

tx_name
(gene_id)

Regions

Mean coverage

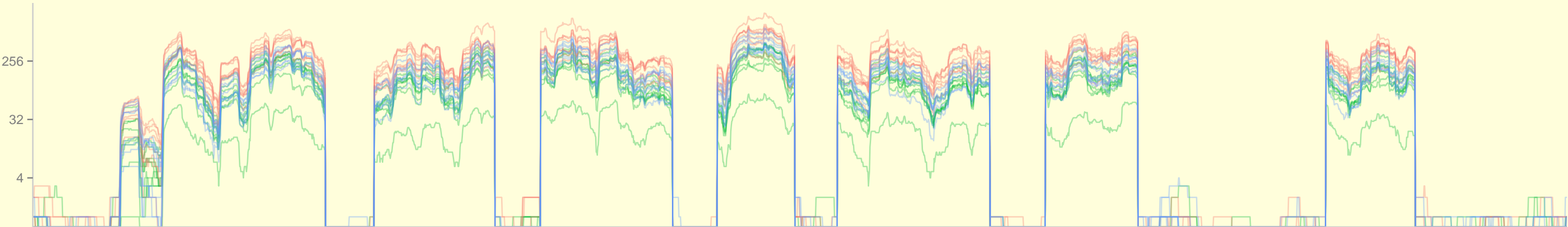
Coverage

Cluster for region with name and q-value 0.05379

chr6

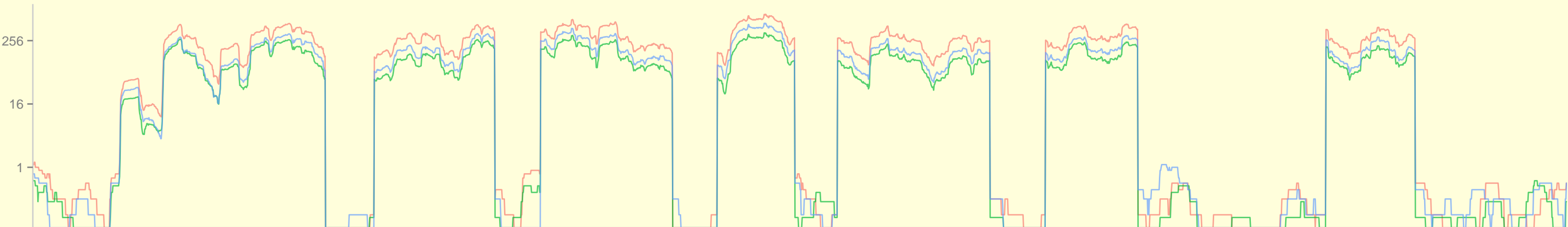


chr6



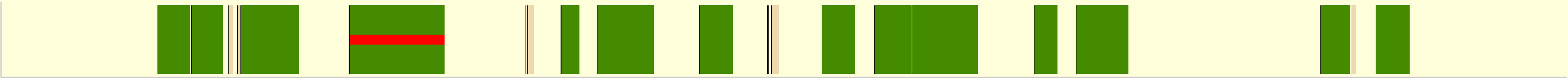
group

- CT
- CO
- ETOH



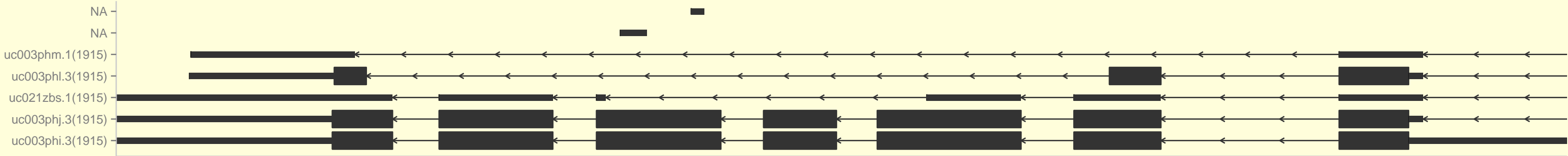
group

- CT
- CO
- ETOH



significantQval

- TRUE
- FALSE



74228000

74229000

74230000