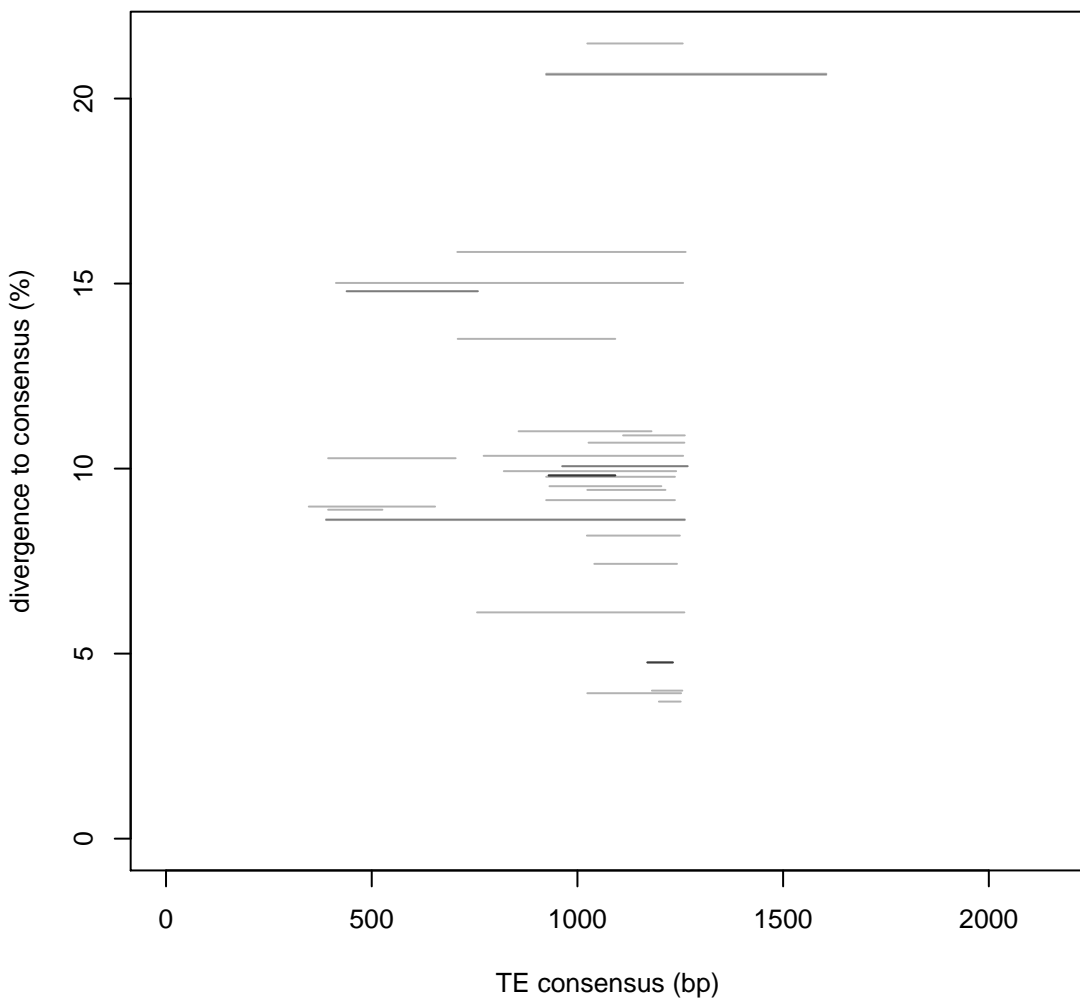
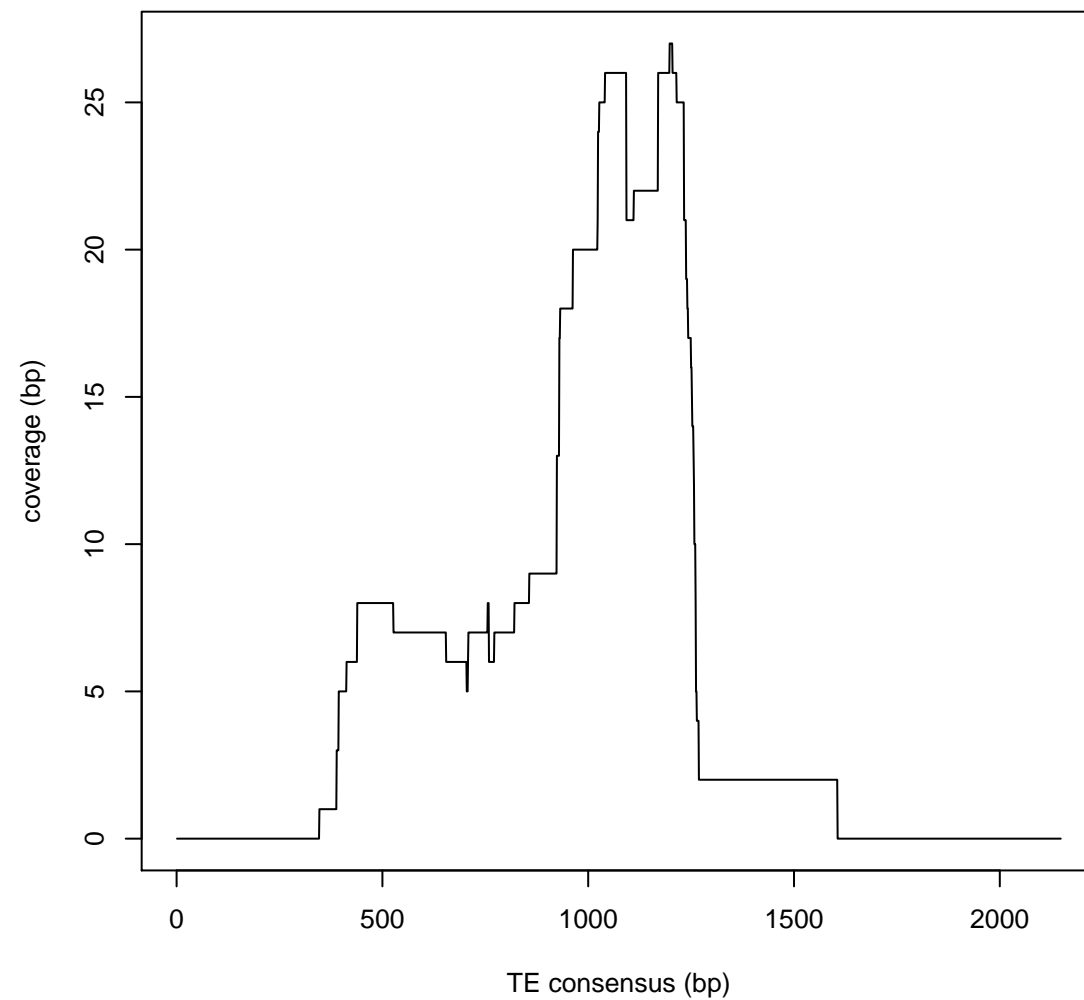


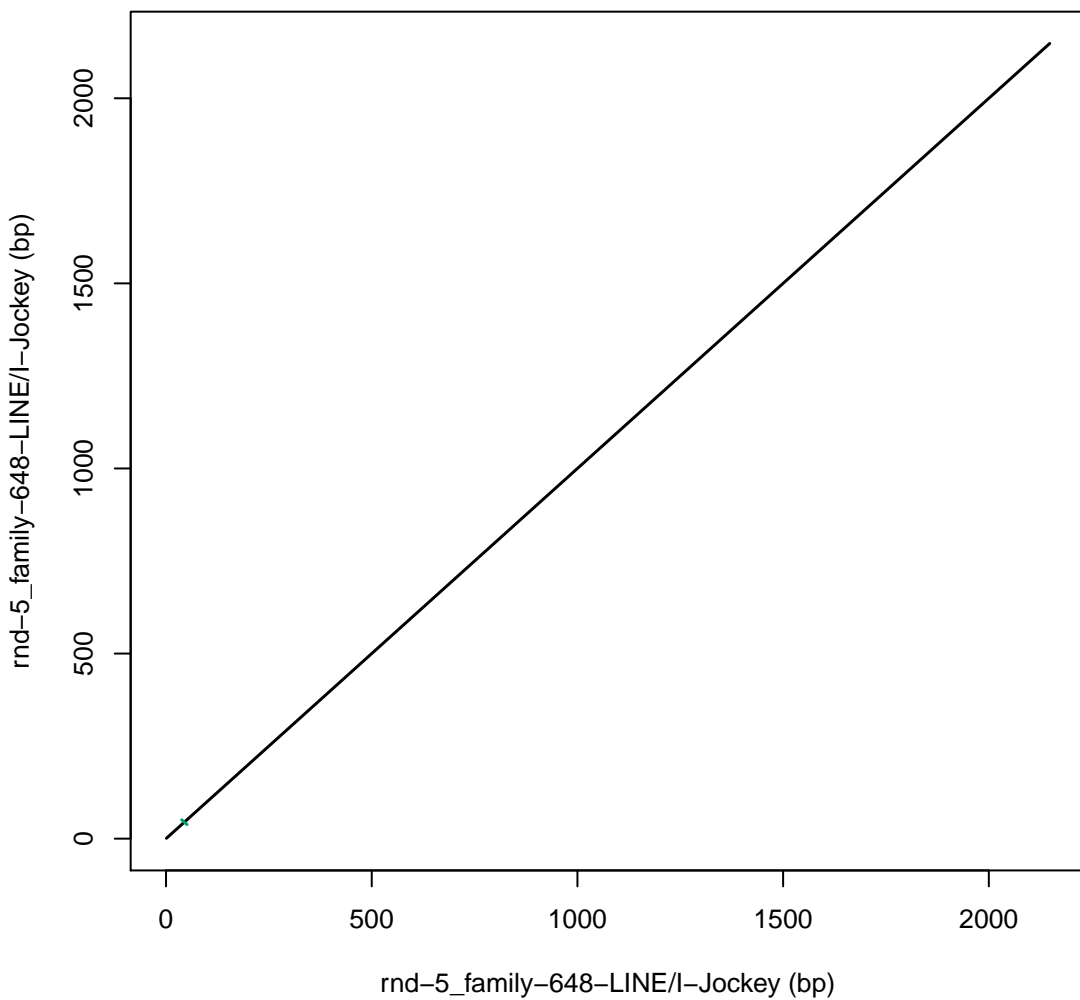
TE: rnd-5\_family-648-LINE/I-Jockey  
consensus size: 2148bp; fragments: 38; full length: 0 (>=1933.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

