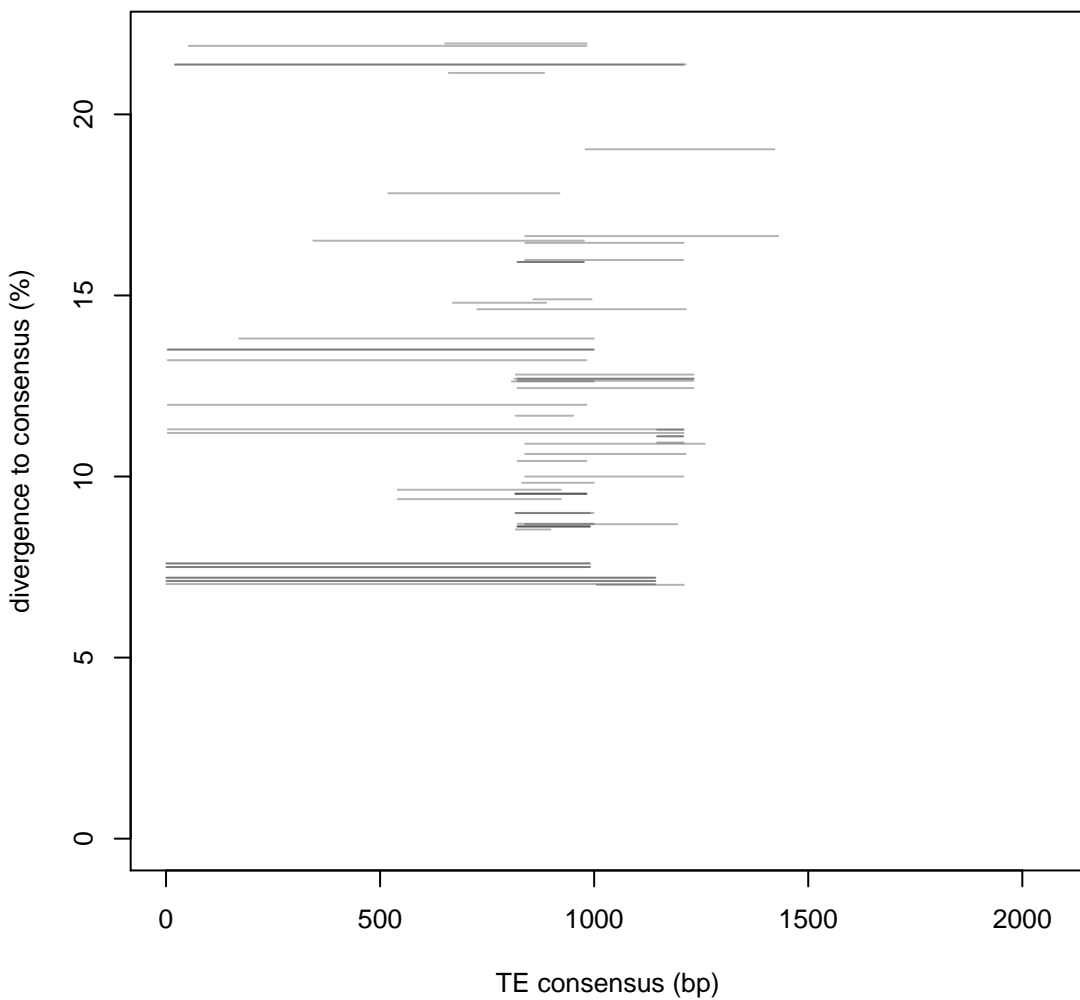
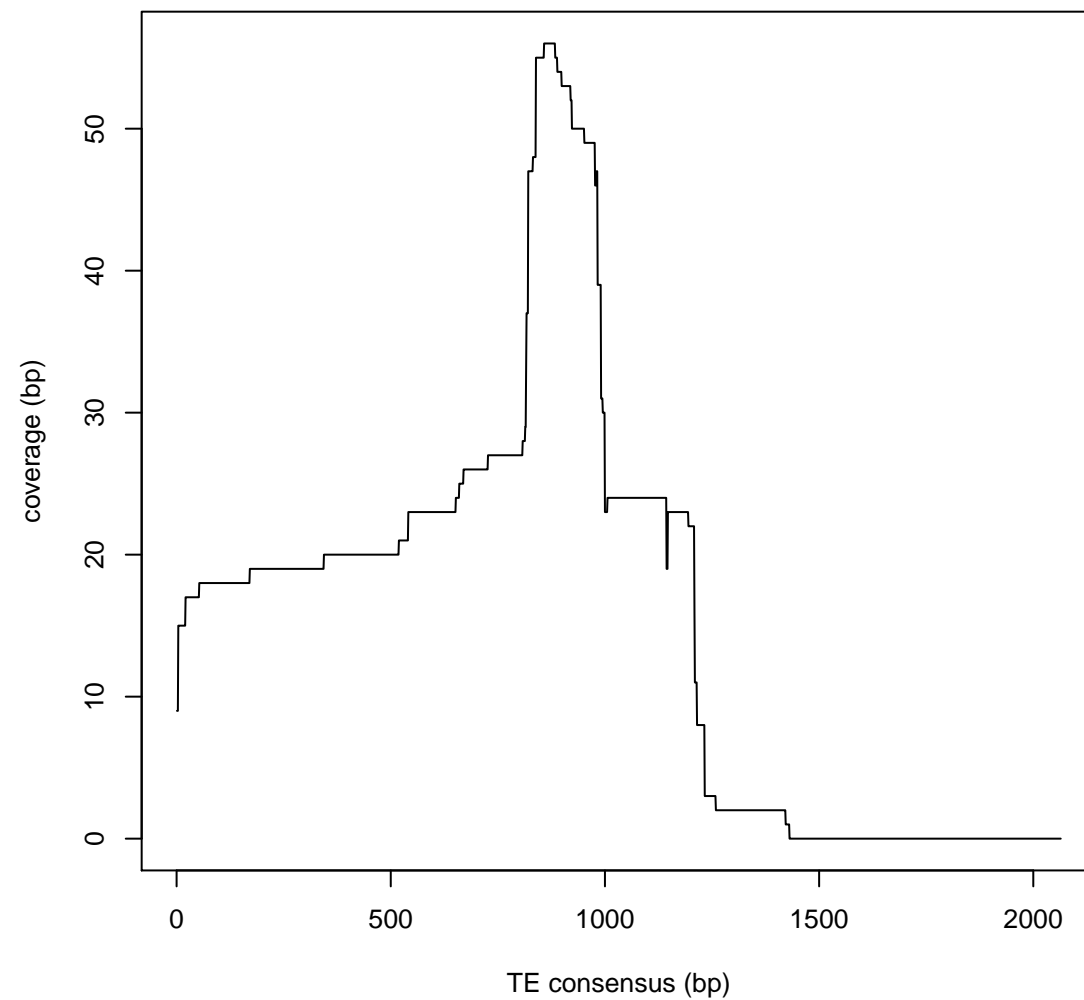


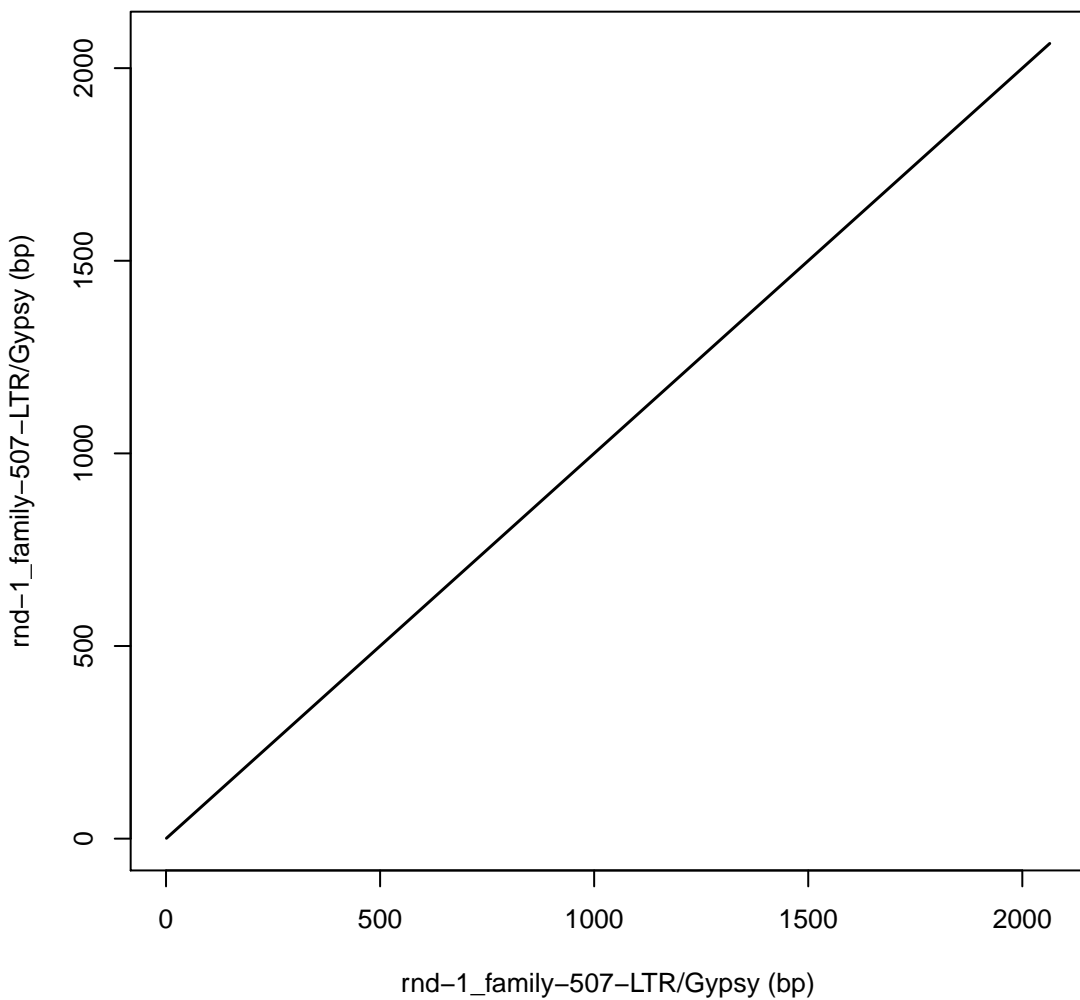
TE: rnd-1\_family-507-LTR/Gypsy  
consensus size: 2064bp; fragments: 62; full length: 0 (>=1857.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

