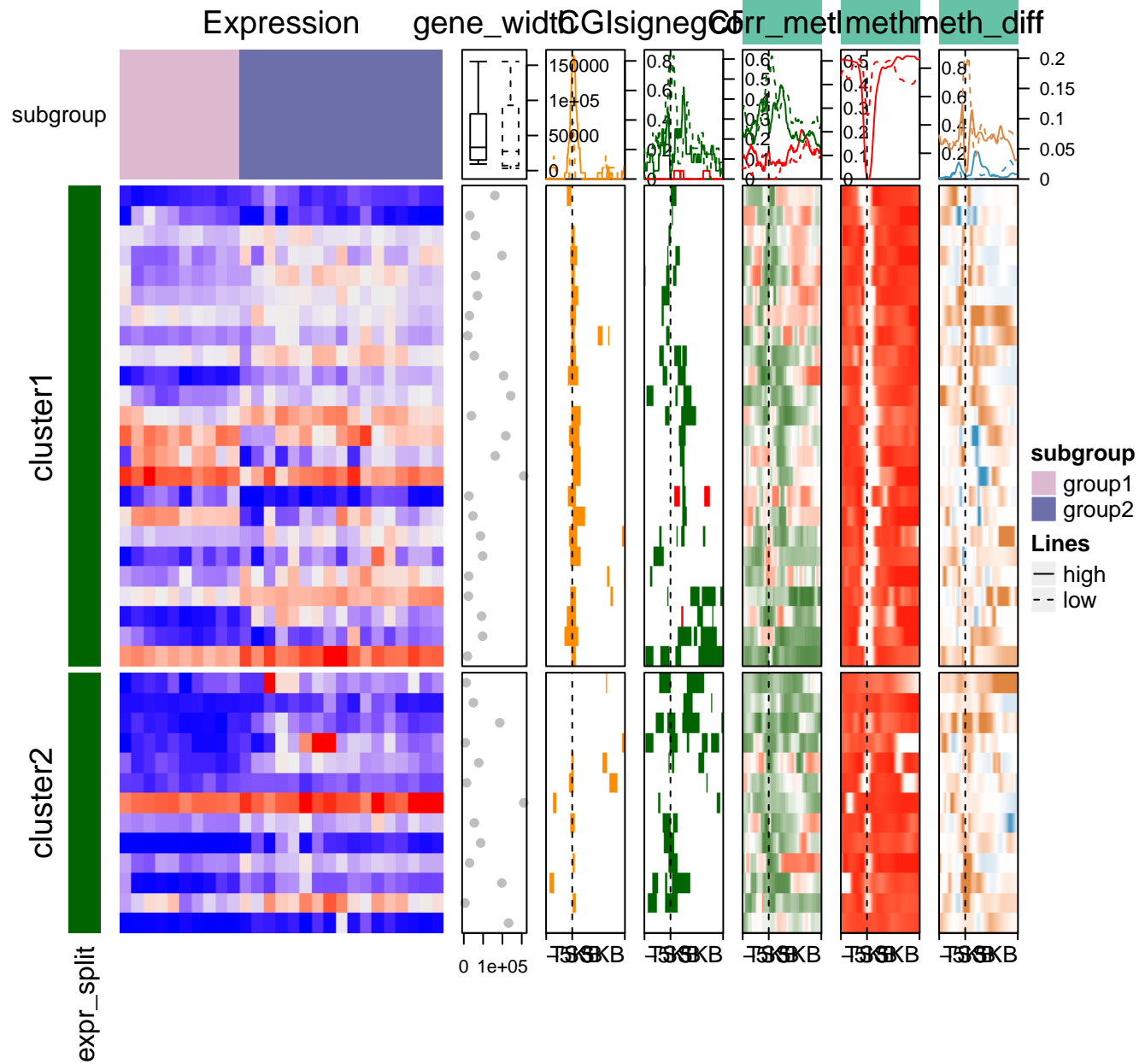




cluster by methylation, 37 rows

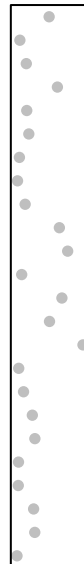
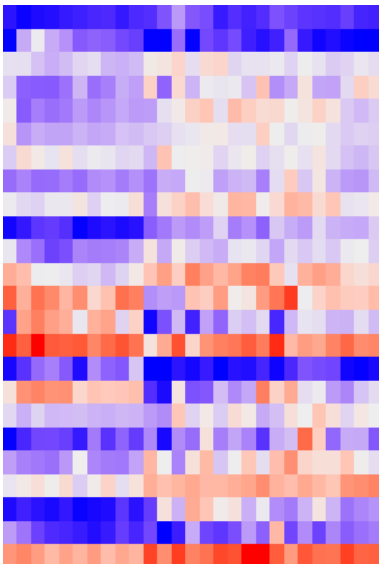


# Expression

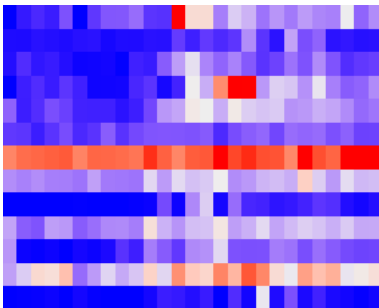
subgroup



cluster1



cluster2



expr\_split

0 1e+05