Billings *et. al.* (2021)

Supplemental Figure 2. Pairwise additive relationship values for 6441 combinations of 114 genotypes in Dataset One and Two. The line of best fit (m = 0.70, b = 0.12, r^2 = 0.77) is plotted in orange. Individuals that show high to moderate genetic differences tend to have the largest overall change in IBS between the two datasets.