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## Data-model comparison using fuzzy logic in paleoclimatology

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**Abstract** Until now, most paleoclimate model-data comparisons have been limited to simple statistical evaluation and simple map comparisons. We have applied a new method, based on fuzzy logic, to the comparison of 17 model simulations of the mid-Holocene (6 ka BP) climate with reconstruction of three bioclimatic parameters (mean temperature of the coldest month, MTCO, growing degree-days above 5°C, GDD5, precipitation minus evapotranspiration,  $P - E$ ) from pollen and lake-status data over Europe. With this method, no assumption is made about the distribution of the signal and on its error, and both the error bars related to data and to model simulations are taken into account. Data are taken at the drilling sites (not using a gridded interpolation of proxy data) and a varying domain size of comparison enables us to make the best common resolution between observed and simulated maps. For each parameter and each model, we compute a Hagaman distance which gives an objective measure of the goodness of fit between model and data. The results show that there is no systematic order for the three climatic parameters between models. None of the models is able to satisfactorily reproduce the three pollen-derived data. There is larger dispersion in the results for MTCO and  $P - E$  than for GDD5. There is also no systematic relationship between model resolution and the Hagaman distance, except for  $P - E$ . The more local character of  $P - E$

has little chance to be reproduced by a low resolution model, which can explain the inverse relationship between model resolution and Hagaman distance. The results also reveal that most of the models are better at predicting 6 ka climate than the modern climate.

### 1 Introduction

One of the aims of the Paleoclimate Modeling Inter-comparison Project (PMIP) (Joussaume and Taylor 1995) is to assess the ability of atmospheric general circulation models (AGCMs) to represent a climate different from the present day. Most of the models involved in PMIP are used in simulation of anthropogenic climate change (Kattenberg et al. 1995), and we therefore need to know if those models are able to successfully represent a climate different from the present one. This can be achieved using paleoclimate data. The 6000 yr BP climate has been chosen as a key period for PMIP because it is a simple experiment from a modelling point of view, and it is data rich. Indeed, for this period, several paleoclimatic data sets are now available for different regions of the globe (Wright et al. 1994), and considerable effort has been made to synthesise available data and building comprehensive data sets that can be used for model-data comparisons (Wright et al. 1994; Prentice et al. 1996; Cheddadi et al. 1997; Harrison et al. 1996; Jolly et al. 1998).

Up to now, most model-data comparisons using paleoclimate data (Liao et al. 1994; Prentice et al. 1998; Dong and Valdes 1995; Texier et al. 1997; Masson et al. 1999) have been limited to simple maps or data comparison in a few regions, and only simple statistics have been used. The biome model of Prentice et al. (1992) has also proved efficient in translating model outputs into biomes that can be directly compared with pollen data reconstructions of past vegetation (Texier et al. 1997; Harrison et al. 1998). In these cases, kappa

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statistics (Monserud and Leemans 1992) are used to measure how different the reconstructed vegetation is from the present-day one. As long as only one model is considered and model-data differences are large, obvious mismatches between observations and model output can help identify weaknesses in the model's ability to reproduce natural events. However when several models are considered or when model-data differences are small, there arises the need for synthesising subtle differences at many spatial points, which can only be achieved with an objective measure of the goodness of fit between model and reality. The present study extends the work of Masson et al. (1999) in comparing PMIP model results over Europe with bioclimatic variables (temperature of the coldest month, growing degree days and precipitation minus evaporation) reconstructed from pollen data (Cheddadi et al. 1997), by introducing a new objective measure of the goodness of fit between model and data and using it to classify the model performances.

Preisendorfer and Barnett (1983), have pointed out that a measure of the goodness of fit between model and reality should be adjustable to allow both a local and global intercomparison, and that it is also needed to assign a measure of the significance to that measure of the fit. Frankignoul et al. (1989) and Braconnot and Frankignoul (1993) demonstrated that it was also very important to include all sources of uncertainties arising either from measurement errors, data sampling, small-scale variability of model forcing in the comparison. There is indeed no need to discuss differences in regions where neither data nor the model outputs are reliable. Evaluation of model performances are intrinsically linked to a specific model or problem, which explains why there is not a universal method for model testing. Instead, different approaches using either parametric or non parametric statistics have been developed in atmospheric and oceanic sciences (e.g. Mielke et al. 1981; Preisendorfer and Barnett 1983; Willmott et al. 1985; Zwiers and Storch 1989; Frankignoul et al. 1989; Braconnot and Frankignoul 1993, 1994). These methods deal with data generated at a large number of grid cells which are spatially autocorrelated, and small samples. This is why prior to model-data comparison, data compression is usually performed using principal component analysis or its derivatives, such as common empirical orthogonal function (EOF) analyses (Duchêne and Frankignoul 1991; Braconnot and Frankignoul 1993, 1994; Frankignoul et al. 1995). In all these above mentioned applications, model outputs and observations were interpolated on a common grid. This procedure is not a problem for present day climate when observations are quite numerous, and when it is possible to have an estimate of the uncertainties in poorly covered data, but it may be one for the sparser networks of paleoclimatic data.

For paleoclimate studies, even with the most complete data set available, the data coverage is poor and not regularly spaced, and is characterised by different spatial scales of variability from local to synoptic. The data used for comparison in this study (Cheddadi et al. 1997) have been interpolated by the authors on a regular grid. Because this interpolation could be influenced by remote points in regions of inadequate sampling, it is difficult to define the reliability of the gridded reconstruction. We have therefore decided to define a method that can work without interpolation on a common grid. Pollen data are taken at the location of drilling, and each model is tested against the data on its own grid.

Ideally, we would also like the goodness-of-fit measure to reflect (i.e. to be larger than) those situations when a model correctly simulates a particular structure in the data but shifted in location, as opposed to those situations when the structure is not simulated at all (e.g. getting an enhanced monsoon, but not exactly the right place versus not getting an enhanced monsoon at all). Point-by-point similarities in the former case (structure present, but geographically shifted) are likely to be disappointingly small. Finally, the method must also be able to take into account the uncertainties of both the pollen-derived variables and model outputs. In our case, the main source of data uncertainties comes from the possibly large tolerance of plants to various climates. For the atmospheric models, the uncertainties arise from initial conditions, internal variability and surface boundary conditions (which includes incomplete experimental design that fixes SSTs and leaves 6 ka vegetation distribution at modern levels). The errors of the initial conditions can be neglected, since each model simulation is 15 y long and the atmosphere loses the memory of the initial conditions in one year. The surface boundary conditions introduce systematic biases but they cannot be considered here and will be part of the mismatch between model and data. The only uncertainty that can be explicitly considered is the internal variability, which can be estimated from the 15-y experiments.

In summary, we will develop a distance-type goodness-of-fit measure able to take into account the difference of resolution between AGCM and data maps, the possible irregularity in the network of data, and the uncertainties in the data and AGCM simulations. Moreover, this measure must not be limited to point-to-point comparisons, but must also allow area-to-area comparisons so that a slight shift in latitude or/and in longitude will not be penalised too strongly in the distance values.

We have adopted a method based on fuzzy logic (Zimmermann 1985) which is well adapted to our problem. The method is described in Sect. 2 and applied to the comparison of PMIP simulations with pollen data over Europe in Sect. 3.

## 2 Method based on fuzzy logic

Fuzzy logic differs from conventional logic in that it aims to provide techniques for approximate rather than precise reasoning. Unlike classical statistics which is based on frequency distributions of random populations, fuzzy logic deals with describing the characteristics of properties by associating intervals of continuous variables with semantic labels. The power of this approach comes from the fact that these semantic intervals can or even should overlap.

As an example, consider the parameter “air temperature”. This parameter can be broken down into four subsets: “cold”, “cool”, “warm”, “hot”. The domain from the smallest to the largest allowable value is called the universe of discourse and is denoted  $X$ . In Fig. 1, we assume that  $X$  is the real interval from 5 to 35°C. Assume that the actual air temperature, say  $x_0$ , is 25°C. We can see that  $x_0$  is included in both the “warm” and the “hot” subset. The difference lies in the degree of membership. This particular value of air temperature  $x_0 = 25^\circ\text{C}$  belongs to the subset “warm” with a degree of membership of 0.8 while it is only 0.2 for the subset “hot”. Moreover,  $x_0$  does not belong either to the subset “cold” or to the subset “cool”. So, the degree of membership of  $x_0$  is equal to zero for both.

Subsets like “cold”, “cool”, “warm” and “hot” are called fuzzy subsets of  $X$  because each element  $x$  of  $X$  may belong to any subset with a degree of membership varying between zero and one. In others words, fuzzy logic is a graduated logic based on the idea of membership function from the universe of discourse  $X$  to the real interval  $[0,1]$ . Each element  $x$  of the universe of the discourse  $X$  is associated with a real number between zero and one giving its degree of membership fulfilment to the subset being considered.

As a contrast, in binary logic, any element  $x$  of  $X$  cannot be at the same time warm and hot and the corresponding degree of membership is one or zero. In fuzzy logic, an ordinary set is called a “crisp” set. To distinguish between fuzzy and crisp concepts, fuzzy subsets will be always denoted with a tilde ( $\tilde{\cdot}$ ).

Assume  $\tilde{A}$  is a fuzzy subset of  $X$  with membership function  $\mu_{\tilde{A}}(x)$  (Fig. 1 illustrates particular triangular cases). It must be emphasised that the universe of discourse  $X$  is a crisp set, often the set of real numbers. The open interval that ranges from the smallest to the largest value of the universe of the discourse is called the “support” of  $\tilde{A}$ . The closed interval consisting of all elements with a degree of membership equal to 1 is called the “core” of the fuzzy subset  $\tilde{A}$ .

Some fuzzy subsets have the empty set as a core. If the core contains only one element this element is called the “pivot” of the fuzzy subset. Thus any fuzzy subset  $\tilde{A}$  is completely defined by its membership function  $\mu_{\tilde{A}}(x)$  which involves both support and core.

### 2.1 Fuzzy numbers

Fuzzy numbers are special fuzzy subsets.  $\tilde{A}$  is a fuzzy number if and only if: (1) the universe of discourse  $X$  is the set of real numbers; (2) at least one element  $x$  of the support has its degree of membership equal to 1 (normal assumption, i.e. the core exists); and (3) the membership function does not have local extrema (i.e. it is assumed to be convex).

Two latter properties limit the shape that a fuzzy number can take: it is always non-decreasing to the left of the core and non-increasing to the right of the core. So, a real number can be seen as a fuzzy number whose support comprises only one element which has a degree of membership exactly equal to 1.

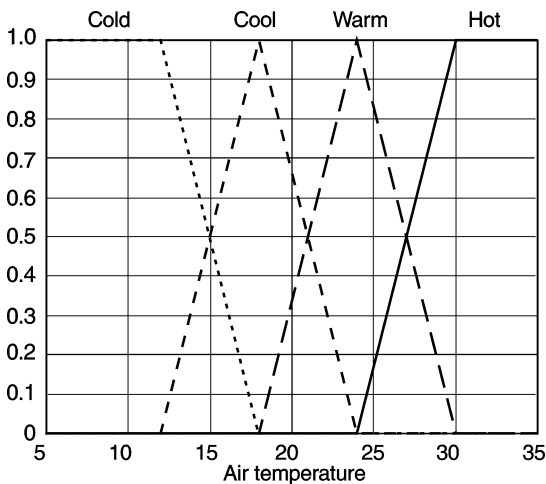
### 2.2 The left-right fuzzy numbers or LRFN

The simplest type of fuzzy number has a triangular or trapezoidal membership function (Fig. 1). A more general class of fuzzy numbers is the left-right fuzzy numbers with curvilinear membership functions (Dubois and Prade 1980).

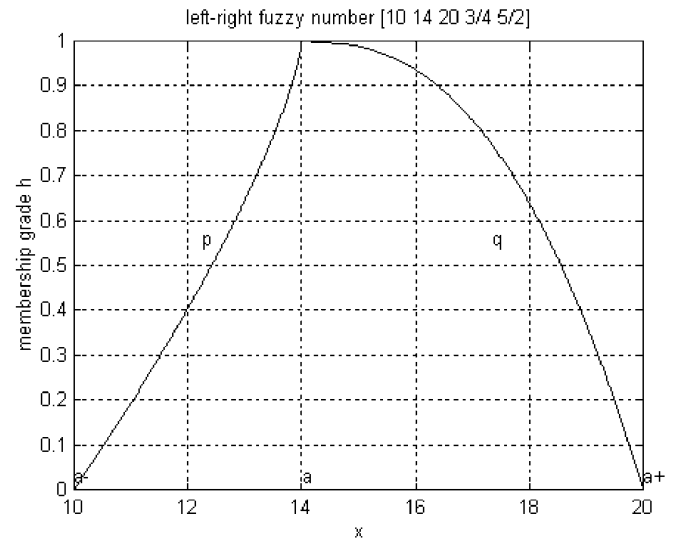
Assume a fuzzy number  $\tilde{A}$  with support  $]a_-, a_+[$  (this kind of bracket indicates an open interval), pivot  $\{a\}$  and membership function  $\mu_{\tilde{A}}(x)$ . This latter can be broken down into a left function denoted  $L_{\tilde{A}}(x)$  and a right function denoted  $R_{\tilde{A}}(x)$  which have a simple analytic form:

$$\mu_{\tilde{A}}(x) = \begin{cases} L_{\tilde{A}}(x) = 1 - ((a - x)/(a - a_-))^{p_A} & x \in ]a_-, a] \\ R_{\tilde{A}}(x) = 1 - ((x - a)/(a_+ - a))^{q_A} & x \in [a, a_+[ \\ 0 & \text{otherwise} \end{cases} \quad (1)$$

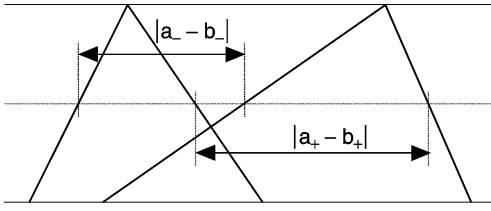
where both exponents  $p_A$  and  $q_A$  are positive real numbers, that determine how sharply the membership function curves are (Fig. 2). The left bracket  $]$  indicates an interval open at left and the right bracket  $[$  indicates an interval open at right. The subscript “ $A$ ” in



**Fig. 1** Four fuzzy subsets to describe air temperature (°C). An element  $x$  in the universe  $X$ , here the real interval 5°C to 35°C, belongs to each fuzzy subset with a degree of membership between zero and one



**Fig. 2** An example of left-right fuzzy number where  $a = 10$ ,  $a = 14$ ,  $a_+ = 20$ ,  $p = 3/4$  and  $q = 5/2$



**Fig. 3** The Hageman distance between two triangular fuzzy numbers

Eq. (1) and anywhere in the current text refers to the fuzzy number  $\tilde{A}$ . Thus relationship Eq. (1) defines left-right fuzzy numbers (LRFN).

The left-right fuzzy number concept allows us to represent not only the data but also the uncertainty in the data. A large value for both exponents gives a “fat” fuzzy number because each  $x$  of its support is thought highly possible (almost total uncertainty). In the opposite case, values less than one produce a “slim” fuzzy number because only the elements  $x$  near the pivot are thought highly possible (almost total certainty). The linear case deals with  $p = q = 1$  (triangular fuzzy number or TFN). Therefore, any LRFN  $\tilde{A}$  is completely defined with five real numbers:  $\tilde{A} \equiv \{a_-, a, a_+, p_A, q_A\}$ .

### 2.3 Distance between two fuzzy numbers (Hageman’s distance)

A degree of membership of a fuzzy number,  $\tilde{A}$ , is denoted by  $h$  and is defined, for all real  $x$ , by  $\mu_{\tilde{A}}(x)$ . Conversely, for each  $\tilde{A}$ , given a level  $h$  in  $[0,1]$ , the corresponding abscissa is  $x = a_-(h)$  for the left side and  $x = a_+(h)$  for the right side. So, we define the squared distance between two fuzzy numbers (Fig. 3) as follows (Bardossy et al. 1990, 1993):

$$D^2(\tilde{A}, \tilde{B}) = \int_0^1 \{ (a_-(h) - b_-(h))^2 + (a_+(h) - b_+(h))^2 \} h dh \quad (2)$$

In short, relationship (2) is a weighted average squared distance between  $\tilde{A}$  and  $\tilde{B}$ . It is very easy to verify that in case of real numbers ( $A$  and  $B$ ). The distance  $D$  is just the usual Euclidean distance,  $|A - B|$ .

Using left-right fuzzy numbers, say  $\tilde{A} \equiv \{a_-, a, a_+, p_A, q_A\}$  and  $\tilde{B} \equiv \{b_-, b, b_+, p_B, q_B\}$ , and denoting  $\delta_A = a - a_-$  and  $\eta_A = a_+ - a$ , the corresponding abscissas follows from Eq. (1):

$$x = \begin{cases} a_-(h) = a - \delta_A(1-h)^{1/p_A} & (\text{left}) \\ a_+(h) = a + \eta_A(1-h)^{1/q_A} & (\text{right}) \end{cases} \quad (3)$$

and similarly for the fuzzy number  $\tilde{B}$ .

From Eqs. (2) and (3), the Hageman distance between two LRFN  $\tilde{A}$  and  $\tilde{B}$  is:

$$D(\tilde{A}, \tilde{B}) = \sqrt{(a-b)^2 - 2I(a-b) + J} \quad (4)$$

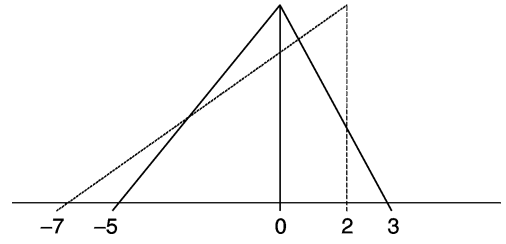
with

$$\begin{aligned} I &= \delta_A f\left(\frac{1}{p_A}\right) - \eta_A f\left(\frac{1}{q_A}\right) - \delta_B f\left(\frac{1}{p_B}\right) + \eta_B f\left(\frac{1}{q_B}\right) \\ J &= \delta_A^2 f\left(\frac{2}{p_A}\right) - 2\delta_A \delta_B f\left(\frac{1}{p_A} + \frac{1}{p_B}\right) + \delta_B^2 f\left(\frac{2}{p_B}\right) + \eta_A^2 f\left(\frac{2}{q_A}\right) \\ &\quad - 2\eta_A \eta_B f\left(\frac{1}{q_A} + \frac{1}{q_B}\right) + \eta_B^2 f\left(\frac{2}{q_B}\right) \end{aligned} \quad (5)$$

while the function  $f$  is defined for any positive real number  $r$  as:

$$f(r) = \frac{1}{(r+1)(r+2)} \quad (6)$$

If we fix the first fuzzy number  $\tilde{A}$ , the pivot and the membership function of  $\tilde{B}$ , then the distance Eq. (4) depends only on the left ( $\delta_B$ ) and right ( $\eta_B$ ) spreads of  $\tilde{B}$ . The minimum of the distance is reached



**Fig. 4** The minimum distance between triangular fuzzy number  $(-5, 0, 3, 1, 1)$  and triangular fuzzy number of pivot 2 is reached for left spread = 9 and right spread 0, giving the TFN equal to  $(-7, 2, 2, 1, 1)$ : representation of these numbers (in fact the right spread cannot be negative)

for  $(\delta_B^*)$  and  $(\eta_B^*)$  by solving:

$$\frac{\partial D}{\partial \delta_B} = 0 \text{ and } \frac{\partial D}{\partial \eta_B} = 0 \quad (7)$$

Assuming  $b > a$ , the corresponding fuzzy number  $\tilde{B}^*$  is then  $\{b^*, b, b^*, p_B, q_B\}$  with spreads (constrained to be positive or null) given as:

$$\begin{aligned} \delta_B^* &= \frac{\delta_A f\left(\frac{1}{p_A} + \frac{1}{p_B}\right) - (a-b)f\left(\frac{1}{p_A}\right)}{f\left(\frac{2}{p_B}\right)} \\ \eta_B^* &= \frac{\eta_A f\left(\frac{1}{q_A} + \frac{1}{q_B}\right) + (a-b)f\left(\frac{1}{q_B}\right)}{f\left(\frac{2}{q_B}\right)} \end{aligned} \quad (8)$$

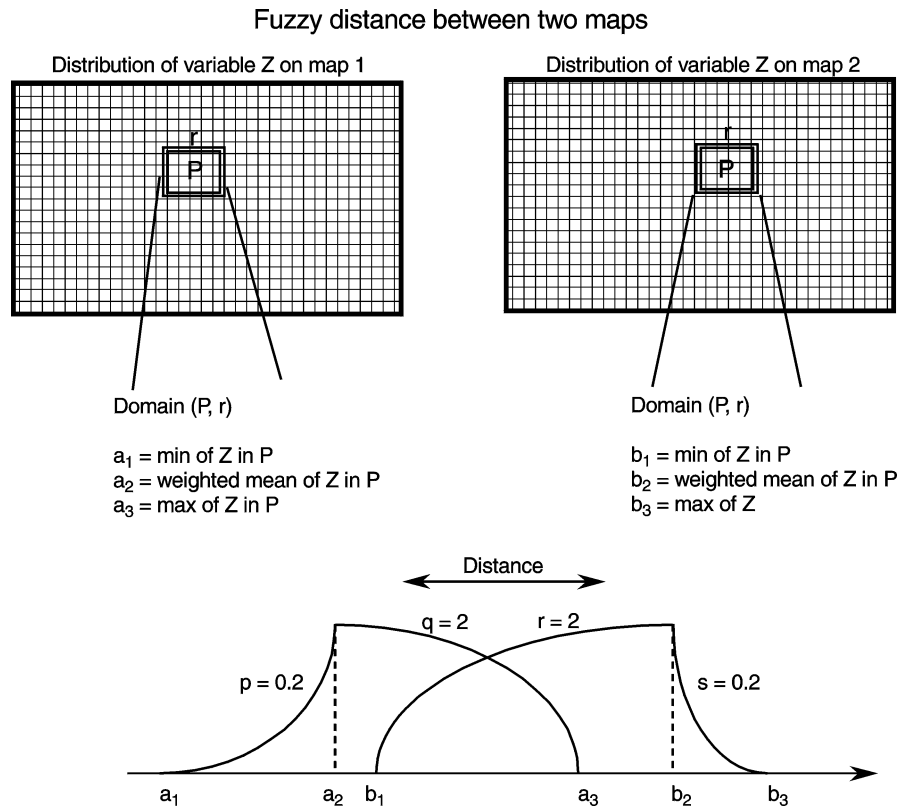
In the triangular case (TFN)  $p_A = q_A = p_B = q_B = 1$ , and we find from Eqs. (4), (5) and (6) that  $D(\tilde{A}_{TFN}, \tilde{B}_{TFN}^*) = (|a-b|)/\sqrt{3}$ . That means that, by taking into account of the uncertainty in the variables, the minimum fuzzy distance reduces to the  $1/\sqrt{3}$  of the classical Euclidean distance. An example is given in Fig. 4: the Euclidian distance is 2 and the minimum Hageman distance is  $2/\sqrt{3}$ . In brief, the Hageman distance tells us that two numbers are closer than it appears in first approximation if we take into account the degree of fuzziness associated with them.

### 2.4 Implementation of the method

The distance between two maps with different resolution, distribution of grid points, and errors, is defined using the iterative approach described (Fig. 5).

1. For each point  $P$  of coordinates  $(x, y)$  on the first map (assumed to be the map to check) representing variable  $z$ , we delimit a square of dimension  $r$ ; we determine the minimum and maximum of  $z$  in this domain; the support of the corresponding fuzzy number is delimited by the minimum and the maximum reached on this square domain; the pivot is defined as the weighted average of the variable  $z$  on this domain, the weights being the inverse Euclidian distance between the point  $(x, y)$  of the checked map and the points belonging to the domain (if this distance is zero, we arbitrary set it to 0.01).
2. We delimit in the second (reference) map the same square of dimension  $r$  around the point of the same coordinates; the pivot will be the weighted average of all the points within the square, the weights being given by the inverse Euclidian distance between the points of the domain and  $(x, y)$ ; the support is given by the minimum and maximum values of  $z$  over the domain.
3. The left and right spreads can be enlarged to account for the uncertainties of  $z$  on one or both maps. In this case, the support is given by the difference between the maximum of the higher values of the confidence intervals and the minimum of their lower values.
4. In order to emphasise the situation where the two fuzzy numbers overlap even partly, in opposition to the situation where they are

Fig. 5 Diagram of the map comparison method



not joined, we define non symmetric membership functions where the interior spreads are convex and the exterior ones are concave as indicated in Fig. 5; this is done to give more importance to the part of the spreads which are consistent. This method of selecting the membership functions is rather subjective, but the important point is to use the same procedure for all the models.

5. The distance is calculated between the two fuzzy numbers and the operation is repeated for each point of the first map.

The method is objective and does not require any special hypotheses to be implemented. In particular decisions about size  $r$  and related distance are taken on the basis of Monte-Carlo simulations (see Sect. 3.2). The only subjective point is the choice of the membership functions which rest on the final goal of the analysis (for example, the decision to emphasise matching instead of discrepancy), but the consequence is rather limited as long as the same procedure is applied to all the models, the main objective being here to compare these models together.

2.5 A simple application

A simple example is provided here to help to understand the behaviour of the Hagan’s distance. Consider two variables  $z_1$  and  $z_2$ , defined on the dimension  $x$  as follows:

$$z_1 = |x + 3|^{0.5}$$

The second variable  $z_2$  is obtained by offsetting  $z_1$  by 6 units:

$$z_{21} = |x - 3|^{0.5}$$

These two variables are defined over a domain  $[-20, 20]$  and are plotted in Fig. 6. The Euclidian distance between these two curves increases from 0 (for  $x = 0$ ) to a maximum of 6.0 for  $x = \pm 3$  and decreases then towards 0 when  $x$  becomes arbitrarily large in absolute value. The latter situation corresponds to the case of null domain size in Fig. 7.

If we define the fuzzy numbers as indicated in Fig. 5, with a progressively larger domain, the support of the fuzzy numbers increase with the size of the domain. The support averaged over all points of the curves is shown in Fig. 7. The mean distance between the curves (calculated for all the points of the curves) has a more complex behaviour: it decreases from a maximum 1.04 to a minimum of 0.18 for domain size between 10 and 12 and increases again to stabilise at about 0.66 from size 24 (this value is linked to the maximum spread of  $x$  used here).

Figure 8 represents the dependence of these distances on  $x$ . For domain sizes lower than 6, the shape of distance profiles are similar (roughly M-shaped like the Euclidian distance). For domain sizes between 8 and 20, the distance has a peak for  $x = 0$  and decreases towards the extremities and for domain sizes higher than 20, we have spurious secondary peaks at the extremities due to the  $x$  interval limited to  $[-20, 20]$ .

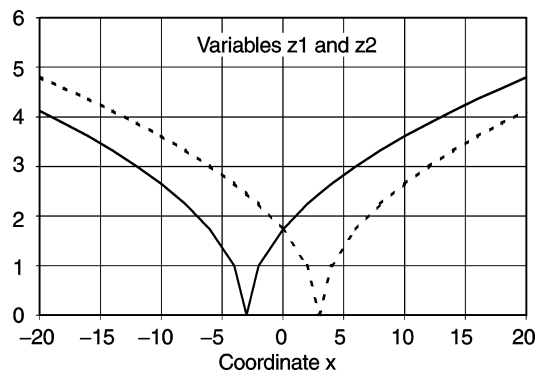
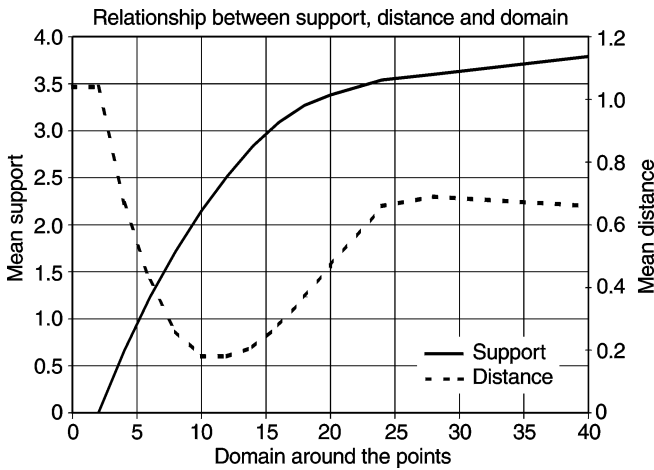


Fig. 6 Variable  $z_1$  and  $z_2$  used for the Hagan distance



**Fig. 7** Evolution of the support and the mean distance in function of the domain size

These profiles show that for small domains, Hagaman distance has a behaviour similar to the Euclidean distance, but for domains comparable to or larger than the typical scale of the phenomena to be represented, Hagaman distance smooths the effects of the pattern shifts smaller than this scale. Domains close to the entire range of the coordinates induce boundary perturbations, so the domain size should be limited to no larger than half the range.

### 3 The application to model-data comparison

#### 3.1 Data and model simulations

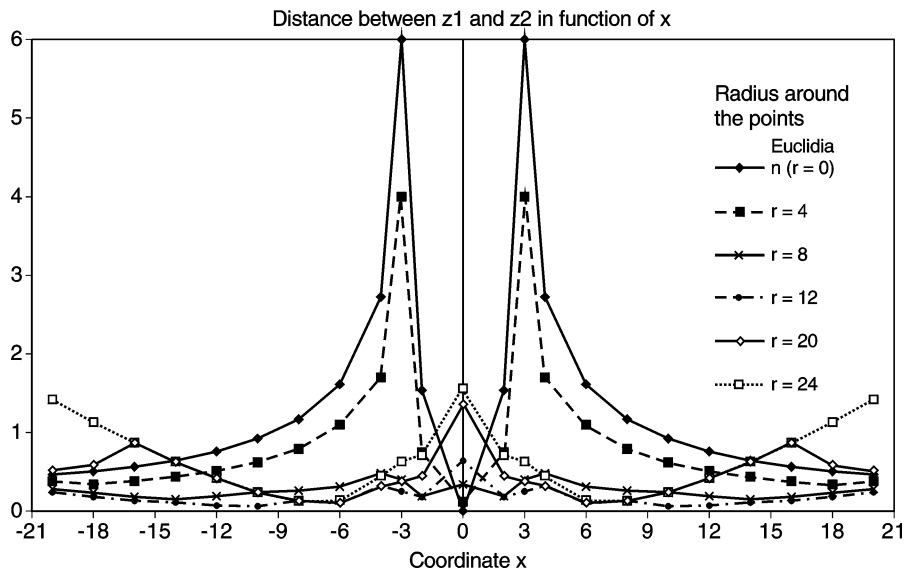
The data used for the comparison are the climatic reconstructions of Cheddadi et al. (1997) for 6000 y BP obtained from pollen data constrained by lake-status indices. The climatic parameters reconstructed are: the temperature of the coldest month (MTCO), the grow-

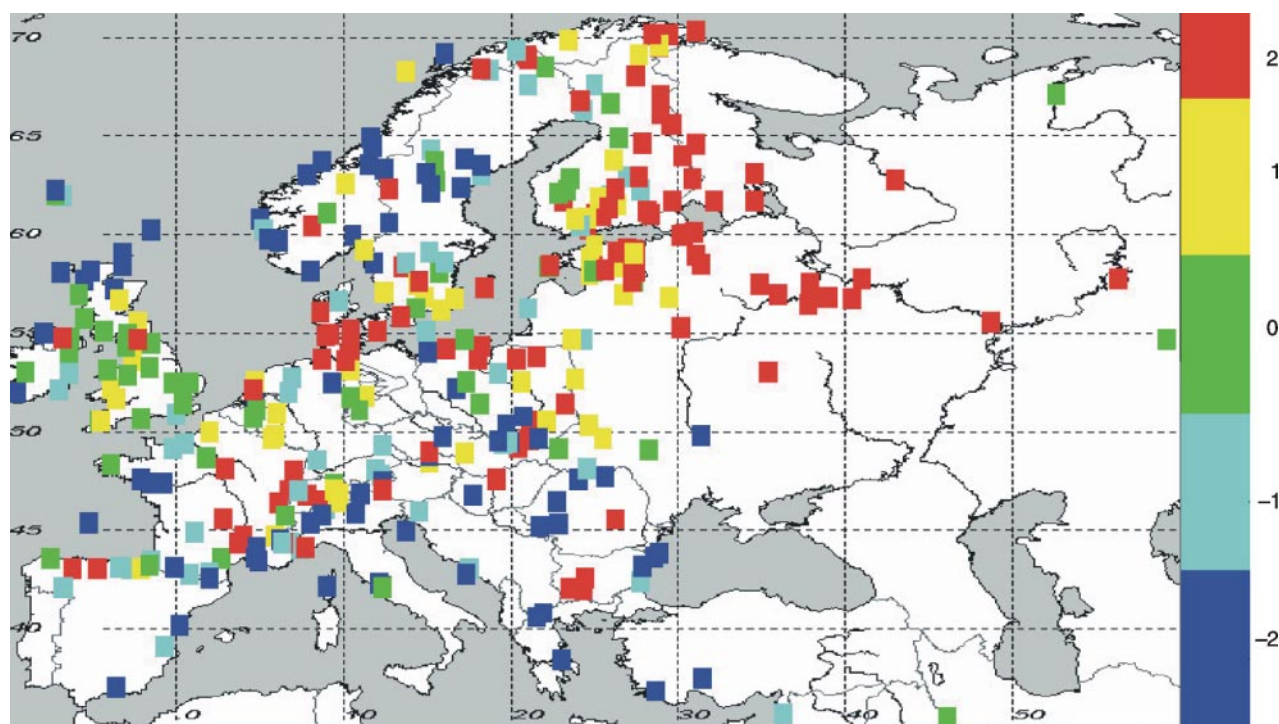
ing-degree days above  $5^{\circ}\text{C}$  (GDD5) and the annual precipitation minus evapotranspiration ( $P - E$ ). The method used for these reconstructions can be summarised as follows: (1) for each 6 ka site, a set of nearest modern samples (analogues) is determined, (2) the value of  $P - E$  anomaly (i.e. deviation from the modern value in the 6 ka site) for each analogue is compared with the status anomaly (i.e. deviation between 6 ka status and modern status) of the closest lake available and the analogues which have a  $P - E$  anomaly with an opposite sign of the lake status are rejected; (3) the reconstructed climate is the average of the climate of the selected analogues; and (4) the error bars of the reconstructions are given by the variability among the climate of the different analogues.

Following Masson et al. (1999), we compare model results with the reconstructed MTCO, GDD5 and  $P - E$ . These parameters can easily be computed from model output. As permitted by the fuzzy approach, we do not use the gridded maps presented in Cheddadi et al. (1997) but rather the reconstructions at the 371 sites (see e.g. in Fig. 9).

Seventeen climate models (Table 1) involved in the Paleoclimate Modelling Intercomparison Project (PMIP) are compared to these data in the region  $35^{\circ}\text{N}$  to  $90^{\circ}\text{N}$  and  $10^{\circ}\text{W}$  to  $60^{\circ}\text{E}$ . The models have a wide range of resolutions and consequently the number of gridpoints belonging to the region studied ranges from 72 (for LMD4) to 525 (GEN2 and ECHAM3). All the simulations of the 6000 BP climate have been performed with exactly the same prescribed changes in boundary conditions relative to the present ones. Orbital parameters have been changed according to Berger (1978), the atmospheric  $\text{CO}_2$  content has been decreased from 345 for modern to 280 for 6000 BP (Raynaud et al. 1993). Seasonal sea-surface temperature and sea-ice extent were held to present-day values. Past

**Fig. 8** Distance between the two variables defined in Fig. 6 for several values of the domain size





**Fig. 9** Anomalies of the temperature of the coldest month (6 k - modern) in Europe, as reconstructed from pollen and lake-levels (Cheddadi et al. 1997)

**Table 1** Characteristics of the 17 models used for the comparison, including the number of gridpoints belonging to the region studied (35°N–90°N, 10°W–60°E)

Model	References	Resolution	Number of points
BMRC	BMRC 12b2 (Australia): Colman and McAvaney (1995)	R21 L9	204
CCC2.0	CCMA version 2 (Canada): Mc Farlane et al. (1992)	T32 L10	270
CCM3	CCM3 (USA): Hack et al. (1994), Bonan (1996)	T42 L18	500
CCSR1	CCSR/NIES 5.4.02 (Japan): Numaguti et al. (1995)	T21 L20	120
CNRM	Arpege-climat version 2 (France): Déqué et al. (1994)	T31 L19	270
CSIRO	CSIRO V4-7 (Australia): Gordon and O'Farrel (1997)	R21 L9	204
ECHAM3	ECHAM 3.6 (Germany): Lorenz et al. (1996)	T42 L19	525
GISS-IIP	GISS AGCM Model II prime: Hansen et al. (1983)	72*46 L9	224
GEN2	GENESIS 2 (USA): Thomson and Pollard (1997)	T31 L18	524
GFDL	GFDL Climate Dynamic Group	R30 L20	486
LMCELM4	LMD4ter (France): Masson and Joussaume (1997)	48*36 L11	72
LMCELM5	LMD5.3 (France): Masson and Joussaume (1997)	64*50 L11	169
MRI2	MRI2 (Japan): Kitoh et al (1995)	72*46	195
UGAMP	UGAMP UGCM version 2 (UK): Valdes and Hall (1994)	T42 L19	500
UIUI11	UI11-PMIP (USA): Schlesinger et al. (1997)	72*46 L14	285
UKMO	UKMO-Unified Model 3.2 AGCM (UK): Hewitt and Mitchell (1996)	96*73 L19	450
YONU	YONU AGCM Tr7.1.1 (Korea): Takioka et al. (1984), Oh et al. (1994)	72*46 L8	210

changes in land surface cover have also been neglected, as requested by PMIP, which can be a major reason for discrepancies between data and models. As shown by Texier et al. (1997), the replacement of tundra by boreal forest in the north of Eurasia contributes to change the effect of surface conditions towards a warming.

Complete maps for all the model simulated changes in the three bioclimatic parameters can be found in

Masson et al. (1999), as well as a discussion of the simulated present-day patterns of these parameters. All the models simulate qualitatively the observed southwest/northeastern temperature gradient of modern MTCO. The largest differences between model and observations are found in the northeastern part of Europe where interannual variability is important. They exceed 5°C only for two simulations.

The north-south gradient of modern GDD5 is also reproduced by all the models. Over northern Europe, the departure between models and observations do not exceed 300 °C days, except for three simulations with a warm bias (more than 2 °C in average) during the growing season. The annual water budget (precipitation minus evaporation) is more difficult to be simulated because it involves sub-grid scale processes. Most of the models overestimate the water budget, as estimated from observation by Masson et al. (1999), up to a factor 4 to 9 in northeastern Europe. Here, to illustrate the model behaviour, we only present data and simulations of MTCO as 90%-level box plots according to the latitude (Fig. 10). The error bars account for the dispersion in longitude. This representation has been chosen because the 6000 BP climate change reconstructed from pollen exhibits a prominent north-south gradient (Fig. 9). All the reconstructed temperatures are negative below 40 °N latitude and most of them are positive above 50 °N. The models with the most similar gradient, at least for the southern part of the map, are ECHAM3, LMD4, YONU, CCSR1 and GFDL. We present also models with opposite gradient (CCC2) or no significant gradient (BMRC, CCM3). Although there is a large diversity of the model results, none of these models are really able to reproduce the magnitude of the changes reconstructed from the data, especially the negative anomalies of the south.

### 3.2 Results

The method developed in Sect. 2 has been applied to compare each model with data over the whole region. The exponents  $p$  and  $q$  are selected as illustrated in Fig. 5, i.e. the right spread of the smallest number and the left spread of the greatest number receive exponent 2 and the two other spreads receive exponent 0.2. From the Hagaman distance computed for each model grid-point, a mean Hagaman distance is obtained for the entire map.

This procedure is repeated from various domain sizes from 1° to 20° (note that low sizes tend toward the classical Euclidian distance and higher sizes tend to smooth the variations in the maps), and at each iteration, a mean distance is calculated. Figure 11 illustrates two examples of the evolution of the distance according to the domain size for MTCO for two models: LMD4, with the lowest resolution, and UKMO, with one of the highest resolutions. The mean distance MD is represented with its 5%-upper limit MD<sub>+</sub>, defined by 2 standard errors around MD (the standard error being the standard deviation divided by square root of the number of data points, say 371). To assess the confidence interval of these distances, the latter is compared to a Monte-Carlo simulation, obtained as follows:

1. We randomly mix the coordinates of the MTCO within both maps and we calculate the mean distance between the two maps;
2. We repeat the process 100 times;
3. Over these 100 pseudo-samples so obtained, we calculate the mean distance and its standard error;
4. We consider the mean minus twice the standard error as a lower random limit (at the 5%-level), denoted RL<sub>-</sub>;
5. Any mean distance is significant at the 95%-level when its MD<sub>+</sub> is less than RL<sub>-</sub>.

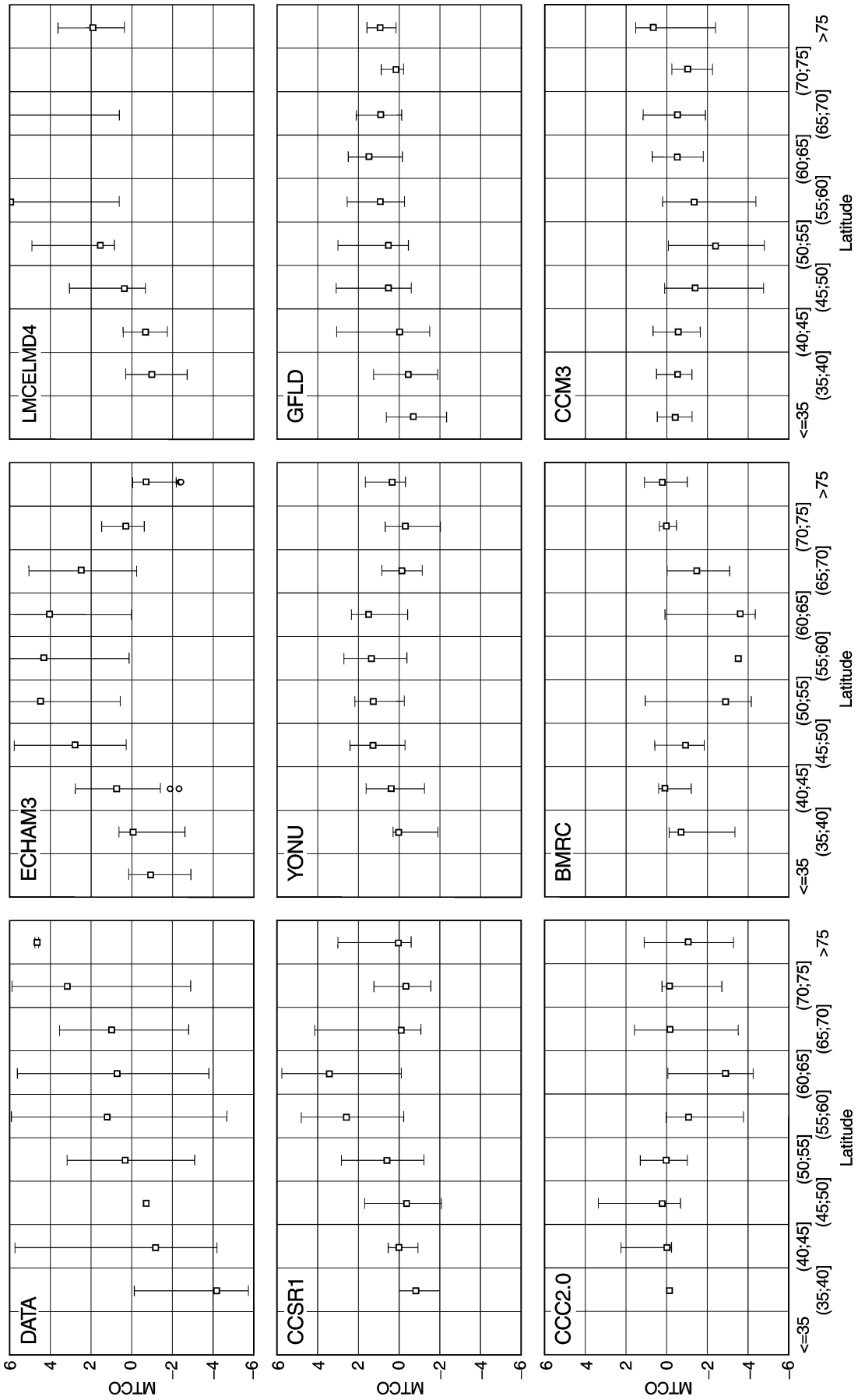
We can see on Fig. 11 that, for LMD4 and UKMO, the first distance significant at the 95%-level is found at a domain size of 8°. However, the distance evolution is chaotic for the low-resolution model whereas it is smoother, with a smaller difference between the mean and the upper-limit for the high-resolution model. For the latter, the signal is clearer.

Finally we retain, for each model and each climatic parameter, the lowest significant domain size and the corresponding mean distance. To facilitate the comparison, we standardise the distances by dividing them by the variance of the spatial distribution of the paleodata. They are represented in Fig. 12 as scatter plots where the ordinate is the distance and the abscissa is the domain size. This figure provides a classification from the closest model to the most distant. All models with distance below 1.1 produce a warming between 50 and 65 north and a gradient towards colder temperature to the south. Models with the largest distance exhibits a reverse north/south gradient. An interesting feature is that the domain size is correlated to the distance: a close model is already close for low domain size, even though the resolution also gives a constraint on this size. Therefore the domain size for the first significant distance comes from a combination of model resolution (the better the resolution the smaller the domain size) and of model data-agreement (the better the agreement the smaller the domain size). For the two other parameters (Fig. 12), the results are somewhat different:

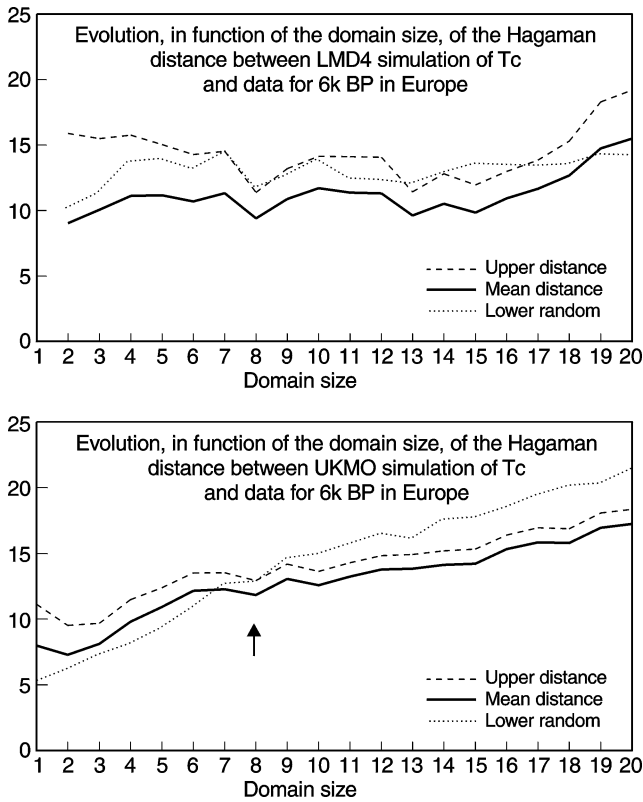
- A. For GDD5, the models are more consistent with a distance ranging between 0.7 and 1.1 (three models have no significant distance and are plotted at domain size 18: LMD4, LMD5, YONU, which are among the lowest resolution ones) and, for the significant models, there is no relationship between the distance and the domain;
- B. For  $P - E$ , the distance is lower (between 0.3 and 1) and, as for MTCO, the domain size tends to increase with the distance.

Considering the relationship between distance and spatial structure of the simulation, the conclusions are obvious. For GDD5, data suggest a cooling south of 50 °N and a warming north of 50 °N (Fig. 10). Most of the models produce a warming over all of Europe, with the exception of LMD4 and LMD5, for which a maximum of warming extends towards the Mediterranean.





**Fig. 10** Box plots of the data of Cheddadi et al. (1997) and simulation by eight selected models of the MTCO parameters, in function of the latitude. The vertical bars represent the 90th percentile interval and the square represents the median calculated on all the gridpoints included in the zonal interval

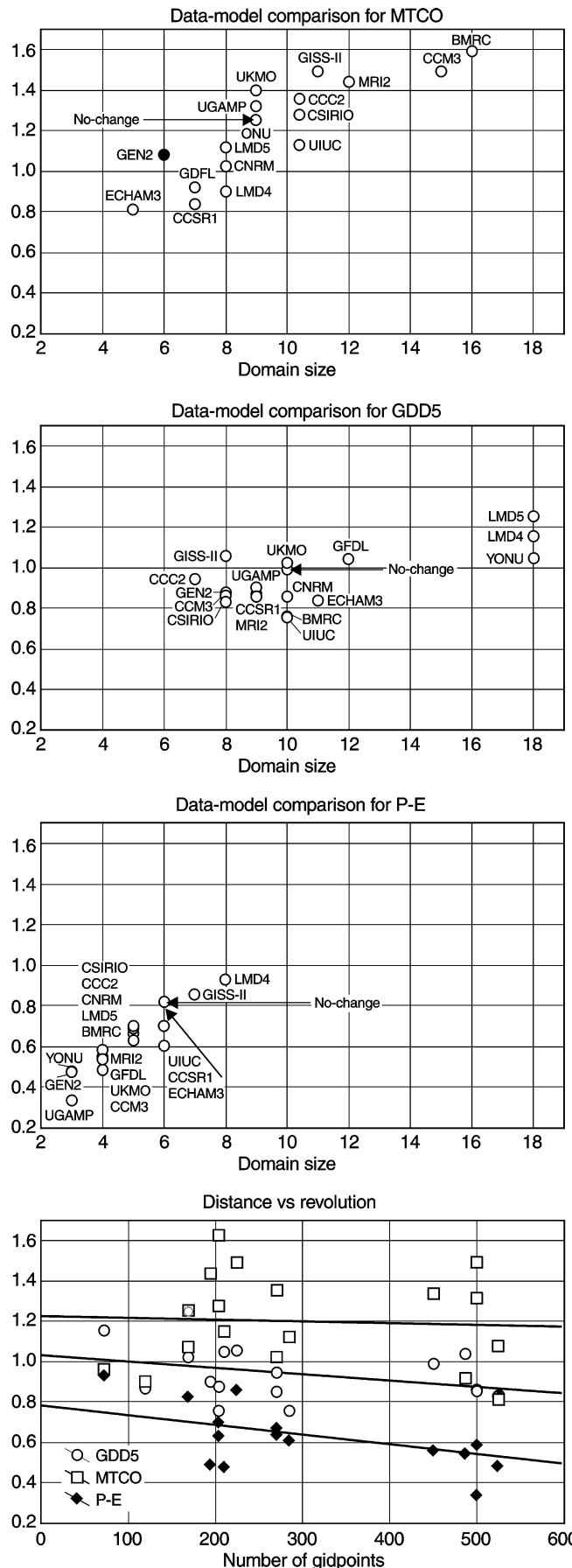


**Fig. 11** Evolution of the distance between data (Cheddadi et al. 1997) and two simulations of two models (with very different resolutions) for the mean temperature of the coldest month (MTCO); the lines with squares represent the mean distance and mean + 2\* standard error calculated on all the gridpoints of the maps between north of 35°N and 10°W 60°E . The dashed line is calculated by Monte-Carlo simulation and represent the 95% significance level

Thus, they have similar Hagan distances. For  $P - E$ , the data show more humid conditions over Europe except in the northwest where dryer conditions prevail. Several models, like UGAMP, GEN2 and MRI2, show a similar pattern: their distance is lower than 0.5 in Fig. 12. Three models (ECHAM3, GISS-II, LMD4) simulate the opposite pattern. These models are located above distance 0.8 in Fig. 12.

If we plot the distance versus the resolution of the model for  $P - E$  (Fig. 12d), it appears, that there is a significant relationship between distance and model resolution ( $r^2 = 0.24$ ): the highest resolution models all have a distance lower than 0.6. For GDD5, the

**Fig. 12** Distance between model simulations and reconstructed data of Cheddadi et al. (1997); three variables are considered: MTCO (the mean temperature of the coldest month), GDD5 (growing-degree days above 5°C) and  $P - E$  (precipitation minus evapotranspiration); The first three graphs represent first significant distance according to domain size and the last one shows the relationship between distance and the resolution of the models (number of gridpoints in the region considered); the distances are standardized by division by the reconstructed data variance



relationship is not significant ( $r^2 = 0.13$ ) and for MTCO, the relationship is null ( $r^2 = 0.01$ ).

Considering model scores for the three different climatic parameters, we cannot conclude that if a model “good” for one parameter it is necessarily also “good” for another one. As an example, ECHAM3 is close for MTCO, intermediate for GDD5 and distant for  $P - E$ .

It is difficult to understand from these distances whether a model is really demonstrating something about the 6 kyr BP climate, or whether in fact any of these models are better than no model at all. In other words, is it better to use a climate model to simulate the 6 ka climate than the present-day climatology? To test this hypothesis, we calculate the distance between the data and a map of null anomalies, called “0-change” in Fig. 12. It appears that the “0-change” is in the middle of all the models for MTCO and, if we exclude the three non-significant models for GDD5, there are only two models poorer than the “0-change” for the two other climatic parameters. Therefore, we conclude that majority of the models analysed are able to simulate the 6 ka climate better than they do for the present climatology.

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#### 4 Conclusions

We have applied a new method based on fuzzy logic to the comparison of PMIP model simulations of the mid-Holocene with a climatic reconstruction from pollen and lake-status data over Europe (Cheddadi et al. 1997). This work extends the comparison of Masson et al. (1999). With the fuzzy logic approach, no assumption is made regarding the distribution of the signal and its error. Also a novelty is that data were taken at the drilling sites and a varying domain size of comparison allows us to work at the best common resolution between observed and simulated maps. In practice, the larger domain sizes for a given parameter are found when the model-data agreement is poor. Three parameters have been tested: MTCO, GDD5 and  $P - E$ . For each parameter and each model, we compute a Haganman distance which gives an objective measure of the goodness of fit between model and data. The mean distance for a map is the global score for the model. On this basis, the different models can be classified. It is however possible to have access to a regional view of the model-data agreement by a map providing the Haganman distance at each grid point of the model grid for the first significant domain size at global scale.

The fuzzy logic implies the use of a membership function of which the shape can be modulated to take into account some *a priori* information. Here, this shape has been chosen to weigh more heavily the part of the intervals of data and model which overlap, but it is easily possible to emphasise, for example, where the part discrepancy is maximum. It is an important prop-

erty of the fuzzy logic to be able to include some *a priori* knowledge in the analysis.

Our results show that there is no systematic order for the three climatic parameters between models. None of the models satisfactorily reproduce the three pollen-derived parameters. To evaluate the quality of these simulations compared to the use of a modern climatology in the model-data comparisons, we have also calculated the distance between the data and a map of null anomalies for the three parameters. Most of the models are better than this climatology for GDD5 and  $P - E$  and half of them also for MTCO data. There is larger dispersion in the results for MTCO and  $P - E$  than for GDD5. GDD5 is an integral of temperature and has less variability than  $P - E$  which is generally noisy in model simulations or than MTCO, which is a winter time temperature highly variable in middle latitudes.

For the temperature parameters, there is also no relationship between model resolution and the Haganman distance. For instance, results of ECHAM3, UGAMP and CCM3, the three T42 resolution model involved in PMIP, cover the whole range of values for MTCO. But they are rather closer to data for GDD5 than the other models. On the other hand, there is an inverse relationship between distance and resolution for  $P - E$  (except ECHAM3 which is the only T42 model which poorly simulates that parameter). That can, perhaps, be explained by the more local character of  $P - E$  which has little chance to be reproduced by a low-resolution model. Integrated parameters such as GDD5 have a smoother spatial distribution and show better consistency between the models. More local parameters such as  $P - E$  need high-resolution models to be simulated adequately.

As far as data-model comparison is concerned, the method described here has the advantage when working on data which have different resolutions and which are not even gridded. It can help to avoid all form of bias induced by spatial interpolation. Another interesting point is the fact that the error bars associated with such maps can be considered. In the absence of probabilistic distributions associated with the Haganman distance, a Monte-Carlo method has been used to assess the significance of this distance.

As this method is based on the distance between intervals, many other applications involving proximity analyses can be envisaged. In particular all the methods based on the research of analogues in the present to explain past assemblages (or in the recent past to forecast the future) can be improved by such Haganman distances. In problems of classification (or cluster analysis) the errors in the data can be taken into account by the replacement of the classical Euclidian distance by the Haganman distance.

When enough data are available for a larger part of the globe, it will be possible to provide an overall evaluation of the models, by averaging the standardised distances. At this stage, the region is too limited and

does not reflect the world situation (some regions have a stronger and more interesting signal, like the East African monsoon). It is too early for that evaluation and we want to avoid the impression of a definitive scoring.

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