

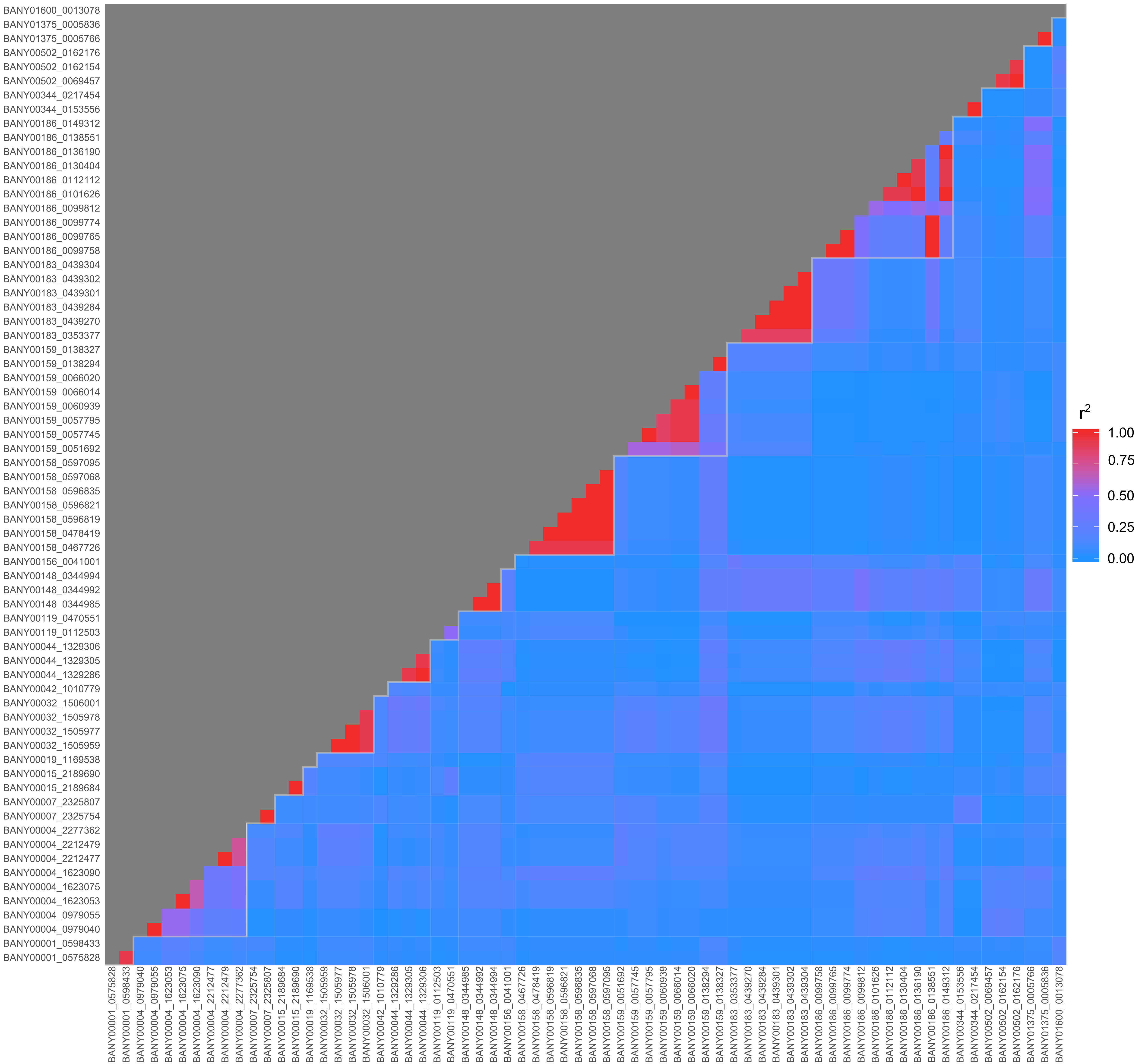


Figure S1. Inter-scaffold linkage disequilibrium matrix for all outlier SNPs. Color scale represents r^2 values product of a pairwise comparison across all individual SNPs. Each SNP is annoated as "BANY00001_0575828", meaning a SNP at position 575,828 bp of the BANY.1.2 scaffold 1, and are ordered according to the BANY.1.2 assembly. Grey lines show the scaffold boundaries in the matrix, with thick grey lines used to show within-scaffold comparisons.