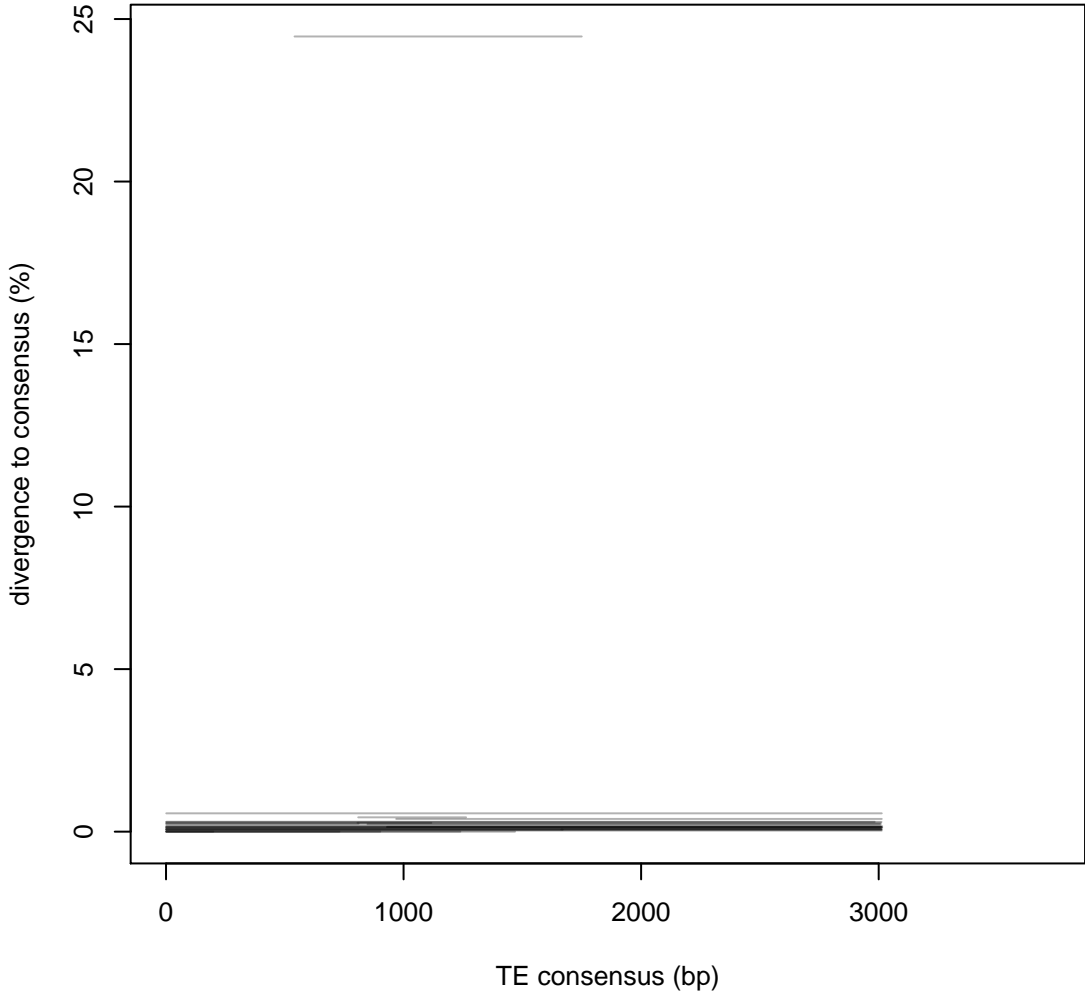
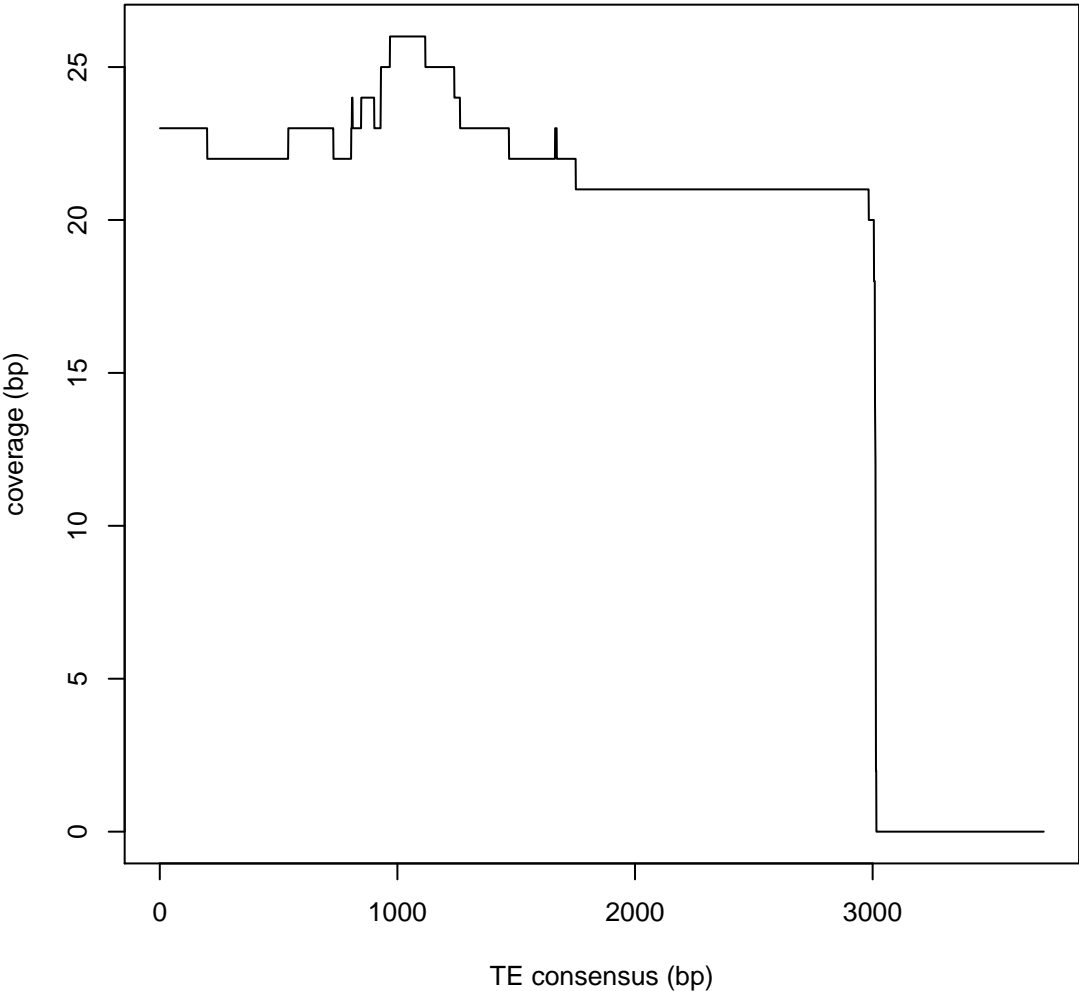


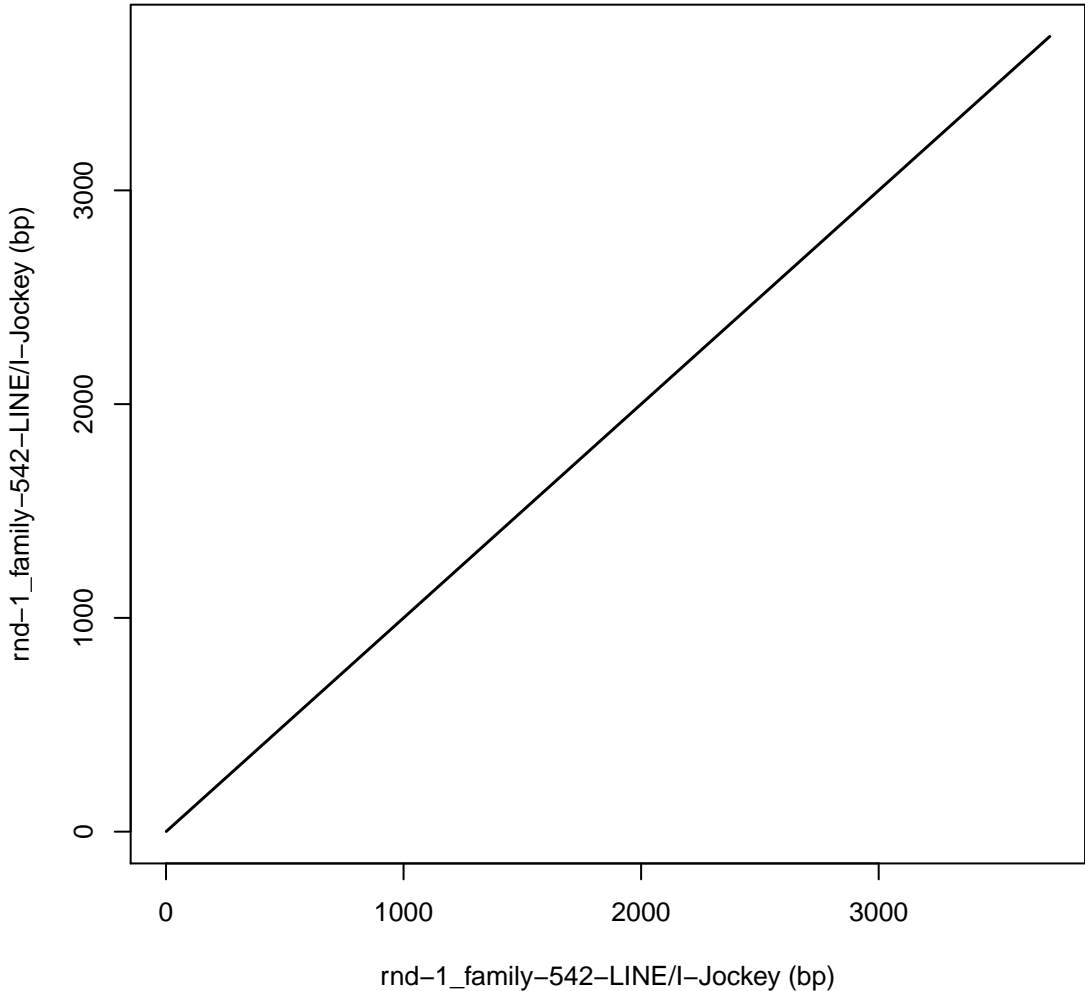
TE: rnd-1\_family-542-LINE/I-Jockey  
consensus size: 3720bp; fragments: 31; full length: 0 (>=3348bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

