



Concordance between MAF in SL Tamils and STU after merging all datasets

This plot shows the correlation between minor allele frequencies (MAF) of SNPs in SL Tamils and 1000 genomes STU, who are expected to be genetically identical. SNPs with $F_{ST} > 0.05$ between Sri Lanka and STU were flagged for re-genotyping, including those circled in red (See Supplemental Digital Contents 1 & 5).