

Figure S1

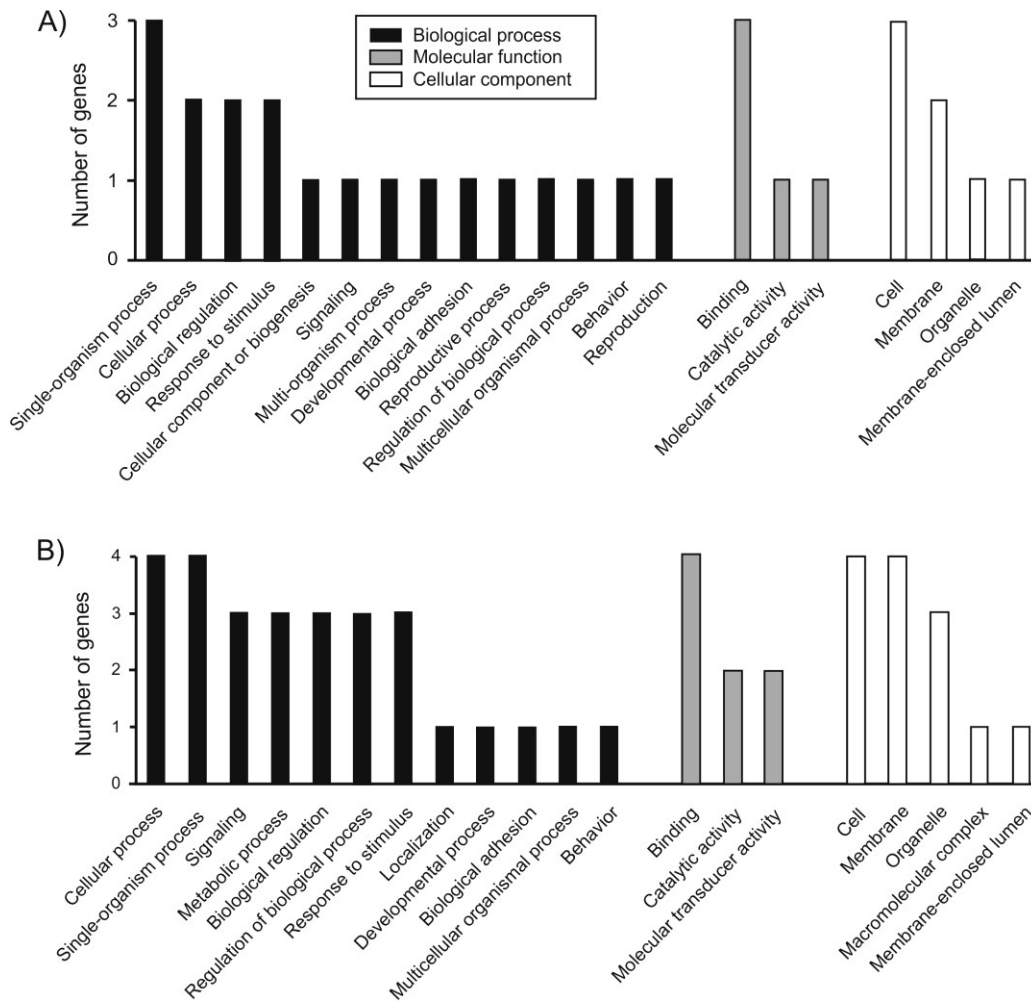


Figure S1. Enriched Gene Ontology terms for candidate genes under selection between (A) East Asian and Arctic breeds and FBDs, (B) modern European breeds and FBDs. The presented terms are significantly enriched at $P < 0.05$. This analysis included genes located within 50 kb distance upstream or downstream of outlier SNPs.