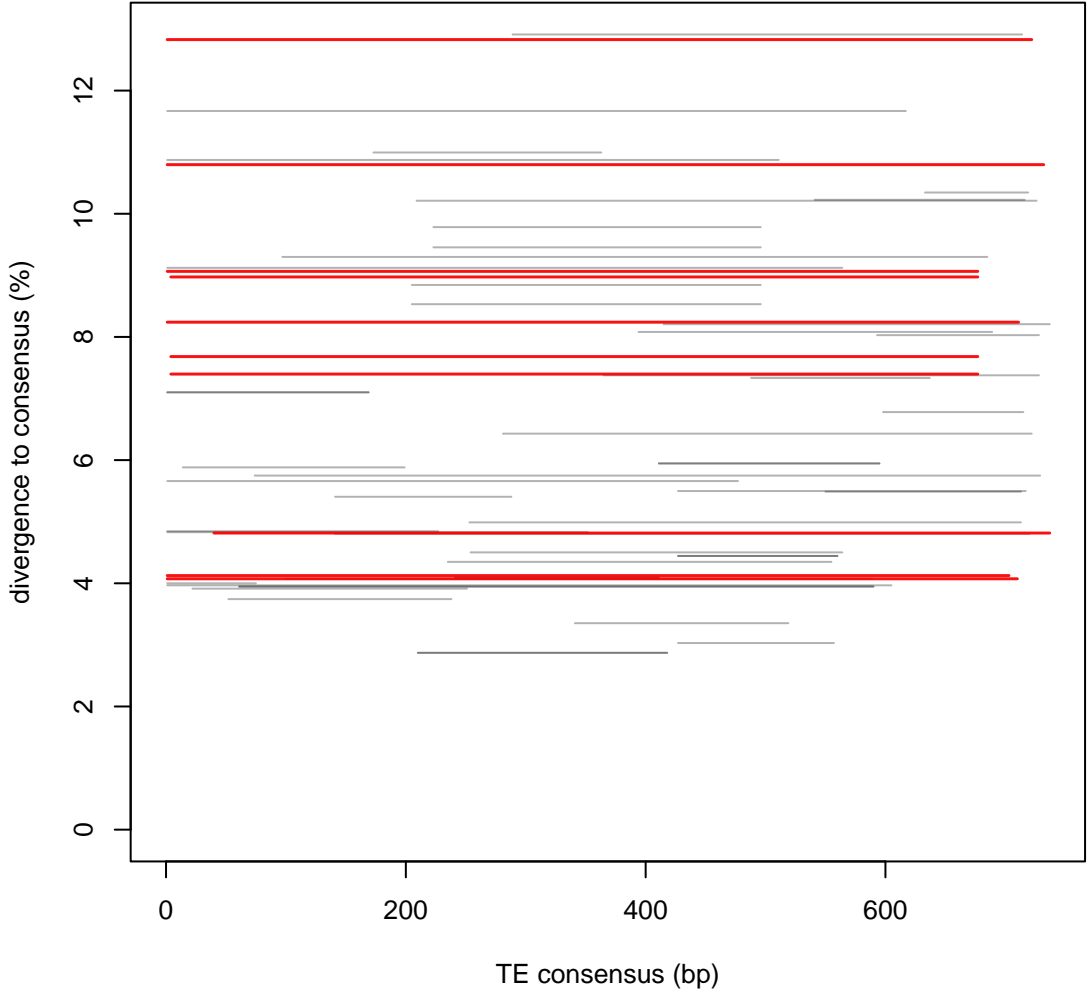
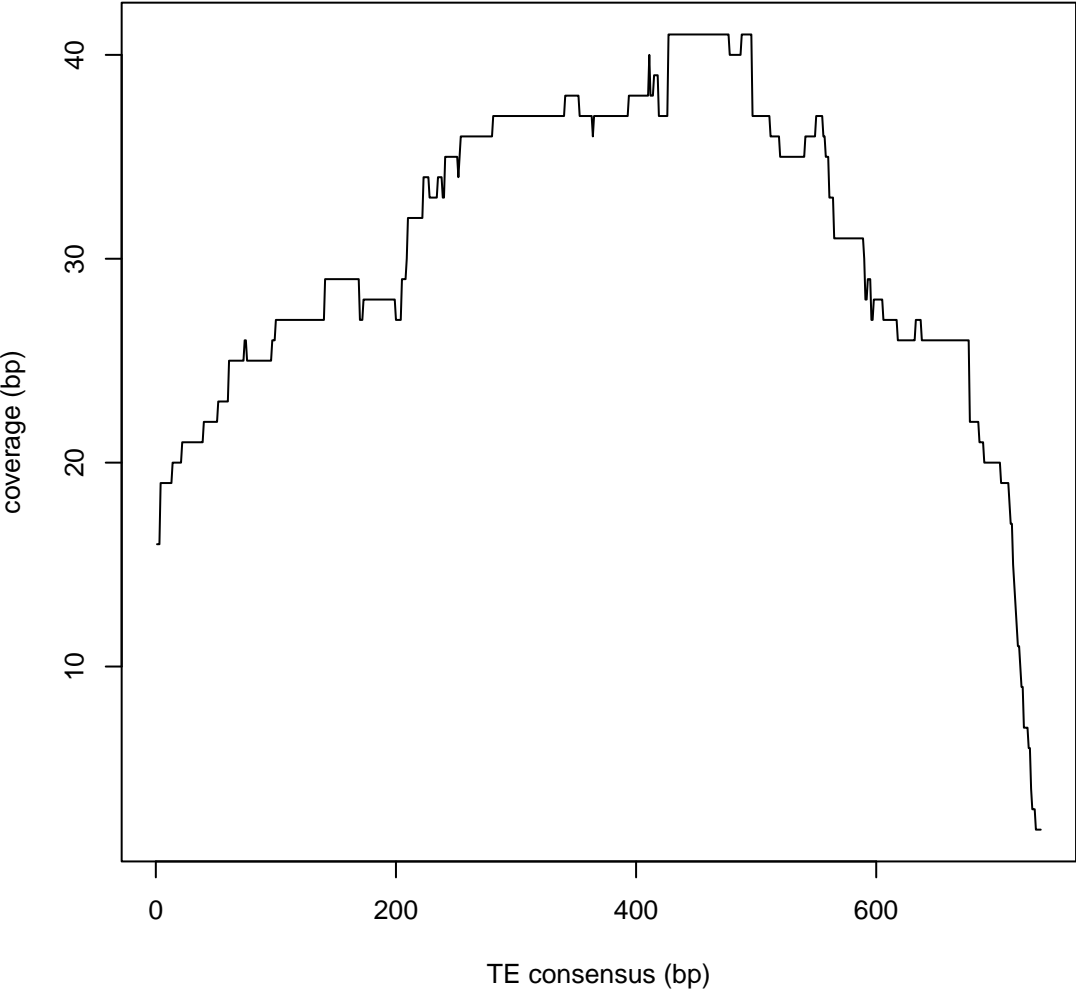


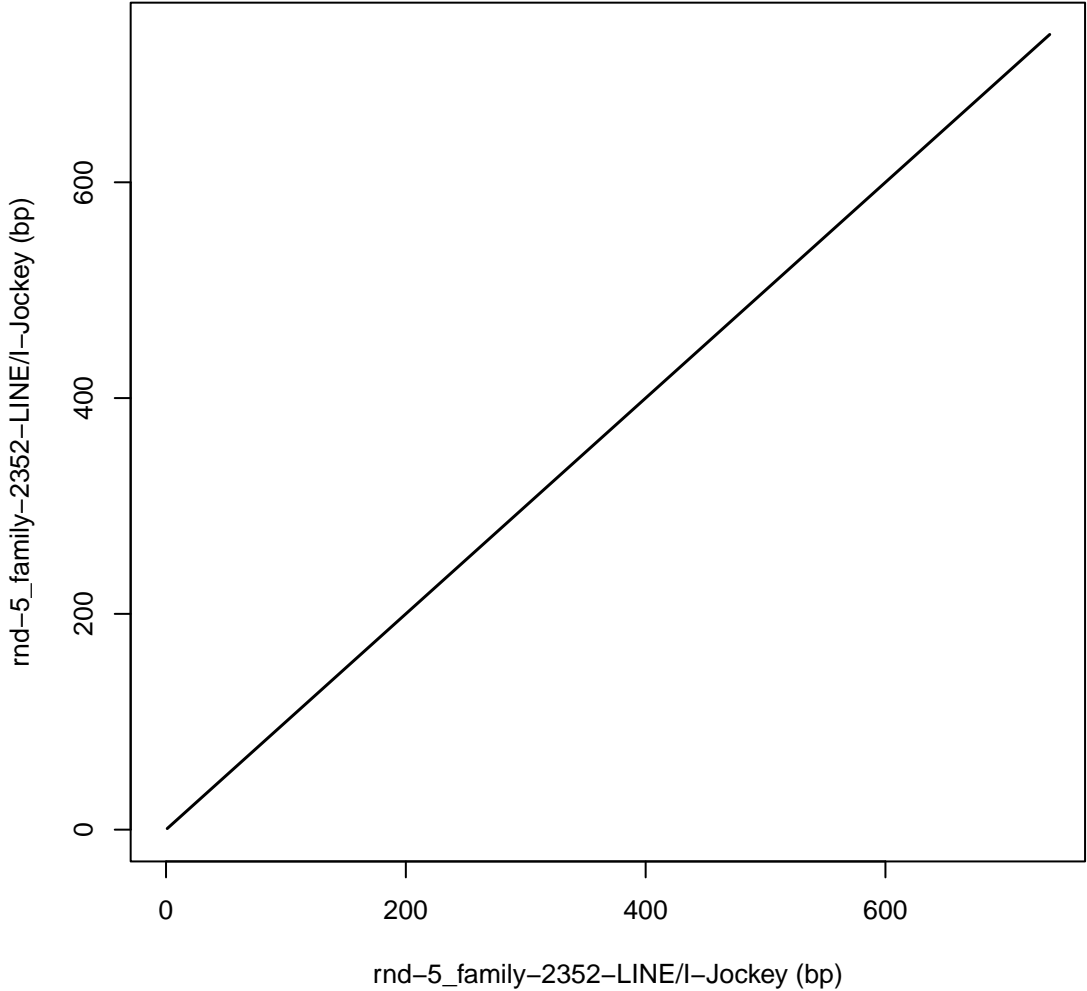
TE: rnd-5_family-2352-LINE/I-Jockey
consensus size: 737bp; fragments: 61; full length: 10 (>=663.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

