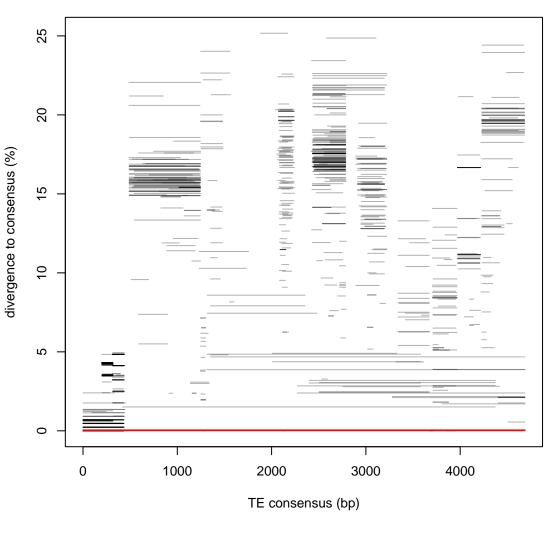
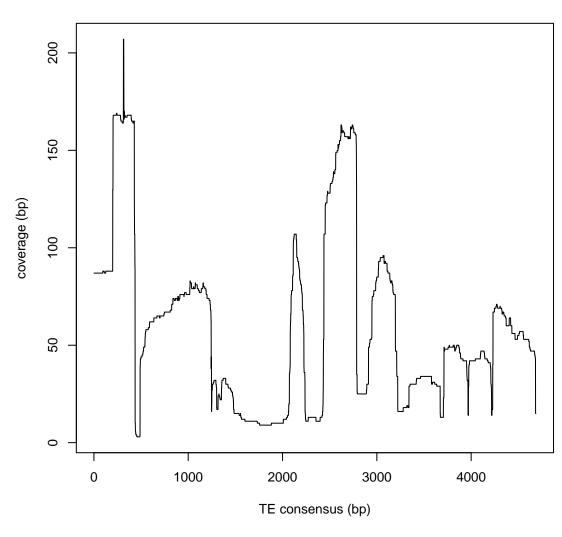
TE consensus genomic coverage





TE consensus self dotplot (blastn)

0 1000 2000 3000 4000 rnd-1_family-64-LTR/Gypsy (bb)

TE consensus structure and protein hits

