



Figure S15 Histograms of similarities between the genotypes and the expression arrays (the proportion of matches between observed and inferred eQTL genotypes) for each tissue. The panels on the left include self-self similarities (along the diagonal of the similarity matrices); the panels on the right include all self-nonself similarities (the off-diagonal elements of the similarity matrices). Self-self values < 0.8 and self-nonself values > 0.8 are highlighted with red tick marks.