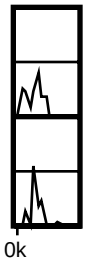
sample_.fa - mlagan

Alignment 1
  Seqs: sample1/sample2
  Criteria: 75%, 100 bp
  Regions: 0

Alignment 2
  Seqs: sample1/sample3
  Criteria: 75%, 100 bp
  Regions: 0

X-axis: sample1
Resolution: 25
Window size: 40 bp



gene
exon
UTR
CNS