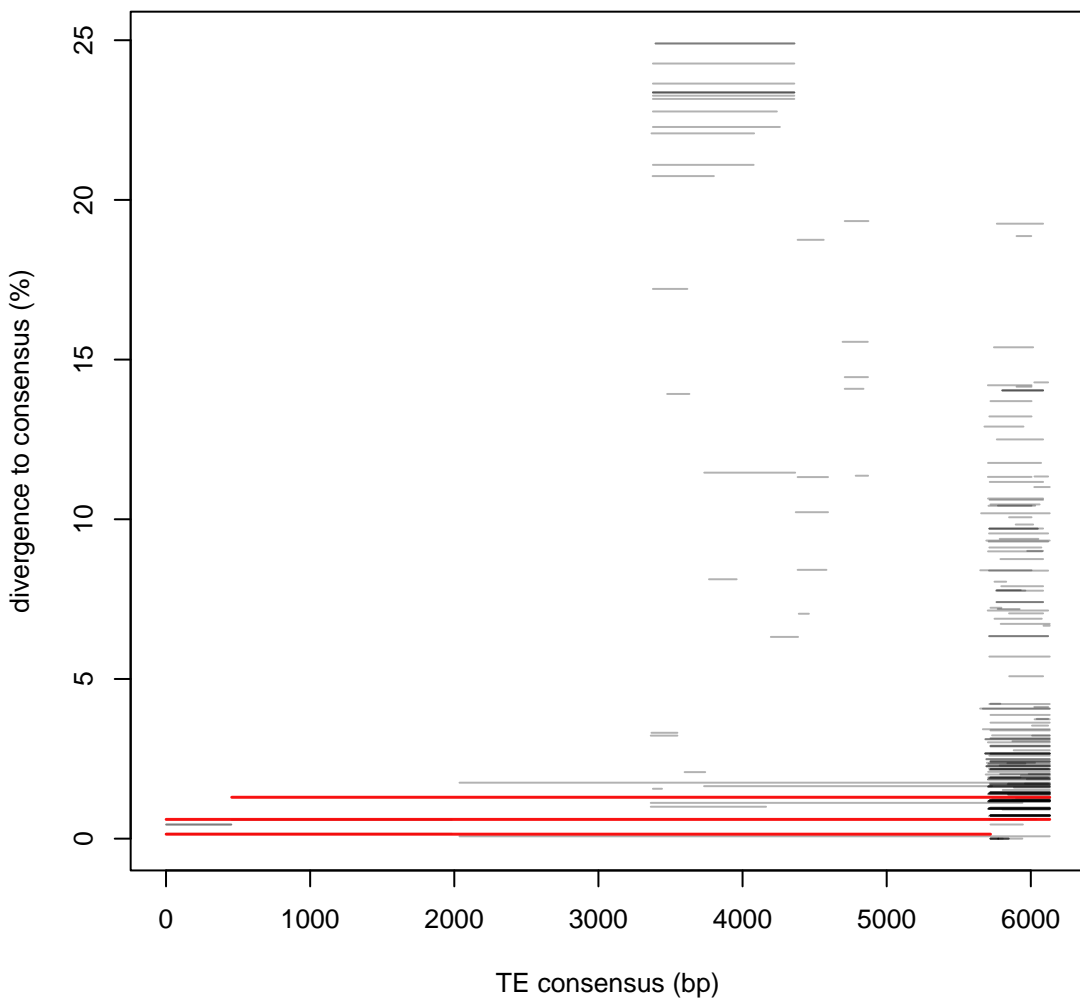
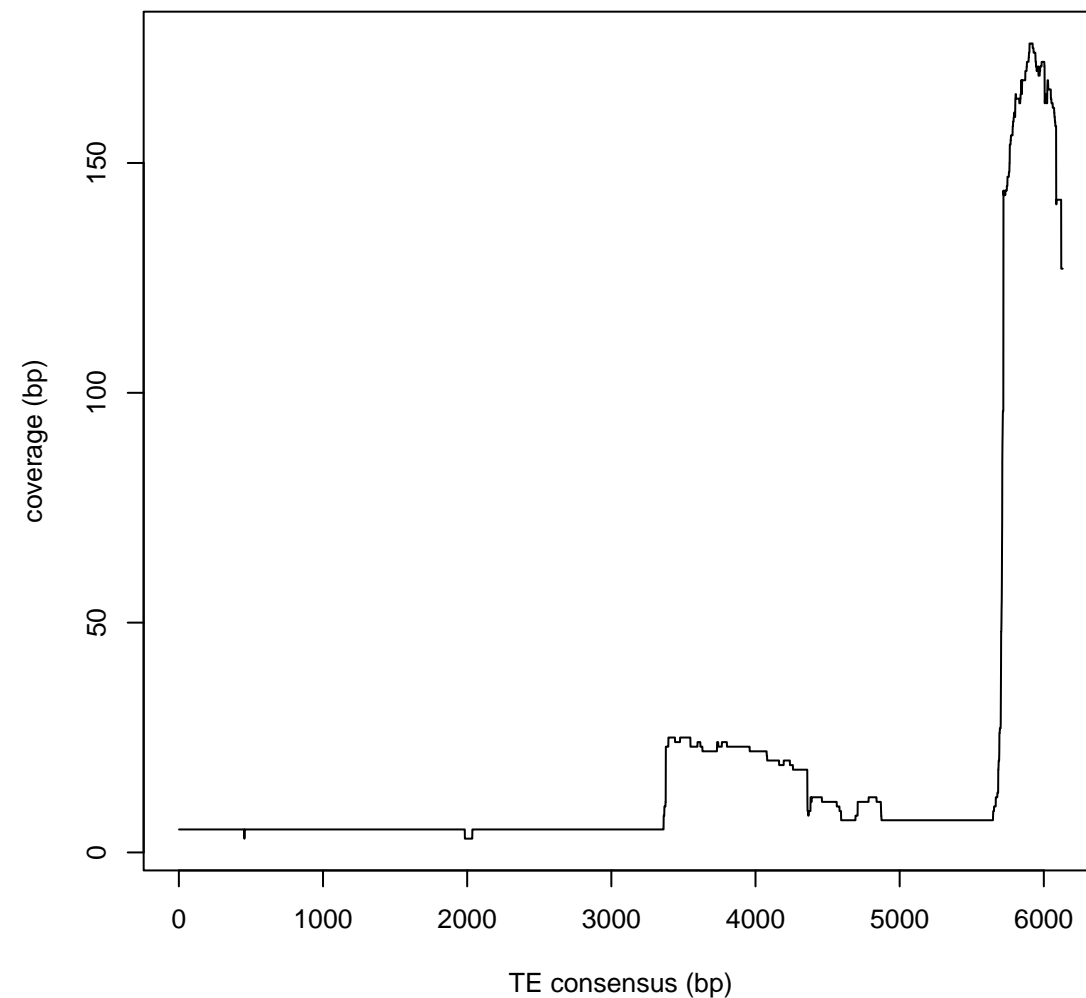


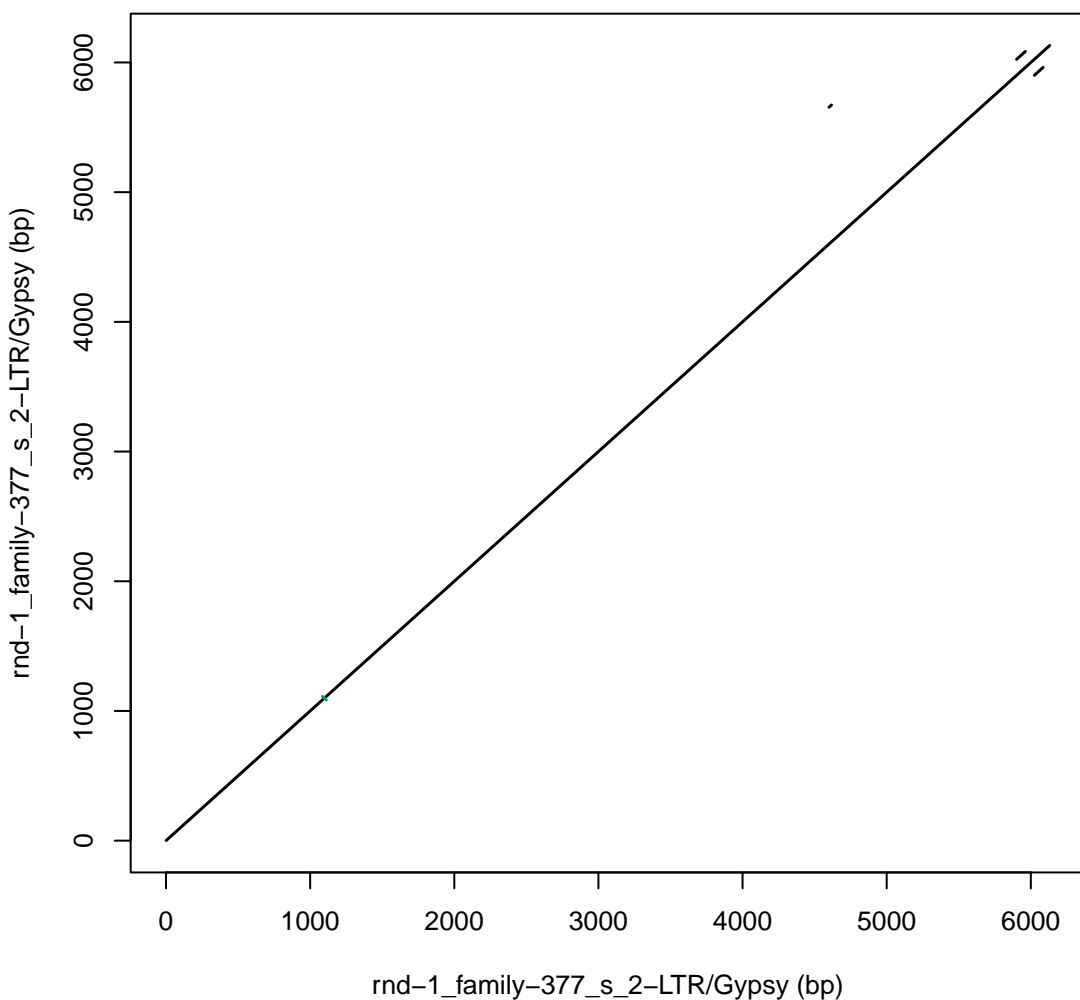
TE: rnd-1_family-377_s_2-LTR/Gypsy
consensus size: 6131bp; fragments: 240; full length: 3 (>=5517.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

