Figure S3  Overlapping QTL region on chromosome 17 for freezing to context, freezing to cue and distance traveled in the periphery. LOD scores quantify support for association at SNPs, accounting for differences in genetic sharing using marker-based estimates of pairwise relatedness. Red lines represent 1.5-LOD support intervals, and the dashed orange line shows the significance threshold (at $\alpha = 0.05$) obtained using the simple permutation test.