Fibrosis Progression in Chronic Hepatitis C: Morphometric Image Analysis in the HALT-C Trial

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Computer-assisted morphometry can provide precise measurement of hepatic fibrosis on a continuous scale. Previous morphometric studies of large cohorts of patients with treatment refractory chronic hepatitis C have shown a mean increase in fibrosis of 30% to 58% in 1 year. The aim of the present study was to quantify fibrosis progression in biopsy specimens obtained over 1.5 to 5 years from three groups of patients with baseline bridging fibrosis or cirrhosis (Ishak stages 3-6) enrolled in the Hepatitis C Antiviral Long-term Treatment Against Cirrhosis Trial. The main group of 346 lead-in nonresponders (viremic after 24 weeks of peginterferon-ribavirin therapy) had a mean fibrosis increase of 61% over pretreatment baseline after 2 years and 80% after 4 years. In contrast, the 78 breakthrough/relapse patients (undetectable serum hepatitis C virus RNA after 24 weeks of peginterferon-ribavirin and receiving antiviral therapy for 48 weeks) showed a mean increase in fibrosis of 48% when biopsied 36 months from pretreatment baseline but no further increase at 60 months. Finally, the 111 express patients with baseline biopsies following unsuccessful peginterferon-ribavirin outside the trial had significantly more baseline fibrosis than the others but an increase of only 21% after 21 months and a slight decrease at 45 months. Maintenance therapy with low-dose peginterferon had no effect on fibrosis changes in any of the groups. Conclusion: Morphometry demonstrated complex, nonlinear changes in fibrosis over time in this heterogeneous cohort of patients with interferon-refractory chronic hepatitis C. (HEPATOLOGY 2009;50:1738-1749.)

Progression of hepatic fibrosis is the mechanism by which chronic hepatitis C leads to cirrhosis and decompensated liver disease. Consequently, interest is high in developing therapeutic options for patients who are not candidates for antiviral treatment as well as those who do not achieve a sustained virological response to antiviral therapy; these patients remain at risk for con-

Received April 22, 2009; accepted July 28, 2009.

tinuing fibrosis progression and the clinical consequences of hepatic decompensation and hepatocellular carcinoma. 1-3 Because chronic hepatitis C takes decades to evolve into cirrhosis, and even longer to progress toward decompensation, demonstration of survival benefit resulting from antiviral therapy would require large treatment trials of very long duration; therefore, treatment benefit in

Abbreviations: CI, confidence interval; HALT-C Trial, Hepatitis C Antiviral Long-term Treatment Against Cirrhosis Trial; HCV, hepatitis C virus; IFN, interferon. From the ¹Armed Forces Institute of Pathology, Division of Hepatic Pathology and Veterans Administration Special Reference Laboratory for Pathology, Washington, DC; ²New England Research Institutes, Watertown, MA; the ³Departments of Medicine and Molecular & Structural Biology and The Liver-Biliary-Pancreatic Center, University of Connecticut Health Center, Farmington, CT; the ⁴Department of Internal Medicine, University of Michigan Medical Center, Ann Arbor, MI; the ⁵Liver Diseases Branch and ⁷Office of the Director, National Institute of Diabetes and Digestive and Kidney Diseases, National Institutes of Health, Department of Health and Human Services, Bethesda, MD; the ⁶Division of Gastroenterology, University of California Irvine, Irvine, CA; the ⁸Gastroenterology Service, VA Long Beach Healthcare System, Long Beach, CA; the ⁹Department of Pathology and Immunology, Washington University, St. Louis, MO; the ¹⁰Laboratory of Pathology, National Cancer Institute, National Institutes of Health, Department of Health and Human Services, Bethesda, MD; the ¹¹Hepatology Section, Virginia Commonwealth University Medical Center, Richmond, VA; the ¹²Section of Hepatology, Division of Gastroenterology and Hepatology, Department of Internal Medicine, University of Colorado Denver, Anschutz Medical Campus, Aurora, CO; the ¹³Division of Gastrointestinal and Liver Diseases, Keck School of Medicine, University of Southern California, Los Angeles, CA; the ¹⁴Gastrointestinal Unit (Medical Services), Massachusetts General Hospital, Boston, MA; the ¹⁵Department of Medicine, Harvard Medical School, Boston, MA; and the ¹⁶Virology Division, Department of Laboratory Medicine, University of Washington, Seattle, WA.

Supported by the National Institute of Diabetes and Digestive and Kidney Diseases, the National Institute of Allergy and Infectious Diseases, the National Cancer Institute, the National Center for Minority Health and Health Disparities, the General Clinical Research Center, and Clinical and Translational Science Center grants from the National Center for Research Resources, National Institutes of Health. Additional support was provided by Hoffmann-La Roche, Inc., through a Cooperative Research and Development Agreement with the National Institutes of Health. (See Appendix for contract and grant details.)

The opinions and assertions expressed in this study are the private views of the authors and do not represent the views of the National Center for Research Resources, the National Institutes of Health, the Department of the Army, or the Department of Defense.

many clinical trials has been demonstrated reliably by assessing fibrosis stage in paired liver biopsies obtained before and after a course of therapy.

In most studies that have relied upon liver biopsy to evaluate changes in fibrosis, semiquantitative scoring systems have been used that have a limited range of categories of severity, such as the Metavir score⁴ (five stages) or the Ishak score⁵ (seven stages). An alternative to semiquantitative fibrosis scores is direct measurement of the amount of fibrosis in the biopsy specimen by computerassisted morphometric image analysis. In contrast to semiquantitative fibrosis scores that rely on quantity and location of, and architectural distortion caused by fibrosis, morphometry determines the proportion of collagenous tissue in a specimen, regardless of location, and provides precise measurements on a continuous scale. Within any given semiquantitative stage, a wide range of collagen may be present, and considerable overlap exists within Metavir and Ishak stages.^{6,7} Accurate morphometric studies, however, require high-quality, unfragmented liver biopsy specimens of sufficient size, because fibrous tissue is underrepresented in small, fragmented specimens of fibrotic livers. Morphometry is also subject to sampling variability,8 but when used to show differences between treatment groups in an adequately powered study, morphometry has the potential to be the most sensitive means of showing changes in the quantity of fibrosis.⁷

Even though clinical and histologic progression is usually very slow, in two previous morphometric analyses of antifibrotic agents tested in randomized controlled trials, investigators found that the actual quantity of fibrous tissue increased at a rapid rate in patients with treatment-resistant chronic hepatitis C. In one study of 245 patients with advanced fibrosis and cirrhosis (Ishak stage 4-6) treated with interferon (IFN)-γ1b or placebo, the mean collagen content of the liver biopsies increased by 58% in only 48 weeks.⁷ Similarly, in a trial of the peroxisome proliferator-activated receptor agonist antifibrotic agent

farglitazar compared with placebo in patients with moderate (Ishak stage 2-4) precirrhotic fibrosis, the mean hepatic collagen content increased by 30% in 52 weeks. Such changes may not have direct clinical significance, especially in patients with relatively little baseline fibrosis, but the ability to detect the change suggests that morphometry is a more useful tool than histologic staging for clinical trials of antifibrotic agents.

The Hepatitis C Antiviral Long-term Treatment Against Cirrhosis (HALT-C) Trial provided a unique opportunity to evaluate morphometry for assessing changes in liver fibrosis over several years in patients with chronic hepatitis C and moderate to advanced fibrosis. In the trial, we enrolled 1,050 prior nonresponders to pegylated IFN and ribavirin with compensated liver disease and bridging fibrosis or cirrhosis. Subjects were randomized to receive either weekly injections of 90 μg peginterferon alfa-2a or no therapy for 3.5 years. Over the course of the trial, we found no difference between the peginterferon-treated and nontreated groups in the development of clinical outcomes or in progression from precirrhotic fibrosis to cirrhosis on liver biopsy. 10 In this paper, we report results of histologic and morphometric analyses of fibrosis in liver biopsies obtained at baseline (up to 18 months before randomization), at approximately 18 months after randomization (study month 24), and at approximately 42 months after randomization (study month 48). We hypothesized that morphometric image analysis would demonstrate a linear increase in collagen content over time and might show an effect of maintenance therapy on fibrosis progression, even if standard histologic scoring did not.

Patients and Methods

The design and results of the HALT-C Trial have been reported.^{10,11} To be eligible for entry into the trial, patients at 10 United States Medical Centers had to be

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Published online in Wiley InterScience (www.interscience.wiley.com).

DOI 10.1002/hep.23211

Potential conflict of interest: Dr. Goodman received grants from Schering-Plough, and Pharmasset. Dr. Bonkovskyi is a consultant for and advises Boehringer-Ingelheim and Novartis. He also advises and is on the speakers' bureau of Lunbeck Pharma. He received grants from Hoffman-LaRoche, Merck, and Vertex. Dr. Fontana is on the speaker's bureau of Roche. Dr. Lindsay received grants from Schering-Plough, Valeant, Vertex, and Wyeth. She is also a consultant for and advises Bristol-Myers Squibb, Clinical Care Options, Kendle International, King, Peregrine, Roche Diagnostics, Valeant, Zymogenetics. She received grants from, is a consultant for, and advises, Hoffman-LaRoche, Human Genome Sciences, and Idenix Pharmaceuticals. Dr. Morgan is a consultant for, is on the speaker's bureau of, and received grants from Roche. He advises and received grants from Vertex. He received grants from Schering-Plough and March. Dr. Shiffman advises and received grants from Biolex, Conatus, Human Genome Sciences, Romark, Valeant, Vertex, and Zymogenetics. He advises Bristol-Myers Squibb, Anadys and Novartis. He received grants from GlaxoSmith Kline, Globeimmune, Idenix, Johnson and Johnson, and Wyeth. He is a consultant for, advises, is on the speakers' bureau of, and received grants from Roche. He also advises, is on the speakers' bureau of, and received grants from Schering-Plough. Dr. Stoddard owns stock in Elan Corp., Johnson and Johnson, Stryker, Proctor and Gamble, and Bristol-Myers Squibb.

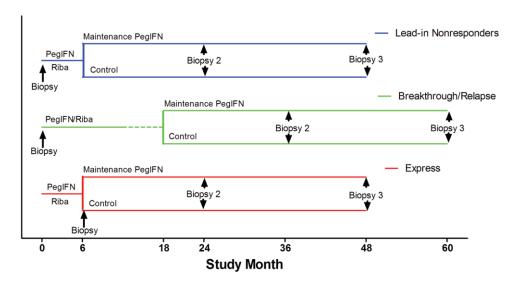


Fig. 1. Design of the HALT-C Trial, showing the source of the three cohorts of patients used in morphometric analyses of fibrosis. Lead-in nonresponder patients (blue lines) were those who remained viremic during the lead-in phase of peginterferon/ribavirin for 6 months and were then randomized to maintenance peginterferon or the untreated control group. These patients underwent baseline liver biopsies before the lead-in phase and approximately 24 and 48 months later. Breakthrough/relapse patients (green lines) developed undetectable HCV RNA during the lead-in phase of peginterferon/ribavirin and completed 12 months of therapy and 6 months of follow-up. Those who became viremic while still receiving therapy (breakthrough) or during follow-up (relapse) were randomized to maintenance or untreated control groups. These patients had baseline biopsies before the lead-in phase and approximately 36 and 60 months later. Express patients (red lines) received the equivalent of the lead-in phase outside the trial but remained viremic. Most of these patients had baseline biopsies immediately before randomization to maintenance peginterferon or the control group. Subsequent biopsies were at 18 and 42 months after randomization. PegIFN, peginterferon.

nonresponders to prior IFN therapy and had to have bridging fibrosis or cirrhosis on liver biopsy. Potential participants were excluded if they had another concurrent liver disease, a history of hepatic decompensation or hepatocellular carcinoma, any contraindication to IFN therapy, or any other medical or psychiatric condition that could interfere with their completion of the trial.

Recruitment for the trial began in August 2000 and ended in August 2004. Initially, all patients underwent a 24-week lead-in phase of treatment with combination, full-dose peginterferon alfa-2a and ribavirin. Patients who were still viremic after 20 weeks of treatment (lead-in nonresponders) were randomized to receive no further therapy (controls) or low-dose (90 µg/week) peginterferon (maintenance) for the next 3.5 years (Fig. 1). Patients with undetectable serum hepatitis C virus (HCV) RNA after 20 weeks of treatment were categorized as responders and continued to receive full-dose peginterferon and ribavirin for a total of 48 weeks. Those who were found thereafter to have detectable HCV RNA while receiving treatment (breakthrough patients) or after completing the 48 weeks of therapy (relapse patients) were eligible for randomization for the following 3.5 years (breakthrough/relapse patients). After combination peginterferon and ribavirin was approved by the U.S. Food and Drug Administration, patients who had received the equivalent of lead-in treatment outside the trial were also allowed to enter the randomized phase (express

patients). Of the 1,050 patients enrolled in the randomized phase of the study, 662 were lead-in nonresponders, 151 were breakthrough/relapsers, and 237 were express patients.

Liver biopsies were evaluated for necroinflammatory changes, fibrosis, steatosis, and other findings by consensus of a committee of 12 pathologists, at least 8 of whom were present at each scoring session. The Ishak fibrosis score⁵ was used as a primary endpoint in the trial. All patients were required to have a baseline liver biopsy showing bridging fibrosis (Ishak stages 3-4) or cirrhosis (Ishak stages 5-6) within 12 months of enrollment in the trial. Exceptions were allowed for patients with stage 2 (portal fibrosis) if a previous biopsy had shown unequivocal bridging fibrosis. For express subjects, baseline biopsies were obtained following unsuccessful peginterferonribavirin outside the trial. Repeat biopsies were performed at 18 months after entering the randomized phase (biopsy 2) and after another 24 months (biopsy 3) unless the patient refused, had a contraindication to biopsy, had a study-terminating clinical outcome, or had withdrawn from the trial.

Morphometric Image Analysis of Hepatic Fibrosis. We used liver biopsy sections stained with Sirius red for assessment of hepatic collagen content as described with Image Pro Plus 6.0 imaging software (Media Cybernetics, Silver Spring, MD) after editing of the digitized images to remove technical artifacts and extraneous tissue, such as

skin or muscle as well as Glisson's capsule or hepatic hilum if present. This method takes advantage of the fact that the degree of Sirius red staining measured by the degree of saturation of the red hue (-0.1 to + 0.04 of the)hue-saturation-intensity color model) correlates well with chemically determined collagen content and morphometrically determined estimates of the area of hepatic fibrosis. This approach relies on the analysis of identical brightfield images and images obtained with polarizing filters to subtract background mathematically at each pixel without interactive thresholding. The collagen content per unit area of the specimen is expressed in arbitrary units calculated as the sum of the pixel-wise collagen measurements divided by the number of summed pixels, corrected for the section thickness as measured by a confocal laser scanning microscope (Zeiss LSM 310) and multiplied by 5 (a constant) to yield a range of values very similar to those obtained by methods that measure the fractional area of fibrous tissue. The surface area of each specimen was computed as the number of pixels multiplied by the pixel size (8.2 \times 10⁻⁶ mm²). Because the method uses the degree of saturation of color to estimate collagen content, a pixel that is darker red (i.e., more saturated), has greater weight than one that is paler red, and although the values obtained do not have a physical dimension, the relative amount of collagen can be compared between specimens and used as a continuous function in statistical analyses. All collagen in the specimen is measured by this method, including the small amount normally present in portal tracts and central veins; therefore, biopsy specimens with relatively little abnormal fibrous tissue are not readily distinguished by this method. Similarly, because only collagen is measured, expansion of portal tracts by inflammation but not fibrosis does not affect the result. Reproducibility, assessed by 10 collagen measurements on the same slide, yielded a coefficient of variation of 7.1% for biopsy specimens with high collagen content (>0.1 U) and 12.1% for those with low collagen content (<0.05 U).

Selection of Biopsy Specimens for Morphometric Analysis. This dataset was limited to 535 HALT-C Trial patients with at least two adequate liver specimens suitable for computerized morphometry. Fragmented specimens (19% overall, 23% of baseline biopsies) were excluded because of inherent inaccuracy, and biopsy specimens with <10 mm² of tissue were excluded to reduce sampling variability. One hundred ninety-nine patients met the criteria for all three biopsy specimens; another 135 patients had adequate specimens at baseline and biopsy 2; an additional 136 patients had adequate specimens for biopsy 2 and biopsy 3 but not at baseline; and an additional 65 had adequate specimens for morphometry

at baseline and biopsy 3 but not in biopsy 2. Overall 1,269 specimens from 535 of the 1,050 randomized patients were used in the analyses of morphometric data. These included 1,267 needle biopsies and tissue from two explanted livers. The 1,267 needle biopsies had a mean \pm standard deviation length of 2.0 \pm 0.7 cm (median 1.9 cm) and a mean \pm standard deviation area of 17.5 \pm 5.8 mm². There were 324 (31%) that were less than 15 mm long. The sections from the two explant livers had 89 and 251 mm² of tissue.

Statistical Analysis. To evaluate the change in morphometric collagen over time, we used repeated-measures analysis of variance assuming an autoregressive covariance structure. Time (baseline, biopsy 2, biopsy 3), randomization stratum (lead-in nonresponders, breakthrough/relapse patients, express patients) and the development of a clinical outcome (yes/no) were included as fixed effects. Correlation was measured by way of Spearman rank correlation. Mann-Whitney and Kruskal-Wallis tests were used for assessing differences in patient characteristics between groups. Contingency tables were analyzed with the chi-square test. P values were two-sided. All analyses were performed with SAS version 9.1 (SAS Institute, Inc., Cary, NC) or Instat version 3.06 (Graphpad Software, San Diego, CA).

Results

Patient Population. The baseline characteristics of the 535 patients whose biopsy specimens were used for morphometric analysis were similar to those of the entire randomized cohort of the HALT-C Trial (Table 1). The mean age was 50, the majority were male (72%), and 95% were infected with HCV genotype 1. Baseline liver biopsy revealed cirrhosis in 182 (34%) of the 535 morphometry patients compared with 41% in the overall HALT-C Trial cohort, reflecting the fact that cirrhotic patients were more likely to have fragmented or small biopsies.

Among the 535 patients in whom morphometric analyses were performed, 346 (65%) were randomized at the end of the lead-in phase, 78 (15%) were breakthrough/relapse patients, and 111 (21%) were express patients. A noteworthy difference among these three randomization strata was in the interval between the baseline biopsy and biopsy 2 (Table 2). The majority of the lead-in nonresponder subgroup had the baseline liver biopsy shortly before the start of the lead-in phase, with a median of 6.3 months (range 5.2-17.5 months) between biopsy and randomization. Most of the breakthrough/relapse patients were also biopsied just before entering the lead-in phase, but nearly all had finished 48 weeks of combination therapy and 24 weeks of follow-up before being randomized;

Table 1. Characteristics of Patients with Two Consecutive Unfragmented Liver Biopsies Analyzed for Morphometry Compared with All Randomized Patients in the HALT-C Trial

	Analyzed for Morphometry (n = 535)	All HALT-C Patients (n = 1,050)
Age		
Mean ± SD	50 ± 7	50 ± 7
Range	19-77	19-80
Sex		
Male	72%	71%
Female	28%	29%
Race		
White	71%	72%
Black	21%	18%
Hispanic	5%	8%
Body mass index		
Mean ± SD	30 ± 5	30 ± 5
Range	18-56	18-58
Diabetes	24%	24%
Genotype 1	95%	94%
Log ₁₀ HCV RNA (mean)	6.5 ± 0.5	6.4 ± 0.5
Randomization stratum		
Lead-in nonresponder	65%	63%
Breakthrough/relapse	15%	14%
Express	21%	23%
Treatment group		
Control	51%	51%
Long-term peginterferon	49%	49%
Baseline liver biopsy	n = 399	n = 1,050
Fragmented	0	24%
Length (mean \pm SD)	$21 \pm 7 \text{ mm}$	18 ± 9 mm
Area (mean \pm SD)	$18\pm6~\text{mm}^2$	$14\pm 8~\text{mm}^2$
Inflammation score (Ishak)		
Mean	7.74 ± 1.91	7.54 ± 2.06
Range	3-12	1-12
Fibrosis score (Ishak)		
Stage 2	11%	7%
Stage 3	39%	34%
Stage 4	17%	18%
Stage 5	21%	22%
Stage 6	13%	19%
Steatosis (<5% of tissue)	39%	42%

therefore, the median time between baseline biopsy and entry into the randomized phase was 18.1 months (range 0.8-25.9 months). Finally, most of the express patients had baseline biopsies after they had failed peginterferon-ribavirin therapy, with a median of 1.5 months (range 0.3-19.1 months) from biopsy to randomization. For all patients, biopsy 2 was scheduled to be at 18 months after entering the randomized phase; therefore, for the lead-in

Table 3. Changes in Ishak Fibrosis Scores During the HALT-C Trial for the 535 Patients Used in Morphometric Analyses

	n	Improved	Same	Worsened
Baseline to biopsy 2	532	30%	43%	27%
IFN-treated	261	31%	40%	29%
Control	271	30%	44%	26%
Month 24 to biopsy 3	531	26%	41%	33%
IFN-treated	261	28%	40%	32%
Control	270	24%	42%	34%
Baseline to biopsy 3	533	32%	34%	34%
IFN-treated	262	33%	34%	33%
Control	271	31%	34%	35%

Three patients were missing biopsy 2 and two were missing biopsy 3. A score was considered improved if it decreased by one or more stages and worsened if it increased by one or more stages.

nonresponder patients, biopsy 2 was a median of 25 months from baseline, whereas for breakthrough/relapse patients and for express patients, biopsy 2 was obtained a median of 37 months and 21 months after baseline, respectively. Biopsy 3, on the other hand, was obtained at a median of 24 months following biopsy 2 for all patients.

Changes in Fibrosis Stage. All liver specimens from the 535 morphometry patients were scored for Ishak fibrosis stage by the pathology committee in blinded review sessions. Three patients had no biopsy 2, and two were missing biopsy 3, but all others had fibrosis stages assigned at each of the three time points. Overall, between baseline and biopsy 2, 30% of patients had an improvement (i.e., decrease) of 1 or more stages in Ishak fibrosis score, 43% remained the same, and 27% worsened, with an increase in stage (Table 3). Between biopsy 2 and biopsy 3, 26% improved, 41% remained the same, and 33% worsened. From baseline to biopsy 3, 32% improved, 34% remained the same, and 34% had worsening of Ishak fibrosis score. No significant differences emerged between patients assigned to the long-term peginterferon treatment arm and those assigned to the untreated control arm, but there were differences among the three randomization strata. The distribution of fibrosis scores at the three time points for each stratum and for the entire cohort is shown in Fig. 2. For the entire cohort, the number of patients with stage 3 fibrosis decreased, and the number with stages 2 and 1 increased, but the total proportion of patients with rela-

Table 2. Timing of Liver Biopsies for the Three Randomization Strata

	Lead-in onresponder $(n = 346)$	Breakthrough/Relapse (n = 78)	Express (n = 111)
Baseline to randomization, months	6.3 (5.2-17.5)	18.1 (0.8-25.9)	1.5 (0.3-19.1)
Baseline to biopsy 2, months	25.3 (22.6-40.5)	36.8 (19.7-48.9)	21.4 (14.2-40.5)
Biopsy 2 to biopsy 3, months	24.2 (15.7-37.9)	23.7 (13.1-32.3)	23.9 (14.0-29.6)

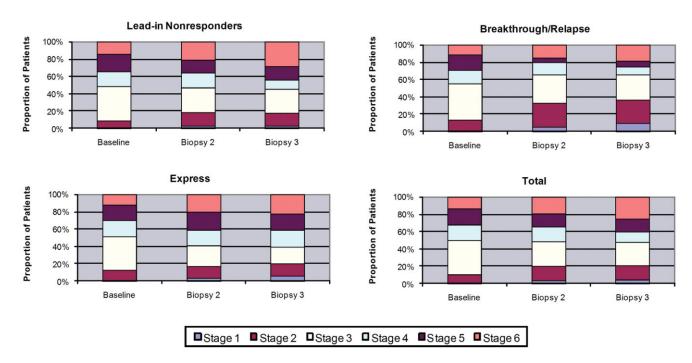


Fig. 2. Distribution of Ishak fibrosis stages at each biopsy among 535 patients used for morphometric analyses, showing an increasing proportion of patients with stage 6 cirrhosis (P < 0.0001). Panels show the three randomization strata with lead-in nonresponders (upper left [n = 346]), breakthrough/relapse (upper right [n = 78]), express (lower left [n = 111]), and the total cohort (lower right [n = 535]). There were no significant differences among the three strata at baseline. All three strata showed a significant increase in cirrhosis, but there were significantly fewer breakthrough/relapse patients with advanced stages at biopsy 2 (P = 0.01) and biopsy 3 (P = 0.004) than in the other two strata.

tively mild fibrosis (stages 1-3) remained constant (47%-49%), whereas patients with stage 6 (established cirrhosis) increased from 14% at baseline to 20% in biopsy 2 to 26% in biopsy 3, with a corresponding decrease in stages 4 and 5 at each time point. Closer inspection of the three randomization strata revealed an increasing proportion of low scores (stages 1-3) as well as stage 6 and a decreasing proportion of stages 4 and 5 in the breakthrough/relapse group, suggesting that some patients may have benefitted from partial virologic suppression that was achieved in this group. The lead-in nonresponders and express patients tended to have worsening fibrosis scores. Differences between breakthrough/relapse patients and the other two strata were significant at biopsy 2 (P = 0.01) and biopsy 3 (P = 0.0004).

Morphometry. Morphometric determination of collagen content was limited to the 1,269 unfragmented biopsies with at least 10 mm^2 of tissue, including 399 baseline, 470 biopsy 2, and 400 biopsy 3 specimens. Overall, a similar relationship and significant correlation occurred (Spearman r = 0.57) between fibrosis stage and collagen content as reported in other studies (6,7); however, considerable overlap was observed in the amount of collagen at the various stages (Fig. 3), indicating that, while correlated, morphometry, which quantifies the amount of fibrous tissue, and Ishak fibrosis score, which

relies primarily on location of fibrosis and architectural distortion, measure different things.

Using repeated measures analysis of variance, we found no differences in mean morphometrically determined collagen content over time between the peginterferontreated patients and the untreated controls (Fig. 4). Surprisingly, however, the mean collagen content increased by 49% between baseline and biopsy 2 (95% confidence interval [CI] 35%-63%) but then further increased only slightly to 54% above baseline at biopsy 3 (95% CI 41%-68%). Because we had anticipated that fibrosis progression would be more linear over time, we undertook additional analyses.

A potential explanation for the unexpected results in overall collagen change with time was found in the striking differences among the three randomization strata (Fig. 5). Patients randomized at the end of the lead-in phase behaved closest to expectation, with an increase in mean collagen by 61% over baseline (95% CI 47%-76%) in biopsy 2 (median 25.3 months after baseline) and 80% in biopsy 3 (95% CI 61%-99%; median 49.5 months after baseline), although the difference between biopsy 2 and biopsy 3 was not statistically significant. In contrast, the breakthrough/relapse patients had slightly less mean collagen at baseline, with an increase of 48% in biopsy 2 (95% CI 8%-89%; median 36.8 months after baseline),

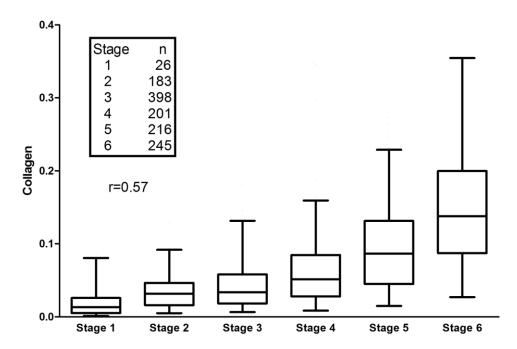


Fig. 3. Distribution of measurements of collagen content at each Ishak fibrosis stage for the 1,269 liver biopsies in the study. The boxes represent the median, 25^{th} , and 75^{th} percentiles; the lines represent the 2.5 and 97.5 percentiles (outliers not shown). Spearman correlation coefficient = 0.57 (P < 0.0001).

but no further increase in biopsy 3 (median 60.5 months after baseline). The express patients were quite different; their baseline biopsies followed an unsuccessful course of peginterferon/ribavirin, rather than preceding the lead-in course of therapy that the other two groups received. Compared with the lead-in nonresponder patients, express patients had a mean of 47% more collagen at baseline (P < 0.0001). Collagen in biopsies 2 and 3 were not significantly different from baseline levels, with an increase of only 21% in biopsy 2 (95% CI 1%-40%; median 21.4 months after baseline) and then a shift back to

the baseline value in biopsy 3 (median 45.3 months after baseline). The greater mean collagen content at baseline in the express patients was not explained by differences from the other groups in age, serum alanine aminotransferase, level of HCV RNA, alcohol history¹² or by inflammation or fat in the biopsy (Table 4). Within each of the three randomization strata, no significant differences occurred in mean collagen content at the various time points between those randomized to long-term peginterferon and the untreated controls (data not shown). The lead-in nonresponder patients had an increase in mean collagen

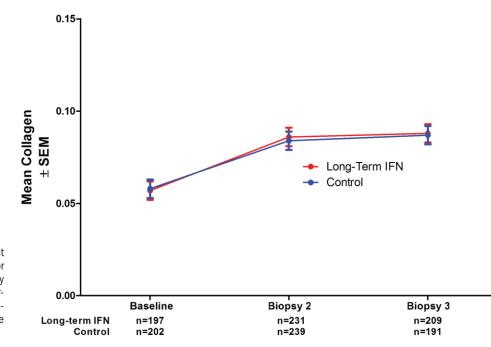
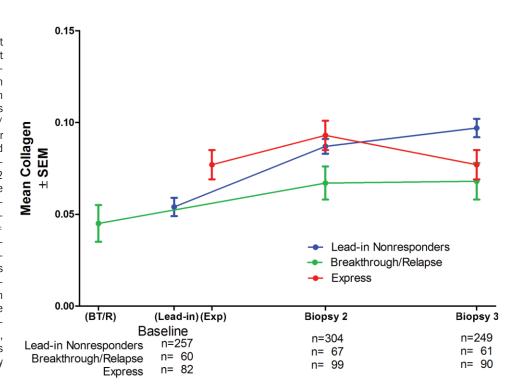


Fig. 4. Mean collagen content (\pm standard error of the mean) for the 535 patients in the study by treatment group, showing no difference between long-term IFN and untreated controls at any of the three time points.

Fig. 5. Mean collagen content (± standard error of the mean) at different time points for the 535 patients in the study by randomization strata. The median interval between baseline biopsy and biopsy 2 was 36.8 months for the breakthrough/ relapse patients, 25.3 months for the end of lead-in patients, and 21.4 months for the express patients. Mean collagen in biopsy 2 was significantly greater than in the baseline biopsy for lead-in nonresponders (P < 0.0001) and breakthrough/relapse patients (P =0.0029), but biopsy 3 was not statistically significantly greater than biopsy 2 for either group. Express patients had significantly more collagen in the baseline biopsy than lead-in and breakthrough/relapse patients (P < 0.0001), but the differences between baseline biopsy, biopsy 2, and biopsy 3 for express patients were not statistically significant.



of 20% to 30% per year, the breakthrough/relapse patients had an increase of 10% to 16% per year, and the express patients had an increase of 0% to 13% per year.

We also examined the possibility that patients with the most substantial fibrosis progression may have suffered clinical outcomes (death, ascites, variceal hemorrhage, spontaneous bacterial peritonitis, Child-Turcotte-Pugh score >7, eligibility for liver transplantation, or hepatocellular carcinoma)¹¹ and failed to have biopsy 3, resulting in a selection bias that could account for the apparent lack of progression between biopsy 2 and biopsy 3. In fact, however, all but one of the 535 patients in this analysis had a biopsy 3, even though only 400 were unfragmented with at least 10 mm² of tissue, a requirement for inclusion in this analysis. Next, we asked whether fragmented, small biopsies, which were excluded in this analysis, were more likely to occur in biopsy 3, another potential source of selection bias. Patients who suffered outcomes had a 119% increase in mean collagen between base-

line and biopsy 3 (95% CI 79%-159%) and significantly more mean collagen at each of the three time points (P =0.0137 at baseline; P = 0.0030 for biopsy 2; P = 0.0031 for biopsy 3) (Fig. 6) than patients without outcomes, even though only 19 patients with outcomes had biopsy 3 specimens adequate for morphometric analysis. Thus, patients with clinical outcomes had both the most fibrosis at baseline and the largest increase in fibrosis progression over time; however, because patients with outcomes had such elevated levels of fibrosis, they were less likely to have a biopsy suitable for morphometric analysis. Although the number of patients with clinical outcomes and fragmented/small biopsies is small, exclusion of their biopsy 3 specimens from morphometric analysis could have been partly responsible for the apparent deceleration in fibrosis between biopsy 2 and biopsy 3.

Finally, we examined the effect of baseline collagen on changes in morphometric fibrosis determinations (Fig. 7).

Table 4. Baseline Characteristics of the Three Randomization Strata

	Lead-in Nonresponder $(n = 346)$	Breakthrough/Relapse $(n = 78)$	Express (n = 111)	P Value
Age*	50.3 ± 7.6	48.7 ± 4.9	50.2 ± 6.4	0.21
Alanine aminotransferase*	110 ± 71	113 ± 94	101 ± 74	0.51
Log HCV RNA*	6.46 ± 0.47	6.58 ± 0.44	6.38 ± 0.59	0.02
Total lifetime alcoholic drinks (thousands)*12	15.5 ± 23.3	22.9 ± 43.0	17.7 ± 24.7	0.09
Histology at baseline				
Inflammation score ⁵	7.6 ± 1.9	7.3 ± 2.0	7.6 ± 2.0	0.40
Steatosis (<5% of tissue)	n = 145 (42%)	n = 22 (28%)	n = 43 (57%)	0.08

Values are expressed as the mean \pm standard deviation.

^{*}At randomization.

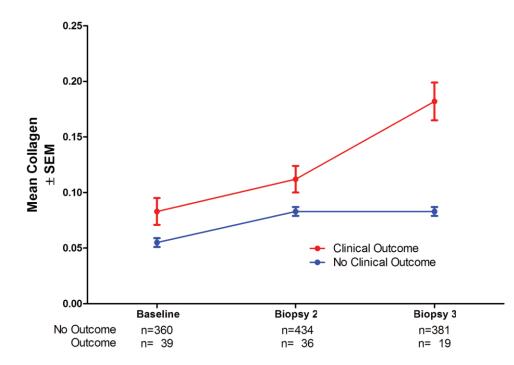


Fig. 6. Mean collagen content (\pm standard error of the mean) at different time points showing significantly more fibrosis at each time (baseline, P=0.0137; biopsy 2, P=0.0030; biopsy 3, P=0.0031) in the 42 patients who had clinical outcomes than in the 493 patients who did not have an outcome.

We chose ranges to approximate the highest (>0.055 units, n = 181), middle (0.024-0.055 units, n = 158), and lowest (<0.024 units, n = 185) third of baseline values. The mean collagen of the cohort with medium baseline collagen behaved closest to expected, increasing by 110% above baseline at biopsy 2 and 158% at biopsy 3. For those with low baseline collagen, the mean increased by approximately the same absolute value, but because the initial mean value was very low, the percentage increase was high (307%), but there was no further increase at biopsy 3. Finally, the mean of the cohort with

high collagen actually fell by 10% at biopsy 2, but remained the same at biopsy 3. This may be explained in part by loss of patients with the most collagen to clinical outcomes (Fig. 6), although a complete explanation remains elusive.

Discussion

In this study, we found that with morphometric image analysis, the most sensitive histologic method for measuring fibrosis, long-term low-dose maintenance peginterferon

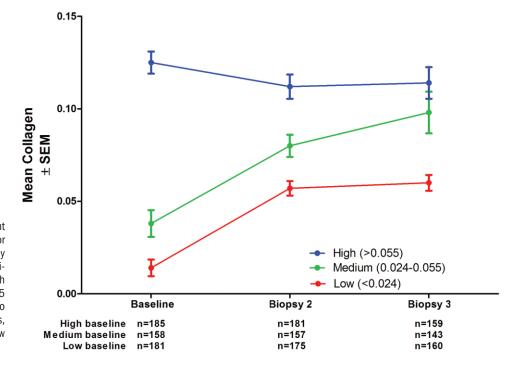


Fig. 7. Mean collagen content (\pm standard error of the mean) for the 535 patients in the study by baseline collagen levels (approximately one-third of patients in each group). The group with high (>.055 units) baseline collagen tended to have less in subsequent biopsies, whereas those with medium or low collagen increased.

therapy had no effect on fibrosis progression in this group of patients with chronic hepatitis C. Our data, however, did suggest that partial virologic suppression by full-dose peginterferon/ribavirin during the lead-in phase may retard fibrosis progression. Moreover, our observations raise the possibility that when full-dose peginterferon/ribavirin therapy is ineffective in producing virologic suppression, fibrosis progression may actually be accelerated.

In two previously completed short-term clinical trials in patients with chronic hepatitis C, collagen was found to increase at a relatively rapid rate that could be detected and quantified by morphometry in large cohorts. In neither of these trials did the study drug have any effect on changes in fibrosis; therefore, data from placebo-treated controls and patients receiving study drugs were combined for analyses of fibrosis progression. In one study, 245 patients with advanced fibrosis (Ishak stage 4-6) had a 58% increase in mean collagen in 1 year. In another, 209 patients with moderate (stage 2-4) fibrosis had a 30% increase in mean collagen of 30% in 1 year.

The HALT-C Trial provided the opportunity to study longer-term changes in fibrosis; however, because all patients received at least 6 months of ribavirin with full-dose peginterferon, the trial did not include a true placebotreated control cohort. Nevertheless, this study was anticipated to provide information about the long-term progression of fibrosis, but the overall results (Fig. 4) were surprising. The mean fibrosis of the cohort increased by nearly 50% between baseline and biopsy 2, similar to results reported in previous studies; thereafter, almost no change occurred over the next 2 years of the study. To some extent, we could explain these observations by invoking the differences in hepatic collagen results among the three randomization strata (lead-in nonresponder, breakthrough/relapse, and "express) (Fig. 5), and the inclusion of fewer biopsy 3 specimens resulting from the loss of patients with the most fibrosis to clinical outcomes (Fig. 6).

The three randomization strata differed from one another in three important ways. First, a considerable difference existed in the time interval between the baseline biopsy and biopsy 2, which averaged 25.3 months for the lead-in nonresponder group, 36.8 months for the breakthrough/relapse group, and 21.4 months for the express group. Second, the breakthrough/relapse patients achieved complete virologic suppression for several months, which did not occur in the other two groups. Third, the express patients underwent their biopsies after 6 months of full-dose peginterferon/ribavirin, whereas the other patients underwent baseline biopsies before receiving therapy.

The cohort of patients randomized at the end of lead-in therapy had fibrosis progression closest to that reported previously, with an increase of 80% in hepatic fibrosis over 4 years. Although this approximately 20% per year is less than the 30% to 58% per year observed in other studies, the current study is the first to include longterm follow-up biopsies. The fibrosis progression we observed in this group was not linear, however; the increase was greater during the first 2 years than during the second 2 years. This observation may have resulted in part from the loss of patients who had outcomes (Fig. 6). The breakthrough/relapse patients had documented virologic suppression for at least part of their treatment courses, which may have contributed to the accumulation of considerably less fibrosis, 50% over 5 years or 10% per year, and relatively fewer patients with cirrhosis (Fig. 2). Within the breakthrough/relapse cohort, however, no significant difference emerged between patients who received long-term low-dose peginterferon and untreated controls; therefore, the retardation they had in fibrosis progression appears to have resulted from the initial viral suppression rather than from maintenance therapy. Additionally, preliminary analyses of clinical outcomes and histologic improvement suggest that the impact of lead-in therapy was more substantial than the impact of maintenance therapy.¹³ Similar to the lead-in nonresponder cohort, the most substantial increase in mean fibrosis for the breakthrough/ relapse group occurred between the baseline biopsy and biopsy 2.

Fibrosis changes observed in the express patients are not readily explained but accounted for much of the apparent lack of fibrosis progression between biopsy 2 and biopsy 3 in the entire cohort (Fig. 4). Even though express subjects did not differ significantly in baseline Ishak fibrosis stages from that in the lead-in nonresponder stratum, mean fibrosis as determined by morphometry was 47% higher at baseline in the express group. Measured collagen levels in the express group and the lead-in nonresponder group were nearly the same in biopsy 2 but fell inexplicably thereafter in the express group. What distinguished the express group from the lead-in nonresponder group was the fact that the express group had baseline biopsies shortly (a median of 1.5 months) after failing to respond to treatment with peginterferon/ribavirin. This observation raises the possibility that ineffective peginterferon and ribavirin therapy might accelerate hepatic fibrosis transiently and might also explain why the patients randomized at the end of the lead-in phase (lead-in nonresponders) had a greater increase in fibrosis during the first 2 years (during which they received ineffective full-dose peginterferon and ribavirin therapy for the initial 6 months) than during the second 2-year interval. Cur-

rently, however, a possible role of peginterferon and ribavirin in promoting transient fibrosis progression remains speculative.

A limitation of the current study is the fact that fewer than half of the patients enrolled in the HALT-C Trial had liver biopsies suitable for morphometric analysis, raising the possibility of selection bias. This observation demonstrates the need for a standardized biopsy approach in studying changes in hepatic fibrosis. Noninvasive techniques may become available in the future but will need to be validated against the natural history of the disease and to be correlated with liver biopsy. If biopsies are to be used as a study endpoint or as a standard for validation of noninvasive markers, then the techniques and requirements for adequacy of biopsy specimens need to be standardized. For studies of fibrosis, specimens obtained with cutting biopsy needles are superior to those obtained with suction needles¹⁴; in addition, larger-gauge needles (14to 16-gauge) and biopsies at least 2 cm long are most likely to provide adequate specimens for assessing histology¹⁵ if this is to be used as a clinical trial endpoint and to correlate with noninvasive markers. Biopsy sampling error will always be present to some extent, so it must be emphasized that morphometry is not appropriate for evaluation of changes in individual patients but should be reserved for statistical analysis and comparison of changes in large cohorts.

The availability of serial liver biopsy specimens during the HALT-C Trial provided an opportunity, for the first time, to analyze fibrosis progression by way of quantitative image analysis over 3.5 to 5 years in a large number of patients with advanced chronic hepatitis C. Our findings show promise for this approach in future studies and clinical trials, but at present the natural history of fibrosis progression remains an area of great uncertainty. Therefore, in the future, carefully designed prospective long-term studies such as the HALT-C Trial will be needed to define the true natural history of fibrosis in chronic hepatitis C.

Appendix

The following individuals and institutions were instrumental in the planning, conduct, and/or care of patients enrolled in this study: Gyongyi Szabo, Barbara F. Banner, Maureen Cormier, and Donna Giansiracusa, University of Massachusetts Medical Center, Worcester, MA (contract N01-DK-9-2326); Gloria Borders and Michelle Kelley, University of Connecticut Health Center, Farmington, CT (grant M01RR-06192); Adrian M. Di Bisceglie, Bruce Bacon, Brent Neuschwander-Tetri, Elizabeth M. Brunt, and Debra King, Saint Louis University School of Medicine, St Louis, MO (contract N01-DK-9-2324); Raymond T.

Chung, Andrea E. Reid, Atul K. Bhan, Wallis A. Molchen, and Cara C. Gooch, Massachusetts General Hospital, Boston, MA (contract N01-DK-9-2319; grant M01RR-01066; grant 1 UL1 RR025758-01, Harvard Clinical and Translational Science Center); S. Russell Nash, Jennifer DeSanto, and Carol McKinley, University of Colorado School of Medicine, Denver, CO (contract N01-DK-9-2327; grant M01RR-00051; grant 1 UL1 RR 025780-01); John C. Hoefs, John R. Craig, M. Mazen Jamal, Muhammad Sheikh, and Choon Park, University of California Irvine, Irvine, CA (contract N01-DK-9-2320; grant M01RR-00827); William M. Lee, Thomas E. Rogers, Peter F. Malet, Janel Shelton, Nicole Crowder, Rivka Elbein, and Nancy Liston, University of Texas Southwestern Medical Center, Dallas, TX (contract N01-DK-9-2321; grant M01RR-00633; grant 1 UL1 RR024982-01, North and Central Texas Clinical and Translational Science Initiative); Sugantha Govindarajan, Carol B. Jones, and Susan L. Milstein, University of Southern California, Los Angeles, CA (contract N01-DK-9-2325; grant M01RR-00043); Anna S. Lok, Joel K. Greenson, Pamela A. Richtmyer, and R. Tess Bonham, University of Michigan Medical Center, Ann Arbor, MI (contract N01-DK-9-2323; grant M01RR-00042; grant 1 UL1 RR024986, Michigan Center for Clinical and Health Research); Richard K. Sterling, Melissa J. Contos, A. Scott Mills, Charlotte Hofmann, and Paula Smith, Virginia Commonwealth University Health System, Richmond, VA (contract N01-DK-9-2322; grant M01RR-00065); T. Jake Liang, David Kleiner, Yoon Park, Elenita Rivera, and Vanessa Haynes-Williams, Liver Diseases Branch, National Institute of Diabetes and Digestive and Kidney Diseases, National Institutes of Health, Bethesda, MD; James E. Everhart, Leonard B. Seeff, Patricia R. Robuck, Jay H. Hoofnagle, National Institute of Diabetes and Digestive and Kidney Diseases, Division of Digestive Diseases and Nutrition, Bethesda, MD; David R. Gretch, Minjun Chung Apodaca, Rohit Shankar, and Natalia Antonov, University of Washington, Seattle, WA (contract N01-DK-9-2318); Kristin K. Snow, Linda Massey, and Margaret C. Bell, New England Research Institutes, Watertown, MA (contract N01-DK-9-2328); Fanny Monge, and Michelle Parks, Armed Forces Institute of Pathology, Washington, DC; and the following Data and Safety Monitoring Board Members: Gary L. Davis (Chair), Guadalupe Garcia-Tsao, Michael Kutner, Stanley M. Lemon, and Robert P. Perrillo.

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