

tx_name
(gene_id)

Regions

Mean coverage

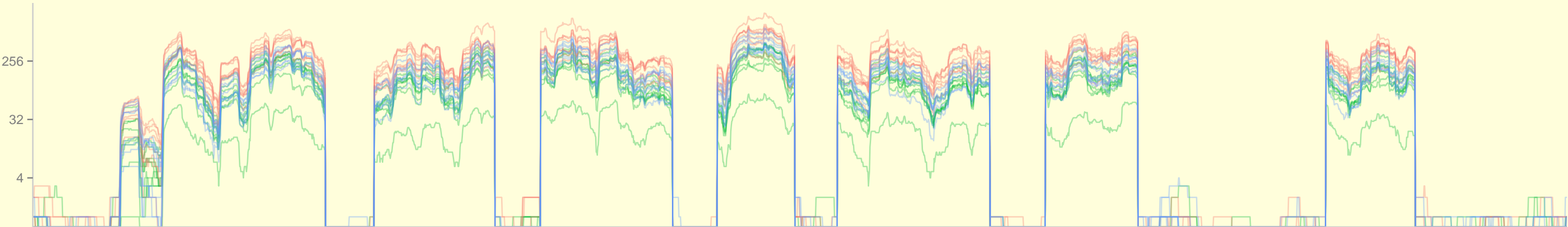
Coverage

Cluster for region with name and q-value 0.05379

chr6

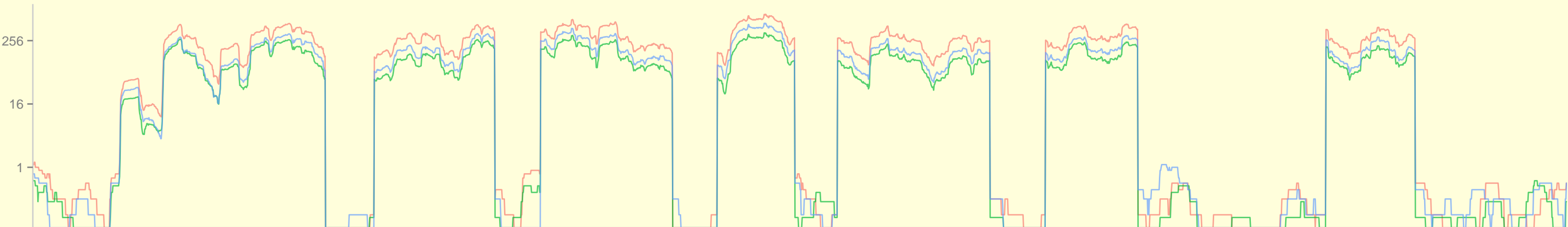


chr6



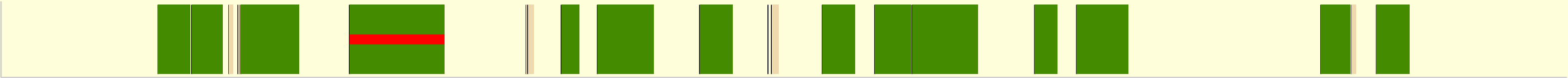
group

- CT
- CO
- ETOH



group

- CT
- CO
- ETOH



significantQval

- TRUE
- FALSE

