GSA DATA REPOSITORY 2011333

Humid periods in southern Arabia - windows of opportunity for modern human dispersal

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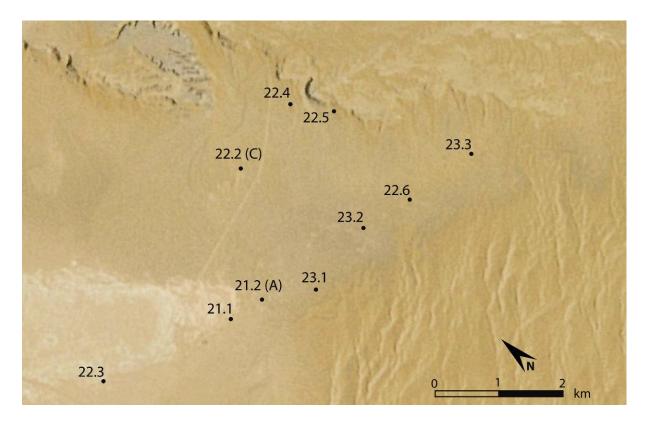
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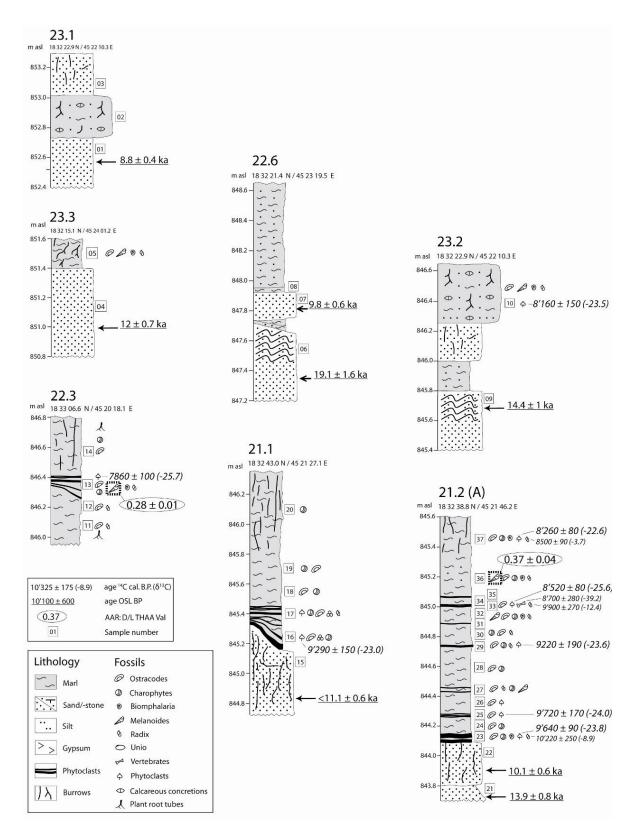
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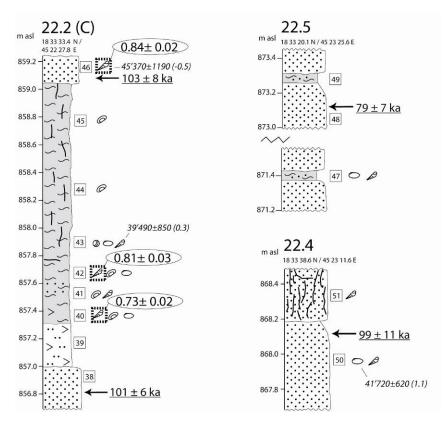
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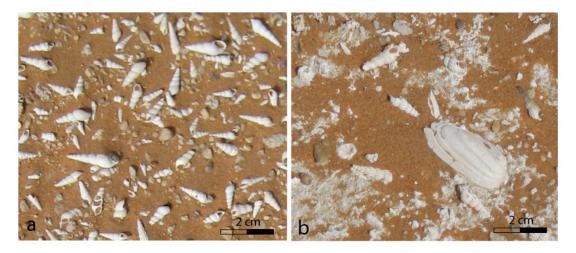
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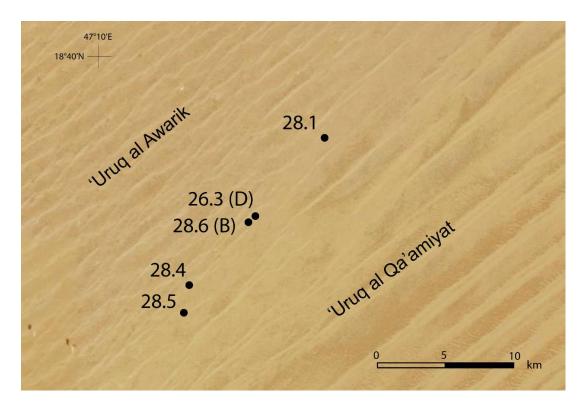
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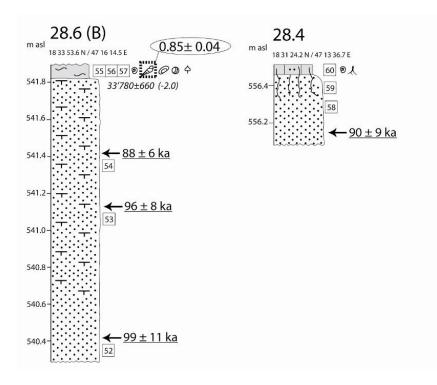
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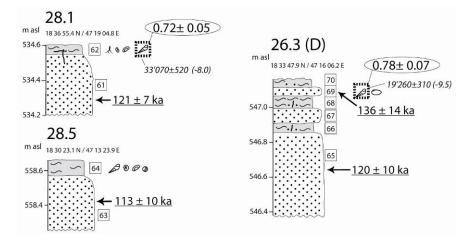
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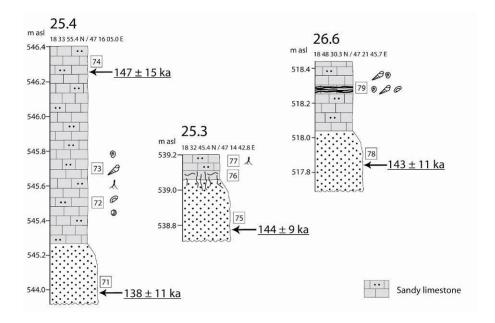
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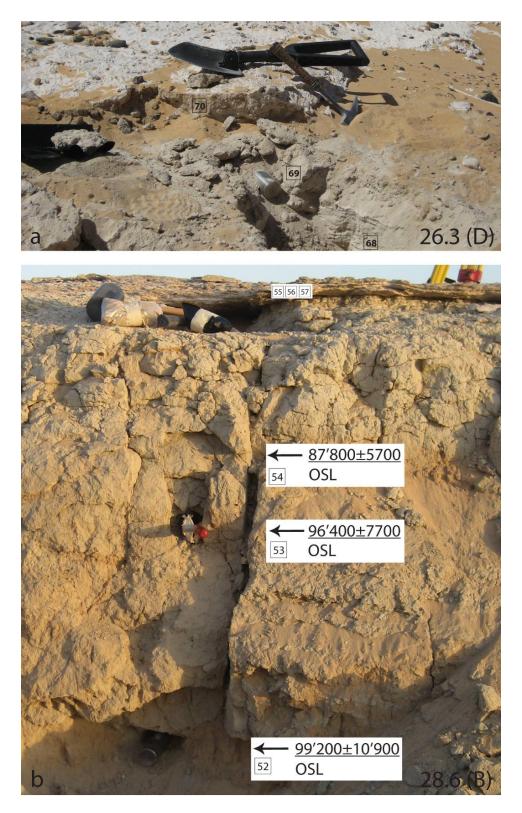
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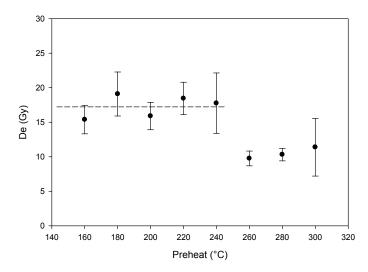
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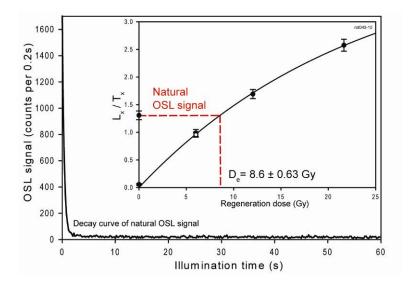
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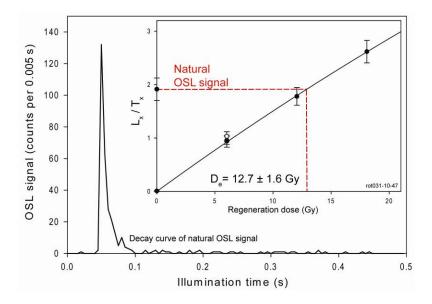
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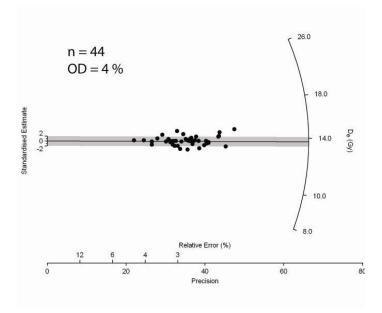
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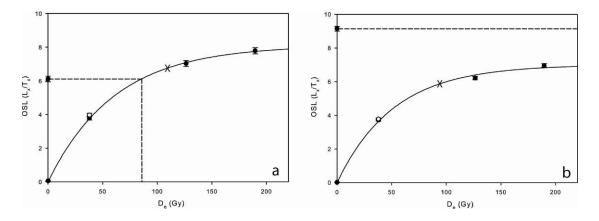
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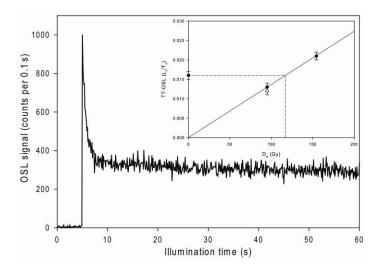
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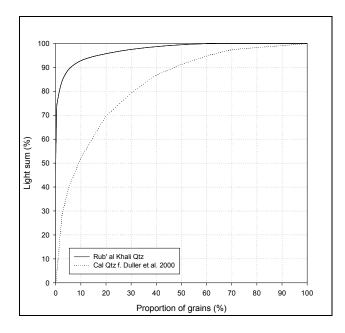
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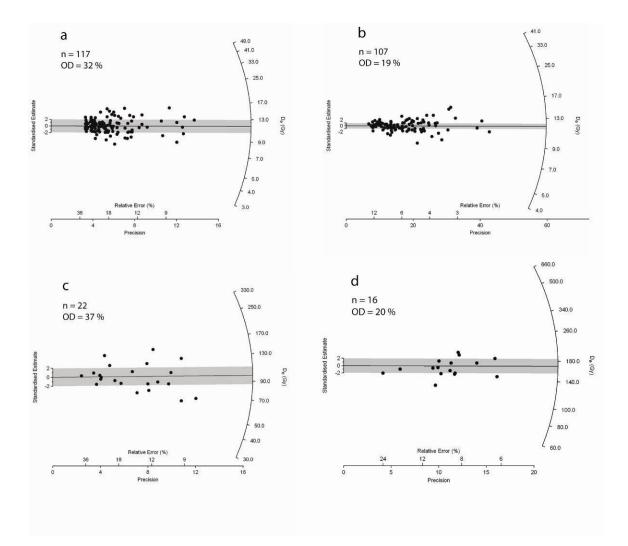
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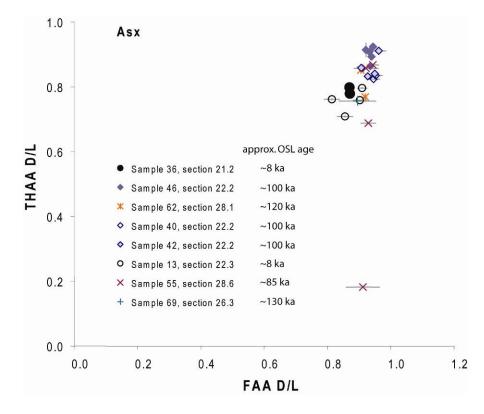
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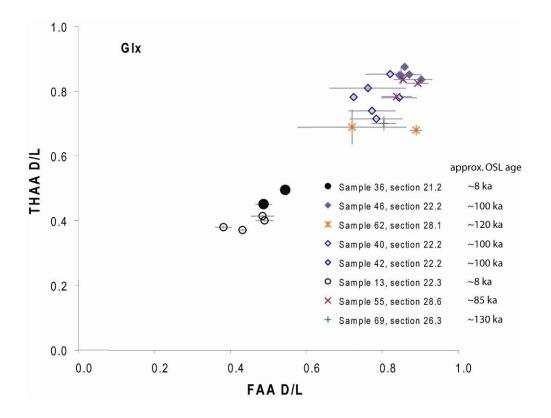
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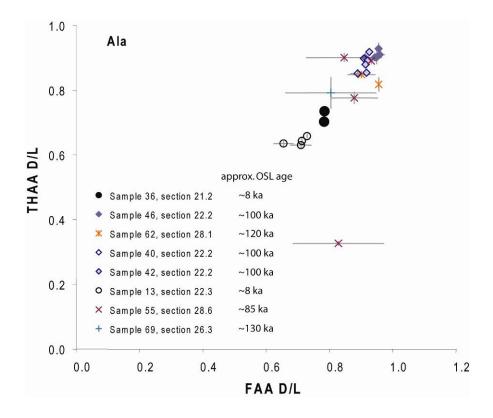
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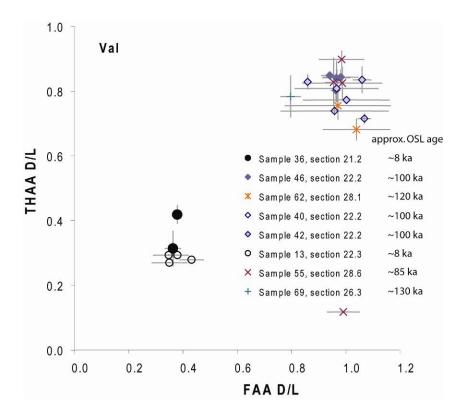
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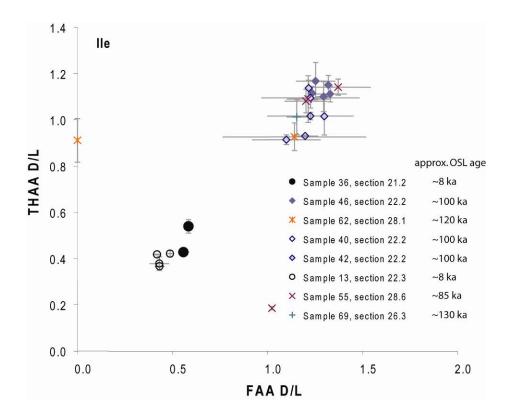
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2. Supplementary Methods

OSL Dating Method

Optically stimulated luminescence (OSL) dating is a method to determine burial ages of sediments. Energy supplied by naturally occurring ionizing radiation in the sediment increases the number of trapped electrons in the crystal lattice of certain mineral grains (e.g. quartz). Sunlight exposure releases light-sensitive trapped electrons. As a consequence, the intensity of the OSL signal is a measure of the radiation dose absorbed by the sediment grains since they were last exposed to daylight. For purposes of dating, the OSL signal is measured and the dose producing an equivalent signal is determined. The equivalent dose divided by the dose rate of the surrounding sediment gives the OSL age of a sample (Huntley et al., 1985).

Common problems when sampling paleolake deposits are limited sediment thickness, low quartz content and sometimes vertical inhomogeneities. The spatial and temporal (water table changes, bioturbation) homogeneity is important as it affects the dose rate, and a homogenous dose rate is generally an assumption for age estimations. As a consequence, our sampling strategy mainly relied on sampling homogenous sand beneath the lake deposits for a maximum age, combined with further samples within and/or above the lake deposits to narrow the age range of deposition. We collected samples by forcing a steel tube into a freshly cleaned exposure. The material from the tubes was then carefully discharged into opaque plastic bags for transport to the lab. Dose rate samples were collected with a small shovel from the same holes. Block samples of lithified lake deposits were wrapped with two layers of black opaque foil. Documentation of outcrops included a detailed stratigraphic sketch (Supplementary Fig. DR2-9).

All samples were opened in the laboratory under the appropriate lighting conditions (dim red/orange illumination). In the case of block samples the outer surface was removed carefully using a diamond saw with water cooling. All dating was carried out on quartz grains. The samples were treated with HCl and H_2O_2 to remove carbonates and organic matter before the grain size 200-250 µm was isolated. A two-step heavy-liquid separation using LST Fastfloat (2.70 and 2.58 g cm⁻³) was applied and the quartz fraction was subsequently etched in HF (40%) for 60 min to reduce remaining feldspars and the outer part of the quartz grains influenced by alpha irradiation during burial. After washing with HCl to dissolve fluorides, the samples were re-sieved at 150 µm to remove remaining feldspar grains. The coarse-grained quartz was then mounted on steel discs using silicone spray or placed in special single grain discs.

The **dose rate** represents the rate of ionizing radiation to which the samples were exposed during burial and is mostly derived from the radioactive decay of ²³⁸U, ²³⁵U, ²³²Th (plus their daughter products) and ⁴⁰K, with lesser contributions from cosmic rays. Radioactive elements (K, Th, U; Supplementary Table DR4) were measured by high-resolution gamma spectrometry

by S. Szidat, Department of Chemistry and Biochemistry, University of Bern (Preusser and Kasper, 2001). These measurements imply some evidence for radioactive disequilibrium in the Uranium decay chain for some of the samples and these have been additionally measured using Instrumental Neutron Activation Analysis (INAA) by J. Kučera, Nuclear Physics Institute Řež, Czech Republic. U-238 from INAA compared to Ra-226 from Gamma spectrometry (Supplementary Table DR4) shows limited U-238 deficiency with a negligible effect on the Pleistocene ages. For example, the age of sample 38 with extreme scenario using the values determined U-238 or Ra-226 differs not more than 3 %. Dose rates were calculated assuming average water content between 1% and 3% during burial. The contribution of cosmic rays to the total dose rate was calculated using present day depth considering full uncertainties due to actual position below surface (Prescott and Hutton, 1994). Total dose rates and resulting OSL ages are summarized in Supplementary Table DR4.

Two of the samples are rather problematic with regard to dose rate determination, as these have been taken from near surface layers. Although problematic, these samples are important as they directly date the lake deposits. Sample 74 (section 25.4) is an almost pure carbonate with little aeolian sand content. This sample shows a deficit of the mother isotope U in the decay chain and extreme low K and Th concentrations. As the layer is highly porous it appears that rain water may have reduced the concentration of radioactive elements after deposition. As a consequence, the resulting age is regarded as maximum estimate. The second problematic sample 69 (section 26.3) is from a thin aeolian sand layer bracketed by lake marls. Unfortunately, we could not measure the gamma radiation field on site, which leaves some uncertainties with regard to homogeneity. Additionally, this sample also shows low K and Th values and therefore we also regard this age as a maximum estimate.

Luminescence measurements were made in a Risø TL/OSL-DA-20 reader with blue diodes ($470 \pm 30 \text{ nm}$; ~ 41 mW cm^{-2}) and with a calibrated 40 mCi 90 Sr/ 90 Y beta source delivering ~ 0.13 Gy s^{-1} to the grains. Luminescence signals were measured with an EMI 9635Q photomultiplier tube through a 7.5 mm Hoya U340 filter. For optical stimulation of single sand-sized grains (SG) a laser attachment was used with a beam from a 10 mW Nd:YVO₄ diode-pumped green laser (532 nm) focused down to a 10 µm diameter spot with a power density of ~50 W/cm² (Bøtter-Jensen et al., 2000).

Early saturation of the OSL signals with $2D_0$ values between 90 - 150 Gy made it necessary to use two different protocols. Equivalent doses (D_e) were either determined with a standard SAR protocol for OSL (Murray and Wintle, 2000) or a SAR protocol for thermally transferred OSL (TT-OSL, Porat et al., 2009).

For the **Holocene samples** the single aliquot regenerative dose (SAR) protocol (Supplementary Table DR1 and DR2) of Murray and Wintle (2000) was applied to the samples

initially characterized using preheat plateau (Supplementary Fig. DR12) and dose recovery tests (Supplementary Fig. DR19).

For multi grain D_e determination a preheat of 220°C for 10 s was used and also applied to the test dose. For single grain D_e determination the conditions were the same except for the much shorter illumination of 0.5 s due to the high power density of the laser. The test dose (D_t) size was always 6 Gy. The dose response curve was constructed by plotting L_x/T_x versus dose, where L_x is the OSL signal from the regenerative dose and T_x is the response to the test dose. For multi grain the initial 0.4 s OSL signal minus a background estimated from the 50-60 s integral was used for OSL dose determinations while the much shorter intervals for single grains were 0.05-0.07 s for the signal minus a background from 0.4-0.5 s. Supplementary Figure DR13 and DR14 show representative decay and dose response curves. Aliquots were rejected if the recycling ratio was outside the range 1.0 ± 0.2 or the OSL IR depletion ratio (Duller et al., 2003) was less than 0.8. The recuperation threshold was 5 %. All heating was ramped at 5°C s⁻¹. Data analysis included a measurement error of 1 % for single aliquots and of 2.5 % for single grains. The total light sum figure (Supplementary Figure DR18) from a typical sample (48) shows that only 3 % of the grains contribute 85 % of the total light sum. We crossed-checked different aliquot sizes using single grains as well as 2 mm and 6 mm (spray mask size) multiple grain approaches (Supplementary Table DR4). The number of grains per aliquot is ~80 (2 mm) and ~600 (6 mm). The most intensively investigated samples (6 and 7) show almost identical results for the different approaches. For consistency, all ages related to the Holocene deposits shown in the figures are those determined for 2 mm aliquots (see: 4. Supplementary Discussion).

Most of the **Pleistocene samples** were out of range for the OSL SAR protocol of Murray and Wintle (2000) due to early signal saturation (Supplementary Fig. DR15). In order to date dune sands with greater expected doses, a previously published (Supplementary Table DR3, Porat et al. (2009)), single-aliquot regenerative-dose protocol (SAR) for TT-OSL was applied. A test dose of 10 Gy was used. The initial 0.4 s TT-OSL signal minus a background estimated from the 0.4-10.4 s integral, as suggested by Ballarini et al. (2007), was used for TT-OSL dose determinations (Supplementary figure DR17). For all measurements the aliquot size was 6 mm. Dose dependent dose recovery tests and recycling ratios met the standard performance criteria (Porat et al., 2009) and dose response curves demonstrated the potential to extend the dose range to beyond 600 Gy. Multiple aliquot additive dose (MAAD) TT-OSL protocols were used to test for sensitivity changes in the SAR TT-OSL protocol up to doses of 1200 Gy. A strong dose dependent deviation of the SAR TT-OSL relative to the MAAD TT-OSL dose response is observed. Comparison of the TT-OSL and OSL sensitivity data obtained from the MAAD and SAR data sets show a lack of proportionality between TT-OSL and OSL for the SAR data which will result in a problem when SAR dose response curves are constructed using many regeneration points with doses above 300 Gy. As a consequence for the samples under consideration the SAR TT-OSL protocol of Porat et al. (2009) can only be used when the

expected doses are below 300 Gy (Supplementary Fig. DR16). A detailed description of the TT-OSL testing results is given in Rosenberg et al. (2011).

AAR Dating Method

A technique of amino acid analysis has been developed for geochronological purposes (Penkman et al., 2008), combining a Reverse-Phase High Pressure Liquid Chromatography method of analysis (Kaufman and Manley, 1998) with the isolation of an "intra-crystalline" fraction of amino acids by bleach treatment (Sykes et al., 1995). This combination of techniques results in the analysis of D/L values of multiple amino acids from the chemically-protected protein within the biomineral; enabling both decreased sample sizes and increased reliability of the analysis. Amino acid analyses were undertaken on Melanoides tuberculata shells from Mundafan and Khujaymah. This involved isolating the intra-crystalline protein fraction of the gastropod Melanoides tuberculata, which is a novel species for amino acid studies. The rate of racemization is governed by a variety of factors, but the intra-crystalline amino acid fraction appears to operate as a closed system (Penkman et al., 2008; Penkman et al., 2007). Environmental factors in the intra-crystalline fraction of a particular species (e.g. water content, concentration of cations, pH) are constant and therefore the extent of racemization is a function of time and temperature. Over a small geographical area, such as that represented in this study, it can be assumed that the integrated temperature histories are effectively the same. Any differences in the extent of decomposition of protein within the sample are therefore agedependent. The pattern of protein degradation with time is slightly different for each species and biomineral. Amino acid racemization is governed by the original protein sequence and conformation. Whilst developing the research into closed-system protein degradation it became clear that the reaction rates were species-specific, even in the intra-crystalline fraction. This necessitates the comparison of amino acid data only within a single species (Penkman et al., 2007). As Melanoides tuberculata has not been characterized for amino acid geochronology before, this study provides the first data as to its usefulness.

Amino acid racemization (AAR) analyses were undertaken on *Melanoides tuberculata* shells from 8 samples:

4 subsamples from sample 13, section 22.3 (Supplementary Fig. DR2), NEaar 6009-6012, ~8 ka

² subsamples from sample 36, section 21.2 (Supplementary Fig. DR2), Neaar 5774, 5807, ~8 ka

² subsamples from sample 40, section 22.2 (Supplementary Fig. DR4), NEaar 6003-6004, ~100 ka

⁴ subsamples from sample 42, section 22.2 (Supplementary Fig. DR4), NEaar 6005-6008, ~100 ka

⁵ subsamples from sample 46, section 22.2 (Supplementary Fig. DR4), NEaar 5775, 5808-5811, ~100 ka

² subsamples from sample 69, section 26.3 (Supplementary Fig. DR8), NEaar 6017-6018, ~130 ka

² subsamples from sample 62, section 28.1 (Supplementary Fig. DR8), NEaar 5776. 5812, ~120 ka

⁴ subsamples from sample 55, section 28.6 (Supplementary Fig. DR7), NEaar 6013-6016, ~85 ka

All samples were prepared using the procedures of (Penkman et al., 2008) to isolate the intra-crystalline protein by bleaching. Two subsamples were then taken from each shell; one fraction was directly dematerialized and the free amino acids analyzed (referred to as the 'Free' amino acids, FAA, F), and the second was treated to release the peptide-bound amino acids, thus yielding the 'total' amino acid concentration, referred to as the 'Total Hydrolysable amino acid fraction' (THAA, H*). During preparative hydrolysis both asparagine and glutamine undergo rapid irreversible deamination to aspartic acid and glutamic acid respectively. It is therefore not possible to distinguish between the acidic amino acids and their derivatives and they are reported together as Asx and Glx.

Carbon Isotope $\delta^{13}C_{org}$

Carbon isotope composition (Supplementary Table DR7) were determined using a ThermoFisher Flash-EA 1112 coupled with a Conflo IV interface to a ThermoFisher Delta V plus isotope ratio mass spectrometer (IRMS). Samples were combusted in the presence of O₂ in an oxidation column at 1030°C. Combustion gases were passed through a reduction column (650°C), and produced CO₂ gases transferred to the IRMS via an open split for on-line isotope measurements. Isotope ratios are reported in the conventional δ -notation with respect to V-PDB (Vienna Pee Dee Belemnite) standards. The methods were calibrated with NBS22 ($\delta^{13}C =$ -30.03‰) and IAEA CH-6 ($\delta^{13}C =$ -10.46‰) for carbon. Reproducibility of the measurements is better than 0.2‰.

3. Supplementary Tables

Supplementary Table DR1. OSL SAR protocol (Murray and Wintle, 2000) as used for 2 mm and 6 mm aliquots of Holocene age. For the natural sample, x = 0, $D_0 = 0$ Gy and the subscript n is used (L_n , T_n).

Step	Procedure
1	Dose, $D_{\rm x}$
2	Preheat 1 at 220°C for 10 s
3	Blue stimulation at 125°C for 60 s; L_X
4	Test dose, $D_t = 6$ Gy
5	Preheat 2 at 220°C for 10 s
6	Blue stimulation at 125°C for 60 s; T_X
	Return to 1

Supplementary Table DR2. OSL SAR protocol (Murray and Wintle, 2000) as used for single grains of Holocene age. For the natural sample, x = 0, $D_0 = 0$ Gy and the subscript n is used (L_n , T_n).

Step	Procedure
1	Dose, $D_{\rm x}$
2	Preheat 1 at 220°C for 10 s
3	Green laser stimulation at 125°C for 0.5 s; L_X
4	Test dose, $D_t = 6 \text{ Gy}$
5	Preheat 2 at 220°C for 10 s
6	Green laser stimulation at 125°C for 0.5 s; T_X
	Return to 1

Supplementary Table DR3. Simplified TT-OSL SAR protocol as suggested by Porat et al. (2009). For the natural sample, x = 0, $D_0 = 0$ Gy and the subscript n is used (L_n , T_n).

Step	Procedure
1	Dose, $D_{\rm x}$
2	Preheat at 200°C for 10 s
3	Blue stimulation at 125°C for 300 s
4	Transfer preheat at 260°C for 10 s
5	Blue stimulation at 125°C for 100 s; L_X
6	Test dose, $D_{\rm t}$
7	Preheat at 220°C for 10 s
8	Blue stimulation at 125°C for 100 s; T_X
9	Heat at 300°C for 100 s
	Return to 1

Supplementary Table DR4. OSL and TT-OSL results. The grain size was always 200-250 μ m and the water content 2±1 %. Doserates were calculated from high resolution Gamma spectrometry results. The highlighted data sets are the ones of the ages used.

		Activities (Bq kg ⁴)															
Z	Section	Region	Depth (m)	Altitude (masl)	Cosmic-dose rate (mGy/ka)	к	Th	U-238 from 186 keV and Ra- 226	LL238 from INAA	Ra-226	Doserate (Gy/ka)	De (Gy)	n	GO	Age (ka)	Aliquot size (mm)	Protocol
1	23.1	Mund.	0.9	860	196	271.0 ± 5.7	8.0 ± 0.4	5.0 ± 3.8		7.2 ± 0.2	1.31 ± 0.06	11.51 ± 0.27	52	16	8.8 ± 0.4	2	OSL
1	23.1	Mund.	0.9	860	196	271.0 ± 5.7	8.0 ± 0.4	5.0 ± 3.8		7.2 ± 0.2	1.31 ± 0.06	11.28 ± 0.97	7	18	8.6 ± 0.8	SG	OSL
4	23.3	Mund.	0.6	860	203	207.7 ± 4.5	4.3 ± 0.2	9.2 ± 1.7		4.2 ± 0.1	1.00 ± 0.05	12.05 ± 0.41	28	17	12.0 ± 0.7	2	OSL
6	22.6	Mund.	1.3	86 0	184	98.0 ± 2.2	3.0 ± 1.1	3.2 ± 1.2		3.7 ± 0.2	0.62 ± 0.05	11.88 ± 0.24	107	19	19.1 ± 1.6	2	OSL
6	22.6	Mund.	1.3	860	184	98.0 ± 2.2	3.0 ± 1.1	3.2 ± 1.2		3.7 ± 0.2	0.62 ± 0.05	11.73 ± 0.40	117	32	18.9 ± 1.7	SG	OSL
6	22.6	Mund.	1.3	860	184	98.0 ± 2.2	3.0 ± 1.1	3.2 ± 1.2		3.7 ± 0.2	0.62 ± 0.05	12.21 ± 0.45	23	17	19.6 ± 1.8	6	OSL
7	22.6	Mund.	0.9	86 0	195	248.5 ± 5.2	6.4 ± 0.2	9.6 ± 1.9		6.6 ± 0.1	1.20 ± 0.05	11.72 ± 0.38	46	21	9.8 ± 0.5	2	OSL
7	22.6	Mund.	0.9	860	195	248.5 ± 5.2	6.4 ± 0.2	9.6 ± 1.9		6.6 ± 0.1	1.20 ± 0.05	12.05 ± 0.52	86	30	10.0 ± 0.6	SG	OSL
7	22.6	Mund.	0.9	860	195	248.5 ± 5.2	6.4 ± 0.2	9.6 ± 1.9		6.6 ± 0.1	1.20 ± 0.05	14.09 ± 0.31	38	13	11.7 ± 0.5	6	OSL
9	23.2	Mund.	1.0	860	192	207.9 ± 4.4	5.4 ± 0.1	7.6 ± 2.7		5.2 ± 0.2	1.03 ± 0.04	14.81 ± 0.71	28	25	14.4 ± 0.9	2	OSL
9	23.2	Mund.	1.0	860	192	207.9 ± 4.4	5.4 ± 0.1	7.6 ± 2.7		5.2 ± 0.2	1.03 ± 0.04	16.77 ± 1.02	10	9	16.3 ± 1.2	SG	OSL
15	21.1	Mund.	1.4	860	182	221.4 ± 4.6	6.0 ± 0.2	19.0 ± 2.6	27.1 ± 0.4	8.5 ± 0.1	1.13 ± 0.05	12.52 ± 0.40	47	15	<11.1 ± 0.6	2	OSL
15	21.1	Mund.	1.4	860	182	221.4 ± 4.6	6.0 ± 0.2	19.0 ± 2.6	27.1 ± 0.4	8.5 ± 0.1	1.13 ± 0.05	12.03 ± 1.08	8	23	<10.6 ± 1.0	SG	OSL
21	21.2	Mund.	1.8	860	172	172.2 ± 3.7	5.9 ± 0.3	39.9 ± 8.8		6.8 ± 0.2	0.94 ± 0.04	13.10 ± 0.42	53	22	<13.9 ± 0.8	2	OSL
21	21.2	Mund.	1.8	860	172	172.2 ± 3.7	5.9 ± 0.3	39.9 ± 8.8		6.8 ± 0.2	0.94 ± 0.04	14.33 ± 0.52	22	16	<15.3 ± 0.9	6	OSL
22	21.2	Mund.	1.6	860	177	172.4 ± 3.8	5.2 ± 0.3	21.4 ± 2.5	22.5 ± 0.4	6.1 ± 0.1	0.95 ± 0.04	9.51 ± 0.24	92	23	<10.1 ± 0.5	2	OSL
22	21.2	Mund.	1.6	860	177	172.4 ± 3.8	5.2 ± 0.3	21.4 ± 2.5	22.5 ± 0.4	6.1 ± 0.1	0.95 ± 0.04	9.73 ± 0.19	52	14	<10.3 ± 0.5	6	OSL
38	22.2	Mund.	2.5	86 0	157	433.8 ± 9.0	20.5 ± 0.1	10.6 ± 4.4	20.9 ± 0.4	16.2 ± 0.3	1.98 ± 0.05	201.05 ± 10.66	15	19	101 ± 6	6	TT-OSL
38	22.2	Mund.	2.5	860	157	433.8 ± 9.0	20.5 ± 0.1	10.6 ± 4.4	20.9 ± 0.4	16.2 ± 0.3	1.98 ± 0.05	165.65 ± 11.87	17	18	>83 ± 7	SG	OSL
38	22.2	Mund.	2.5	860	157	433.8 ± 9.0	20.5 ± 0.1	10.6 ± 4.4	20.9 ± 0.4	16.2 ± 0.3	1.98 ± 0.05	130.70 ± 9.52	21	30	>65 ± 5	2	OSL
38	22.2	Mund.	2.5	860	157	433.8 ± 9.0	20.5 ± 0.1	10.6 ± 4.4	20.9 ± 0.4	16.2 ± 0.3	1.98 ± 0.05	141.30 ± 6.77	24	23	>71 ± 4	6	OSL
46	22.2	Mund.	0.3	860	212	182.3 ± 4.0	5.9 ± 0.4	6.4 ± 3.2	11.9 ± 0.3	10.3 ± 0.1	0.97 ± 0.05	109.30 ± 6.76	12	18	103 ± 8	6	TT-OSL
46	22.2	Mund.	0.3	860	212	182.3 ± 4.0	5.9 ± 0.4	6.4 ± 3.2	11.9 ± 0.3	10.3 ± 0.1	0.97 ± 0.05	87.60 ± 6.55	30	33	>83 ± 7	SG	OSL
46	22.2	Mund.	0.3	860	212	182.3 ± 4.0	5.9 ± 0.4	6.4 ± 3.2	11.9 ± 0.3	10.3 ± 0.1	0.97 ± 0.05	108.53 ± 7.21	23	30	>103 ± 8	2	OSL
46	22.2	Mund.	0.3	860	212	182.3 ± 4.0	5.9 ± 0.4	6.4 ± 3.2	11.9 ± 0.3	10.3 ± 0.1	0.97 ± 0.05	92.02 ± 3.28	26	18	>87 ± 5	6	OSL
48	22.5	Mund.	0.8	86 0	198	429.8 ± 8.9	11.5 ± 0.6	12.4 ± 1.9		8.8 ± 0.3	1.79 ± 0.06	150.18 ± 11.40	11	24	79±7	6	TT-OSL
48	22.5	Mund.	0.8	860	198	429.8 ± 8.9	11.5 ± 0.6	12.4 ± 1.9		8.8 ± 0.3	1.79 ± 0.06	154.61 ± 15.00	15	29	>82 ± 9	SG	OSL
48	22.5	Mund.	0.8	860	198	429.8 ± 8.9	11.5 ± 0.6	12.4 ± 1.9		8.8 ± 0.3	1.79 ± 0.06	137.11 ± 12.54	24	43	>72 ± 7	2	OSL
48	22.5	Mund.	0.8	860	198	429.8 ± 8.9	11.5 ± 0.6	12.4 ± 1.9		8.8 ± 0.3	1.79 ± 0.06	164.08 ± 6.82	24	16	>87 ± 5	6	OSL
50	22.4	Mund.	8.0	860	198	363.1 ± 7.5	9.1 ± 0.1	8.4 ± 2.0		9.0 ± 0.2	1.81 ± 0.12	161.63 ± 16.45	14	36	99 ± 11	6	TT-OSL
50	22.4	Mund.	0.8	860	198	363.1 ± 7.5	9.1 ± 0.1	8.4 ± 2.0		9.0 ± 0.2	1.81 ± 0.12	145.52 ± 23.45	16	59	>89 ± 15	SG	OSL
50	22.4	Mund.	8.0	860	198	363.1 ± 7.5	9.1 ± 0.1	8.4 ± 2.0		9.0 ± 0.2	1.81 ± 0.12	149.43 ± 13.67	19	37	>92 ± 9	2	OSL
50	22.4	Mund.	0.8	860	198	363.1 ± 7.5	9.1 ± 0.1	8.4 ± 2.0		9.0 ± 0.2	1.81 ± 0.12	154.26 ± 5.72	18	15	>95 ± 5	6	OSL

ž	Section	Region	Depth (m)	Altitude (masl)	Cosmic-dose rate (mGy/ka)	к	Th	U-238 from 186 keV and Ra- 226	U-238 from INAA	Ra-226	Doserate (Gy/ka)	De (Gy)	n	QO	Age (ka)	Aliquot size (mm)	Protocol
52	28.6	Khuj.	1.5	560	172	303.7 ± 6.3	5.4 ± 0.1	5.0 ± 1.7		6.3 ± 0.2	1.23 ± 0.03	128.86 ± 13.32	8	18	99±11	6	TT-OSL
52	28.6	Khuj.	1.5	560	172	303.7 ± 6.3	5.4 ± 0.1	5.0 ± 1.7		6.3 ± 0.2	1.23 ± 0.03	63.49 ± 11.74	11	57	>49 ± 9	SG	OSL
52	28.6	Khuj.	1.5	560	172	303.7 ± 6.3	5.4 ± 0.1	5.0 ± 1.7		6.3 ± 0.2	1.23 ± 0.03	75.90 ± 7.50	25	47	>58 ± 6	2	OSL
53	28.6	Khuj.	1.2	560	179	337.7 ± 7.1	11.3 ± 0.3	11.7 ± 3.2		15.0 ± 0.4	1.68 ± 0.06	161.51 ± 11.34	8	19	96±8	6	TT-OSL
53	28.6	Khuj.	1.2	560	179	337.7 ± 7.1	11.3 ± 0.3	11.7 ± 3.2		15.0 ± 0.4	1.68 ± 0.06	144.00 ± 18.00	23	39	>86 ± 11	SG	OSL
53	28.6	Khuj.	1.2	560	179	337.7 ± 7.1	11.3 ± 0.3	11.7 ± 3.2		15.0 ± 0.4	1.68 ± 0.06	137.69 ± 12.59	14	32	>82 ± 8	2	OSL
53	28.6	Khuj.	1.2	560	179	337.7 ± 7.1	11.3 ± 0.3	11.7 ± 3.2		15.0 ± 0.4	1.68 ± 0.06	131.03 ± 4.67	23	16	>78 ± 4	6	OSL
54	28.6	Khuj.	0.5	560	197	406.4 ± 8.4	12.8 ± 0.4	15.7 ± 4.3		16.2 ± 0.2	1.94 ± 0.06	170.10 ± 9.36	16	20	88±6	6	TT-OSL
54	28.6	Khuj.	0.5	560	197	406.4 ± 8.4	12.8 ± 0.4	15.7 ± 4.3		16.2 ± 0.2	1.94 ± 0.06	99.92 ± 11.83	18	46	>51±6	SG	OSL
54	28.6	Khuj.	0.5	560	197	406.4 ± 8.4	12.8 ± 0.4	15.7 ± 4.3		16.2 ± 0.2	1.94 ± 0.06	120.62 ± 11.22	9	10	>62 ± 6	2	OSL
58	28.4	Khuj.	0.4	577	200	311.3 ± 6.5	5.4 ± 0.2	5.8 ± 2.7		5.2 ± 0.2	1.30 ± 0.01	119.76 ± 11.18	16	34	90± 9	6	TT-OSL
58	28.4	Khuj.	0.4	577	200	311.3 ± 6.5	5.4 ± 0.2	5.8 ± 2.7		5.2 ± 0.2	1.30 ± 0.01	46.60 ± 7.00	13	54	>35 ± 5	2	OSL
61	28.1	Khuj.	0.3	55 5	203	315.8 ± 6.6	6.5 ± 0.4	22.5 ± 5.6		15.4 ± 0.2	1.48 ± 0.07	188.88 ± 8.81	8	9	121 ± 7	6	TT-OSL
61	28.1	Khuj.	0.3	555	203	315.8 ± 6.6	6.5 ± 0.4	22.5 ± 5.6		15.4 ± 0.2	1.48 ± 0.07	120.70 ± 15.50	14	46	>77 ± 10	2	OSL
63	28.5	Khuj.	0.3	580	203	285.8 ± 6.0	6.1 ± 0.2	7.1 ± 1.9		7.7 ± 0.1	1.03 ± 0.04	147.69 ± 11.44	8	17	113 ± 10	6	TT-OSL
63	28.5	Khuj.	0.3	580	203	285.8 ± 6.0	6.1 ± 0.2	7.1 ± 1.9		7.7 ± 0.1	1.03 ± 0.04	143.60 ± 4.65	13	8	>110 ± 6	2	OSL
65	26.3	Khuj.	0.8	570	189	253.2 ± 5.4	5.5 ± 0.3	8.1 ± 3.0	7.7 ± 0.3	6.7 ± 0.2	1.12 ± 0.01	141.14 ± 10.01	8	15	120 ± 10	6	TT-OSL
65	26.3	Khuj.	8.0	570	189	253.2 ± 5.4	5.5 ± 0.3	8.1 ± 3.0	7.7 ± 0.3	6.7 ± 0.2	1.12 ± 0.01	107.74 ± 6.33	17	23	>92 ± 7	2	OSL
65	26.3	Khuj.	8.0	570	189	253.2 ± 5.4	5.5 ± 0.3	8.1 ± 3.0	7.7 ± 0.3	6.7 ± 0.2	1.12 ± 0.01	113.50 ± 4.82	18	17	>97±6	6	OSL
69	26.3	Khuj.	0.1	57 0	210	143.7 ± 3.1	2.5 ± 0.2	3.9 ± 2.4	5.5 ± 0.2	5.9 ± 0.1	0.73 ± 0.01	108.07 ± 9.83	8	20	136 ± 14	6	TT-OSL
69	26.3	Khuj.	0.1	570	210	143.7 ± 3.1	2.5 ± 0.2	3.9 ± 2.4	5.5 ± 0.2	5.9 ± 0.1	0.73 ± 0.01	97.65 ± 8.55	22	34	>123 ± 12	SG	OSL
71	25.4	Khuj.	1.3	57 0	177	275.1 ± 5.8	5.3 ± 0.2	6.1 ± 2.6	10.2 ± 0.3	9.2 ± 0.3	1.3 ± 0.05	175.64 ± 11.68	16	24	138 ± 11	6	TT-OSL
71	25.4	Khuj.	1.3	570	177	275.1 ± 5.8	5.3 ± 0.2	6.1 ± 2.6	10.2 ± 0.3	9.2 ± 0.3	1.3 ± 0.05	136.83 ± 21.03	12	44	>108 ± 17	SG	OSL
74	25.4	Khuj.	0.2	570	206	22.7 ± 0.8	0.7 ± 0.2	11.0 ± 4.9	18.0 ± 0.3	25.0 ± 0.3	0.8 ± 0.04	111.15 ± 9.46	12	32	147 ± 15	6	TT-OSL
74	25.4	Khuj.	0.2	570	206	22.7 ± 0.8	0.7 ± 0.2	11.0 ± 4.9	18.0 ± 0.3	25.0 ± 0.3	0.8 ± 0.04	93.90 ± 3.17	10	9	>124 ± 8	2	OSL
74	25.4	Khuj.	0.2	570	206	22.7 ± 0.8	0.7 ± 0.2	11.0 ± 4.9	18.0 ± 0.3	25.0 ± 0.3	0.8 ± 0.04	103.27 ± 3.56	20	15	>140 ± 8	6	OSL
75	25.3	Khuj.	0.5	560	197	263.9 ± 5.5	4.5 ± 0.2	4.9 ± 2.0		6.0 ± 0.2	1.2 ± 0.05	170.91 ± 8.32	7	26	144 ± 9	6	TT-OSL
75	25.3	Khuj.	0.5	560	197	263.9 ± 5.5	4.5 ± 0.2	4.9 ± 2.0		6.0 ± 0.2	1.2 ± 0.05	155.50 ± 27.30	15	66	>131 ± 24	SG	OSL
75	25.3	Khuj.	0.5	560	197	263.9 ± 5.5	4.5 ± 0.2	4.9 ± 2.0		6.0 ± 0.2	1.2 ± 0.05	157.40 ± 16.20	10	30	>133 ± 15	2	OSL
78	26.6	Khuj.	0.7	540	191	275.7 ± 5.7	4.3 ± 1.2	6.5 ± 1.7		7.7 ± 0.1	1.2 ± 0.06	177.96 ± 9.79	8	7	143 ± 11	6	TT-OSL
78	26.6	Khuj.	0.7	540	191	275.7 ± 5.7	4.3 ± 1.2	6.5 ± 1.7		7.7 ± 0.1	1.2 ± 0.06	157.84 ± 12.40	10	22	>127 ± 12	2	OSL

Activities (Bqkg⁻¹)

Supplementary Table DR5. Amino acid data for *Melanoides tuberculata* from Mundafan and Khujaymah. Error terms represent one standard deviation about the mean. Each sample was bleached, with the free amino acid fraction signified by 'FAA' and the total hydrolysable fraction by 'THAA'.

	FAA						THAA											
		Asx D	/L	GIx D	/L	Ala D	/L	Val D	/L	Asx D	0/L	GIx D	/L	Ala D	a D/L Val D/L			
Sample	Neaar ı	x	σ	x	σ	x	σ	x	σ	x	σ	x	σ	x	σ	x σ		
Sample 13	6009	0.910	0.016	0.485	0.030	0.729	0.002	0.347	0.058	0.796	0.002	0.414	0.003	0.658	0.007	0.293	0.003	
Sample 13	6010	0.903	0.023	0.490	0.024	0.713	0.001	0.350	0.066	0.758	0.001	0.400	0.003	0.643	0.012	0.270	0.002	
Sample 13	6011	0.814	0.025	0.382	0.022	0.655	0.032	0.380	0.042	0.761	0.003	0.380	0.004	0.635	0.008	0.293	0.002	
Sample 13	6012	0.856	0.026	0.433	0.011	0.710	0.034	0.432	0.045	0.708	0.003	0.371	0.004	0.630	0.012	0.279	0.001	
Sample 13		0.871	0.044	0.448	0.051	0.702	0.032	0.377	0.039	0.756	0.036	0.391	0.020	0.642	0.012	0.284	0.011	
Sample 36	5774	0.870	0.008	0.545	0.012	0.785	0.010	0.379	0.006	0.798	0.000	0.495	0.009	0.735	0.004	0.419	0.029	
Sample 36	5807	0.871	0.007	0.488	0.022	0.783	0.004	0.363	0.031	0.778	0.001	0.451	0.007	0.702	0.007	0.314	0.055	
Sample 36		0.870	0.001	0.516	0.040	0.784	0.001	0.371	0.011	0.788	0.014	0.473	0.031	0.718	0.023	0.366	0.074	
Sample 40	6003	0.953	0.022	0.763	0.101	0.909	0.013	1.068	0.024	0.835	0.001	0.809	0.009	0.898	0.003	0.715	0.016	
Sample 40	6004	0.946	0.021	0.785	0.069	0.889	0.024	0.958	0.200	0.824	0.001	0.714	0.002	0.852	0.013	0.739	0.014	
Sample 40		0.949	0.022	0.774	0.085	0.899	0.018	1.013	0.112	0.829	0.008	0.762	0.067	0.875	0.033	0.727	0.017	
Sample 42	6005	0.962	0.024	0.822	0.066	0.926	0.005	1.060	0.034	0.911	0.002	0.853	0.002	0.918	0.004	0.835	0.041	
Sample 42	6006	0.949		0.725		0.922		0.859		0.840	0.003	0.782	0.001	0.892	0.004	0.829	0.023	
Sample 42	6007	0.907	0.017	0.774	0.062	0.917	0.007	0.966	0.155	0.858	0.005	0.739	0.006	0.854	0.006	0.809	0.044	
Sample 42	6008	0.927	0.013	0.845	0.047	0.914	0.014	1.002	0.160	0.832	0.002	0.780	0.006	0.879	0.001	0.773	0.013	
Sample 42		0.937	0.024	0.791	0.053	0.920	0.005	0.972	0.084	0.860	0.035	0.788	0.047	0.886	0.027	0.811	0.028	
Sample 46	5775	0.936	0.005	0.903	0.029	0.943	0.024	0.965	0.052	0.864	0.009	0.836	0.010	0.901	0.001	0.841	0.053	
Sample 46	5808	0.933	0.008	0.872	0.032	0.913	0.000	0.967	0.014	0.908	0.010	0.851	0.012	0.900	0.014	0.839	0.028	
Sample 46	5809	0.922	0.003	0.846	0.002	0.959	0.017	0.940	0.032	0.914	0.022	0.849	0.009	0.910	0.006	0.849	0.012	
Sample 46	5810	0.944	0.011	0.859	0.010	0.955	0.001	0.982	0.064	0.924	0.012	0.875	0.009	0.928	0.019	0.844	0.024	
Sample 46	5811	0.939	0.004	0.850	0.000	0.949	0.000	0.963	0.018	0.893	0.006	0.848	0.027	0.902	0.021	0.802	0.035	
Sample 46		0.935	0.008	0.866	0.023	0.944	0.018	0.963	0.015	0.900	0.023	0.852	0.014	0.908	0.012	0.835	0.019	
Sample 55	6013	0.943	0.019	0.894	0.029	0.932	0.007	0.987	0.148	0.868	0.004	0.825	0.008	0.891	0.011	0.825	0.050	
Sample 55	6014	0.912	0.054	0.836	0.005	0.828	0.144	0.991	0.061	*0.1817	*0.000	*0.138	*0.001	*0.326	*0.007	*0.117	*0.001	
Sample 55	6015	0.922	0.039	0.855	0.025	0.846	0.121	0.984	0.083	0.857	0.000	0.836	0.008	0.901	0.014	0.898	0.027	
Sample 55	6016	0.929	0.025	0.839	0.040	0.879	0.075	0.955	0.031	0.688	0.002	0.783	0.009	0.776	0.019	0.828	0.076	
Sample 55		0.927	0.013	0.856	0.027	0.871	0.045	0.979	0.016	0.804	0.101	0.815	0.028	0.856	0.069	0.850	0.041	
Sample 62	5776	0.920	0.007	0.890	0.016	0.901	0.044	1.039	0.125	0.768	0.001	0.680	0.016	0.848	0.004	0.682	0.034	
Sample 62	5812	0.906	0.006	0.721	0.144	0.956	0.000	0.971	0.196	0.851	0.009	0.689	0.053	0.818	0.022	0.755	0.043	
Sample 62		0.913	0.010	0.805	0.119	0.929	0.038	1.005	0.048	0.810	0.059	0.684	0.007	0.833	0.021	0.718	0.052	
Sample 69	6017	0.895	0.059	0.804	0.033	0.804	0.144	0.797	0.038	0.756	0.016	0.701	0.022	0.792	0.049	0.784	0.065	
Sample 69	6018	0.902	0.052	0.798	0.002	0.801	0.167	0.855	0.007									
Sample 69		0.898	0.005	0.801	0.004	0.803	0.002	0.826	0.041	0.756	0.016	0.701	0.022	0.792	0.049	0.784	0.065	
*not included i	in average d	ue to conta	mination															

Supplementary Table DR6. Radiocarbon dating results. Calibration was done with OxCal 4.10 using the IntCal09 curve (Bronk Ramsey, 2009). Preparation and measurement procedure are described in Hajdas et al. (2004). Samples with the comment "leach+" were subject to additional HCl leaching.

Sec	Nr	Lab Nr.	Material	DeltaC13 (‰)	C14 age BP	Cal (2σ) age (OxCal 4.10) BP	Comment
21.2	37	ETH-38514	Phytoclasts	-22.6 ± 1.1	7420 ± 35	8260 ± 80	
21.2	37	ETH-36454	Shell, Lymnea	-3.7 ± 1.2	7720 ± 55	8500 ± 90	
21.2	33	ETH-36452	Bone, Vertebrate	-39.2 ± 1.1	7810 ± 70	8700 ± 280	
21.2	33	ETH-36453	Shell, Lymnea	-12.4 ± 1.2	8820 ± 65	9900 ± 270	
21.2	33	ETH-37794	Phytoclasts	-25.6 ± 1.1	7755 ± 40	8520 ± 80	
21.2	29	ETH-37793	Phytoclasts	-23.6 ± 1.1	8255 ± 45	9220 ± 190	
21.2	25	ETH-38512	Phytoclasts	-24.0 ± 1.1	8710 ± 35	9720 ± 170	
21.2	23	ETH-38511	Phytoclasts	-23.8 ± 1.1	8685 ± 35	9640 ± 90	
21.2	23	ETH-36567	Phytoclasts (raw)	-2.2 ± 1.2	9300 ± 40	10480 ± 175	
21.2	23	ETH-36451	Shell, Lymnea	-8.9 ± 1.2	9075 ± 60	10220 ± 250	
21.1	16	ETH-37792	Phytoclasts	-23.0 ± 1.1	8300 ± 45	9290 ± 150	
22.3	13	ETH-37906	Phytoclasts	-25.7 ± 1.1	7030 ± 40	7860 ± 100	
23.2	10	ETH-37907	Phytoclasts	-23.5 ± 1.4	7325 ± 50	8160 ± 150	
22.2	43	ETH-36455	Shell, Melanoides	0.3 ± 1.2	31570 ± 420	35930 ± 870	
22.2	43	ETH-36455-	Shell, Melanoides	-4.1 ± 1.2	34350 ± 320	39490 ± 850	leach+
22.2			Shell, Melanoides		38065 ± 895	42830 ± 1350	
22.2	46	ETH-36456-	Shell, Melanoides	-1.9 ± 1.2	41785 ± 770	45370 ± 1190	leach+
22.4	50	ETH-36457	Shell, Unio	1.1 ± 1.2	34065 ± 580	39040 ± 1630	
22.4			Shell, Unio	-2.5 ± 1.2	36755 ± 415	41720 ± 620	leach+
28.6	55	ETH-38709	Marl, bulk	-2.0 ± 1.1	28970 ± 135	33780 ± 660	
26.3			Shell, Unio	-9.5 ± 1.1	16230 ± 50	19260 ± 310	
28.1	62	ETH-38711	Shell, Melanoides	-8.0 ± 1.1	28685 ± 125	33070 ± 520	

Supplementary Table DR7. Carbon isotope $\delta^{13}C_{org}$ ratios

Sample	Weight		δ ¹³ C _{org} VPDB
(Fig. DR2)	(mg)	Run	(%0)
16	4.856	NC0183	-24.830
23	6.077	NC0183	-24.844
29	4.791	NC0183	-25.976
37	9.142	NC0183	-26.141
13	6.489	NC0183	-25.918
10	6.206	NC0183	-23.233

4. Supplementary Discussion

Discussion of OSL dating

Sources of scatter in D_e distributions include the variability in luminescence sensitivity and dose saturation characteristics (Roberts et al., 1999; Duller et al., 2000), incomplete bleaching (Olley et al., 1998), variations in microdosimetry (Murray and Roberts, 1997; Kalchgruber et al., 2003; Vandenberghe et al, 2003), and post-depositional mixing (Roberts et al., 1999). While incomplete bleaching and postdepositional mixing increases the spread of the distribution by skewed distributions or a mixture of different populations, variations in microdosimetry and the variability in sensitivity and saturation characteristics only generally increases the spread of the distribution. The samples from Mundafan and Khujaymah were tested for mixed populations by single grain methodology. Single grain D_e distributions (Supplementary Fig. DR19a,c) show little skewness and only a single population ruling out problems of incomplete bleaching and post-depositional mixing. In order to describe D_e distributions quantitatively, a parameter defining the spread of the distribution was calculated. The over-dispersion parameter (OD) is defined as the spread in a population that is additional to the spread expected from the error on each individual D_e value (Galbraith et al., 1999), which is strongly influenced by differences in signal intensity. The OD of the samples under consideration (Supplementary Table DR4) is on average 20 %, ranging from 15 to 25 % for the Holocene samples (2 mm aliquots). Such broad distributions have previously been observed for dune sands from Southern Australia and have been attributed to variations in microdosimetry (Kalchgruber et al. 2003). Appling the decision process of Bailey and Arnold (2006) shows that none of the distributions from Holocene samples has a skew that indicates a mixture of populations. In order to obtain a central value that best reflects the average D_e of each sample, the central age model (Galbraith et al., 1999) was chosen for calculations.

For the Holocene deposits (Supplementary Fig. DR2), it is shown that all OSL ages for the dune sands below the lake deposits are predating the onset of lake formation, as given by radiocarbon, by not more than a few thousand years. We hence assume that the deposition of aeolian sand below the lake deposits in general usually just predates the onset of lake formation. This assumption has been applied for the Pleistocene deposits and is supported by evidence from sections 22.2 (Supplementary Fig. DR4), 26.2 (Supplementary Fig. DR8) and 25.4 (Supplementary Fig. DR9).

Discussion AAR geochronology

The DL ratios of Asx, Glx, serine (Ser), alanine (Ala) and valine (Val) provide an overall estimate of protein decomposition. These indicators of protein decomposition have been selected as their peaks are cleanly eluted with baseline separation and they cover a wide range of rates of reaction. It is expected that with increasing age, the extent of racemization (D/L) will increase. If the amino acids were contained within a closed system, the relationship between the FAA and the THAA fractions should be highly correlated, with non-concordance enabling the recognition of compromised samples (Preece and Penkman, 2005). The plot of FAA to THAA data from each sample can also be used as a relative timescale, with younger samples falling towards the bottom left corner of the graph and older samples falling towards the upper right corner, along the line of expected decomposition. The data from the Mundafan samples have been plotted in this way for each of the amino acids. As the site of Mundafan has experienced a different temperature history to that of previous studies of intra-crystalline protein degradation, correlation to absolute dates requires the development of an aminostratigraphic framework for this area.

Asparagine (Asx) is one of the fastest racemizing of the amino acids discussed here (due to the fact that it can racemize whilst still peptide bound). This enables good levels of resolution at younger age sites, but decreased resolution with older material. The D/L Asx data from the Holocene samples have the lowest values of the samples analyzed, as expected. However, most of the other horizons are nearing equilibrium, meaning that Asx is not particularly useful for age resolution for sites of this age. The subsamples of sample 55 are divergent – 6014bH* shows very low D/L values in the THAA fraction, falling away from the line of expected decomposition, along with high concentrations and a differing amino acid composition. This sample shows clear signs of contamination, so although the data is presented in the figures, it should not be used for age determination.

Glutamine (Glx) is one of the slower racemizing amino acids discussed here and so the low levels of racemization do help discriminate between material of the ages of this study. It is noteworthy that Glx has a slightly unusual pattern of racemization in the free form, due to the formation of a lactam. This results in difficulties in measuring Glx in the Free form, as the lactam cannot be derivitized and is therefore unavailable for analysis. The Glx D/L values from sample 13 show the lowest values (Supplementary Fig. DR23), with sample 36 yielding slightly higher D/L values, consistent with their relative radiocarbon dates within the Holocene. Again, the 6014bH* subsamples from sample 55 show that their closed system has been compromised and should be rejected. The other sites show high levels of racemization, nearing equilibrium, although sample 62 and sample 69 exhibit the lower values of the group. Sample 46, sample 51 and sample 40 show similar levels of protein breakdown, consistent with relatively rapid deposition of this sequence.

Alanine (Ala) is a hydrophobic amino acid, whose concentration is partly contributed from the decomposition of other amino acids (notably serine). Ala racemizes at an intermediate rate, so is one of the amino acids that may help distinguishing samples of the timescales under consideration. The results for Ala are broadly similar to that seen in the other amino acids (Supplementary Fig. DR24). The alanine data supports the interpretation that sample 13 is the youngest in age, with sample 36 slightly older. The 6014bH* subsamples from sample 55 again show divergence from closed system behavior. The replication of sample 55 and sample 69 samples is not particularly good, so further samples are being run to confirm this. Sample 62 tend to show a slightly younger age, with sample 46, sample 42 and sample 40 showing tight clustering at almost equilibrium levels.

Valine (Val) has extremely low rates of racemization, and as the concentration of Val is quite low, the difficulty of measuring the D/L accurately results in higher variability. It does however still prove useful for age discrimination within material of Pleistocene age. The Val D/L in the FAA and THAA fractions again support the other amino acid data (Supplementary Fig. DR23). The Holocene samples are demonstrably younger, while the one subsample from sample 55 is compromised. Data from sample 62 is more variable, but still consistent with a younger age than sample 46, sample 42 and sample 40 (but significantly older than the Holocene samples). Again, the older samples show values nearing equilibrium; it is therefore likely that AAR dating has reached its temporal limit for this area.

Isoleucine (Ile) also has extremely low rates of racemization, and baseline separation is not very accurate using the RP-HPLC technique, so is usually separated by ion exchange chromatography. However, results are presented here as they support the data obtained from the other amino acids. Isoleucine has two chiral centers, but the important geochronological reaction is the breakdown of L-isoleucine to its diastereomer D-alloisoleucine, which is reported to equilibrate at D/L values of 1.3 (often reported as an A/I value). Again, the Holocene samples are clearly distinguishable, and the other sites are tending towards equilibrium.

The Melanoides tuberculata samples seem to show relatively good closed system behavior, although a higher degree of variability is seen within sample 62, and there is clear contamination in sample 55. While optical thin section analysis showed that the structure of the shells is well preserved, there were indications of epitaxial calcite growth. Our preparation isolates the intra-crystalline material within a shell, therefore we are theoretically analyzing a closed system of protein. However, if any re-crystallization has occurred, then this compromises the intra-crystalline fraction, meaning it has not been closed system for the history of the sample, and therefore cannot be used for dating. It may be that the variability is due to re-crystallization, but the relative consistency between the amino acids would argue against any significant effect. The two samples of well-constrained Holocene age: Sample 13 (~ 8000 years BP) and sample 36 (~9000 years BP) show levels of protein degradation consistent with these ages. The rest of the samples show levels of racemization nearing equilibrium, limiting their relative resolution. At what point equilibrium is achieved with these species in this tropical environment is unknown. Further study will inform on this. Using AAR, it is not possible to discriminate between the three

samples analyzed which have been excavated within a sequence (sample 46, sample 42 and sample 40).

In this study, the amino acid data has been used as a relative dating technique to present a pilot aminostratigraphy for the area in question. The conversion of relative sequences into absolute dates and accurate correlation between different areas requires further studies on the patterns of breakdown in *Melanoides tuberculata*. However, it is concluded that sample 13 and sample 36 are the youngest within the sequence, consistent with their radiocarbon ages. Very high levels of protein degradation are observed in the other samples, inconsistent with a Holocene age. Amino acid geochronology has the potential to yield useful information on the relative age of these deposits in this area, but it is vital to establish the time point where equilibration occurs to constrain these ages.

Discussion Radiocarbon Dating

For understanding the discrepancy between Pleistocene OSL and radiocarbon ages, it is important to realize that samples (mainly porous mollusk shells; e.g. Melanoides tuberculata) used for radiocarbon dating of pre-Holocene deposits will have strongly been affected by epitaxial calcite growth due to the interaction with water during the Holocene humid period, leading to an incorporation of younger ¹⁴C. In section 22.2 (Supplementary Fig. DR4) radiocarbon ages of Melanoides tuberculata shells (39,490 \pm 850 cal BP (sample 43) and 45,370 \pm 1190 cal BP (sample 46)) are stratigraphically inverted. OSL ages of ~ 100 ka for aeolian deposits from above and below the same lacustrine sequence are considered to reflect the real depositional age. Additional leaching on the same samples (43, 46 and also 50) had only increased the age by 3.5 ka, 2.5 ka and 2.7 ka but was not able to remove all of the epitaxial calcite due to the porous structure of the shells. This problem has been reported on other occasions from similar environments (Fontes and Gasse, 1989). The Holocene radiocarbon ages (Supplementary Fig. DR2) on shells on the other hand show some hard water effect with the ages on shells being up to 1.4 ka older than the ages on phytoclasts.

Discussion Ostracodes

Ostracode species from Mundafan and Khujaymah are typical of fresh- to brackish, shallow water environments. Most prominent species include *Cyprideis torosa* that mainly occurs in marine brackish waters with fluctuating salinities but is also characteristic of a wide range of continental water bodies down to a water depth of 30 m (McClure, 1980; Meisch, 2000). Other species that tolerate an increase in salinity are *Cypris* sp., *Cypris bispinosa*, and *Heterocypris salina* (Meisch, 2000). Freshwater species include *Cypretta* sp., a genus usually found in warm (often subtropical), shallow, vegetated waters (Bronshtein, 1947; Smith and Delorme, 2010) and *Paracandona* sp. that prefers small water bodies with a swampy or boggy bottom and the littoral zone of lakes (Meisch, 2000). A species that needs stable conditions and cannot survive in desiccating environments is *Darwinula stevensoni*. This species lives in aquifers and seeps (Bronshtein, 1947; Griffiths et al., 2001; Martens et al., 1997; Schwalb, 2003; Smith and Delorme, 2010) and also occurs in the littoral of lakes, rivers, bogs and springs (Dumont et al., 1986; Martens et al., 1997). Less characteristic and less abundant are *Candona* sp. and *Limnocythere* sp., both benthonic species as well as *Candonocypris* sp., a nektobenthonic species that is an indicator for running shallow waters (Pérez et al., 2010).

Each humid phase is characterized by a typical species assemblage (number of samples):

Holocene (16), *Paracandona* sp. is accompanied by *Cypris* sp., *Darwinula stevensoni*, and *Heterocypris salina*. This suggests small and shallow, permanent and possibly slightly saline environments.

~80 ka (3), *Darwinula stevensoni* is the dominant ostracode. This species is accompanied by *Paracandona* sp., *Cypris* sp., and a few specimen of *Cyprideis torosa*. This suggests significantly fresher conditions than during the Holocene and at ~100 ka with permanent water bodies.

~100 ka (5), The dominance of *Cyprideis torosa* hints at brackish waters. The presence of *Candonocypris* sp., *Darwinula stevensoni*, and *Limnocythere* sp. suggests the existence of permanent, shallow water bodies with possibly running water.

~125 ka (1), *Darwinula stevensoni* is the dominant species accompanied by *Cypris* sp. and *Paracandona* sp. This suggests significantly fresher conditions than during the Holocene and at ~100 ka with permanent water bodies.

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