

## Supporting information

### Population genetic structure of a centipede species with high levels of developmental instability

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**Figure S1.** Bar plot representing the genetic composition of single individuals (thin vertical columns) as inferred from STRUCTURE with  $K=2$ . We applied the LOCPRIOR with admixture model, and the assumption of correlated allele frequencies among populations. The burn-in period was set to 50000, and the run consisted of  $10^7$  iterations.

