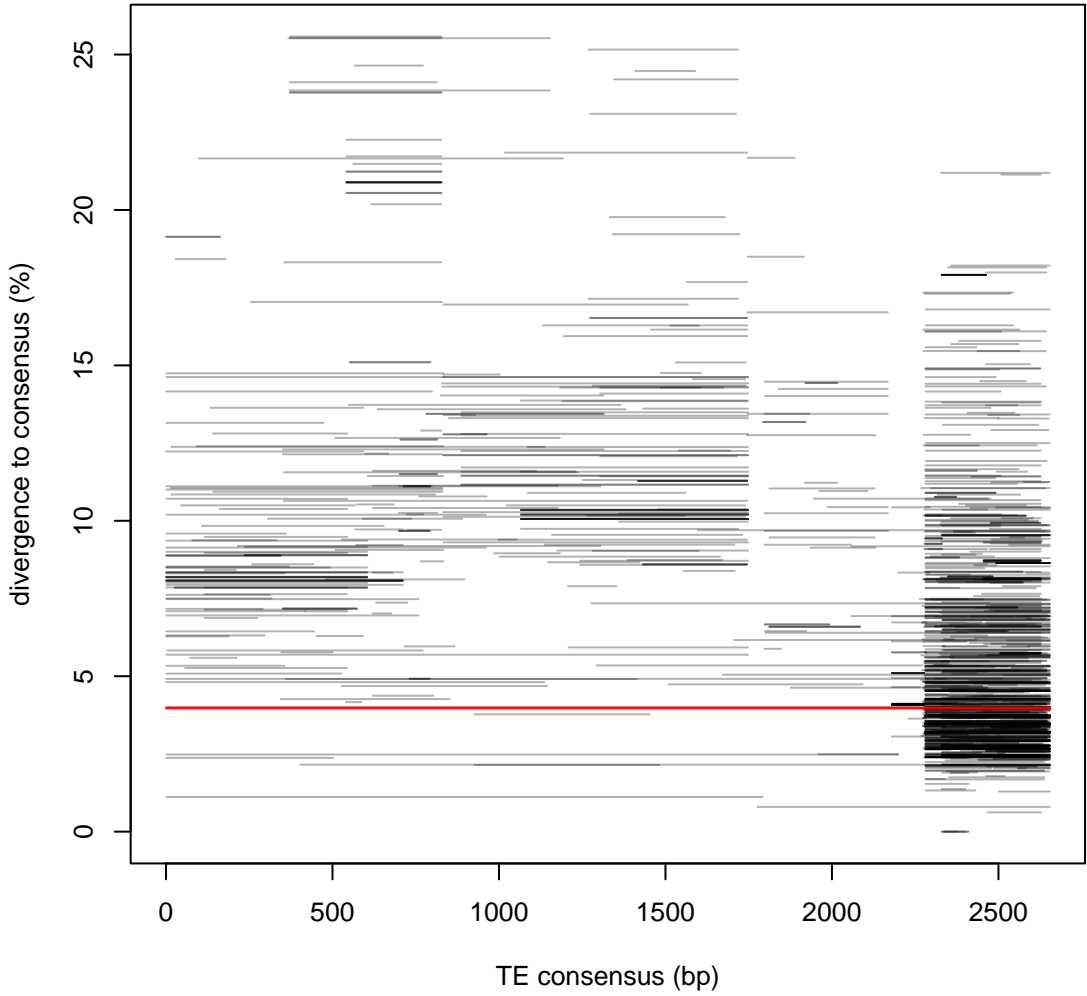
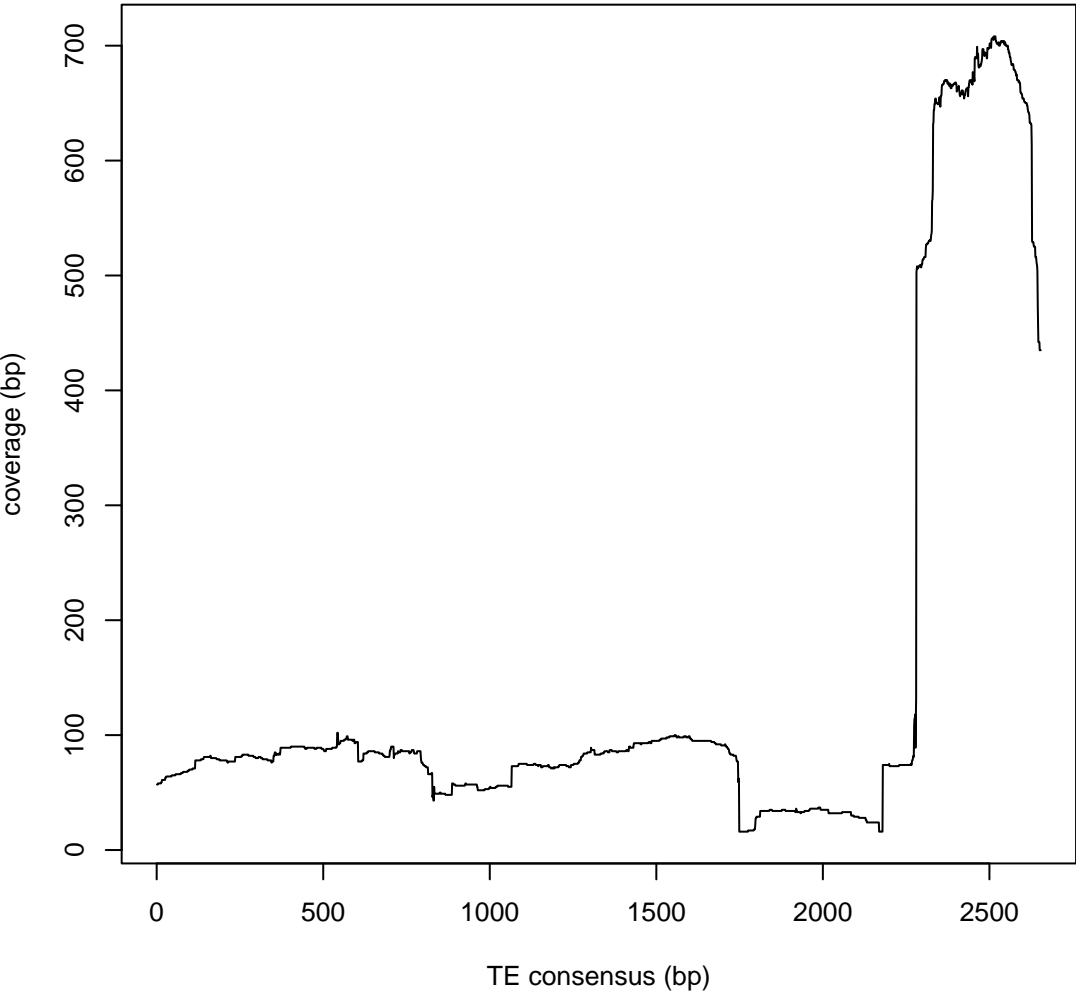


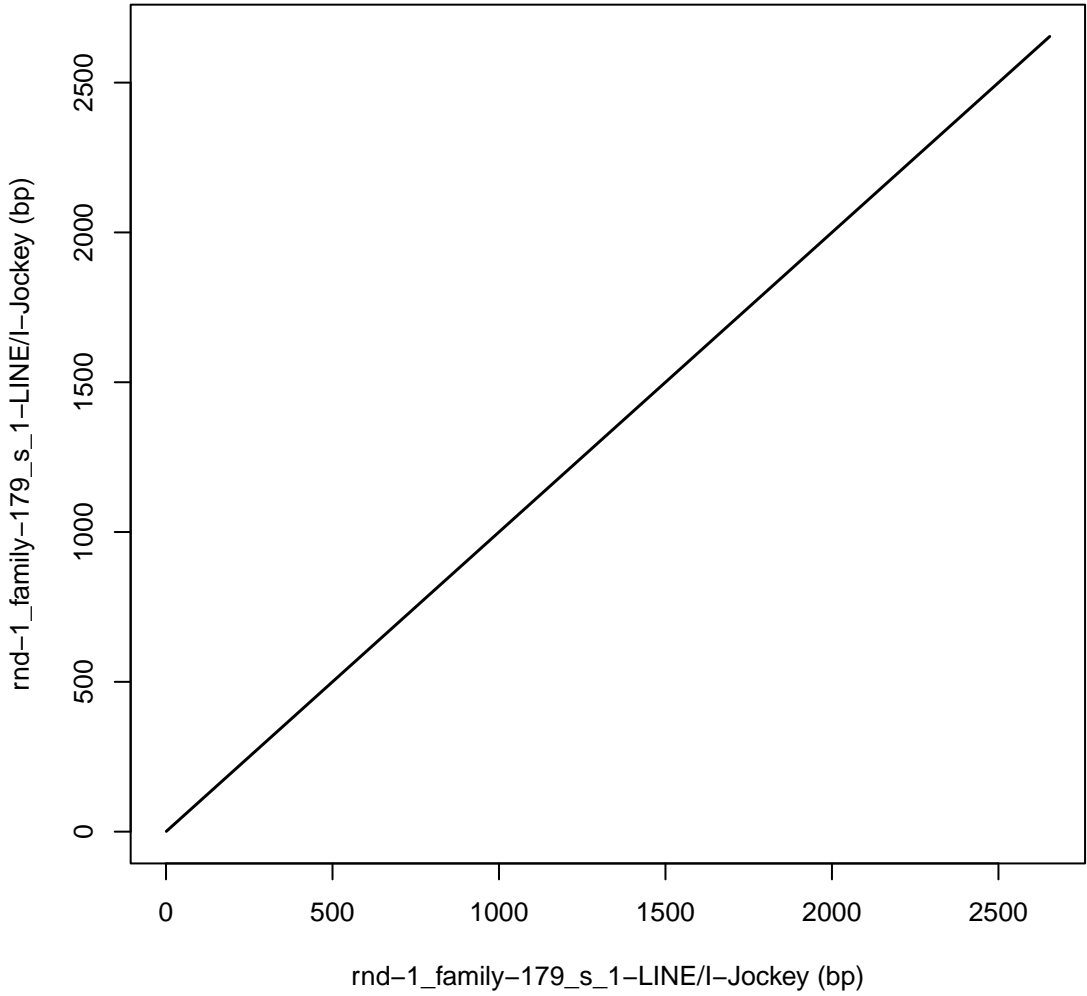
TE: rnd-1\_family-179\_s\_1-LINE/I-Jockey  
consensus size: 2654bp; fragments: 1257; full length: 1 (>=2388.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

