Fig S1A PCA analysis of two clusters based on consensus clustering.

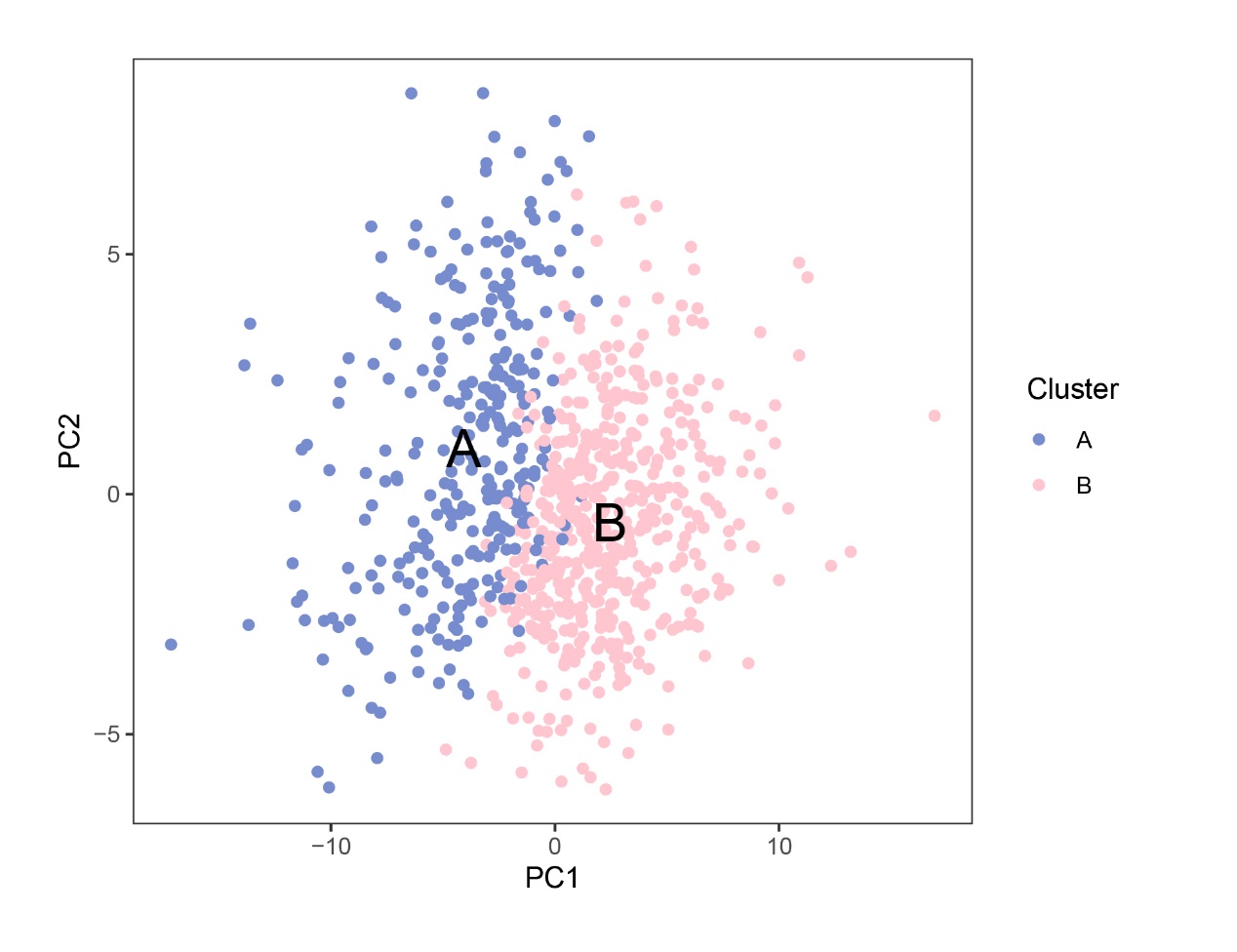


Fig S1B Heatmap shows the relationship between UAS-based molecular clusters and different clinical traits.

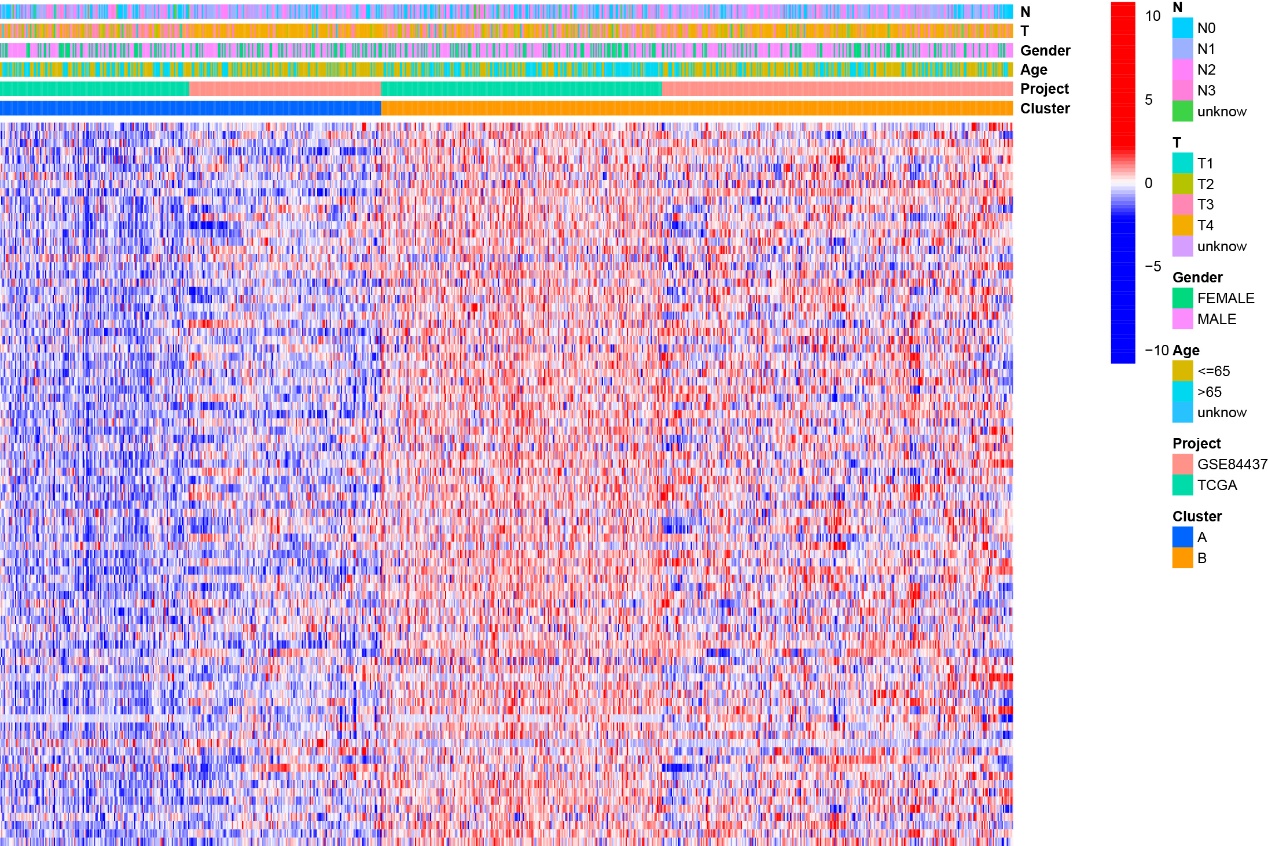


Fig S2A A total of 14 GC samples from GSE167297 were enrolled into single-cell analysis.

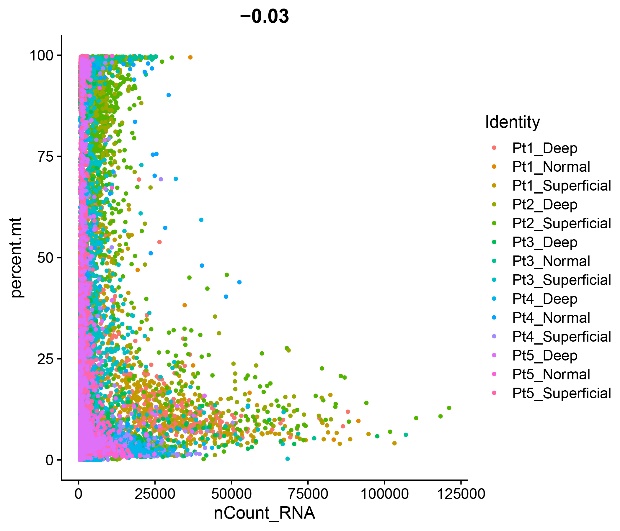
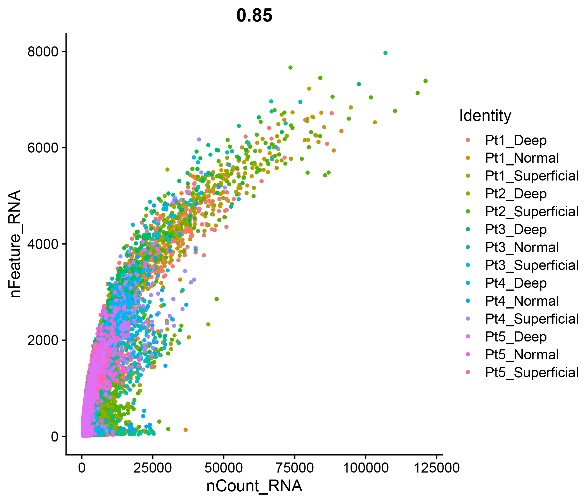
 

Fig S2B PCA analysis unearthed the distribution of all GC case with no significant batch effects.

