

PCA plot of 1000 Genomes Project data. The x-axis represents PC1 (34.8%) and the y-axis represents PC2 (22.4%). The plot shows genetic variation across different populations, with labels for various genomic regions and their associated p-values. Key populations labeled include GCA, TCG, ACC, and GCC. The plot shows a clear separation of populations along the PC1 axis, with labels for various genomic regions and their associated p-values.

- NC\_072398.2
- NC\_072402.2
- NC\_072404.2
- NC\_072407.2
- NC\_072408.2
- NC\_072409.2
- NC\_072410.2
- NC\_072411.2
- NC\_072412.2
- NC\_072413.2
- NC\_072414.2
- NC\_072416.2
- NC\_072418.2
- NC\_072419.2
- NC\_072420.2
- NC\_072421.2