

PCA plot showing genetic variation across populations. The x-axis is PC1 (39.4%) and the y-axis is PC2 (24.2%). Populations are color-coded: European (blue), African (red), East Asian (green), South Asian (orange), and Admixed American (purple). A large cluster of points is visible on the right side of the plot, with many points labeled with accession numbers and coordinates. A small cluster of points is visible on the left side of the plot. A legend at the bottom right identifies the populations by color.

- NC_072423.2
- NC_072424.2
- NC_072425.2
- NC_072426.2
- NC_072427.2
- NC_072429.2
- NC_072430.2
- NC_072431.2
- NC_072433.2
- NC_072435.2
- NC_072436.2
- NC_072437.2
- NC_072439.2
- NC_072440.2
- NC_072441.2
- NC_072443.2
- NC_072447.2
- NC_072448.2