

Paleoceanography

Supporting Information for

Millennial-scale Climate Variability Potentially Shaped the Early Interglacial Optimum in Southern Europe

Stéphanie Desprat¹, Gauthier Guillem¹, Maria Fernanda Sánchez Goñi¹, Teresa

Rodrigues^{2,3}, Qiuzhen Yin⁴, Joan Grimalt ⁵

¹ Environnements et paléoenvironnements océaniques et continentaux (EPOC, UMR 5805), Univ. Bordeaux, CNRS, Bordeaux INP, EPHE, PSL University, F-33600 Pessac, France.

² Divisão de Geologia e Georecursos Marinhos, Instituto Português do Mar e da Atmosfera, Rua Alfredo Magalhães Ramalho, 6, 1495-006 Lisboa, Portugal.

³ Centro de Ciências do Mar (CCMAR), Universidade do Algarve, Campus de Gambelas, 8005-139 Faro, Portugal.

⁴ Earth and Climate Research Center, Earth and Life Institute, Université catholique de Louvain, Louvain-la-Neuve, Belgium.

⁵ Institute of Environmental Assessment and Water Research (IDAEA), CSIC, Jordi Girona, 18, 08034, Barcelona, Spain

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							(1.3 interpolation)

Figure S1: Sensitivity test to assess the influence of the interpolation step, windows and smoothing methods used to extract the residuals representing the millennial-scale variations. Black curve with dots: raw MF percentages.

Black bold lines in all plots: MF_orb and MF_hi using a 7 ky Gaussian filter on interpolated data using a 0.3 ky step.

Grey area in all plots: global envelope of the smoothed data and residuals using variable filters (Gaussian, moving average, Lowess and Taner low-pass filter with windows varying from 5 to 10 ky and interpolation from 0.1 to 1.3 ky).

Colored lines: individual smoothing and residuals (cf. legends)

Dashed bold lines: Mean filter and residuals calculated from all results (in MF-orb and MF_hi upper plots) and for results obtained using the different methods.



Window	Interpolation		
— 5 ky — 7 ky — 9 ky	— 0.1 ky — 0.4 ky	— 0.7 ky — 1 ky	— 1.3 ky – – Mean
6 ky 8 ky 10 ky	— 0.2 ky — 0.5 ky	— 0.8 ky — 1.1 ky	— 7-ky Gaussian
	0.3 ky 0.6 ky	0.9 ky 1.2 ky	(0.3 ky Interpolation)

Figure S2: Sensitivity test to assess the influence of the interpolation step on the filters with fixed 7 ky window and of the window on data interpolated at 0.3 ky on the orbital component of the MF data.



Window	Interpolation		
<u> </u>	0.1 ky0.4 ky	0.7 ky1 ky	—1.3 ky Mean
6 ky8 ky10 ky	0.2 ky0.5 ky	0.8 ky1.1 ky	7-ky Gaussian
	0.3 ky0.6 ky	0.9 ky1.2 ky	(0.3 ky Interpolation)

Figure S3: Sensitivity test to assess the influence of the interpolation step on the filters with fixed 7 ky window and of the window on data interpolated at 0.3 ky on the millennial components of the MF data.



Figure S4. Pollen percentages diagram of the interval encompassing MIS 9e from IODP Site U1385. It shows the main pollen taxa only. Pollen zones were determined using constrained hierarchical clustering of the pollen percentage data. Clustering was performed using the CONISS method and implemented with the *chclust* function of the package *rioja* (Juggins, 2009) in the R environment (R Core Team, 2022). Only pollen taxa included in the main sum were selected to construct the Bray-Curtis distance matrix used for clustering analysis. The distance matrix was calculated using the R package *vegan* (Oksanen et al., 2022). The diagram was drawn with the *stratplot* function of the package *rioja*.



Figure S5: Sampling resolution, distance of the pollen assemblages and RoC for the MIS 9 pollen record from Site U1385.



Figure S6. Results of the LOVECLIM simulations presenting the differences in annual, JJA and DJF temperature (°C) and precipitation (cm.year⁻¹) between the MIS 5 and MIS 9 peak simulations (cf. Yin & Berger, 2015). The grey shaded color indicates that difference is significant at more than 95% confidence level calculated using student t-test.