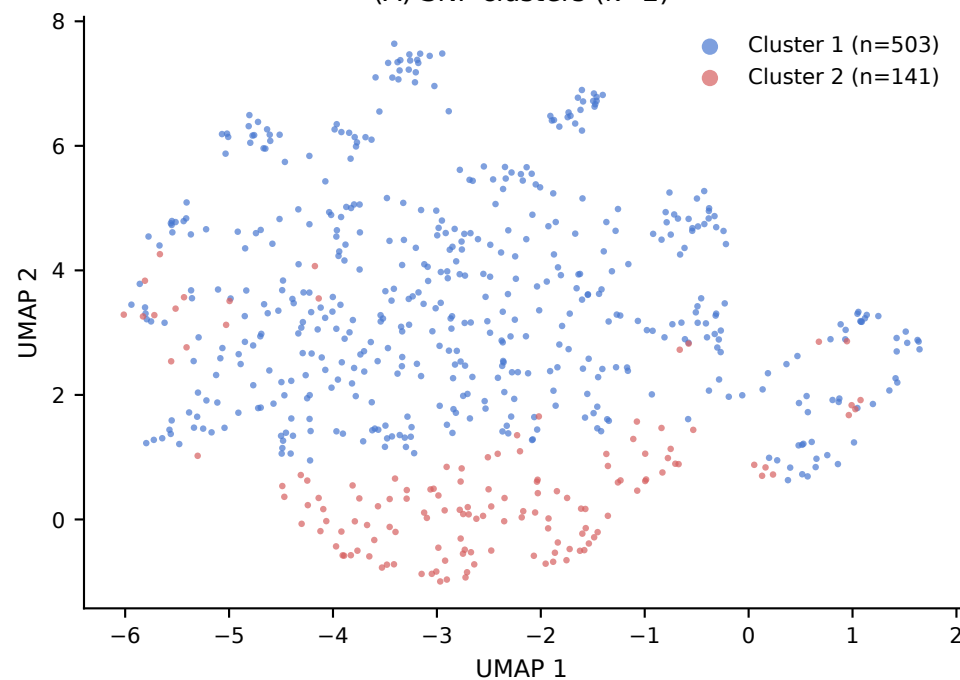
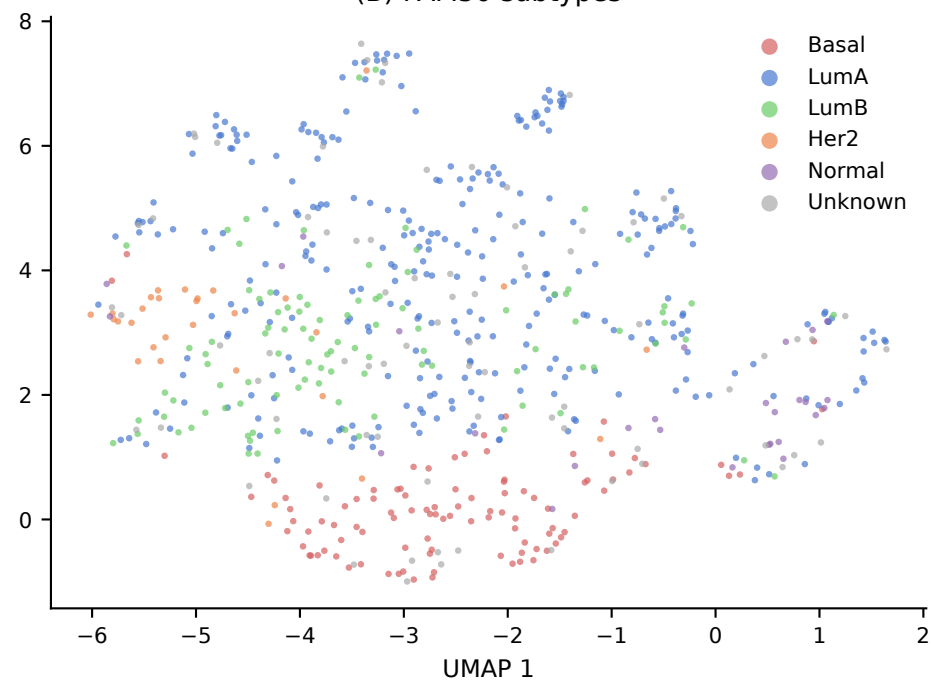


Figure 1 - SNF multi-omics clustering of TCGA-BRCA (n=644)

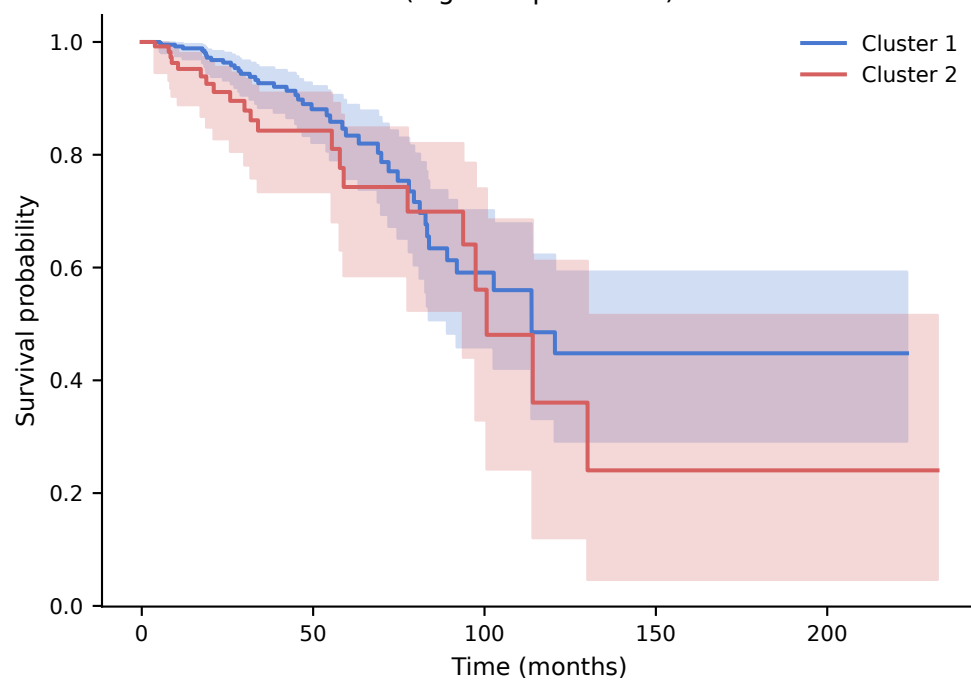
(A) SNF clusters (k=2)



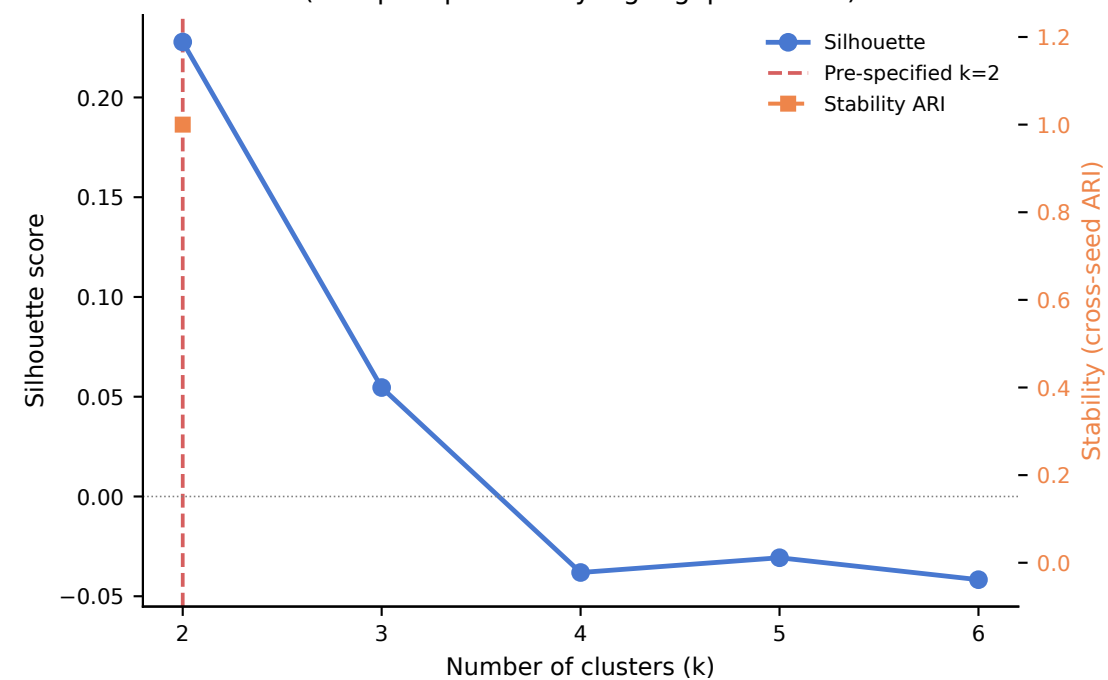
(B) PAM50 subtypes



(C) Overall survival: SNF k=2  
(log-rank p=0.1438)



(D) k-selection: silhouette & stability  
(k=2 pre-specified by eigengap heuristic)



Cluster 2					
At risk	503	384	19	27	21
Censored	0	386	144	145	118
Events	0	123	180	244	2144