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1 Holocene temperature and hydrological changes  
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4

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16

17 **Abstract**

18 To achieve a sufficient understanding of the spatial dynamics of terrestrial climate  
19 variability, new proxies and networks of data that cover thousands of years and run up  
20 to the present day are needed. Here we show the first Gram-negative bacterial  
21 3-hydroxy fatty acid (3-OH-FA) based temperature and hydrological records from any  
22 palaeoclimate archive globally. The data, covering the last 9 ka before present (BP),  
23 are generated from an individual stalagmite, collected from Heshang Cave, located on  
24 a tributary of the Yangtze River, central China (30°27'N, 110°25'E; 294 m). Our  
25 results indicate a clear early-to-middle Holocene Climatic Optimum (8.0-6.0 ka BP)  
26 followed by a long-term monotonic cooling and increasing variability over the last 0.9  
27 ka BP. The hydrological record shows two relatively long wet periods (8.8-5.9 ka BP  
28 and 3.0-0 ka BP) and one relative dry period (5.9-3.0 ka BP) in central China. We  
29 show that 3-OH-FA biomarkers hold promise as independent tools for palaeoclimate  
30 reconstruction, with the potential to deconvolve temperature and hydrological signals  
31 from an individual stalagmite.

32

33 **Keywords:**

34 Holocene; Paleoclimatology; Novel proxy; 3-hydroxy fatty acid; China; Monsoon;  
35 Speleothems

36

37 **1. Introduction**

38 Nearly half of the Earth's population live within the influence of the modern  
39 monsoon and its importance to terrestrial eco-systems, societal wellbeing and the  
40 global economy can not be overstated ([Webster et al., 1998](#)). Records of past  
41 Holocene rainfall and temperature, which extend the relatively short instrumental  
42 record, can constrain natural monsoon variability and are particularly important for  
43 the Asian monsoon region where prediction of future changes in rainfall using climate  
44 models has proven challenging ([IPCC, 2014](#)). Such records can also illustrate the  
45 influence of the monsoon on prehistoric cultures and settlements ([Xie et al., 2013](#)).

46 Stalagmites have become a key archive in Quaternary palaeoclimatic  
47 reconstruction due to their ability to yield continuous and undisturbed records, precise  
48 and absolute chronologies, and their global terrestrial distribution ([Blyth et al., 2016](#);  
49 [Fairchild et al., 2006](#); [Fairchild and Baker, 2012](#); [McDermott, 2004](#); [Wong and](#)  
50 [Breecker, 2015](#)). Oxygen isotopes are effectively the 'master' or standard approach  
51 for speleothem analysis, but inherently encode a mix of climatic signals ([Lachniet,](#)  
52 [2009](#); [McDermott, 2004](#)), including, at the regional scale: temperature changes, the  
53 isotopic composition of source waters and precipitation amount. In addition, complex  
54 site-specific factors must be taken into account, such as drip rate ([Dreybrodt and](#)  
55 [Scholz, 2011](#)) and CaCO<sub>3</sub> precipitation ([Fairchild and Baker, 2012](#)). Many previous  
56 studies have focused on the interpretation of oxygen isotopes in speleothems, but  
57 deconvolving independent temperature and precipitation signals from speleothem  
58 CaCO<sub>3</sub> remains highly challenging, as evidenced by the paucity of such deconvolved  
59 records ([Hu et al., 2008b](#); [Yuan et al., 2004](#)).

60 Biomarker based proxies are now firmly established in the fields of  
61 paleoceanography and paleolimnology (Castañeda and Schouten, 2011; Eglinton and  
62 Eglinton, 2008; Schouten et al., 2013). Recently attention has turned to the potential  
63 of organic matter and biomarker techniques for speleothem research (Blyth et al.,  
64 2008; Blyth et al., 2016). A number of biomarkers with known paleoclimatic utility  
65 have been discovered and measured in speleothems, including glycerol dialkyl  
66 glycerol tetraethers (GDGTs) (Blyth et al., 2014; Blyth and Schouten, 2013; Yang et  
67 al., 2011), plant derived biomarkers (Blyth et al., 2007; Blyth et al., 2011; Blyth et al.,  
68 2010; Bosle et al., 2014; Xie et al., 2003), branched fatty acids and hydroxy fatty  
69 acids (Blyth et al., 2006; Huang et al., 2008; Wang et al., 2012). Furthermore, Blyth  
70 and Schouten (2013) recently proposed a novel GDGT calibration, based on samples  
71 derived from 33 globally distributed speleothems from caves with a range of average  
72 air temperatures.

73 Biomarkers in stalagmites may originate from the overlying vegetation,  
74 overlying soil ecosystem, limestone aquifer and cave fauna (Blyth et al., 2008).  
75 Moreover, different biomarker classes may have different sources, for example, Yang  
76 et al. (2011) found that the majority of the archaeal isoprenoid and bacterial branched  
77 GDGTs measured in stalagmite samples from Heshang cave were likely produced *in*  
78 *situ*. Most recently, Blyth et al. (2014) found that GDGTs preserved in stalagmites in  
79 the UK and Australia likely originated from the *in situ* microbial communities within  
80 cave systems. An artificial irrigation experiment conducted in Cathedral Cave,  
81 Australia, found different GDGT distributions among speleothem, soil and drip water

82 samples (Baker et al., 2016). In contrast, a 2 year monitoring experiment of drip  
83 waters in Heshang Cave found that fatty acids in drip waters were most likely derived  
84 from the overlying soil and/or groundwater system via particulate entrainment and  
85 deposition (Li et al., 2011). It is noteworthy that the fatty acid ratios (ratios of  
86  $nC_{16:1}/nC_{16:0}$  and  $nC_{18:1}/nC_{18:0}$ ; the prefix  $n$  indicates normal, the number before the  
87 colon specifies the number of C atoms, and the number after the colon gives the  
88 number of double bonds) showed a strong negative relationship with the external  
89 temperature recorded in Yichang meteoric station (located ca. 100 km east of Heshang  
90 Cave), whereas the two ratios displayed no relationship with internal cave  
91 temperatures recorded at the HS4 site, which suggests that *in situ* cave microbes are  
92 probably not the predominant source for  $C_{16}$  and  $C_{18}$  acids in drip water collected in  
93 Heshang Cave. Li et al., (2011) concluded that, based the distributional patterns of the  
94 fatty acids, microbes living in the overlying soils and/or groundwater system are the  
95 dominant source of fatty acids to the Heshang Cave drip waters. We note that  
96 Vaughn et al. (2011) discovered microbia living on speleothem surfaces in Kartchner  
97 Caverns, USA. Such consortia of microbes are an inevitable source of *in situ* fatty  
98 acids. Thus fatty acids measured in stalagmites may be derived from mixed sources,  
99 including overlying soils/ sediments (Li et al., 2011), the ramifying network of  
100 conduits and reservoirs in the limestone and *in situ* microbes (Vaughan et al., 2011).  
101 However, even though the origin and pathways of inclusion into speleothems of  
102 biomarkers may be complex (Blyth et al., 2008; Blyth et al., 2016), it doesn't hinder  
103 the utilization of biomarkers in paleoclimate reconstruction. Site specific

104 interpretation and ground truthing is required, but this is also true for established  
105 paleoclimate techniques, as outlined above. In summary lipid biomarkers preserved in  
106 speleothems show clear potential for paleoclimate reconstruction. However, very few  
107 such biomarker based paleoclimatic reconstructions have been published (Blyth et al.,  
108 2011; Huguet et al., 2018; Li et al., 2014; Xie et al., 2003).

109 Gram-negative bacterial 3-hydroxy fatty acids (3-OH-FAs) are abundant in  
110 stalagmites (Blyth et al., 2006; Huang et al., 2008; Wang et al., 2016; Wang et al.,  
111 2012) and are characteristic compounds of Lipid A, the lipid component of the  
112 lipopolysaccharides (LPS) located in the outer membrane of Gram-negative bacteria  
113 (Szponar et al., 2003; Szponar et al., 2002; Wollenweber and Rietschel, 1990). Based  
114 on strong relationships with environmental pH and temperature from an altitudinal  
115 transect of soils on Shennongjia Mountain (Mt.) central China, a number of novel  
116 3-OH-FA based proxies have been proposed (Wang et al., 2016). For example, the  
117 ratio of *anteiso* to *normal* C<sub>15</sub> 3-hydroxy fatty acid (RAN<sub>15</sub>) was propounded to be a  
118 novel temperature proxy, and the ratio of the total sum of *iso* and *anteiso* 3-OH-FAs to  
119 the total amount of *normal* 3-OH-FAs (Branching Ratio) and RIAN (negative  
120 logarithm of Branching Ratio) were propounded to be novel pH proxies (Wang et al.,  
121 2016).

122 In this study we present inferred temperature and hydrological records, spanning  
123 the last 9 ka BP, based on 3-OH-FA derived proxies from a single stalagmite  
124 collected from Heshang Cave, central China (Fig. 1). This work is the first  
125 demonstration of the application of 3-OH-FA based proxies for paleoclimatic

126 reconstruction and suggests that such approaches may be used to derive independent  
127 quantitative temperature and qualitative hydrological signals from an individual  
128 stalagmite.

129

## 130 **2. Materials and methods**

### 131 **2.1 Sampling site and sample information**

132 Heshang Cave is located at 294m above sea level (a.s.l.), on the Qing River, a  
133 tributary in the middle reaches of the Yangtze River, central China (30°27'N,  
134 110°25'E) (Fig. 1A). Heshang Cave is one of several caves which characterize the  
135 regional karst landscape. The overlying dolomite is ca. 400 m thick and is capped  
136 with a mature layer of soil (20-40 cm-thick) and reasonably dense vegetation (Fig.  
137 1B). The regional climate is strongly impacted by the East Asian Monsoon, with a hot  
138 and moist summer, but relatively cold and dry winter (An, 2000). Regional average  
139 annual precipitation is 1161 mm, based on the recent 65 years (1951-2014) of  
140 meteorological data from Yichang station. The seasonal temperature ranges, inside  
141 and immediately outside the cave, were constrained by 2-hour resolution logging  
142 between 2004 and 2007 using HOBO H8 Pro T loggers (Hu et al., 2008a). The  
143 modern temperature immediately outside the cave varies seasonally from 3°C to 30°C,  
144 with an annual average of 18°C and is statistically identical to that of the nearest  
145 government meteorological station in Changyang county (Hu et al., 2008a). The  
146 annual mean temperature inside the cave is identical to the outside measurements.  
147 However, the amplitude of the internal temperature range is about one fifth of the



148 external cycle and lags the external temperatures by about 10 days (Hu et al., 2008a).  
149 Heshang cave extends a distance of  $\approx 250$  m, roughly horizontally from its opening  
150 (see Fig. 1C) and is well decorated with stalagmites, rimstone pools, and less frequent  
151 stalactites (including an exquisite ‘Lotus Flower’ stalactite).

152 The HS4 stalagmite is 2.5 m long, and was actively growing when collected  
153 from ca. 150 m within Heshang Cave in 2001 (Fig. 1C). It shows clear annual banding  
154 throughout its growth axis, generated by the strong seasonal cycle at this site (Johnson  
155 et al., 2006). Highlights of previous work on this stalagmite include a quantitative  
156 Holocene Asian monsoon rainfall record (Hu et al., 2008b) and high resolution 8.2 ka  
157 event record (Liu et al., 2013; Owen et al., 2016). The HS4 stalagmite was divided  
158 longitudinally into 4 sections. Each section was dedicated to a different branch of  
159 analyses (e.g.  $\delta^{18}\text{O}$ , trace elements, organic geochemistry etc.). 206 subsamples were  
160 taken from the organic geochemistry section along the stalagmite growth axis and  
161 73 subsamples were selected at intervals for biomarker analysis. All the outer layers  
162 of the subsample were removed during sampling to avoid any potential contamination.  
163 Based on annual layering each sample has a resolution of several decades to >100  
164 years.

165 In 2013 seventeen cave sediment samples were collected within Heshang cave  
166 from the entrance to the deepest accessible part of the cave (Fig. 1C) and nine  
167 overlying soil samples were collected from the land-surface immediately above the  
168 cave (which slopes upwards from an altitude of 457 m to 489 m) (Fig. 1B). The  
169 sediment inside the cave is oligotrophic with < 6 g/kg total organic carbon (Gong et

170 [al., 2015](#)).

## 171 **2.2 Chronology**

172 The chronology of HS-4 was established independently by U-Th dating and layer  
173 counting. Twenty-one subsamples were sampled and prepared in a class-1000 clean  
174 lab before being analyzed by multi-collector inductively coupled plasma mass  
175 spectrometry (MC-ICP-MS) at Oxford University (Nu Instruments), following the  
176 techniques of Robinson et al. (2002). Layer counting was used for the uppermost 150  
177 years and U-Th dating for the period ca.  $378 \pm 57$  to  $9446 \pm 146$  years BP. The  
178 chronology of each sample is based on linear interpolation between the  $^{230}\text{Th}$  dates,  
179 the average age uncertainty is 67 yrs. Further details on the chronological techniques  
180 and model are reported by Hu et al. (2008b).

## 181 **2.3 pH measurement**

182 The pH of cave surface sediments and overlying soils was measured following  
183 the methods of Yang et al. (2014). Samples were mixed with ultrapure water in a ratio  
184 of 1:2.5 (g/mL). After standing for 30 min, the pH of the supernatant was measured  
185 using a meter with a precision of  $\pm 0.01$ .

## 186 **2.4 Lipid extraction and work-up**

187 The stalagmite samples were treated with an optimized acid digestion method  
188 following Wang et al. (2012). In brief, 10 grams of stalagmite sample were digested  
189 with 3M HCl, then re-fluxed at 130°C for 3 hours with a condenser/ heating mantle  
190 assembly. An internal standard (pregn-5-en-3.beta.-ol) was quantitatively added to  
191 each sample to quantify the amount of lipids in the stalagmite. After cooling, the

192 residue was extracted by dichloromethane (15mL×4) and the extracts combined.  
193 Solvents were removed by rotary evaporation (Buchi R210) under reduced pressure.  
194 Soil samples and cave surface sediments underwent the same work-up protocol as the  
195 stalagmite samples. The condensed lipids were further derivatized by BF<sub>3</sub>-methanol  
196 (14% BF<sub>3</sub>/methanol, Sigma) and BSTFA (N, O-bis(trimethylsilyl) trifluoroacetamide,  
197 Supelco) before undergoing gas chromatography-mass spectrometry (GC-MS). In  
198 order to minimize contamination, all glassware was soaked in a decontamination  
199 solution, rinsed with ultra purified water, and heated for 6 h at 500 °C. The HCl was  
200 pre-extracted with dichloromethane (DCM, ×4), and all other reagents were tested for  
201 background contaminants.

## 202 **2.5 Instrumental analysis**

203 All the samples were analyzed using GC-MS with a Hewlett Packard 6890 gas  
204 chromatograph coupled to a Hewlett Packard 5973 mass selective detector. Separation  
205 was performed on a ZB-5MS fused silica capillary column (60 m×0.25 mm id.; 0.25  
206 µm film thickness). The GC oven temperature was programmed from 70°C to 200°C  
207 at 10°C per min, then from 200°C to 300°C at 2°C per min, and finally held at 300°C  
208 for 27 min. The carrier gas was He (1 mL/min). The spectrometers were operated in  
209 electron-impact (EI) mode, the ionization energy was set at 70 eV and the scan range  
210 was from 50 to 550 aum.

## 211 **2.6 Proxy calculation**

212 The RAN<sub>15</sub> and RIAN in the HS4 stalagmite samples were calculated using the  
213 relative abundances of the 3-OH-FAs with carbon numbers from C<sub>10</sub> to C<sub>18</sub>, which are

214 derived from Gram-negative bacteria. Standard deviations for the RAN<sub>15</sub>-MAAT and  
215 RIAN were calculated by a duplicate extraction and analyses on 9 randomly selected  
216 samples.

217

### 218 **3. Results and Discussion**

#### 219 **3.1 Distribution and source of 3-OH-FAs**

220 Below we discuss the distributional characteristics of 3-OH-FAs in Heshang  
221 cave sediments and overlying soils and, with consideration of recent bacterial  
222 monitoring of the cave environment and drip waters (Liu et al., 2010; Yun et al.,  
223 2016b), constrain their possible sources and pathways.

224 The average distributions of 3-OH-FAs in the overlying soils, cave surface  
225 sediments and the HS4 stalagmite samples are illustrated in [Figure 2](#). There is an  
226 overall similarity in the distribution patterns of the three sample sets (with some  
227 differences discussed below), with the C<sub>10</sub>, C<sub>12</sub>, C<sub>14</sub>, C<sub>16</sub> and C<sub>18</sub> homologues being  
228 typically most abundant. The carbon number of the detectable 3-OH-FA homologues  
229 varies from C<sub>8</sub> to C<sub>30</sub> ([Fig. 2](#)), however only the overlying soil samples contain the  
230 lowest carbon numbers of the 3-OH-FAs (C<sub>8</sub>, *i*-C<sub>9</sub>, *a*-C<sub>9</sub>, C<sub>9</sub>) ([Fig. 2A](#)). The  
231 distribution of 3-OH-FAs in the HS-4 stalagmite reported here agrees with previous  
232 studies of the HS4 stalagmite from Heshang cave ([Huang et al., 2008](#); [Wang et al.,](#)  
233 [2012](#)). The only other reports of 3-OH-FAs in speleothem samples comes from an  
234 Ethiopian stalagmite ([Blyth et al., 2006](#)) and a British stalagmite ([Blyth et al., 2011](#)).  
235 The molecular distributions reported by Blyth et al. ([2006](#)) and ([2011](#)) are similar

236 (maxima at C<sub>12</sub>, C<sub>14</sub>, C<sub>16</sub> etc.) to the HS4 distributions, but the higher molecular  
237 weight 3-OH-FAs (>C<sub>20</sub>) we detected were not previously reported.

238 Normal and branched 3-OH-FAs homologues of C<sub>10</sub> to C<sub>18</sub> chain length are  
239 abundant constituents of Lipid A, a constituent of Lipopolysaccharide (LPS), the main  
240 component of the outer membrane of Gram-negative bacteria (Lee et al., 2004;  
241 Szponar et al., 2003). 3-OH-FAs have been used to quantify and characterize the  
242 Gram-negative bacterial community in atmospheric aerosols (Lee et al., 2004), marine  
243 dissolved organic matter (DOM) (Wakeham et al., 2003) and snow samples (Tyagi et  
244 al., 2016; Tyagi et al., 2015), and have recently been utilized to define a number of  
245 novel terrestrial paleoclimate proxies (Wang et al., 2016). The fractional abundance of  
246 the individual 3-OH-FA homologues varies from 0 to 30% in the studied samples. The  
247 3-OH-FAs generally show a strong even/odd predominance (Fig. 2). In the HS4  
248 stalagmite, this general even/ odd predominance is accentuated for the homologues in  
249 the range from C<sub>11</sub> to C<sub>15</sub>, but is reversed for the *i*-C<sub>17</sub> 3-OH-FA which is notably  
250 higher than the *n*-C<sub>16</sub> 3-OH-FAs (Fig. 2C). Furthermore, the *n*-C<sub>12</sub> 3-OH-FAs account  
251 for ca.18% in the stalagmite samples, but only ca.11% in both the overlying soils and  
252 cave surface sediments. The higher proportion of *n*-C<sub>12</sub> and *i*-C<sub>17</sub> homologues in the  
253 stalagmite may derive from *in situ* bacterial production or possibly better preservation  
254 of these compounds in stalagmites.

255 Culturable bacteria in drip waters from Heshang Cave are dominated by  
256 Gram-negative heterotrophs, derived from Proteobacteria with the dominance of  
257 Gamma proteobacteria (Liu et al., 2010). A two-year drip water monitoring

258 experiment in Heshang Cave likely demonstrates a pathway for transporting  
259 3-OH-FAs from the overlying soil microbial community to cave sediments and  
260 speleothem surfaces (Li et al., 2011). Meanwhile, a recent molecular survey of  
261 bacterial communities in Heshang cave drip waters, over the period of 2008 to 2013,  
262 confirms a diverse Gram-negative bacterial community and reveals a seasonal control  
263 on Proteobacteria, whereby Beta-proteobacteria are supplied in the summer and  
264 Gamma-proteobacteria in the winter (Yun et al., 2016b). A latest investigation shows  
265 that Proteobacteria are both abundant in Heshang cave overlying soils and drip waters  
266 (Yun et al., 2016a). This demonstrates that the seasonal signal of changes in the  
267 Gram-negative bacterial community is transmitted readily through the Heshang cave  
268 system to drip waters and to the cave and speleothems. This seasonal cycle is positive  
269 from a paleoclimate perspective, suggesting minimal attenuation of bacterial based  
270 climate signals transmitted from the overlying soils to the HS4 stalagmite, at least  
271 sufficient for centennial to millennial scale paleoclimate studies.

272 As noted above, a distinctive feature of the 3-OH-FA distribution in the HS4  
273 samples, is the greater relative abundance of the *n*-C<sub>12</sub> and *i*-C<sub>17</sub> 3-OH-FAs compared  
274 to both overlying soils and cave sediments (Fig. 2). This may suggest an additional  
275 contribution of 3-OH-FAs, derived from microbes from the cave drip water, which are  
276 sequestered and trapped in the calcite matrix (Supplementary Information). We note  
277 that Paction et al. (2013) report that microbial activity can initiate calcite deposition in  
278 the aphotic zone of caves before inorganic precipitation of carbonates. We also note  
279 the low abundances of long chain 3-OH-FAs (C<sub>20</sub>-C<sub>26</sub>), which might originate from

280 fungi, and/or Gram-positive actinomycetes (Keinänen et al., 2003). However, we  
281 suggest that the broad similarity of 3-OH-FA distributions in the overlying soils and  
282 stalagmites, supported by the site-specific analyses of bacterial diversity and transport  
283 pathways (Liu et al., 2010; Yun et al., 2016b), supports a major contribution of  
284 3-OH-FAs from Gram-negative bacteria dwelling in the overlying soils to the HS4  
285 stalagmite samples. This is consistent with previous findings that lipids preserved in  
286 speleothems are principally derived from the overlying soil ecosystem and vegetation,  
287 having been transported from the surface by percolating groundwater, although a  
288 proportion may be derived from the cave ecosystem (Blyth et al., 2014; Xie et al.,  
289 2005; Xie et al., 2003; Yang et al., 2011).

290

### 291 **3.2 Holocene hydrological reconstruction**

292 Recent work has demonstrated that pH is a key environmental parameter in  
293 controlling soil bacterial community structure and diversity (Bååth and Anderson,  
294 2003; Griffiths et al., 2011; Lauber et al., 2009; Shen et al., 2013; Zhang et al., 2015).  
295 Notably, Giotis et al. (2007) found that a strain of Gram-negative bacterium  
296 increased/decreased the proportion of branched-chain fatty acids in higher pH/lower  
297 pH conditions. Recently, a novel pH proxy RIAN which is based on Gram-negative  
298 bacterial derived 3-OH-FAs in soils from Shennongjia Mountain was proposed by  
299 Wang et al. (2016), with a low RIAN value when pH is high. Based on our finding  
300 that 3-OH-FAs in the HS4 stalagmite are mainly derived from the overlying soils,  
301 here we interpret the HS4 RIAN record as reflecting local changes of pH (Fig. 3A).

302 We note that the HS4 RIAN record is, for the period of mutual overlap, consistent  
303 with hydrological records based on Hopanoids from the Dajihu peatland (Xie et al.,  
304 2013) (Fig. 3B; Supplementary Information), situated 130 km to the NW of Heshang  
305 Cave and the proportion of soil-derived magnetic minerals ( $IRM_{\text{soft-flux}}$ ) incorporated  
306 into the HS4 stalagmite, which Zhu et al. (2017) interpret as a proxy for rainfall  
307 amount and intensity, with sensitivity to extreme precipitation events (Fig. 3C). These  
308 consistent lines of evidence demonstrate RIAN could be used as qualitative  
309 hydrological proxy in stalagmites. We argue below that, for the HS4 record, when  
310 effective precipitation was higher in the past, the RIAN value is lower and likely  
311 indicates leaching of soils and the influence of higher groundwater pH on the  
312 3-OH-FA producing Gram-negative bacteria.

313 In soil environments pH reflects the balance between precipitation and  
314 evaporation (e.g. hydrologically effective precipitation) and water movement through  
315 the soil. Rainwater is naturally acidic due to the reaction with  $CO_2$  in the atmosphere  
316 to form carbonic acid. Excess rainfall leaches base cations increasing the relative  
317 percentage of  $H^+$  and  $Al^{3+}$  ions (and thus acidity) in water. Soil temperatures, pH and  
318 aeration conditions also affect soil microbial activity and diversity which determines  
319 bacterial respiration of  $CO_2$  (and the formation of carbonic acid), further influencing  
320 the pH of soil water and the degree of leaching (Fairchild and Baker, 2012). Thus we  
321 argue that in the soils above Heshang cave, during the Holocene, pH was lower with  
322 higher effective precipitation rates (and vice versa). This is consistent with the  
323 generally observed relationship between effective precipitation and pH in global soils



324 (Slessarev et al., 2016; Yang et al., 2014) and wider evidence of the substantial  
325 influence of pH on bacterial communities at both local and continental scales (Lauber  
326 et al., 2009; Rousk et al., 2010). However, on the contrary, changes of pH in  
327 groundwater systems may display in an opposite trend to that of the overlying soils in  
328 response to increased effective precipitation. In a well-drained karst landscape  
329 increases in precipitation will also be associated with the increased movement of  
330 material (including organic matter and biomarkers), in dissolved or colloidal form to  
331 the groundwater system, which may ultimately percolate or flush into cave systems  
332 (Blyth et al., 2008). The pH of the groundwater is greatly affected by soil-derived  
333 colloids and fine sands flushed into the groundwater system (Fairchild and Baker,  
334 2012). The soil processes outlined above influence the latter source of material and  
335 thus contribute to the pH of the groundwater system. More importantly, increased  
336 rainfall can result in a fall in the total cation content which will lead to an increase of  
337 pH in the groundwater system (Fairchild and Baker, 2012). Thus increased effective  
338 precipitation may lead to antiphased pH variations between the overlying soils and  
339 underlying groundwater systems. Here we interpret Holocene changes in RIAN in the  
340 HS4 stalagmite record as primarily reflecting local changes in precipitation regime  
341 which control the pH of the soil and groundwater systems. Specifically, we suggest  
342 that RIAN records increases in groundwater pH in response to increased rainfall,  
343 whereby microbes either initially derived from the overlying soils or already present  
344 in the groundwater system modified their membrane lipids to adapt to higher pH in  
345 the groundwater system resulting from increased rainfall and soil leaching.

346 From the above, hydrological changes during the last 9 ka BP were qualitatively  
347 reconstructed from the HS4 stalagmite using the RIAN proxy (Fig. 3), which we  
348 interpret as reflecting local changes in pH of the groundwater, originally driven by  
349 changes in precipitation. The RIAN record from HS4 reveals two (relatively long)  
350 wetter periods in central China, between 8.8-5.9 ka BP and 3.0-0 ka BP and one  
351 relative dryer period from 5.9-3.0 ka BP (Fig. 3A). Both the HS4 and the Dajiuhu  
352 peatland records reconstruct a dry middle Holocene with decreased precipitation  
353 centered on ca. 5.5, 4.8 and 3.5 ka BP in central China (Fig. 3B, C) and overlap for  
354 two long wet periods between > 9 ka to 6 ka BP and 3 to 0.6 ka BP. These drying  
355 events occur simultaneously with colder RAN<sub>15</sub>-MAATs and heavier  $\delta^{18}\text{O}$  values in  
356 HS4 (Fig. 4) and coeval cold/dry events recognized in a number of global NH and  
357 regional paleoclimate records (Liu et al., 2013; Ljungqvist, 2010; Mayewski et al.,  
358 2004; Owen et al., 2016; Rohling and Palike, 2005). These proxy-inferred  
359 hydrological reconstructions re-enforce the conclusion of Xie et al. (2013) that the  
360 overturn of distinctive cultures in the Neolithic Period to Iron Age in central China (as  
361 observed in the distributions of >1600 prehistoric settlement sites) correlates with wet  
362 or flood episodes.

363

### 364 **3.3 Holocene temperature reconstruction**

365 Temperature changes during the last 9 ka BP were reconstructed from the HS4  
366 stalagmite using the RAN<sub>15</sub> index (Wang et al., 2016). RAN<sub>15</sub> is a novel temperature  
367 proxy based on 3-OH-FA distributions measured in the soils from an altitudinal

368 transect on Shennongjia Mountain located 120 km to the NW of Heshang Cave (Wang  
369 et al., 2016). Higher/lower RAN<sub>15</sub> values (higher/lower ratio of the *anteiso* to *normal*  
370 C<sub>15</sub> 3-OH-FAs) are obtained in soils with cooler/warmer MAATs. The quantitative  
371 correlation between MAAT and RAN<sub>15</sub> is expressed in the following equation (Wang  
372 et al., 2016):

$$373 \text{ MAAT} = 23.03 - 3.03 \times \text{RAN}_{15} \quad (R^2 = 0.51, p < 0.001, \text{RMSE} = 2.6^\circ\text{C}) \quad (3)$$

374 RAN<sub>15</sub> in the HS4 stalagmite varies from 0.79 to 2.14 during the last 9 ka BP,  
375 with the lowest value at ca. 7.4 ka BP and highest value at ca. 0.5 ka BP (Fig. 4A). By  
376 applying equation (3) to the HS4 samples, we obtain RAN<sub>15</sub>-MAAT reconstructions  
377 over the last 9 ka BP (Fig. 4A). The average RAN<sub>15</sub>-MAAT of 18.4°C over the most  
378 recent part of the record (<0.8 ka BP) overlaps with the range of MAATs, ca. 16.2 to  
379 18.7 (av. 17.5°C) measured since 1952 at the nearest meteorological station (Yichang,  
380 located ca. 100 km away) and is very close to the av. MAAT of 18°C measured  
381 directly outside the cave by a temperature logger between 2004 and 2007 (Hu et al.,  
382 2008a). This agreement between reconstructed temperatures and instrumental  
383 measurements increases our confidence in the potential of the RAN<sub>15</sub> proxy.  
384 RAN<sub>15</sub>-MAATs in HS4 vary from 16.5 to 20.6°C (av. 19°C), during the last 9 ka BP,  
385 and broadly follow a long-term trend of declining temperatures in line with declining  
386 solar insolation at 30°N in July (Laskar et al., 2004) (Fig. 4B). The temperature  
387 variation (4.1°C) in our record is larger than the calibration error of the RAN<sub>15</sub> proxy  
388 (RMSE = 2.6°C; Wang et al., 2016). The Holocene Climate Optimum (HCO) shown  
389 in the RAN<sub>15</sub>-MAAT record is from 8 to 6 ka BP, with the highest temperature at ca.

390 7.0 ka BP (Fig. 4A). Superimposed on the orbital-scale Holocene trend are centennial  
391 to millennial scale climate fluctuations of ca. 1 to 2°C (Fig. 4A). Interestingly, the  
392 most recent 0.9 ka BP is distinguished by greater variability with the highest (20.5°C)  
393 and lowest (16.5°C) RAN<sub>15</sub>-MAATs occurring consecutively at 0.6 ka BP and 0.5 ka  
394 BP.

395 Our reconstructed RAN<sub>15</sub>-MAAT follows a similar trend to the δ<sup>18</sup>O record (Hu  
396 et al., 2008b) from the HS4 stalagmite (Fig. 4C ). The high resolution HS4 δ<sup>18</sup>O  
397 record encodes a mixture of temperature and hydrological signals and clearly defines  
398 a series of centennial scale episodes of heavier δ<sup>18</sup>O (dry/cool events) superimposed  
399 on the longer term Holocene trend (Hu et al., 2008b). Although the novel biomarker  
400 based proxy has a relatively low resolution it's worth noting that a number of cooler  
401 episodes observed in the HS4 RAN<sub>15</sub>-MAAT record, centered on ca. 8.2 ka, 3.4 ka  
402 and 0.5 ka BP (little ice age, LIA) occur simultaneously with heavier values in the  
403 high resolution δ<sup>18</sup>O record in HS4. Coeval cooling events are broadly recognized in a  
404 number of global NH and regional (monsoonal) paleoclimate records (Ljungqvist,  
405 2010; Mayewski et al., 2004; Rohling and Palike, 2005). Notably our RAN<sub>15</sub>-MAAT  
406 record is consistent with globally distal δ<sup>18</sup>O ice-core record from Greenland (Johnsen  
407 et al., 2001) (Fig. 4D) and the Northern Hemisphere Holocene stacked temperature  
408 anomalies record (30° to 90°N) (Marcott et al., 2013) (Fig. 4E). Although our  
409 sampling resolution is necessarily low due to biomarker sampling requirements (at  
410 this stage of analytical development), the HS4 age model is well constrained and we  
411 note that these RAN<sub>15</sub>-MAAT maxima and minima at 0.6 ka BP and 0.5 ka BP

412 coincide with Northern Hemisphere (NH) scale warm and cold episodes during the  
413 late medieval warm periods (MWP) and LIA respectively (Ljungqvist, 2010; Mann et  
414 al., 2008; Moberg et al., 2005).

415

#### 416 **4. Conclusion**

417 Hydrological and temperature changes in the middle reaches of the Yangtze  
418 River during the last 9 ka BP were reconstructed using Gram-negative membrane  
419 lipids extracted from the HS4 stalagmite from Heshang Cave, central China. RAN<sub>15</sub> is  
420 a temperature proxy while RIAN is interpreted as qualitative hydrological proxy.  
421 Temperatures varied from 16.5 to 20.6°C during the last 9 ka BP, with a relatively  
422 warm period in the early to middle Holocene (8.0-6.0 ka BP), and then a relative cool  
423 period in the late Holocene. The hydrological record shows two relatively long wet  
424 periods and one relative dry period in central China, 8.8-5.9 ka BP, 3.0-0 ka BP and  
425 5.9-3.0 ka BP respectively. The HS4 Holocene Climatic Optimum (HCO) between  
426 8.0-6.0 ka BP is warmer and wetter than any other period in the Holocene and  
427 supports a conclusion of the seminal review of Monsoon palaeoclimate by An et al.  
428 (2000) that an early Holocene Optimum in Monsoon strength occurred in the middle  
429 and lower reaches of the Yangtze River, China, centered on ca. 6 ka BP. Moreover,  
430 this agrees with an ensemble of 18 different model simulations (Joussaume et al.,  
431 1999) for 6 ka, which all indicate enhanced low level convergence into the monsoon  
432 low over Eurasia, with the summer monsoon flow extending further inland. The  
433 present study demonstrates both the first paleoclimate application of 3-OH-FA based

434 proxies and how such biomarker tools can record independent hydrological and  
435 temperature signals in speleothems.

436

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## 649 **FIGURE CAPTIONS**

650 **Figure 1** The location of Heshang Cave and sample sites. (A) Schematic map  
651 showing the main regional surface drainage, revised after Hu et al. (2008a). The red  
652 star shows the location of Heshang Cave. (B) The view of Heshang Cave entrance  
653 from the opposite site of the Qing River. (C) Sampling locations of HS4 stalagmite  
654 and cave surface sediments. Black solid triangle denotes the location of HS4 which  
655 was collected in 2001 (Hu et al., 2008a). Black solid circles denote the sampling sites  
656 of stream surface sediments; hollow squares denote the sampling sites of cave surface  
657 sediment.

658

659 **Figure 2** Distribution and fractional abundance of hydroxy fatty acid homologues in  
660 (A) cave overlying soils, (B) cave sediments, and (C) HS4 stalagmites.

661

662 **Figure 3** Comparison of the HS4 RIAN record with other local and regional  
663 palaeo-hydrological records. A) Heshang Cave hydrological record inferred from the  
664 RIAN record during the last 9 ka BP. U–Th dating errors (Hu et al., 2008b) are shown

665 on the top of the RIAN curve as red line segments. B) Hydrological record based on  
666 hopanoids biomarkers from the Dajiuhu peatland (Xie et al., 2013), raw data are  
667 shown as triangle symbols, the black line is a locally weighted scatter plot smoothing  
668 with a quadratic polynomial (lowess) using a span of 5%. C)  $IRM_{\text{soft-flux}}$  in stalagmite  
669 HS4. Peaks in  $IRM_{\text{soft-flux}}$  indicate intervals with increased precipitation events (Zhu et  
670 al., 2017).

671

672 **Figure 4** Comparison of the HS4 stalagmite  $RAN_{15}$ -MAAT record with other  
673 time-series and proxy records. A)  $RAN_{15}$  and  $RAN_{15}$ -MAAT record reconstructed  
674 from the HS4 stalagmite. The standard deviation of the  $RAN_{15}$ -MAAT record is  $\pm$   
675  $0.1^{\circ}\text{C}$ . The green star represents the mean annual air temperature (MAAT)  
676 immediately outside the cave, as by a temperature logger between 2004 and 2007 (Hu  
677 et al., 2008a). U–Th dating errors (Hu et al., 2008b) are shown on the bottom of the  
678  $RAN_{15}$ -MAAT curve as red line segments. B) Solar insolation changes at  $30^{\circ}\text{N}$  in July  
679 during the last 9 ka BP (Laskar et al., 2004). C) The  $\text{CaCO}_3$  oxygen isotope record  
680 from the HS4 stalagmite (Hu et al., 2008b). D) Ice core  $\delta^{18}\text{O}$  record from North GRIP  
681 (Johnsen et al., 2001). E) The Northern Hemisphere stacked temperature anomalies  
682 ( $30^{\circ}$  to  $90^{\circ}\text{N}$ ) for the  $5^{\circ} \times 5^{\circ}$  area-weighted mean calculation with its  $1 \sigma$   
683 uncertainty (Marcott et al., 2013).

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