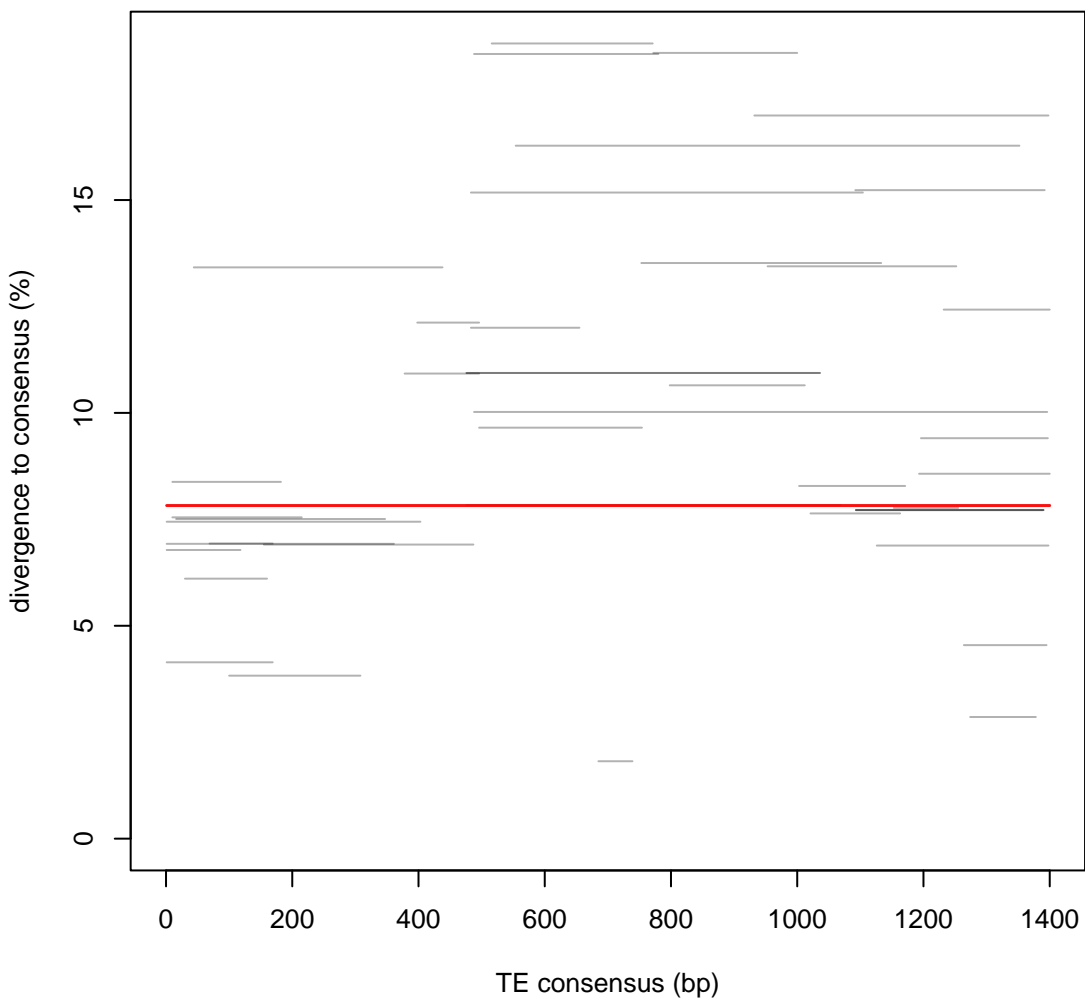
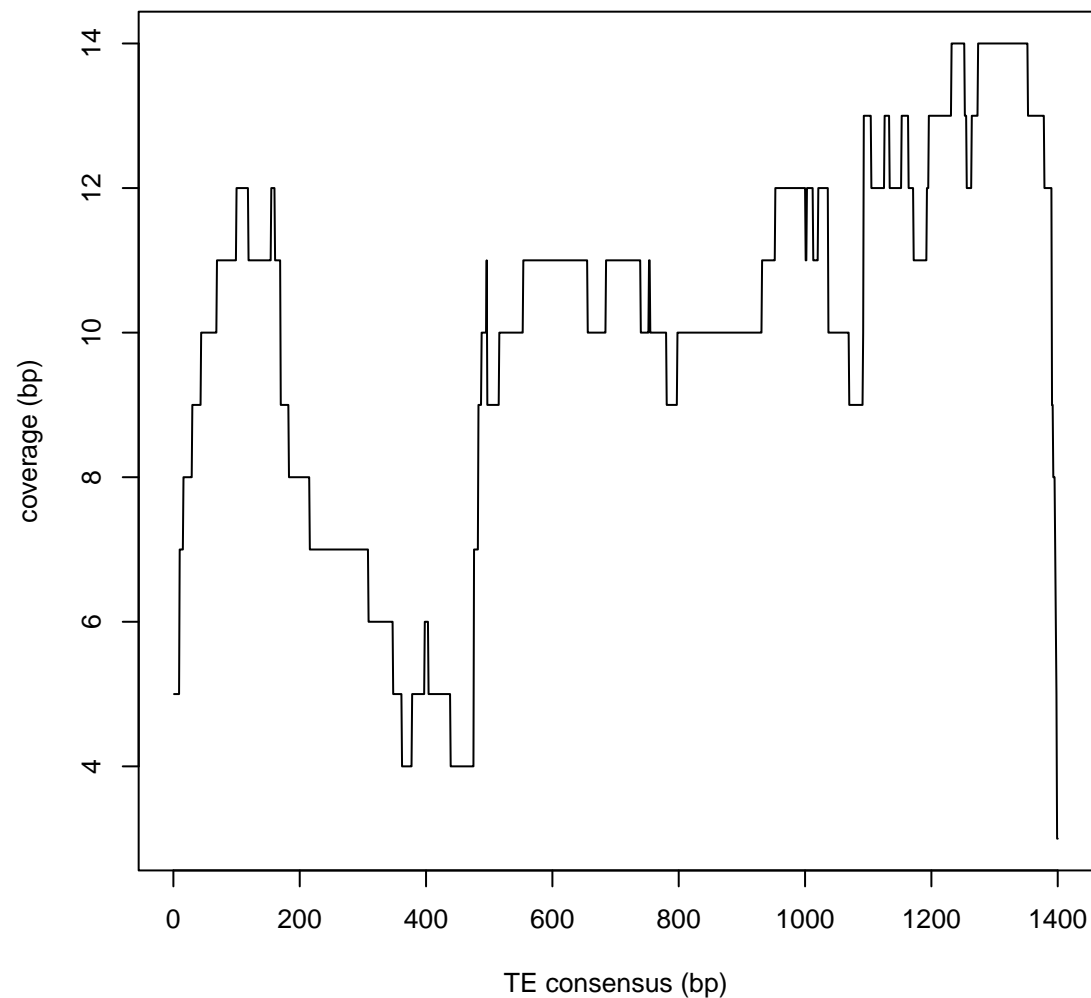


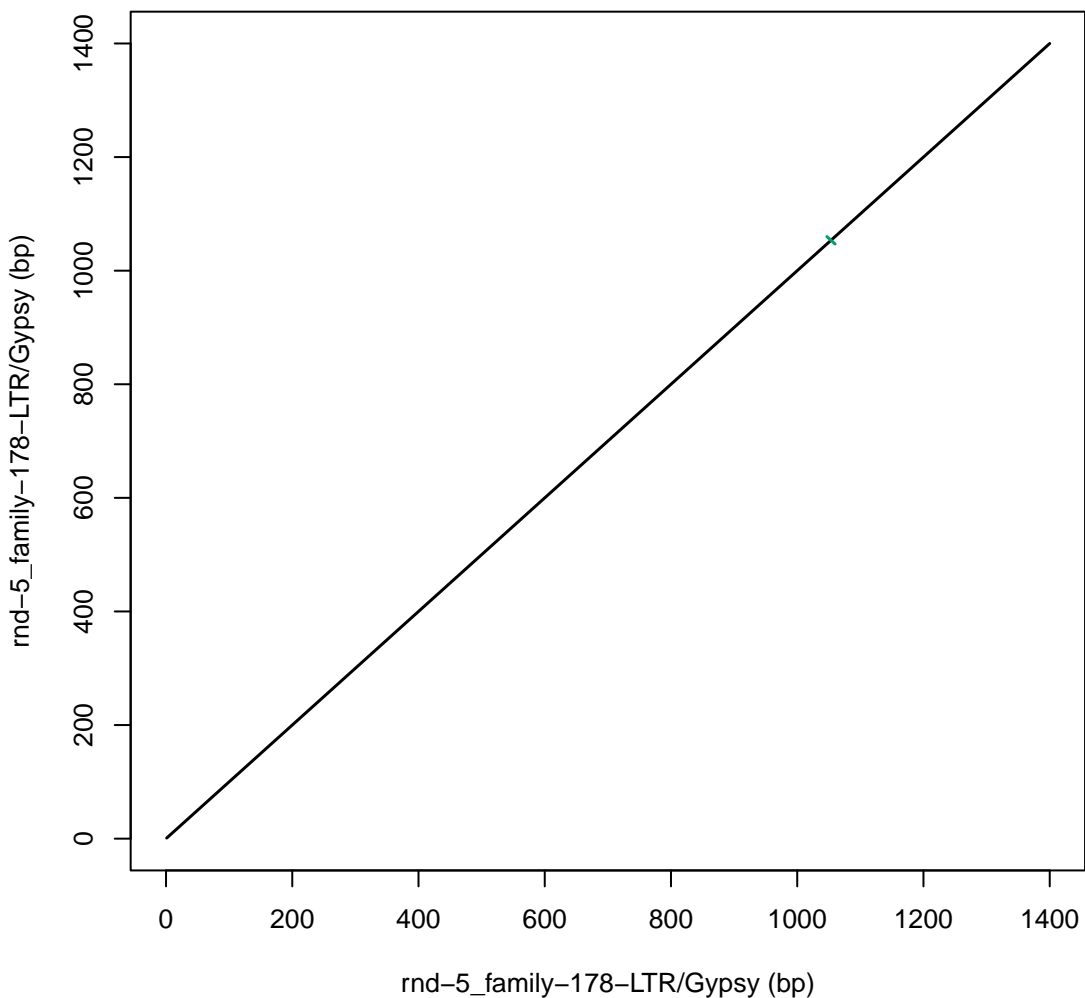
TE: rnd-5\_family-178-LTR/Gypsy  
consensus size: 1400bp; fragments: 44; full length: 1 (>=1260bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

