mitochondrial_variants

PC2 variance: 21.19%

PC1 variance: 33.97%

GB_ISE_004
GB_ISE_003
GB_ISE_006

MHCO3_P0_L3_n200_01
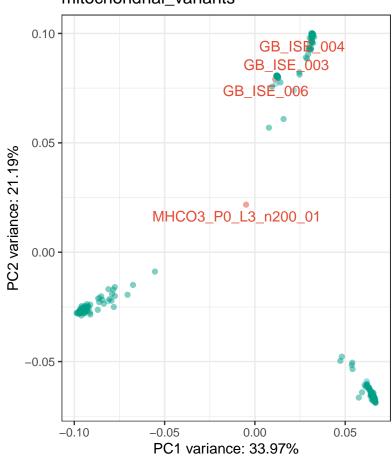
population

a Parent_susceptible

a XQTL_F5_resistant

a XQTL_F5_susceptible