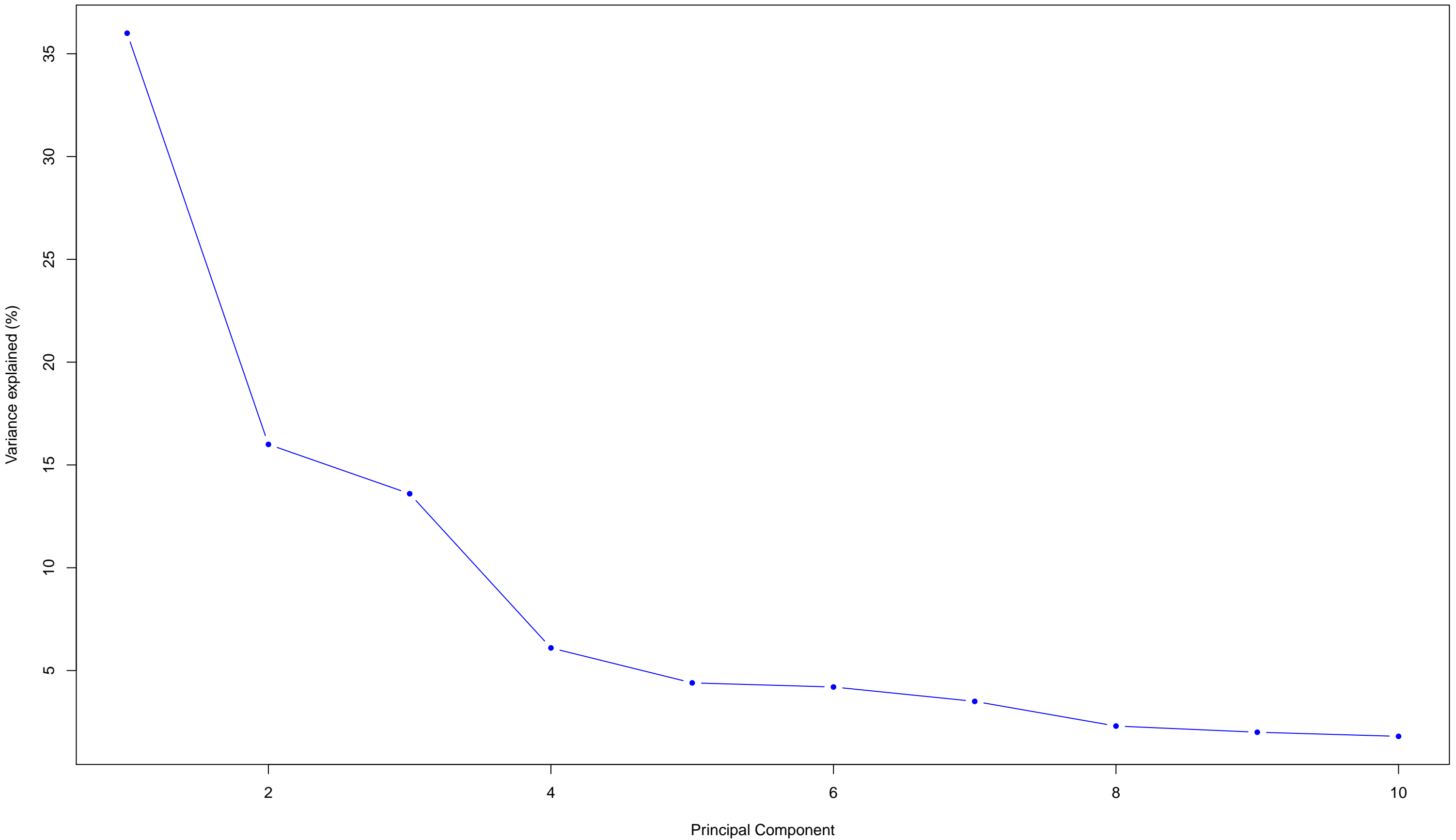


Screeplot



1326Hyp

1336Dtum

1283Dtum

1587Dskin

1587Dtum

1277Dtum

1277Dtum2

1603Gtum

1603Dtum

1603Dtum2

1317Dtum

1317Dtum2

1316Dtum

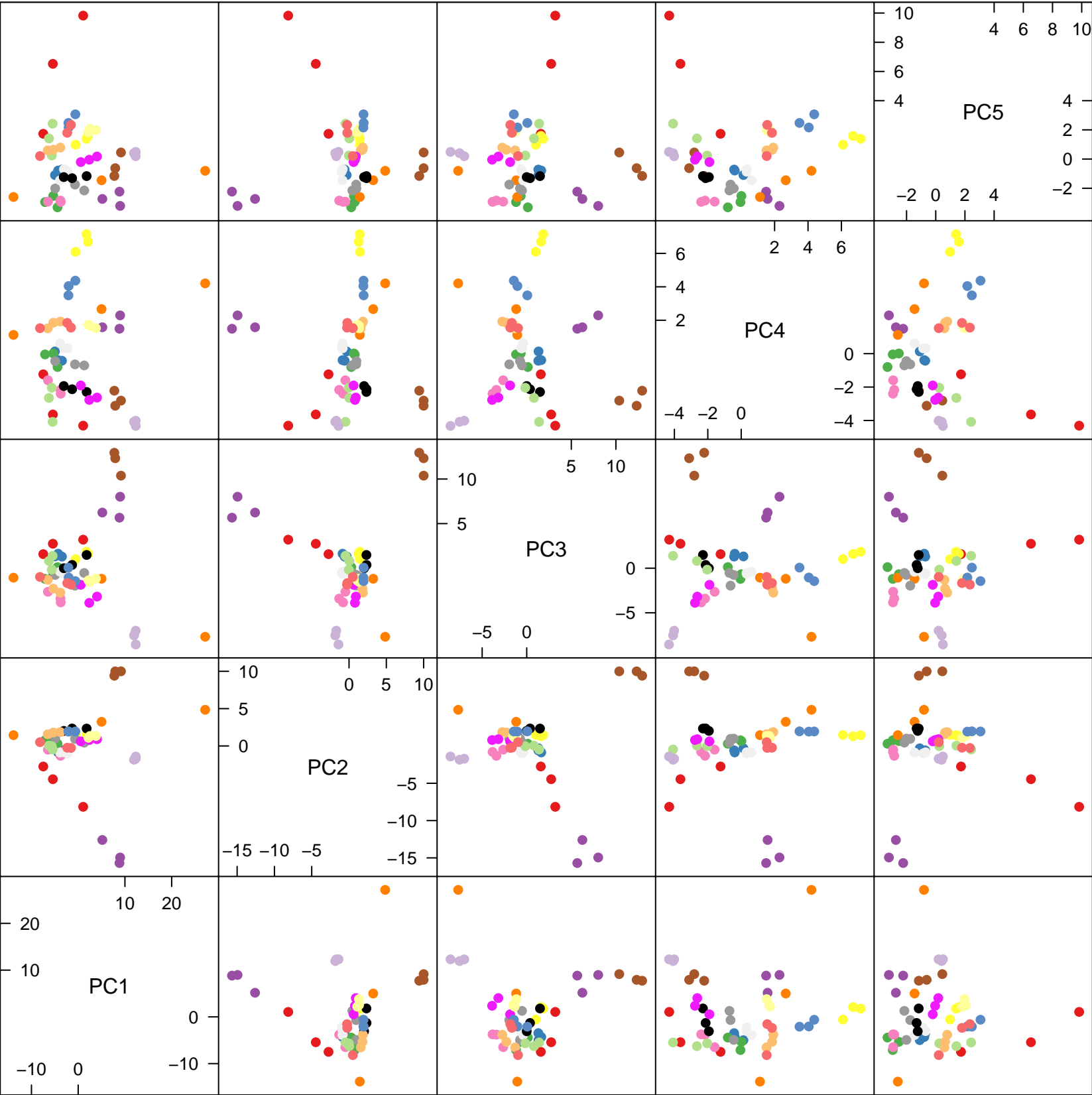
1325Dtum

1325Dtum2

1325Dtum3

1325Dtum4

1325Dtum5



Scatter Plot Matrix

PCA plot showing the first two principal components (PC1 and PC2) of the data. The x-axis is PC1 (36%) and the y-axis is PC2 (22%). The plot displays a large cluster of points representing various samples, with labels indicating specific groups and replicates. The samples are color-coded by group: 1277Dum (orange), 1587Dum (yellow), 1325Dum (purple), 1326Hyp (red), 1587Dskin (purple), and 1316Hyp (blue). The plot shows distinct clustering of samples within each group, with some overlap between groups. The 1277Dum samples are clustered at the top, 1587Dum at the top right, 1325Dum at the bottom right, 1326Hyp at the bottom left, 1587Dskin at the bottom, and 1316Hyp at the top left.

PCA plot showing the first two principal components (PC1 and PC2) of metabolite data. The x-axis represents PC1 (36%) and the y-axis represents PC2 (16%). The plot displays a dense cluster of metabolites, with labels for various compounds such as a-Ketoglutarate, Lysine, Citrulline, Uric Acid, Tyramine, Succinyl-CoA, Glutamine, and many others. The x-axis ranges from -1.0 to 1.0, and the y-axis ranges from -1.0 to 1.0.