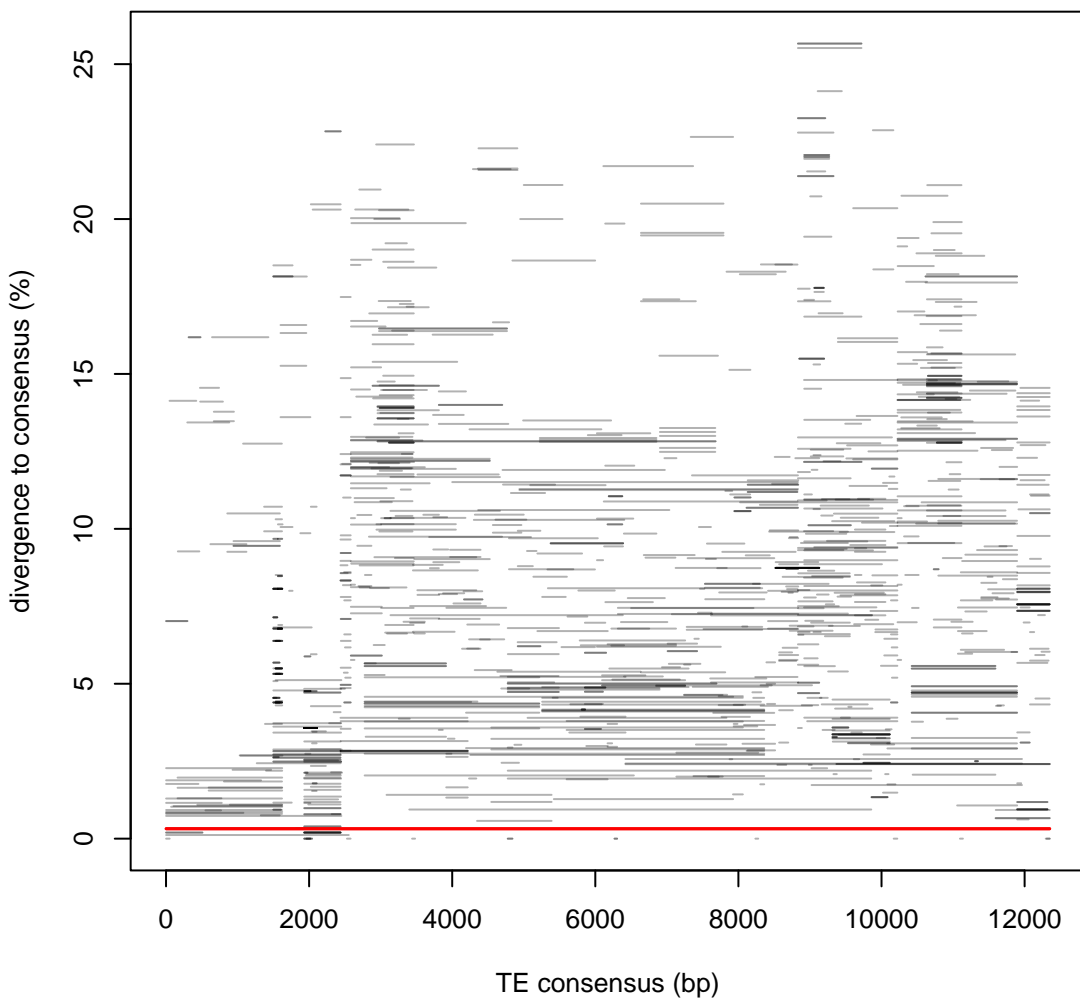
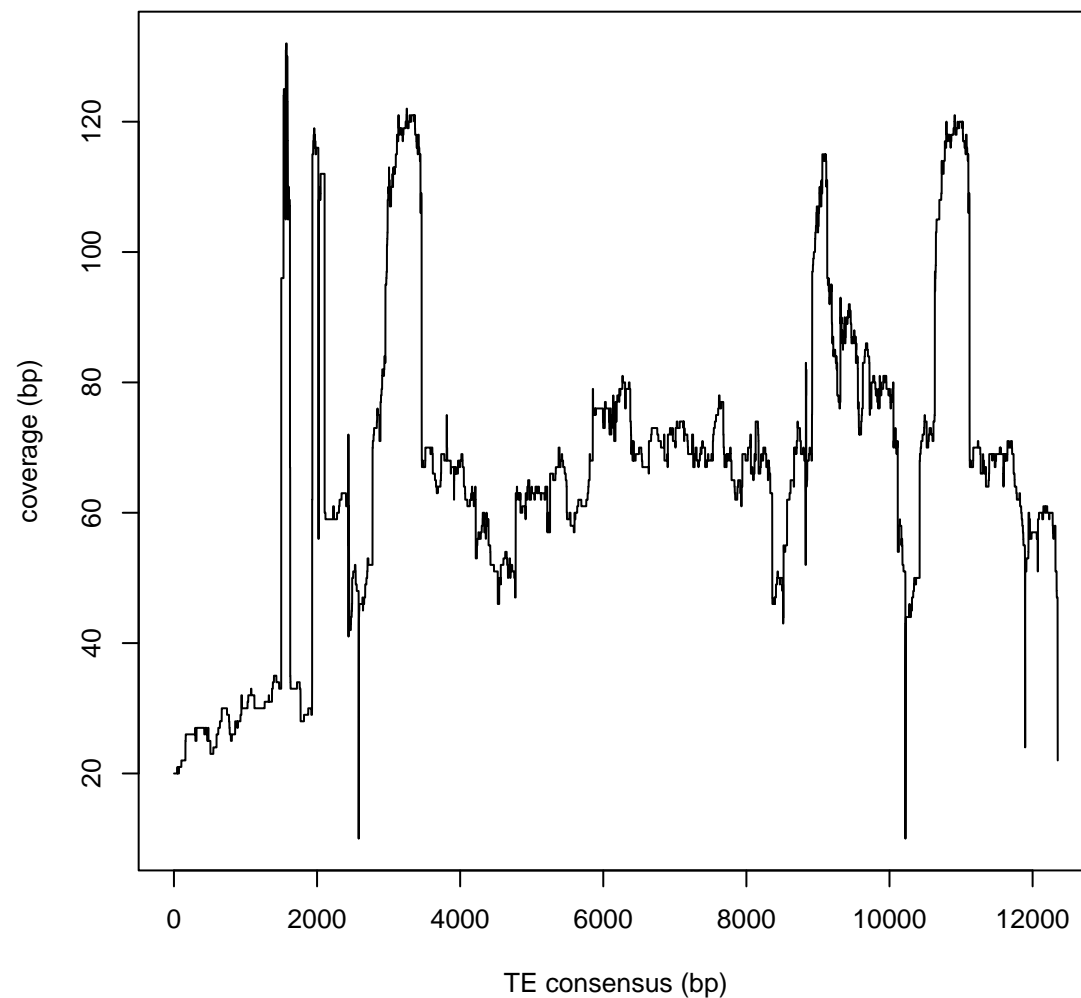


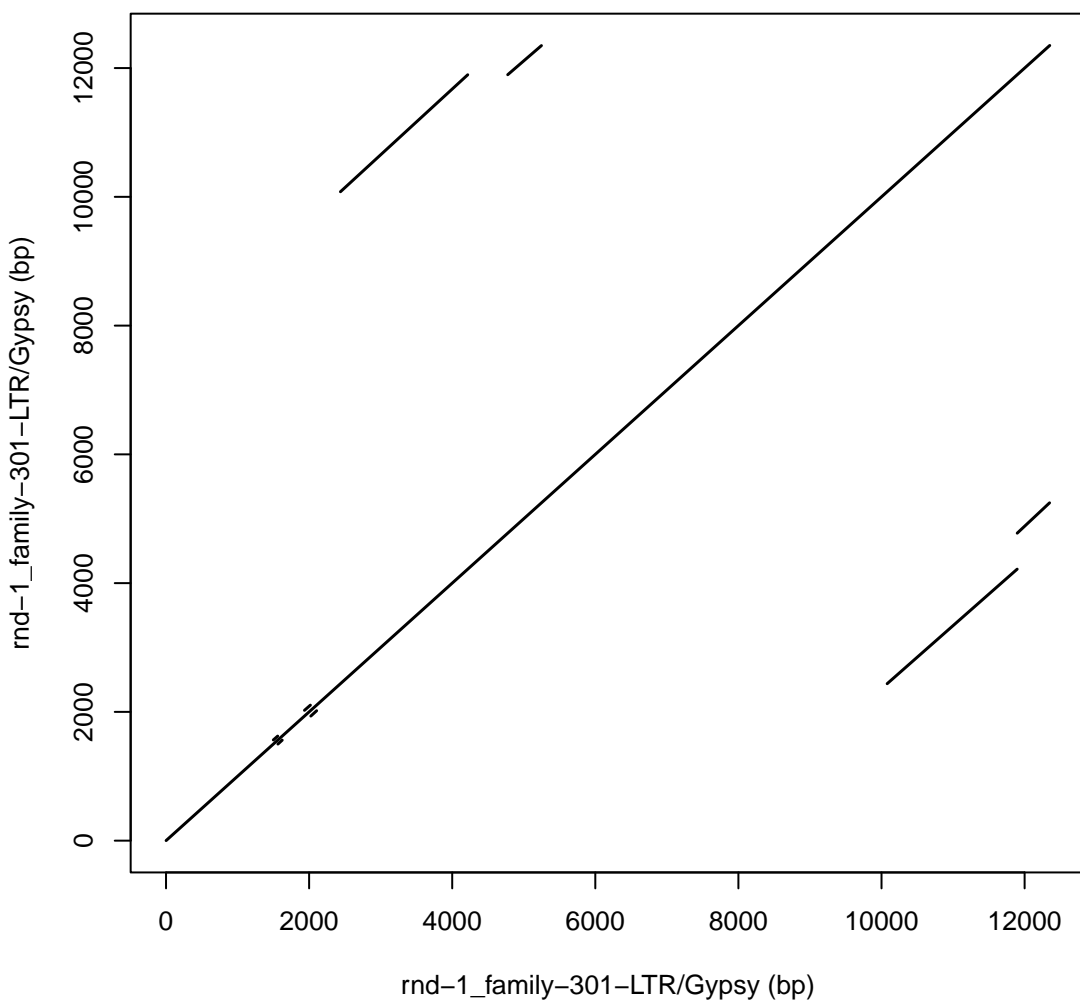
TE: rnd-1\_family-301-LTR/Gypsy  
consensus size: 12351bp; fragments: 1340; full length: 2 (>=11115.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

