 2$ for $\underline{u}$ and $\bar{u}$, respectively. Credible limits for $\hat{a}_{2}-$ $\hat{a}_{1}$ are calculated from the limits $(\underline{u}, \bar{u})$ with $\langle v\rangle$ and Eq. 13. Variance in $\hat{a}_{2}$ - $\hat{a}_{1}$ comes primarily from variance in $u$, and much less by variance in $v$, and therefore a joint credible region is unnecessary.

## APPLICATIONS

Equations 11, 12, and 13 use four parameters $\left(x_{1}, x_{2}, \sigma_{1}, \sigma_{2}\right)$ determining two estimates ( $\hat{a}_{1}, \hat{a}_{2}$ ), obscuring the dependence of the $a_{i}$, and their difference, on the data. The general behavior is shown (Fig. 1) for $\sigma_{1}=\sigma_{2}=\sigma$.

Quantity $u$, and hence $\hat{a}_{2} / \hat{a}_{1}$, is always greater than 1.0 , even if $x_{2} / x_{1}<$ 1.0, agreeing with known age order. As expected, $\hat{a}_{2} / \hat{a}_{1}$ tends to $x_{2} / x_{1}$ as $x_{2} / x_{1}$ increases or $\sigma$ decreases. Age estimates also yield $\hat{a}_{1}=x_{1}$ if $\sigma_{1}=0$, and $\hat{a}_{2}=$ $x_{2}$ if $\sigma_{2}=0$.

Example 1 (Table 1) shows data for a stalagmite dated by Harmon and Curl (1978a,b). Two-sigma standard errors about measured ages overlapped, reducing the significance of classic age difference estimates [as pointed out by Gascoyne, 1978)]. A Bayesian estimate for age difference is positive with positive credible limits.

Example 2 is from Latham et al. (1986) who regressed point age estimates

Fig. 1. Dependence of Bayes point and $75 \%$ credible-limit age-ratio estimates on measured age ratios for equal coefficients of variation.

versus position in a stalagmite, implicitly accepting age-order despite overlap of all adjacent (and many more distant) standard-error intervals. Despite large uncertainty in $x_{1}, \hat{a}_{2}-\hat{a}_{1}$ is nearly "classical." Credible interval limits are positive.

Examples 3 to 7 are hypothetical cases to illustrate effects of relative $x_{1}$ and $x_{2}$, and $\sigma$ for $\sigma_{1}=\sigma_{2}$. Examples 3 and 4 both have $x_{2} / x_{1}=2$, for small or large $\sigma$. Estimated age differences are classical. Examples 5 and 6 have $x_{2}$

Table 1. Examples of Bayesian Estimates of Age Differences

| No. | $i$ | $X_{i}$ | $\sigma_{i}$ | $\hat{a}_{1}, \mathrm{ka}$ | $\hat{a}_{2}-\hat{a}_{1}, \mathrm{ka}$ | $75 \%$ C.I., ka |
| :--- | :---: | :---: | :---: | :---: | :---: | :---: |
| 1 | 1 | 126.0 | 0.209 | 124.0 |  |  |
|  | 2 | 207.0 | 0.241 | 212.0 | 88.0 | $[33.0,147]$. |
| 2 | 1 | 0.045 | 2.30 | 0.024 |  |  |
|  | 2 | 0.525 | 0.64 | 0.550 | 0.526 | $[0.11,1.75]$ |
| 3 | 1 | 100.0 | 0.02 | 100.0 |  |  |
|  | 2 | 200.0 | 0.02 | 200.0 | 100.0 | $[95.2,105]$. |
| 4 | 1 | 100.0 | 0.20 | 99.7 |  |  |
|  | 2 | 200.0 | 0.20 | 200.6 | 100.0 | $[53.5,151]$. |
| 5 | 1 | 100.0 | 0.02 | 101.1 |  |  |
|  | 2 | 100.0 | 0.02 | 98.9 | 2.2 | $[0.40,4.40]$ |
|  | 1 | 100.0 | 0.20 | 112.0 |  |  |
| 7 | 2 | 100.0 | 0.20 | 89.3 | 22.6 | $[4.40,43.7]$ |
|  | 1 | 100.0 | 0.20 | 82.5 |  |  |
|  | 2 | 80.0 | 0.20 | 96.9 | 14.4 | $[2.50,29.0]$ |

$=x_{1}$, and example 7 has reversed age measurements ( $x_{2}<x_{1}$ ). Estimates $\hat{a}_{1}$ and $\hat{a}_{2}$ are ordered, $\hat{a}_{2}-\hat{a}_{1}$ are positive, and $75 \%$ credible intervals are large.

## DISCUSSION

At first, estimating ordered dates from reversed measurements appears startling. However, Bayesian estimates have only reasonable and desirable properties. These include:

1. $\hat{a}_{2}>\hat{a}_{1}: \hat{a}_{1}=\hat{a}_{2}$ cannot be accepted.
2. Classic estimates result for small $\sigma_{i}$ or large $x_{2} / x_{1}$.
3. Credible interval estimates have positive limits, but reflect increased uncertainty when $x_{2}<x_{1}$.

Of course, no guard exists against reversed age measurements resulting from systematic errors in dating. To be properly interpreted, Bayesian estimates msut be based on data subject only to (approximately log-normal) imprecision. However, systematic errors do not lead to contradictory ( $\hat{a}_{2}<\hat{a}_{1}$ ) results, but only to large credible intervals.

In view of these properties of Bayesian estimates of age difference, no reason exists for accepting a null hypothesis that $a_{1}=a_{2}$ when it is known, a priori, to be false.

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