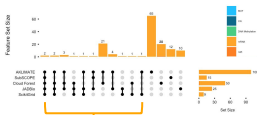
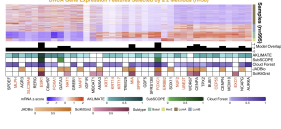


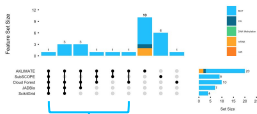
#### A) BRCA Top Model Feature Sets



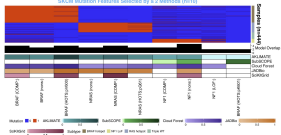
**B)** BRCA Gene Expression Features Selected by a 2 Methods (n=38)



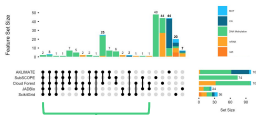
### C) SKCM Top Model Feature Sets



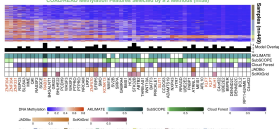
D) SKCM Mutation Features Selected by a 2 Methods (n=10)



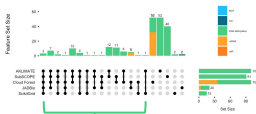
#### E) COADREAD Top Model Feature Sets



**F)** COAD/READ Methylation Features Selected by a 2 Methods (n=58)



#### G) LGGBM Top Model Feature Sets



H) LGG/GBM Methylation Features Selected by  $\alpha_2$  Methods (m62)

