

tx_name
(gene_id)

Regions

Mean coverage

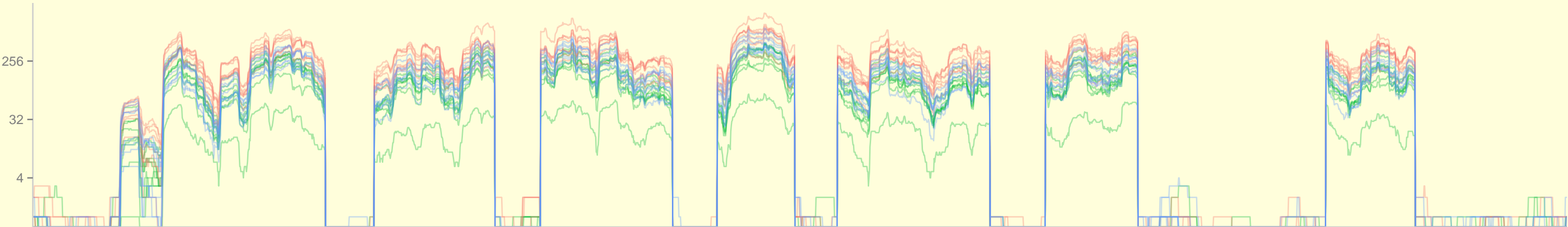
Coverage

Cluster for region with name and q-value 0.05379

chr6



chr6

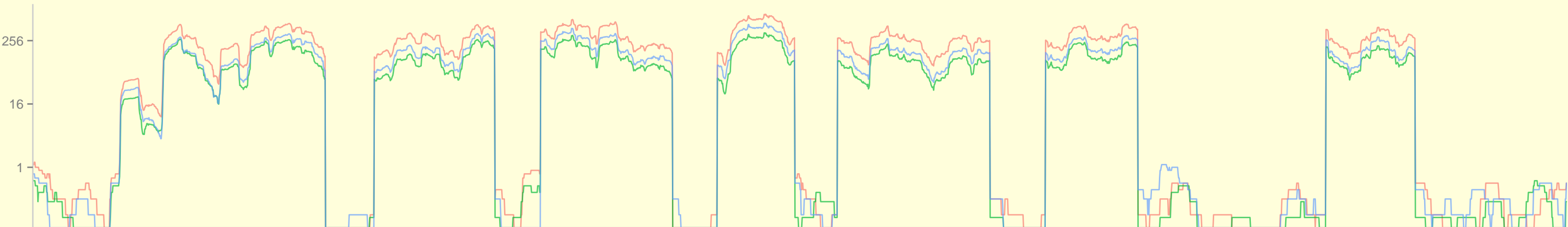


group

CT

CO

ETOH

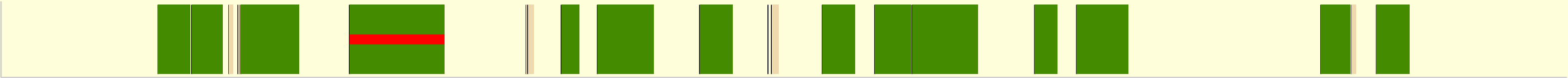


group

CT

CO

ETOH



significantQval

TRUE

FALSE

