

PCA plot showing PC3 (7.8%) on the x-axis and PC1 (12.2%) on the y-axis. The plot displays numerous data points, each representing a gene expression profile. The points are colored and labeled with gene names and coordinates, such as '1::NC\_072411.2:14510607-14518467(+)' and '3::NC\_072411.2:14510603-14518436(+)'.

- NC\_072398.2
- NC\_072401.2
- NC\_072402.2
- NC\_072404.2
- NC\_072405.2
- NC\_072407.2
- NC\_072408.2
- NC\_072409.2
- NC\_072410.2
- NC\_072411.2
- NC\_072412.2
- NC\_072413.2
- NC\_072416.2
- NC\_072417.2
- NC\_072418.2
- NC\_072419.2
- NC\_072420.2
- NC\_072421.2
- NC\_072422.2