

SUPPORTING INFORMATION

Reconstructing hotspots of genetic diversity from glacial refugia and subsequent dispersal in Italian common toads (*Bufo bufo*)

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Supplementary Figure S2 – SDM projection under LGM bioclimatic conditions

Supplementary Figure S2 – Projections of the species distribution model at the last glacial maximum, for the central lineage (left) and the southern lineage (right), according to the three global circulation models used in this study (CCSM4, MIROC, MPI); binary maps, maxTSS thresholds: central lineage, 0.36; southern lineage, 0.46. Maps were generated in ArcGIS 10.1.



