

Temperature oscillations over the last 10,000 years in western Europe estimated from terrestrial mollusc assemblages

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Snail assemblages are used to estimate February and August temperatures during the past 10,000 years in western Europe. We find that a strong warming occurred after the Younger Dryas event, followed by several rapid cooling and warming events. These observations are in agreement with insect and pollen proxy data from the European continent and with estimates of sea-surface temperature from the North Atlantic Ocean as well as with fluctuations of glaciers in western Norway. This study also confirms that terrestrial molluscs can provide reliable climatic data in conjunction with other proxy data.

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Numerous records from the Atlantic Ocean (Duplessy *et al.* 1992; Lehman & Keigwin 1992; Bond *et al.* 1992) and the Greenland ice sheet (Oeschger & Arquit 1991; Johnsen *et al.* 1992) provide a basis for comparison with proxy data from continental sequences in Europe. Those regions seem directly climatically connected, according to Broecker's (1990) theory of the salt conveyor belt. Up to now, the terrestrial data have mainly come from pollen and insect remains found in lacustrine deposits or peat bogs. In other types of environments, molluscs have been shown to yield good ecological and climatic information that can be correlated with biological, physical or chemical climatic signals (Rousseau 1991, 1992; Rousseau *et al.* 1993; Rousseau & Puisségur 1990; Goodfriend 1988, 1991). The purpose of this paper is to present temperature estimates for February and August based on molluscan assemblages from a well-dated composite sequence in Burgundy, France, and to compare the results with independent regional climate data.

Methods and material

Terrestrial molluscs provide environmental information that can be deduced using multivariate analyses (Rousseau 1987, 1990). Molluscan studies have provided quantitative reconstructions of past temperature changes in western Europe using an analogue procedure (Rousseau 1990, 1991; Magnin 1992). Mollusc assemblages collected in samples of about 10 kg of sediment were washed and sieved to extract the shells. The taxa were identified and counted for broken shells applying Lozek's (1964) method.

The modern set of molluscan assemblages corresponds to samples of top soils from which mollusc species were all identified and counted. These modern molluscan assemblages are distributed from northern Scandinavia to southern France (Fig. 1) and correspond to a wide range of vegetation, from high latitude tundra to deciduous temperate mid-latitude forests (Fig. 2). The best modern analogues of the fossil assemblages are selected from the modern database by using the squared chord distance method (Overpeck *et al.* 1985). The meteorological data are taken from a compilation of modern measured values averaged over thirty years (Wernstedt 1972), and from local stations (same time interval) close to modern sampling sites.

The molluscan data set is analyzed using the correspondence analysis (Benzecri & Benzecri 1980). Each element (row or column) I is represented in a J dimensional space by its coordinates on each J variable (k_{ij} ; and $\sum k_{ij} = k_i$), but is also allocated a weight ($= k_i/k$, $k = \sum k_i$, $\sum k = 1$). For example, three assemblages yield the following countings for three species a, b, c :

assemblage 1: $a = 2 - b = 4 - c = 8 - k_i = 14$,
assemblage 2: $a = 4, -b = 8 - c = 16 - k_i = 28$,
assemblage 3: $a = 3 - b = 10 - c = 0 - k_i = 13; k = 55$.

The coordinates (k_{ij}) of each assemblage will be 2/14, 4/14 and 8/14 for the first two and 3/13, 10/13 and 0 for the third. The weight will be 14/55, 28/55 and 13/55.

As this procedure can be used for both rows and columns, superimposable factors are yielded (reference axes of the hyperspace) that allow the simulta-

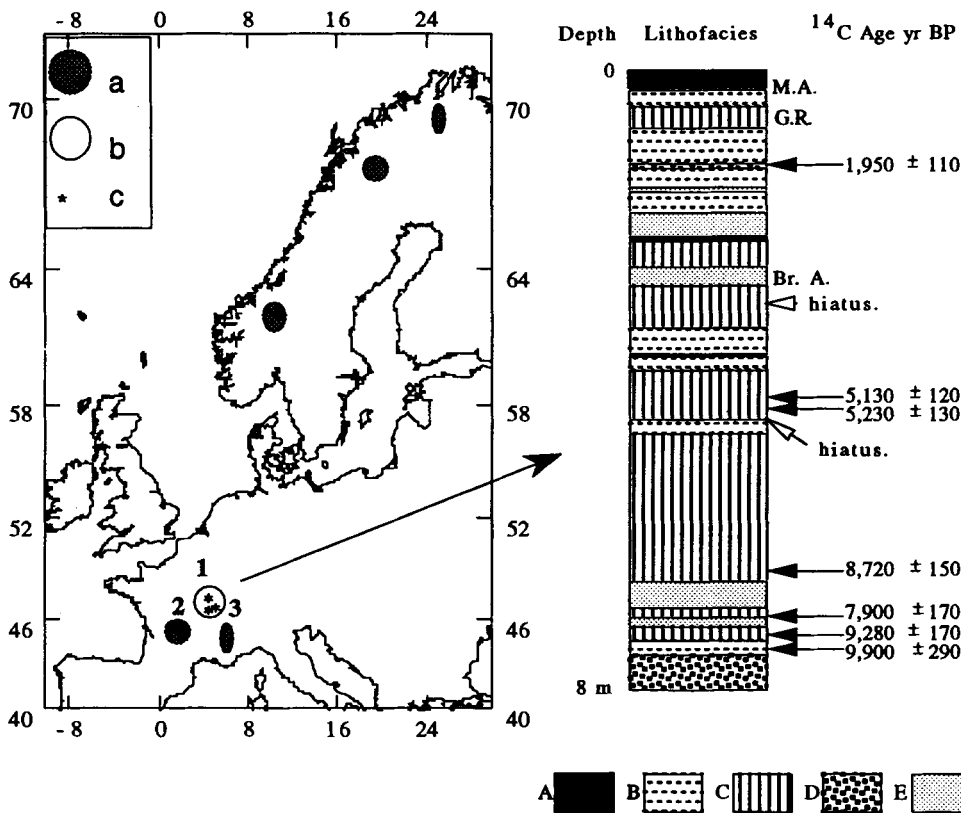


Fig. 1. Location map of (a, b) the modern snail analogue assemblages used for the estimates of palaeotemperatures, and of (c) the fossil sequences. 1: Molesmes, 2: Beaume-lès-Créancey, 3: Cleenay. Lithostratigraphic column: (A) present soil, (B) Clay, (C), (D) indurated tufa, (E) unindurated tufa. M.A. Middle Age, G.R. Gallo-Roman, Br. A. Bronze Age archaeological remains.

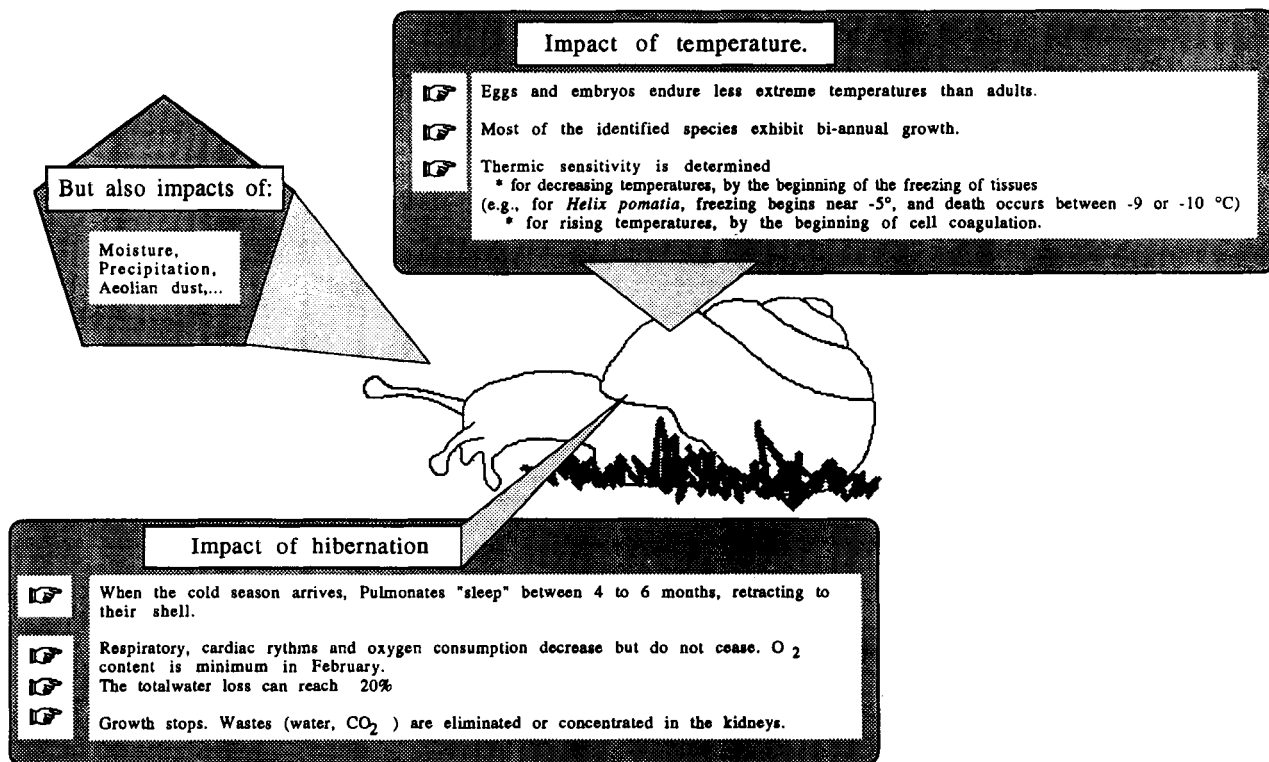


Fig. 2. Main limiting factors constraining the growth of terrestrial molluscs.

neous plotting of both variables and individuals (rows). For our analysis, the mollusc species are columns, and molluscan assemblages, corresponding to a single stratigraphical level, are rows. To avoid too much disturbance in the variability, the data are coded following an abundance class principle (Rousseau 1987).

Correspondence analysis is applied in two steps. The first allows the selection of species which primarily influence the variance of the total data set. If all the species had equal influence on the variance, their contribution would be equal to $1/p$, p representing the total number of species. The selection of predominant species is made using the first seven axes, which usually explain about 50% of the general variability. Species which do not show a high contribution to the variability of the first seven axes are thereby eliminated. The second correspondence analysis (second step of our investigation) is then applied to the reduced data set. Generally, the first 24 factors explain about 90% of the total variance. Stepwise multiple regression between the modern temperature values, used as dependent variables, and loadings of the modern assemblages on these first axes determine the equations of the transfer functions to be applied to the fossil assemblages. The retained coefficients correspond to the highest multiple correlation and the lowest standard error. The analysis of the residuals (real values-estimated values) provides a basis on which to accept or reject the results. Plotting the residuals against the reconstructed values, the scatter plot must show a shape corresponding to a horizontal band if all the conditions necessary for the use of the regression are to be met (Scherrer 1984).

The fossil data come from three ^{14}C dated stratigraphic sequences from the valley bottom in the Burgundy, France. A composite stratigraphy based on these three sites which consists of 42 molluscan assemblages covering the last 10,000 years has been proposed by Rousseau *et al.* 1993 (Fig. 1). All the fossil species analyzed in this Holocene set still exist in assemblages identical to those living during the Pleistocene. The advantage of correspondence analysis is that the modern ecology and distribution of the mollusc species can be used to interpret precisely the general variability of the Holocene data set.

A previous study of this sequence (Rousseau *et al.* 1993) shows that the molluscan assemblages recorded climate events which are in agreement with the classical European Holocene chronostratigraphy. Two steps in Termination I were suggested to occur after the Younger Dryas could event, in agreement with the chronology proposed by Mix & Ruddiman (1985): Termination Ib between 10,000 and 9000 and Ic between 8000 and 6000 years. Termination Ia preceding the Younger Dryas, was however absent from the record. The purpose of the present paper is to provide temperature estimates for the Holocene based on snail assemblages.

Results

Correspondence analysis is first made on a data set comprising 96 species (see list in Table 1) and 132 assemblages (present and fossil). The theoretical contribution of the columns (species) to the variability is then $1/96 = 0.010$. Species with values lower than the threshold on the first seven factors that explain 44.41% of the total variance are eliminated from the data set (Table 1).

The reduced data set comprises 52 columns (species) and 132 rows (assemblages). Using the transfer function procedure for molluscs developed by Rousseau (1991), the first 24 factors are retained. They explain 87.54% of the total variance of the data set. Stepwise regression is then applied to the coordinates of the modern assemblages (total number – fossil number: $132 - 142 = 90$). The largest part of the variance of each predicted variable (February and August temperature) was obtained with partial F ratios equal to 4.

February mean temperature: among the 24 factors, six were selected (Table 2). The transfer function has a high standard error of estimate of 2.18°C and a multiple correlation coefficient of 0.93.

August mean temperature: the standard error of estimates is lower than that of February temperature, 1.31°C, with a multiple correlation coefficient equal to 0.91. Seven factors were selected among the 24 (Table 2).

The equations determined for February and August estimates respectively are indicated in Table 2. The plots of residuals are random, suggesting that the assumptions underlying the application of regression analysis are not violated.

Application of the transfer functions

The composite Burgundian sequence has been time calibrated in accordance with the northwest European chronostratigraphy (Mangerud *et al.* 1974) using ^{14}C dates on organic matter by biostratigraphy and sedimentology. The correspondence between our radiometric dates and the Holocene chronology forms the basis of the proposed time-scale of the sequence (Rousseau *et al.* 1993).

The temperature estimates for the past 10,000 years in Burgundy are plotted, taking into account the stratigraphic position of their corresponding fossil assemblage (Fig. 3). The two main stratigraphic hiatuses previously recognized (Rousseau *et al.* 1993) are also shown. The temperature estimates show less variation relative to present values than those obtained for the pleniglacial interval, i.e. stage 2 (Rousseau 1991; Magnin 1992). At the base of the sequence, the estimates indicate that the Younger Dryas cold event was followed, at 9900 BP, by a rapid warming (Fig. 3). February and August values rose to 1.5 and 17.2°C (Table 3). Cold conditions occurred after this warm

Table 1. List of the species used as variables in the first correspondence analysis. Contributions higher than the theoretical value (10) of the species to the general variability on the first seven factors (F1-F7). The values are expressed in thousands. Variables eliminated for the second analysis are indicated in italics.

| Name | F1 | F2 | F3 | F4 | F5 | F6 | F7 |
|-------------------------------|-------|-----|-----|------|-----|------|------|
| Pomatia elegans | 42 | 14 | 285 | 25 | - | - | - |
| Acicula dupuyi | 18 | - | 119 | 99 | - | 27 | - |
| Carychium minimum | 55 | 287 | 363 | 34 | 31 | 32 | 12 |
| <i>Carychium tridentatum</i> | - | 162 | 19 | - | 51 | - | - |
| Succinea oblonga | - | 32 | 28 | - | 41 | 18 | 11 |
| Succinea putris | 48 | 193 | 517 | 24 | - | 20 | - |
| Succinea sp. | 30 | - | - | 31 | - | 55 | 128 |
| <i>Oxyloma pfeifferi</i> | - | - | - | - | 149 | - | - |
| <i>Azeza goodalli</i> | - | 36 | 48 | - | 82 | - | - |
| Cochlicopa lubrica | 652 | - | 14 | 34 | 31 | 53 | 76 |
| Cochlicopa lubricella | 25 | 40 | - | - | 11 | - | 11 |
| Columella edentula | 50 | 20 | - | 47 | 71 | 45 | - |
| Columella columella | 455 | 176 | 20 | 3641 | - | - | 245 |
| Truncatellina cylindrica | 114 | 288 | 29 | - | 145 | - | 11 |
| <i>Vertigo pusilla</i> | - | 37 | - | 50 | 11 | - | - |
| Vertigo antivertigo | 59 | 155 | 744 | 14 | - | - | - |
| <i>Vertigo substriata</i> | - | 18 | 11 | - | 27 | - | - |
| Vertigo pygmaea | 58 | 70 | 145 | - | 516 | 17 | - |
| Vertigo moulinsiana | 28 | 106 | 140 | 17 | 75 | 11 | - |
| Vertigo genesii | 333 | 12 | 535 | 2239 | - | 1246 | - |
| <i>Vertigo alpestris</i> | 11 | - | - | - | 49 | 209 | - |
| <i>Vertigo angustior</i> | 24 | 73 | 303 | - | - | - | - |
| <i>Orcula dohiolum</i> | 12 | - | 75 | 11 | - | - | - |
| <i>Pagodulina pagodula</i> | - | - | 108 | 14 | - | - | - |
| Abida secale | 293 | 994 | 201 | 13 | 571 | - | - |
| Pupilla muscorum | 120 | 142 | 15 | - | 751 | 18 | - |
| Vallonia costata | 99 | 60 | 68 | - | 279 | 43 | - |
| Vallonia pulchella | 11 | - | 129 | - | 812 | 27 | 16 |
| <i>Vallonia enniensis</i> | 32 | 53 | 500 | - | - | - | - |
| <i>Vallonia excentrica</i> | 65 | 115 | - | - | 266 | - | - |
| <i>Acicula aculeata</i> | - | 39 | 139 | - | 54 | - | - |
| <i>Chondrula tridens</i> | 16 | 37 | - | - | 58 | - | - |
| <i>Ena montana</i> | - | 11 | 21 | - | - | - | - |
| <i>Ena obscura</i> | 23 | - | 191 | 20 | 16 | 17 | - |
| <i>Punctum pygmaeum</i> | - | - | - | 17 | - | - | - |
| Discus ruderas | 235 | 113 | 101 | 368 | - | 22 | 229 |
| <i>Discus rotundatus</i> | - | 159 | 229 | - | 87 | - | - |
| Vitrea pellucida | 1085 | 631 | 268 | 819 | 14 | 41 | 1535 |
| Vitrea subrimata | 52 | 195 | 23 | - | 544 | - | - |
| <i>Vitrea crystallina</i> | - | 162 | 27 | - | 127 | - | - |
| <i>Vitrea contracta</i> | 40 | - | 115 | - | - | - | - |
| Nesovitrea hammonis | 83 | 44 | 114 | 47 | - | - | - |
| Aegopinella pura | - | 42 | 245 | 11 | 127 | - | - |
| <i>Aegopinella nitidula</i> | - | 131 | 77 | - | 89 | - | - |
| Aegopinella minor | - | 52 | - | 11 | 14 | 27 | - |
| Oxychilus cellarius | 39 | - | 299 | 47 | - | 58 | - |
| Oxychilus sp. | - | 95 | - | 16 | 98 | 31 | - |
| Zonitoides nitidus | 41 | 201 | 279 | 26 | 54 | 27 | - |
| Euconulus fulvus | 10007 | 38 | 53 | - | 48 | 34 | 797 |
| <i>Cecilioides acicula</i> | 146 | 138 | - | - | 401 | - | - |
| Cochlodina laminata | 17 | - | 334 | 38 | 50 | 51 | - |
| <i>Macrogastra ventricosa</i> | - | - | - | - | - | - | - |
| <i>Macrogastra rolpheii</i> | 11 | - | 268 | 34 | 18 | 48 | - |
| Clausilia parvula | 228 | 607 | 103 | - | 231 | - | - |
| Clausilia bidentata | 16 | 16 | 223 | 16 | 32 | 22 | - |
| <i>Clausilia dubia</i> | - | - | - | - | - | - | - |
| <i>Clausilia sp.</i> | - | - | - | - | 36 | - | - |
| <i>Clausilia sylv</i> | - | 37 | - | - | 24 | 25 | - |
| <i>Bradybaena fruticum</i> | - | 23 | - | - | 16 | - | - |
| <i>Candidula unifasciata</i> | 91 | 182 | - | - | 368 | - | - |
| Hellicella itala | 144 | 259 | 11 | - | 504 | - | - |
| Hellicella sp. | 28 | 75 | 15 | - | 58 | - | - |
| Trochoidea geyeri | 110 | 399 | 100 | - | 684 | - | - |
| <i>Monacha cartusiana</i> | - | - | - | 42 | - | - | - |

Table 1. (Contd.)

| Name | F1 | F2 | F3 | F4 | F5 | F6 | F7 |
|----------------------------------|------|------|------|------|------|------|------|
| <i>Perforatella incarnata</i> | – | – | 38 | – | – | – | – |
| <i>Hygromia cinctella</i> | – | – | – | – | – | – | – |
| <i>Trichia hispida</i> | 23 | 39 | 35 | – | 102 | – | – |
| <i>Euomphalia strigella</i> | – | – | – | – | – | – | – |
| <i>Helicodonta obvoluta</i> | – | – | 90 | – | 42 | 11 | – |
| <i>Arianta arbustorum</i> | 90 | 23 | 14 | 268 | – | 503 | – |
| <i>Helicigona lapicida</i> | – | 14 | 67 | – | 35 | – | – |
| <i>Cepaea hortensis</i> | – | – | 138 | 17 | – | 16 | – |
| <i>Cepaea</i> sp. | – | 53 | – | – | 20 | – | – |
| <i>Helix pomatia</i> | – | – | 22 | – | 16 | – | – |
| <i>Aegopinella nitens</i> | – | – | 95 | 15 | 12 | 25 | – |
| <i>Cochlostoma obscurum</i> | 36 | – | 351 | 52 | 45 | 76 | – |
| <i>Macrogastra lineolata</i> | – | – | – | – | – | – | – |
| <i>Cepaea nemoralis</i> | – | – | – | – | – | – | – |
| <i>Cochlostoma septemspirale</i> | 133 | 52 | 276 | 47 | – | 37 | – |
| <i>Pupilla triplicata</i> | 184 | 762 | 202 | 14 | 257 | – | – |
| <i>Zebrina detrita</i> | – | – | – | – | – | – | – |
| <i>Chondrina avenacea</i> | 13 | 41 | – | – | – | – | – |
| <i>Pyramidula rupestris</i> | 155 | 663 | 207 | 23 | 1391 | – | 18 |
| <i>Helix aspersa</i> | – | – | – | – | – | – | – |
| <i>Theba plebeia</i> | 13 | – | 66 | 10 | – | 14 | – |
| <i>Phenacolimax major</i> | 56 | 28 | 37 | – | 41 | – | – |
| <i>Lauria cylindracea</i> | – | – | – | – | 31 | – | – |
| <i>Limax</i> sp. | – | 85 | 53 | – | – | – | – |
| <i>Vitrea</i> sp. | – | – | – | – | 14 | – | – |
| <i>Nesovitrea petronella</i> | 1693 | 742 | 256 | 534 | 40 | 311 | 138 |
| <i>Deroceras agreste</i> | 30 | – | 40 | 209 | – | 433 | – |
| <i>Vertigo modesta</i> | 766 | 398 | 225 | 887 | 47 | 5369 | – |
| <i>Vertigo lilljeborgi</i> | 351 | 120 | 13 | 76 | – | 143 | 1793 |
| <i>Euconulus alderi</i> | 101 | 30 | – | 20 | 11 | – | 270 |
| <i>Vertigo ronneybyensis</i> | 145 | 60 | 25 | 78 | – | 99 | 4478 |
| <i>Vertigo geyeri</i> | 63 | 21 | – | – | – | 530 | – |
| % of variance explained | 11.1 | 8.99 | 6.21 | 5.17 | 4.7 | 4.41 | 3.51 |

Table 2. Statistics of transfer functions.

| | Temperature (°C) | |
|----------------------------------|------------------|--------|
| | February | August |
| Multiple correlation coefficient | 0.93 | 0.91 |
| Standard error of estimates | 2.18 | 1.31 |
| Intercept | 1.5 | 17.21 |
| F1 | –3.66 | –1.61 |
| F2 | –3.25 | –2.15 |
| F3 | 0.77 | 0.84 |
| F4 | 0.80 | 0.44 |
| F5 | –2.09 | –1.71 |
| F15 | 1.62 | 0.86 |
| F21 | – | 0.49 |

event during the Boreal chronozone (9000–8000 BP). The temperature estimates for that time go down to 3.5°C below present in February and 3°C in August (Fig. 3). A rapid warming, close to 2°C, took place just after 8000 BP. This warm interval lasted uninterrupted until almost 6000 BP and corresponds to the Atlantic chronozone. At the end of this interval an abrupt and small decrease in temperature is indicated,

marked mainly during winter (Fig. 3). A new cold fluctuation is recorded after the first stratigraphic hiatus at around 5000 BP, with temperatures approximately 3°C less than present-day values for February and August (Fig. 3; Table 3). This event is immediately followed by a new rapid warming of at least 2°C. The interval from 3000 to 2000 BP shows small fluctuations in temperature after the second hiatus (Fig. 3). Estimates for August show values around 2000 BP similar to today (Fig. 3). Both February and August reconstructions show cool values at around 3000 BP (Fig. 3). The reconstructed values for the interval 2000 to around 1000 BP are not taken into account, as they reflect the effect of human impact on the environment in the studied area (Rousseau *et al.* 1993).

Discussion

The trends in the estimates from snail assemblages are in agreement with other proxy data. Temperature estimates from British insect (Coleoptera) assemblages show that an interval as warm as today immediately succeeded the Younger Dryas cold event, and was

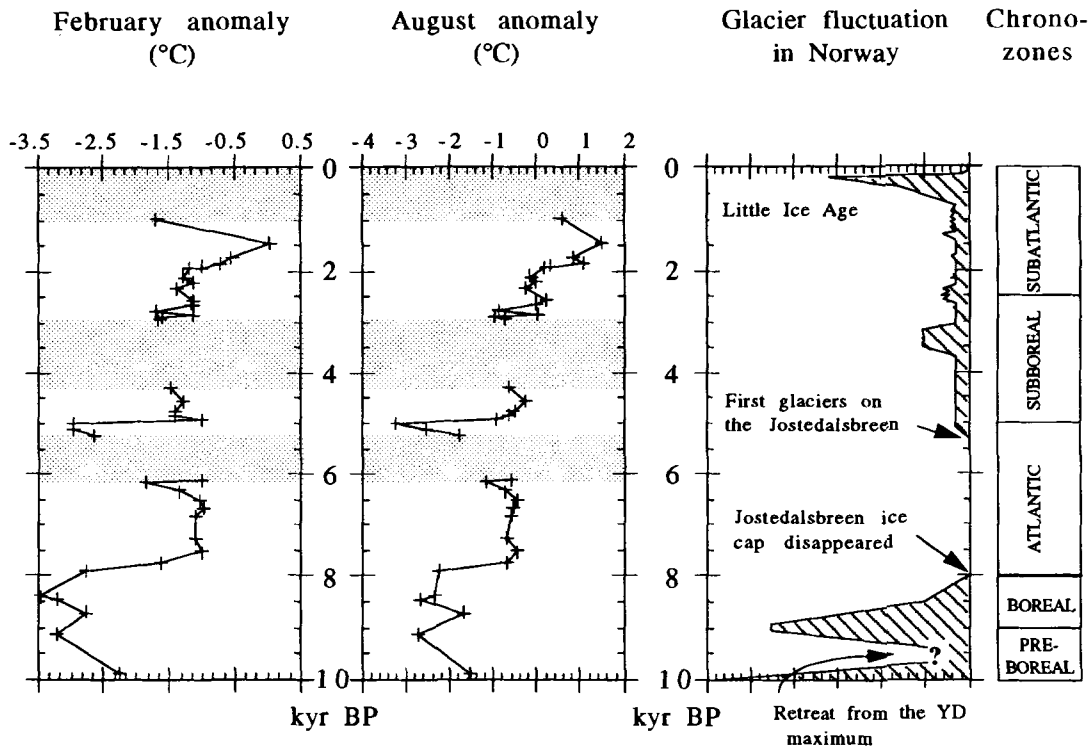


Fig. 3. Reconstructed February and August average temperatures in France expressed as deviations from the modern value. Grey areas correspond to stratigraphic hiatuses previously recognized (Rousseau *et al.* 1993). Fluctuations of Norwegian glaciers (Nesje *et al.* 1991) are also shown.

followed by short cold phases (Atkinson *et al.* 1987; Coope 1987). Estimates based on molluscs also indicate a significant warming, by comparison with values reconstructed for stage 2 in the Achenheim sequence (negative anomalies between -12.7 and -4.2°C in February and -5.3 and -2.7°C in August during the interval lasting from 24,000 to 12,000 BP) which corresponds to late Glacial maximum conditions (Rousseau 1991). This rapid warming, also recorded in Greenland ice cores (Johnsen *et al.* 1992; Taylor *et al.* 1993) has been recognized in both the Antarctic ice sheet (Jouzel *et al.* 1987) and in Southern Ocean (Charles & Fairbanks 1992) cores. This appears to be a global event which is part of the complex Termination I interval, and corresponds to the second major freshwater pulse into the Atlantic Ocean (Fairbanks 1989). Reconstructions from pollen data provide composite proxy data from numerous European localities (Guiot 1987, 1990; Huntley 1990; Pons *et al.* 1987). These studies are mainly centered on the Termination interval. Pons *et al.* (1987), however, show similar general trends to our results with cold fluctuations at around 5000 BP in the annual temperature.

These independent interpretations of terrestrial records are also in agreement with marine records from the Norwegian Sea (Lehman & Keigwin 1992), in which the variation in the percentage of the sinis-

trally coiled *Neogloboquadrina pachyderma* is used to define climatic fluctuations during the last deglaciation. A rapid but short warming of estimated sea-surface temperature occurred after the Younger Dryas cold event, and was immediately followed by a Preboreal cold oscillation (Lehman & Keigwin 1992). Sea-surface temperature then seems to have remained relatively constant from 8800 BP until the present. Moreover, August anomalies from our reconstruction range in the same interval, about 4°C , as do sea surface temperature estimates from core NA 87-22 located off the Rockall Plateau in the North Atlantic (Duplessy *et al.* 1992).

Indirect estimates from geological and paleoecological analyses of lake sediments and terrestrial deposits in western Norway show several glacier fluctuations of the Jostedalsbreen ice cap during the Holocene (Nesje & Kvamme 1991; Nesje *et al.* 1991; Fig. 3). First a district retreat from the Younger Dryas maximum is recorded during the Preboreal. This short warm interval was immediately followed by a new advance of the glaciers between 9500 and 8000 BP, with a maximum at around 9000 BP (Fig. 3). Then, between 8000 and 5000 BP, the ice cap disappeared. The glaciers reappeared just before 5000 BP (Fig. 3). Peaks for extension of glaciers are then recorded between 3000 and 4000 BP, and especially during the Little Ice Age

Table 3. Mean calculated February and August temperatures based on molluscan data from French Holocene sequences. Abbreviations used: MO for Molesmes, B for Beaume-lès-Créancey, CL for Clénay (Fig. 1). Ages come from Rousseau *et al.* (1993) and have been interpolated between ^{14}C dates. Modern February and August temperatures are respectively 3.8° and 18.8°C at Molesmes, 2.8°C and 17.8°C at Beaume-lès-Créancey, and 3.6°C and 18.6°C at Clénay. The ^{14}C date of sample MO4 was used for the time scale of the sequence even if, because of too few terrestrial mollusc individuals, the assemblage itself was not used for the reconstruction.

| Mollusc assemblages | February | August | Age ($\times 1000$ yrs) | ^{14}C dates |
|---------------------|----------|--------|--------------------------|-----------------------|
| M015 | 2.12 | 19.38 | 0.99 | |
| B + 20 | 2.84 | 19.28 | 1.45 | |
| B + 19 | 2.26 | 18.63 | 1.75 | |
| B + 18 | 2.08 | 18.85 | 1.85 | |
| B + 17 | 1.79 | 18.11 | 1.92 | |
| B + 16 | 1.62 | 18.00 | 1.95 | 1950 \pm 110 BP |
| B + 15 | 1.53 | 17.67 | 2.11 | |
| B + 14 | 1.66 | 17.80 | 2.20 | |
| B + 13 | 1.43 | 17.54 | 2.35 | |
| B + 12 | 1.68 | 18.01 | 2.58 | |
| B + 10 | 1.64 | 17.80 | 2.63 | |
| B + 9 | 1.66 | 17.78 | 2.67 | |
| B + 8 | 1.12 | 16.95 | 2.76 | |
| B + 7 | 1.67 | 17.85 | 2.85 | |
| B + 6 | 1.19 | 16.85 | 2.89 | |
| B + 5 | 1.15 | 17.06 | 2.94 | |
| | | | Hiatus | |
| B + 4 | 1.33 | 17.16 | 4.28 | |
| B + 3 | 1.51 | 17.56 | 4.54 | |
| B + 2 | 1.38 | 17.29 | 4.76 | |
| B + 1 | 1.38 | 17.19 | 4.85 | |
| B - 1 | 1.81 | 16.90 | 4.93 | |
| MO11 | 0.84 | 15.60 | 5.02 | |
| MO10 | 0.84 | 16.27 | 5.13 | 5130 \pm 120 BP |
| MO9 | 1.15 | 17.05 | 5.23 | 5230 \pm 130 BP |
| | | | Hiatus | |
| B - 5 | 1.79 | 17.23 | 6.10 | |
| B - 6 | 0.96 | 16.65 | 6.17 | |
| B - 7 | 1.46 | 17.09 | 6.33 | |
| B - 8 | 1.78 | 17.35 | 6.50 | |
| B - 9 | 1.83 | 17.28 | 6.67 | |
| B - 10 | 1.70 | 17.20 | 6.83 | |
| B - 11 | 1.71 | 17.12 | 7.27 | |
| B - 12 | 1.80 | 17.34 | 7.52 | |
| B - 13 | 1.17 | 17.11 | 7.73 | |
| CL15 | 0.85 | 16.35 | 7.90 | 7900 \pm 170 BP |
| MO8 | 0.34 | 16.47 | 8.40 | |
| MO7 | 0.60 | 16.15 | 8.45 | |
| MO6 | 1.05 | 17.12 | 8.72 | 8720 \pm 150 BP |
| MO5 | 0.59 | 16.10 | 9.12 | |
| MO4 | | | | 9280 \pm 170 BP |
| MO3 | 1.54 | 17.25 | 9.90 | 9900 \pm 290 BP |

(Nesje & Kvamme 1991). The re-appearance of glaciers just before 5000 BP implies a sufficient supply in moisture and cool temperatures. This event is in agreement with the small cooling phase determined from mollusc estimates.

Our results demonstrate the complexity of the temperature variation during the past 10,000 years. These observations are also in agreement with the Holocene $\delta^{18}\text{O}$ fluctuations recorded from the Greenland ice

sheet (Oeschger & Arquit 1991; Johnsen *et al.* 1992). The Greenland ice record shows climatic oscillations superimposed on the deglacial trend, the complex Friesland oscillation, following the Younger Dryas, and followed by the so-called climatic optimum.

One can question these interpretations, arguing about the size of standard errors, and if strictly considering these standard errors several oscillations would in fact disappear. But does this mean that they are not real? The comparison of our results with other independent and different proxy data give us confidence, and our purpose is to show general trends in changes of the past environments during the Holocene, and how the molluscan assemblages recorded them.

Conclusions

The quantitative reconstructions of temperature from molluscan communities has been shown to be generally successful in loess environments. The results presented here indicate the usefulness of, records provided by snail communities in another environmental context, and for the relatively short time interval of the Holocene.

The succession of the reconstructed events, as recorded by the molluscs, follows the classical West European framework. After the cold Younger Dryas event, alternating warm and cool events occur just before the climatic optimum of the classical literature (the Atlantic chronozone). These results are in agreement with temperature estimates based on Coleoptera from the British Isles, but disagree with the COHMAP (1988) and other reconstructions indicating warmer summers than today.

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