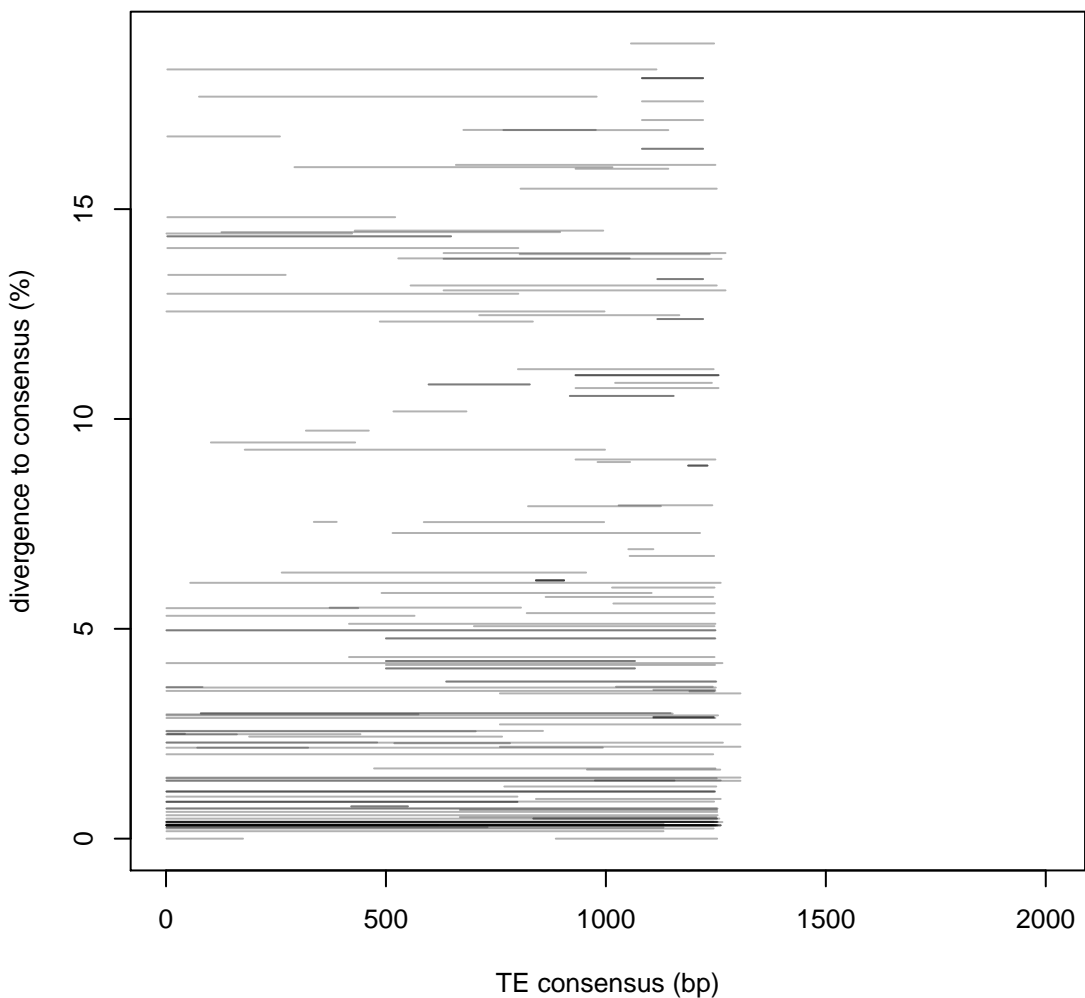
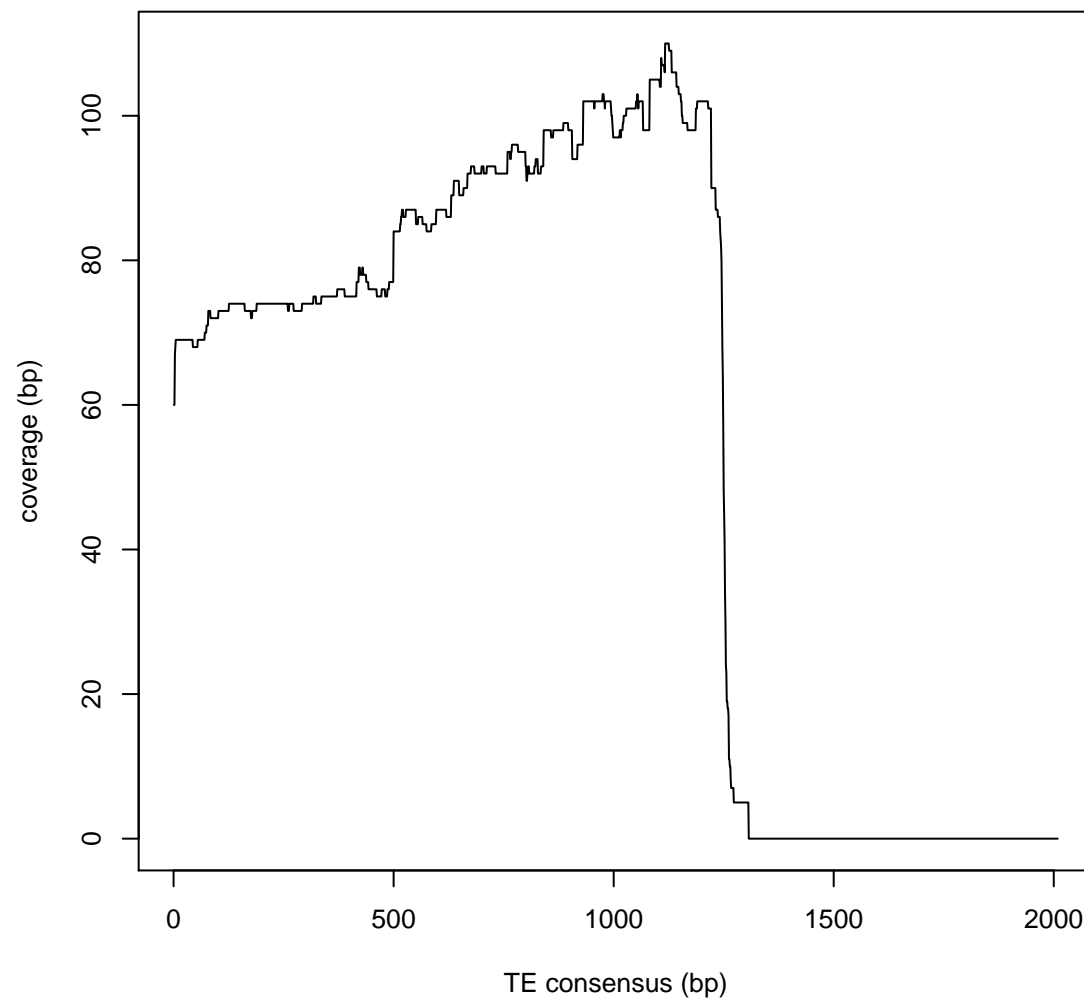


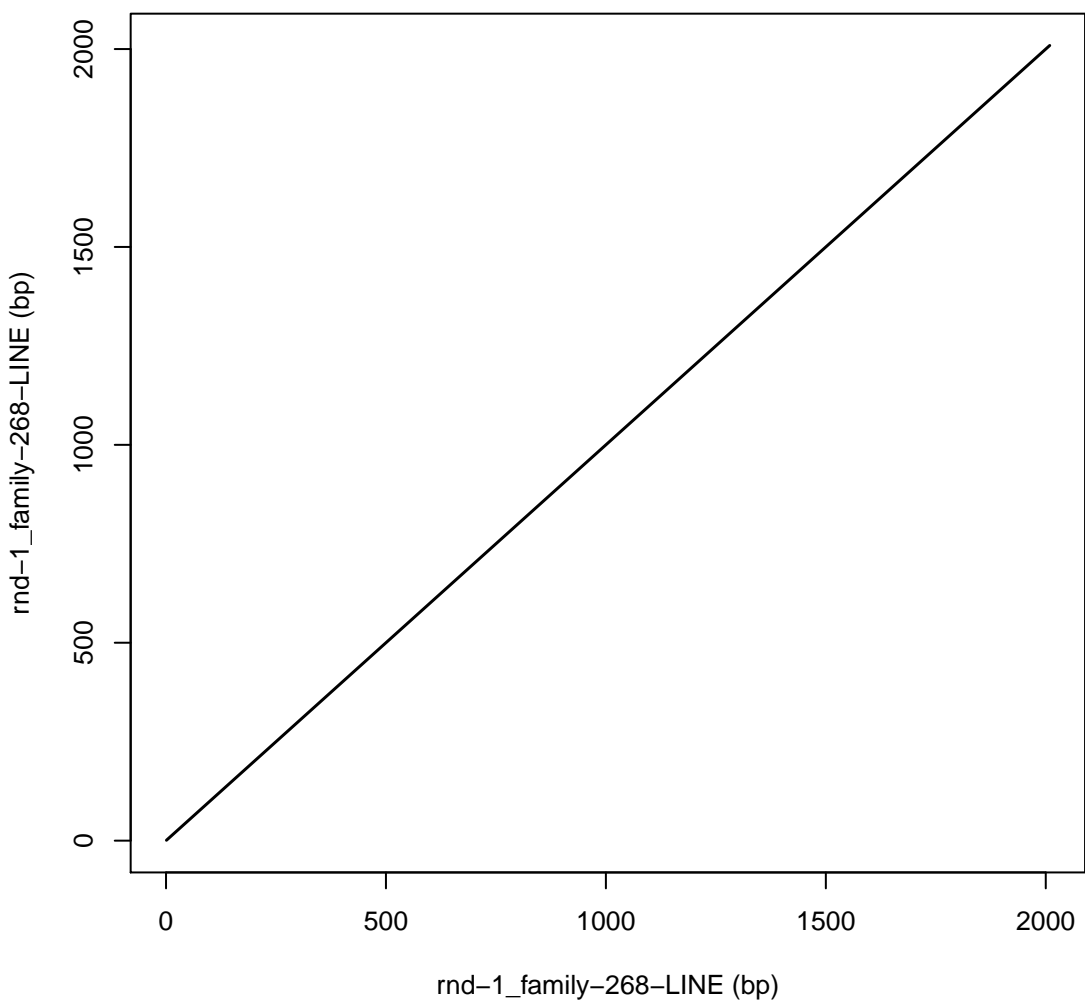
TE: rnd-1\_family-268-LINE  
consensus size: 2009bp; fragments: 174; full length: 0 (>=1808.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

