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Mass-Spectrometric ²³⁰Th-²³⁴U-²³⁸U Dating of the Devils Hole Calcite Vein

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The Devils Hole calcite vein contains a long-term climatic record, but requires accurate chronologic control for its interpretation. Mass-spectrometric U-series ages for samples from core DH-11 yielded ²³⁰Th ages with precisions ranging from less than 1,000 years (2σ) for samples younger than ~140 ka (thousands of years ago) to less than 50,000 years for the oldest samples (~566 ka). The ²³⁴U/²³⁸U ages could be determined to a precision of ~20,000 years for all ages. Calcite accumulated continuously from 566 ka until ~60 ka at an average rate of 0.7 millimeter per 10³ years. The precise agreement between replicate analyses and the concordance of the ²³⁰Th/²³⁸U and ²³⁴U/²³⁸U ages for the oldest samples indicate that the DH-11 samples were closed systems and validate the dating technique in general.

Core DH-11 of the Devils Hole (DH) calcite vein contains a continuous record of stable-isotopic variation in Great Basin ground water for most of the past several hundred thousand years (1). Spectral analyses of ¹⁸O and ¹³C records of DH-11, as well as inferences on timing of glacial-interglacial transitions (1), rely on the accuracy of the chronometric control. To provide this control, we determined mass-spectrometric (MS) uranium-series ages (3) for 21 samples across DH-11, using replicate analyses to validate estimates of uncertainty.

For each analysis, ~ 300 mg of calcite chips were selected for visual purity and freedom from porosity. The chips were ultrasonically cleaned, dissolved in HNO₃, spiked with ²²⁹Th, ²³³U, and ²³⁶U, and purified with conventional ion-exchange methods. The purified U and Th were

*To whom correspondence should be addressed. SCIENCE • VOL. 258 • 9 OCTOBER 1992 loaded with colloidal graphite on separate Re filaments (3) and analyzed in an automated mass spectrometer (4). To evaluate both the precision and accuracy of the dates, we examined possible sources of bias and external variance in the measurements (5) using instrumental checks (6) and two levels of replicate analyses. Thus, we have not relied on precision estimates arising solely from the internal statistics of the mass-spectrometric analyses.

The primary standard for this study was a solution of Precambrian uraninite that has been shown to be in secular equilibrium (7). Analyses of spiked aliquots of this standard were performed after every few samples and provide a test of the reproducibility of single analyses. The mean single-analysis precision of 230 Th/ 238 U for the secular-equilibrium standard was 0.24% (2σ , 12 analyses, no apparent external variance) (5, 8). For 234 U/ 235 U, a proxy for 234 U/ 238 U, the mean single-analysis precision was 0.31% (2σ , 29 analyses), including a resolvable external error of 0.25% (5, 9). Therefore, an additional variance corresponding to the 0.25% external error was added to the internal variance for each 234 U/ 235 U analysis. The weighted-mean

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²³⁴U/²³⁵U ratio of the secular-equilibrium standard was 0.0075690 ± 0.0000045, which corresponds to a ²³⁴U half-life of $245,290 \pm 140$ years (10). This value is in reasonable agreement with the half-life of 244,600 \pm 700 years determined from AS (alpha spectrometric) measurements (11). For age calculations, we normalized our measured ²³⁴U/²³⁵U and ²³⁰Th/²³⁵U ratios for DH-11 samples relative to the corresponding ratios measured for the secularequilibrium standard; this approach in effect results in ²³⁰Th/²³⁸U and ²³⁴U/²³⁸U activity ratios. Advantages of this procedure are that (i) systematic errors from spike miscalibration are eliminated, (ii) errors in calculated ages arising from uncertainties in the ²³⁴U and ²³⁰Th decay constants are demagnified, and (iii) systematic errors arising from uncorrected biases in MS measurement are largely canceled out.

In addition to replicate analyses of the secular-equilibrium standard, we evaluated the real precision of the analyses by replicating most DH-11 sample analyses, starting from a different set of calcite chips. Besides providing another check on the validity of the calculated errors for each analysis, the replicate analyses are also sensitive to small-scale, open-system histories



Fig. 1. Agreement between measured isotopic ratios for replicate DH-11 sample analyses. The *y*-axis indicates deviation, in percent, from weighted-mean value for each sample location. Height of vertical lines indicates 2σ analytical error for each replicate analysis (sample locations for each group of replicates expanded in *x*-direction for clarity: each clump of vertical lines represents replicate analyses for one sample location). Excess scatter of 230 Th/ 238 U age replicates of the youngest samples probably arises from non-reproducibility of sample locations (see text).

of the samples. Replicate $^{234}U/^{238}U$ analyses agreed within analytical error at all sample ages. Replicate $^{230}Th/^{238}U$ analyses agreed within analytical error for samples that formed before 130 ka, but not for those that formed at ~60 and ~122 ka (Table 1 and Fig. 1). This excess error arises largely from the uncertainty in the locations of these samples (~0.7 mm) and the thick-

nesses of the sample bands (1 and 3 mm, respectively). Thus, for samples having analytical age uncertainties less than or comparable to the errors predicted from the uncertainty in sample locations (that is, samples younger than ~ 200 ka), the actual scatter of replicate 230 Th/ 238 U ratios is expected to be somewhat greater than analytical error. These concerns are taken into

Table 1. Mass-spectrometric U-series ages (thousands of years) and calculated initial 234 U/ 238 U activity ratios of individual DH-11 samples versus distance from the free face of the vein (2σ errors). Calculated using half lives for 230 Th of 75,381 ± 590 and 234 U of 244,600 ± 490 years (2σ) (*11, 13*). Median U content of the samples was 0.46 ppm (range = 0.30 to 0.58), median 232 Th was 0.33 ppb (range <.02 to 16). The 230 Th ages are almost unaffected by corrections for initial 230 Th, 234 U, and 238 U, although a small correction corresponding to an initial 232 Th/ 238 U atomic ratio of 3.8 ± 2.0 (230 Th, 234 U, and 238 U are assumed to be in secular equilibrium) was applied. 234 U ages of samples <385 ka are not shown (all agree with the corresponding 230 Th ages) because they are not truly independent ages. Uncertainties in 234 U ages arise almost entirely from the ±0.10 uncertainty in initial 234 U/ 238 U ratio (Fig. 4).

Distance (mm)	²³⁰ Th age	²³⁴ U age	Initial ²³⁴ U/ ²³⁸ U _{act}	Observed ²³⁰ Th/ ²³² Th _{act}
Water			2.7693 ± 0.0066	
			2.7657 ± 0.0082	
1.0	61.5 ± 0.6		2.818 ± 0.010	104
	59.1 ± 0.5		2.816 ± 0.010	135
16.0	81.2 ± 0.7		2.787 ± 0.010	174
	80.2 ± 0.6		2.790 ± 0.010	215
	80.5 ± 0.5		2.792 ± 0.009	228
41.0	119.7 ± 1.0		2.666 ± 0.010	47,200
44.5	122.7 ± 0.8		2.656 ± 0.009	33,400
	121.1 ± 0.8		2.647 ± 0.010	33,800
	123.3 ± 1.0		2.649 ± 0.010	36,700
48.5	131.9 ± 1.2		2.674 ± 0.011	1,370
54.0	149.7 ± 2.5		2.737 ± 0.016	8,280
	149.8 ± 1.4		2.749 ± 0.012	6,630
58.5	156.0 ± 1.2		2.769 ± 0.012	3,710
	156.4 ± 1.5		2.767 ± 0.014	4,100
98.5	212.5 ± 2.9		2.762 ± 0.018	4,370
	209.8 ± 2.4		2.759 ± 0.017	5,460
111.0	234.5 ± 4.9		2.748 ± 0.028	12,900
	235.9 ± 3.8		2.752 ± 0.024	9,730
	236.9 ± 3.9		2.757 ± 0.023	11,200
126.0	266.8 ± 4.2		2.797 ± 0.025	1,010
	274.0 ± 6.6		2.832 ± 0.036	766
137.0	283.3 ± 7.1		2.796 ± 0.038	3,140
	284.1 ± 4.2		2.803 ± 0.025	2,870
	288.6 ± 4.5		2.812 ± 0.026	2,300
171.5	326.1 ± 8.7		2.712 ± 0.044	2,270
	328.7 ± 7.5		2.710 ± 0.039	2,320
188.0	349.2 ± 8.9		2.782 ± 0.047	16,800
	356.9 ± 8.6		2.823 ± 0.046	16,100
	362 ± 18		2.844 ± 0.093	15,500
219.0	390 ± 13	385 ± 20	2.774 ± 0.068	65,400
	401 ± 15	386 ± 20	2.821 ± 0.079	64,600
227.0	404 ± 14	411 ± 20	2.716 ± 0.069	31,300
	400 ± 13	411 ± 20	2.895 ± 0.064	35,000
250.0	439 ± 17	435 ± 20	2.773 ± 0.087	33,500
077.0	437 ± 18	437 ± 20	2.749 ± 0.091	36,400
277.0	$4/3 \pm 43$	459 ± 22	2.82 ± 0.23	71,200
	$4/4 \pm 31$	459 ± 20	2.82 ± 0.16	98,300
210 5	509 ± 34	463 ± 21	2.99 ± 0.19	76,700
318.5	507 ± 35	510 ± 20	2.74 ± 0.17	11,100
0045	555 ± 49	515 ± 20	2.96 ± 0.27	3,250
334.5	535 ± 44	532 ± 21	2.77 ± 0.22	294
0445	531 ± 47	531 ± 21	2.75 ± 0.23	415
344.5	$584 \pm /1$	547 ± 21 545 ± 20	2.94 ± 0.39	73,800
	637 + 192	543 ± 20 548 + 24	32 + 12	63 400
357.5	560 + 53	563 + 21	273 ± 0.26	3 330
007.0	630 ± 109	568 + 21	3.09 ± 0.65	710
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Fig. 2. Ages of DH-11 samples (weighted means of replicate analyses at each location; Table 2) versus distance from the free face of the core. Circles with vertical lines are ²³⁰Th ages and 2σ uncertainties (including effect of nonreproducibility of sample locations; expanded in inset for youngest samples); shaded area represents ²³⁴U ages (each at the same sample location as the ²³⁰Th ages), and vertical dimension of shaded area indicates ²³⁴U-age uncertainties.

Table 2. Weighted-mean ages (thousands of years) of DH-11 replicates (Table 1) versus distance from the free face of the vein. 230Th ages are given for locations from 1.0 mm through 250.0 mm and ²³⁴U ages are given for locations from 277.0 mm to 357.5 mm inclusive (because ²³⁴U ages for these older samples are more precise). Weighting and error-propagation procedure considered uncertainties resulting from location errors (along the core) of each sample and also uncertainties from location errors of replicates within each sample. Thus the (2σ) errors in this table represent the total uncertainty of the ages at these locations within DH-11. Systematic errors arising from half-life uncertainties (14) and biases in the HU-1 secular-equilibrium standard (15) are small and are not included in age errors.

Distance (mm)	Weighted mean age
1.0	60.3 ± 1.7
16.0	80.6 ± 2.5
41.0	119.7 ± 2.1
44.5	122.4 ± 2.6
48.5	131.9 ± 3.0
54.0	149.8 ± 2.6
58.5	156.2 ± 3.1
98.5	211.0 ± 2.9
111.0	235.9 ± 3.1
126.0	269.1 ± 4.2
137.0	285.7 ± 3.5
171.5	327.5 ± 6.4
188.0	354.1 ± 6.2
219.0	395 ± 10
227.0	402 ± 10
250.0	438 ± 13
277.0	461 ± 20
318.5	513 ± 20
334.5	531 ± 20
344.5	546 ± 20
357.5	566 ± 20



Fig. 3. Comparison of AS and weighted-mean MS analyses of DH-11. Jagged, heavy lines are MS analyses (line thickness defined by analytical uncertainties at each location); boxes are AS analyses (solid, $^{234}U/^{238}U$ activity ratios; open, $^{230}Th/^{234}U$ activity ratios; height of boxes indicates 2σ analytical uncertainty).



Fig. 4. Variation in weighted-mean initial 234 U/ 238 U activity ratios versus weighted-mean 230 Th ages of DH-11 samples (2σ uncertainties indicated by dimensions of ellipses). Horizontal dashed lines show approximate limits of variation of <350-ka samples, used to calculate uncertainties for 234 U ages of older samples.

account in Table 2, wherein the data are weighted to account for both analytical and sample-location errors, in Figs. 2 to 4, and in assigning ages to the 18 O and 13 C time series (1).

The ²³⁰Th ages for DH-11 increase monotonically and roughly linearly with distance in from the free face of the vein (Fig. 2 and Tables 1 and 2). All four of the samples having 230 Th ages >500 ka give meaningful ages (based on their consistency with the age-distance trend defined by the younger samples), and we thus conclude that even at an age of more than seven half-lives of ²³⁰Th, the ages remain reliable. The AS analyses done on \sim 5-g samples from DH-11, although less precise than the MS analyses, are in essential agreement (Fig. 3) and validate the earlier AS dates on core DH-2 (12). Except for the cessation of vein growth at ~60,000 ka, the age-distance trend of DH-11 (Fig. 2) confirms the

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Fig. 5. ²³⁴U-²³⁰Th isotopic evolution of DH-11. Curved lines are closed-system isotopic evolution for initial ²³⁴U/²³⁸U activity ratios of 2.6 and 2.9. Open circles are ages on these evolution curves. Solid circles are activity ratios for DH-11 samples (weighted mean of replicate analyses except for youngest sample). Analytical errors are generally smaller than symbol size.

petrographic observations (1, 2, 12) that there were no detectable hiatuses in growth during the previous 500,000 years.

The AS analyses of 61,000- to 296,000year-old samples from DH-2 (12) showed that, within the precision of the measurements, initial $^{234}U/^{238}U$ ratios that were back-calculated for these samples are indistinguishable from a mean value of about 2.70. Our analyses on DH-11 are consistent with this value, but, because of greater analytical precision, a secular variation in initial ²³⁴U/²³⁸U ratios can be readily resolved (Figs. 4 and 5). Weighted-mean initial ²³⁴U/²³⁸U ratios for the younger 360,000 years of DH-11 (where precision of the weighted-mean initial ratios is generally better than ± 0.03) range from 2.651 \pm 0.006 at 122.4 ka to 2.817 ± 0.007 at 60.3 ka. Uranium dissolved in present-day Devils Hole water has a ²³⁴U/²³⁸U activity ratio of 2.768 (weighted mean; Table 1), well within this range. Back-calculated initial ²³⁴U/²³⁸U activity ratios for the older DH-11 samples are necessarily less precise (because the ²³⁰Th ages are less precise), but show no resolvable variation outside the limits defined by the younger samples. We calculated ²³⁴U ages for DH-11 assuming that the average initial ²³⁴U/²³⁸U activity ratio was 2.75 ± 0.10 (Table 1 and Fig. 4). This value falls near the midpoint of the observed range, such that the uncertainty encompasses the extremes of the samples <360,000 years old. The age uncertainty (~20,000 years for all DH-11 samples) corresponding to this uncertainty in the initial²³⁴U/²³⁸U ratio is much larger than the 2σ uncertainty arising from analytical error alone (2000 to 3000 years). Moreover, because the error arising from the uncertain initial ²³⁴U/²³⁸U ratio is likely to be highly correlated for closely spaced samples (the time constant for significant fluc-

tuations in initial ²³⁴U/²³⁸U ratio is probably thousands of years), we are unable to take advantage of replicate analyses to diminish the ²³⁴U age uncertainties. Nonetheless, ²³⁴U ages are more precise than the corresponding ²³⁰Th ages for samples older than 450 ka and provide a useful check on the validity of the oldest ²³⁰Th ages. The agreement between ²³⁴U ages and ²³⁰Th ages (Table 1) for the samples that formed >350 ka (and especially for those that formed >500 ka), independently confirms the ²³⁰Th ages, and also shows that the growth rate of the vein for its first 100,000 years was similar to its long-term average growth rate.

Overall, the U-series ages form a remarkably self-consistent suite of age determinations. Because this consistency is both internal (from replicate samples) and external (from the stability of the overall agedistance trend), it seems highly unlikely that the dates have been significantly corrupted by open-system processes such as uranium gain or loss or alpha-recoil phenomena. The apparent ideality of the U-Th system in the vein material is probably the result of continuous submergence in water that showed limited secular variation of its physical and chemical properties (2).

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- Isotopic analyses were done with a VG-54E sin-4. gle-collector mass spectrometer using an analogue-mode Daly detector, controlled by the program ANALYST [K. R. Ludwig, U.S. Geol. Surv. Open-File Rep. 85-141 (1985)]. U was analyzed in automatic mode with $^{235}U^+$ beams of $\sim 5 \times 10^{-14}$ A (238 U/ 235 U ratio assumed to be 137.88); Th was typically analyzed with 230 Th⁺ beams of ~10⁻¹⁵
- 5. Internal variance is variance in isotopic ratios from within-run, MS measurement only. External vari-ance is variance in isotopic ratios not accounted for by the internal variance.
- 6. Amplifier nonlinearity was calibrated by measurement of fractionation-corrected 235U/238U ratio over an ion-beam range of 5×10^{-16} to 4×10^{-13} A. A slightly nonlinear response of the Daly detector was detected, with ~0.3% less gain at 4 × 10^{-13} A than at 5 × 10^{-16} A. This (softwarecorrected) change in gain was approximately linear with beam size, and corrections were much less than measurement precision. A regression of ²³⁴U/²³⁵U ratio versus ion-beam intensity for hundreds of blocks of standard runs shows no resolvable residual nonlinearity. Mass fractionation during measurements was corrected by normalizing to measured 233U/236U ratios (from the 1:1 ²³³U:²³⁶U spike). No fractionation correction was applied to ²³⁰Th/²²⁹Th or ²³²Th/²²⁹Th ratios, because the average effect on ²³⁰Th/²²⁹Th was canceled out by the similarly uncorrected spike calibrations, and fractionation corrections on the ²³⁰Th/²³²Th ratio were unimportant. Each Re filament was doped with graphite before use, heated to running temperature in the mass spectrometer, and examined for mass-spectral purity. Isobaric interferences were absent during data acquisi-

tion, and blank levels of 230Th were close to zero (average = $-0.8 \pm 4.0 \times 10^6$ atoms). Blanks for ²³⁸U were negligible (median = 1.2×10^{-11} g), although blanks for ²³²Th were significant (median = 4.6×10^{-12} g). Tails under the small peaks from nearby large peaks were insignificant (abundance sensitivity at 1 atomic mass unit offset was 2 to 4 ppm), as were amplifier time-constant effects

- Uraninite standard HU-1, obtained from M. Ivanovich, yielded U-Pb isotopic ages of 620 million years ago, concordant to within 0.5% (K. R. Ludwig, unpublished data). Alpha-spectrometric measurements yielded 234 U/ 238 U activity ratios of 0.9982 ± 0.0088 and 230 Th/ 234 U activity ratios within 0.5% of unity (J. N. Rosholt, personal communication to M. Ivanovich, 1979). The Precambrian age, concordance of U/Pb isotopic ages, and alpha-spectrometric measurements all indicate that the HU-1 standard is in, or very close to, secular equilibrium for ²³⁰Th/²³⁸U and ²³⁴U/²³⁸U ratios.
- 8. MSWD (mean square of weighted deviates) = 0.54

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- 14. For all samples, systematic age errors arising from uncertainty in the ²³⁰Th half-life (*13*) are smaller than those from analytical errors (because of the error-demagnification effect of normalizing to the isotopic ratios of the secular-equilibrium standard). For those samples where ²³⁰Th ages are preferred over 234U ages, the worst-case age error from ²³⁰Th decay-constant uncertainty is 1150 years (for the 235.9-ka samples at 111 mm), compared to the analytical error for that sample of 3100 years. Errors from 230Th decay-constant uncertainties for all other samples <400 ka are less; errors for 400- to 600-ka samples rise exponentially from ~0 to 10 ka. Age errors arising from uncertainty in the 234 U half-life (11) are negligible for all samples
- 15. For example, a 0.2% departure in the ²³⁰Th/²³⁸U ratio of the HU-1 standard from secular equilibrium would result in systematic 230Th age errors that are negligible for the samples <300 ka, and about half the analytical error for the samples with ages from 327.5 to 438 ka.
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Regulation of the Amount of Starch in Plant Tissues by ADP Glucose Pyrophosphorylase

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Starch, a major storage metabolite in plants, positively affects the agricultural yield of a number of crops. Its biosynthetic reactions use adenosine diphosphate glucose (ADPGIc) as a substrate; ADPGIc pyrophosphorylase, the enzyme involved in ADPGIc formation, is regulated by allosteric effectors. Evidence that this plastidial enzyme catalyzes a ratelimiting reaction in starch biosynthesis was derived by expression in plants of a gene that encodes a regulatory variant of this enzyme. Allosteric regulation was demonstrated to be the major physiological mechanism that controls starch biosynthesis. Thus, plant and bacterial systems for starch and glycogen biosynthesis are similar and distinct from yeast and mammalian systems, wherein glycogen synthase has been demonstrated to be the rate-limiting regulatory step.

The α -1,4 glucans (starch and glycogen) are the main storage carbohydrates in practically all living systems (1). In several crops, starch is a major component of the harvest and thus directly has an impact on yield. Within the last 10 years, the demand for starch has dramatically increased for both specialized food and industrial uses (2), primarily as a result of the development of high fructose corn syrups and bio-ethanol. A number of specialty starches (such as amylose and waxy starch) are being

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increasingly recognized for their superior material and nutritional properties as well as biodegradability. Understanding the critical components of the plant starch biosynthetic machinery therefore has a major impact on agriculture and industry. We have used transgenic plants to probe the rate-limiting step in starch biosynthesis.

Starch biosynthesis occurs in the plastids of plant cells, involving ADPGlc pyrophosphorylase (E.C. 2.7.7.27), starch synthase (E.C. 2.4.1.21), and branching enzyme (E.C. 2.4.1.18) (1, 3). In view of its sensitivity to allosteric effectors, ADPGlc pyrophosphorylase (ADPGPP) has been suggested to play a pivotal role in plant starch biosynthesis, as it is in the bacterial pathway for glycogen biosynthesis.

The Escherichia coli ADPGPP, encoded by the glgC gene (4), is a regulated ho-

⁹ MSWD = 2.0.

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