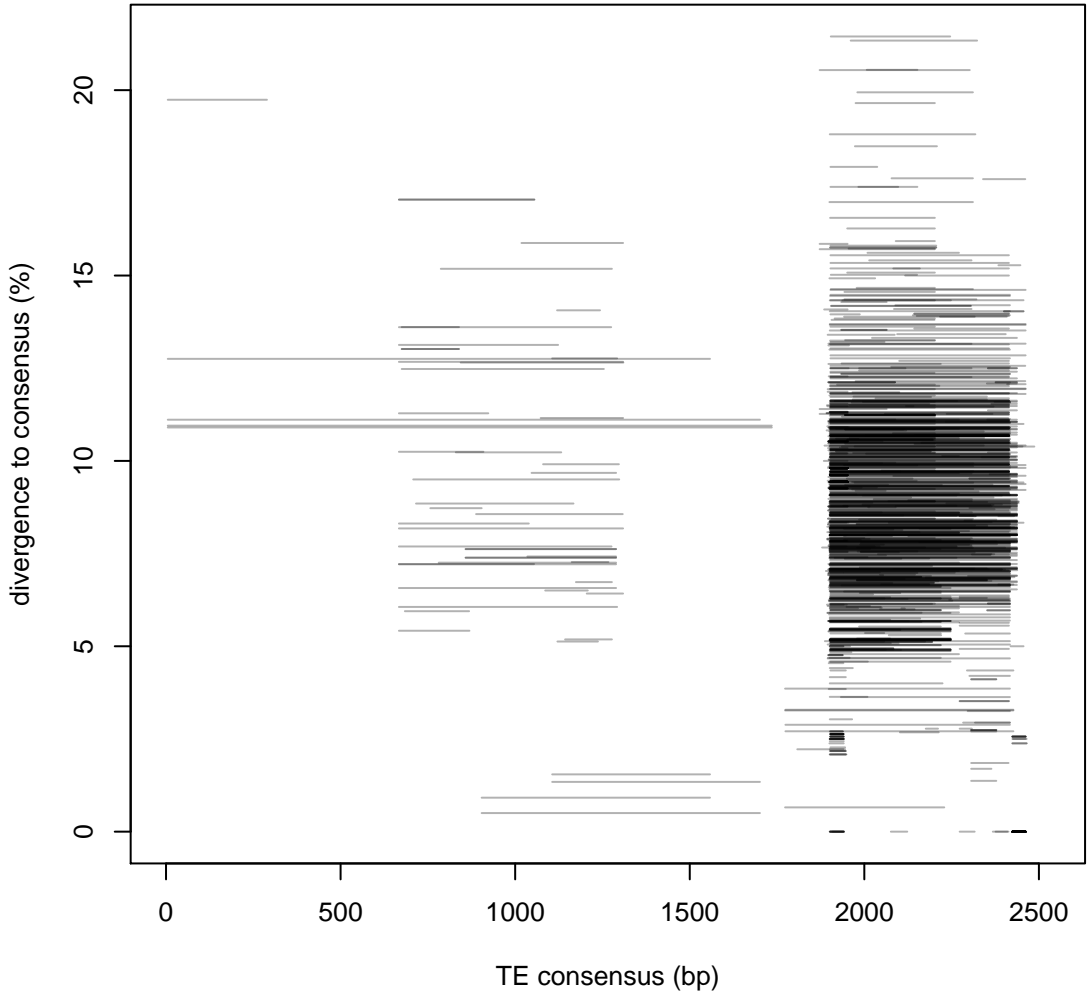
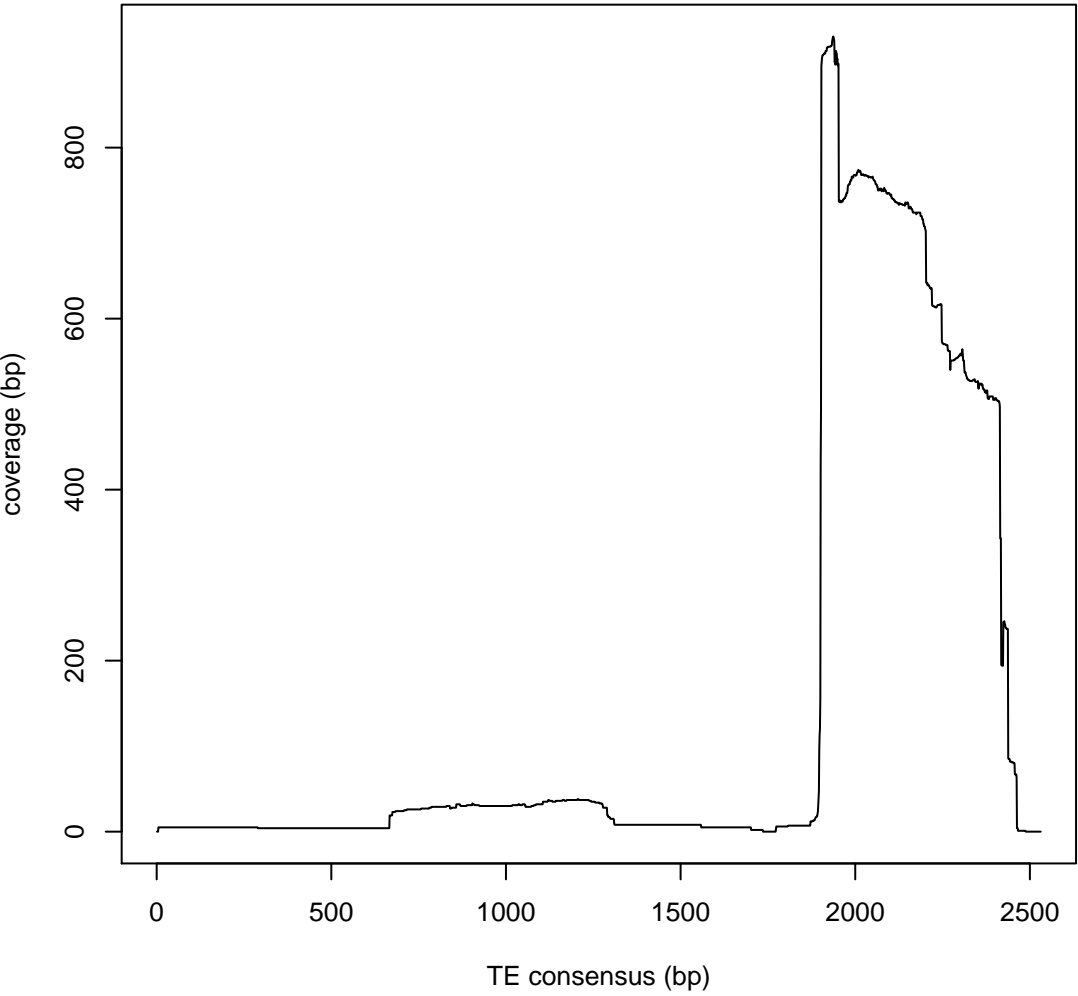


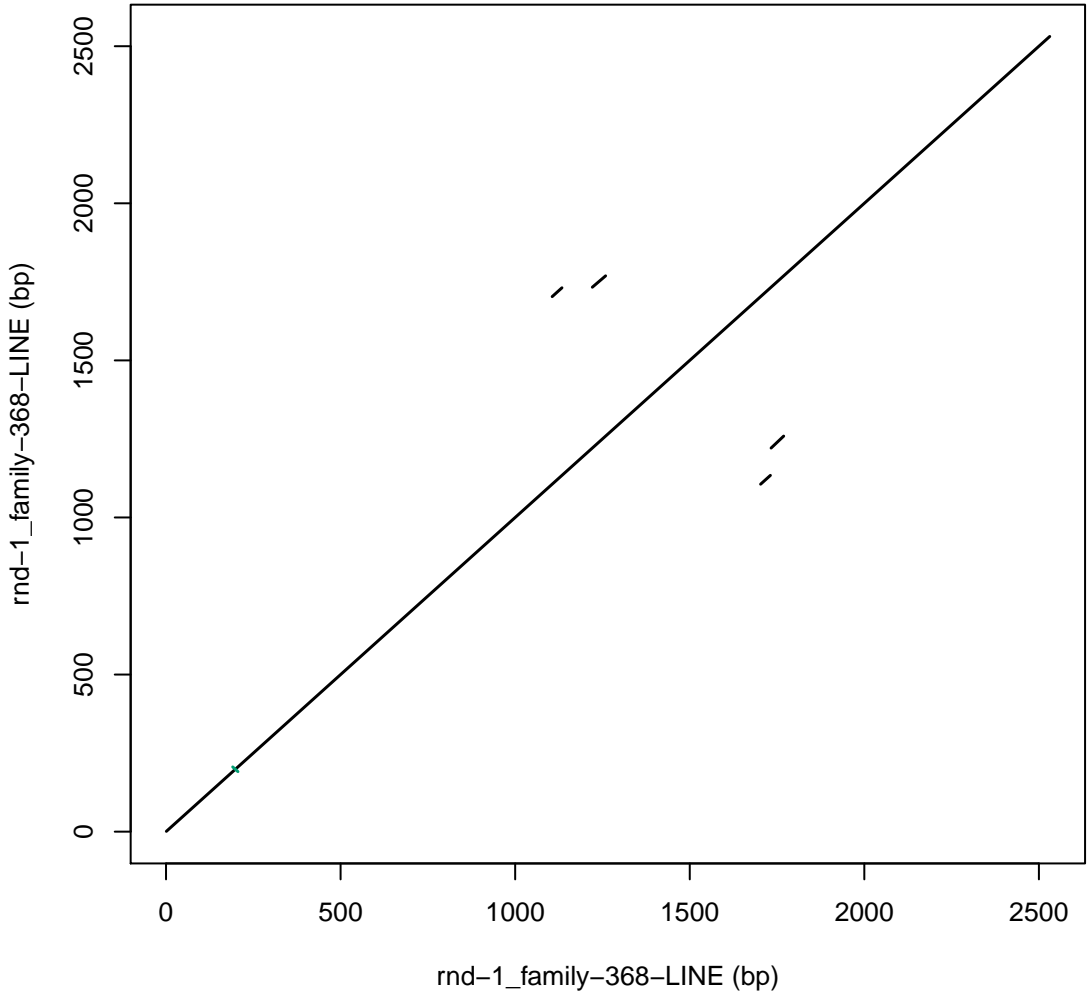
TE: rnd-1\_family-368-LINE  
consensus size: 2531bp; fragments: 1234; full length: 0 (>=2277.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

