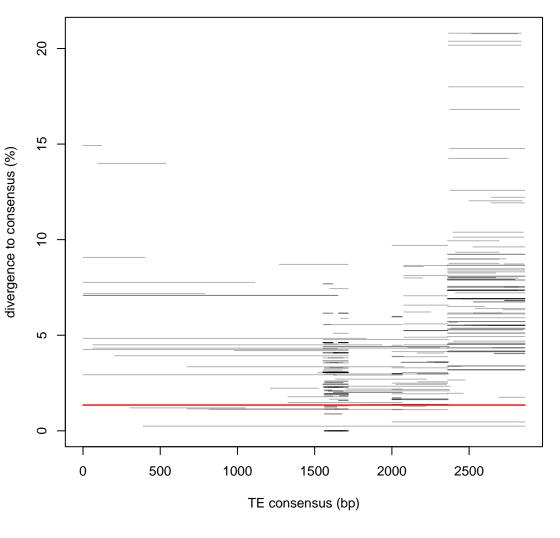
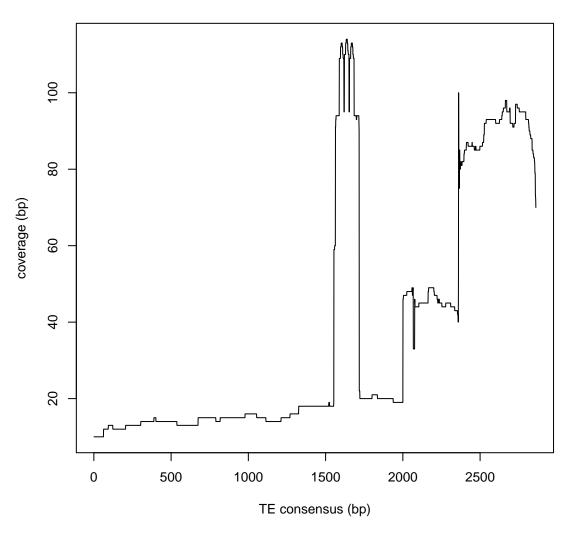
TE consensus genomic coverage





TE consensus self dotplot (blastn)

0 1000 1500 2500 rnd–5_family–487–LTR/Gypsy (bp)

TE consensus structure and protein hits

