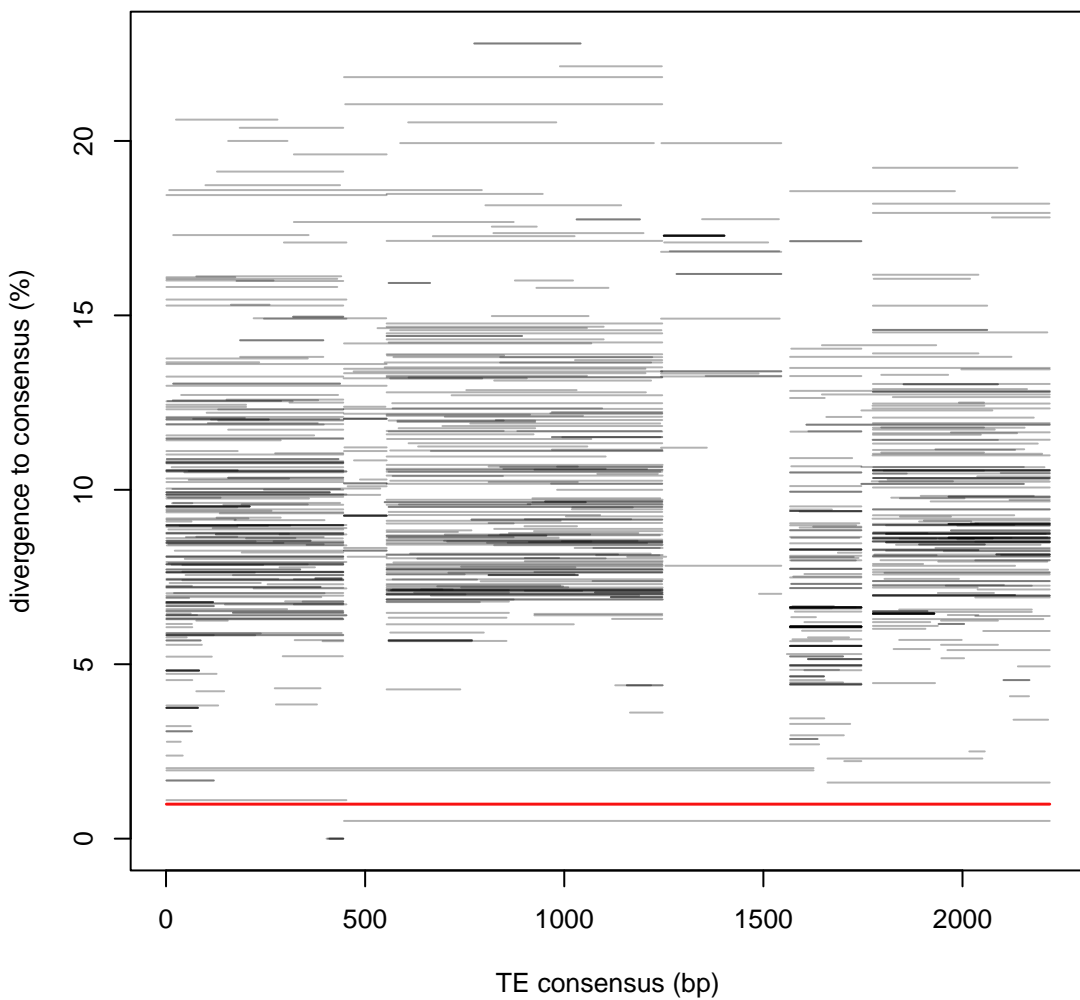
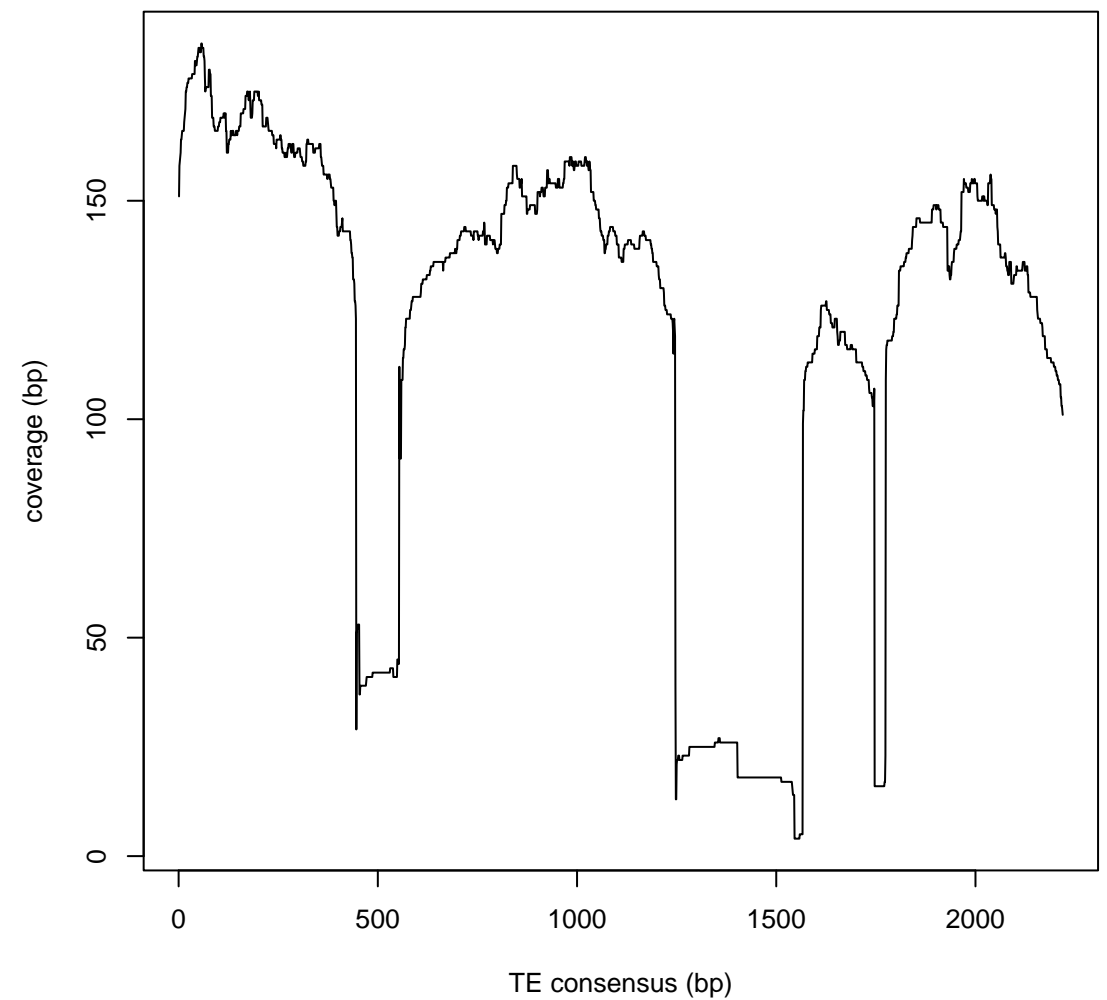


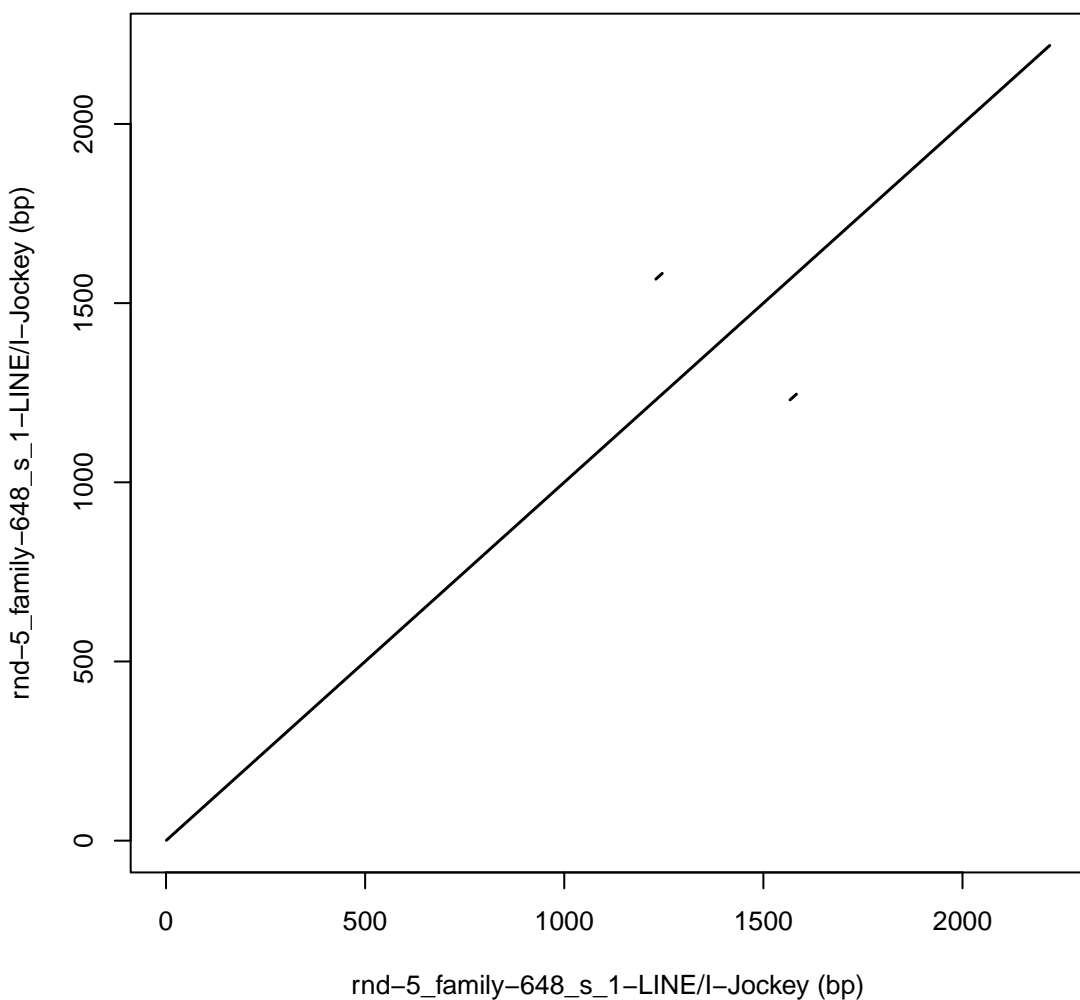
TE: rnd-5\_family-648\_s\_1-LINE/I-Jockey  
consensus size: 2219bp; fragments: 851; full length: 1 (>=1997.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

