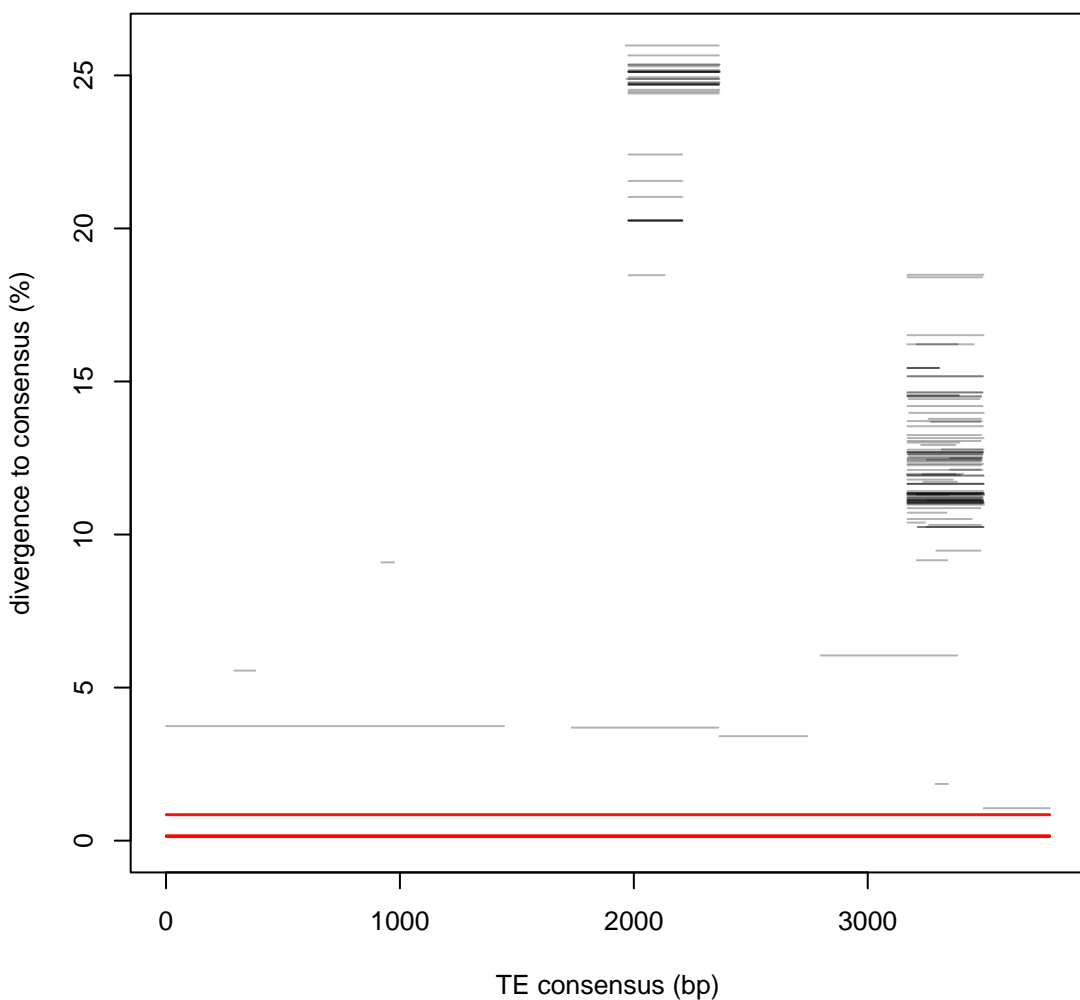
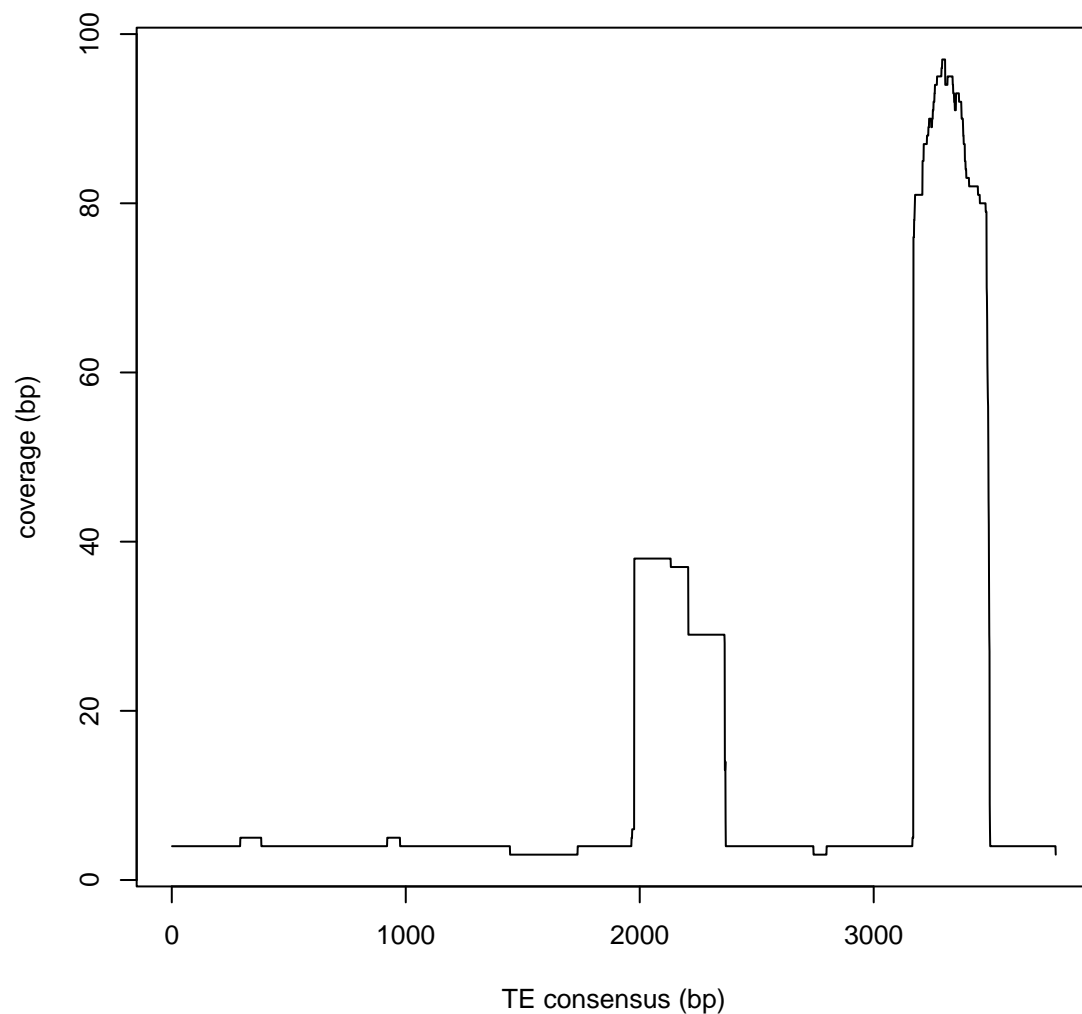


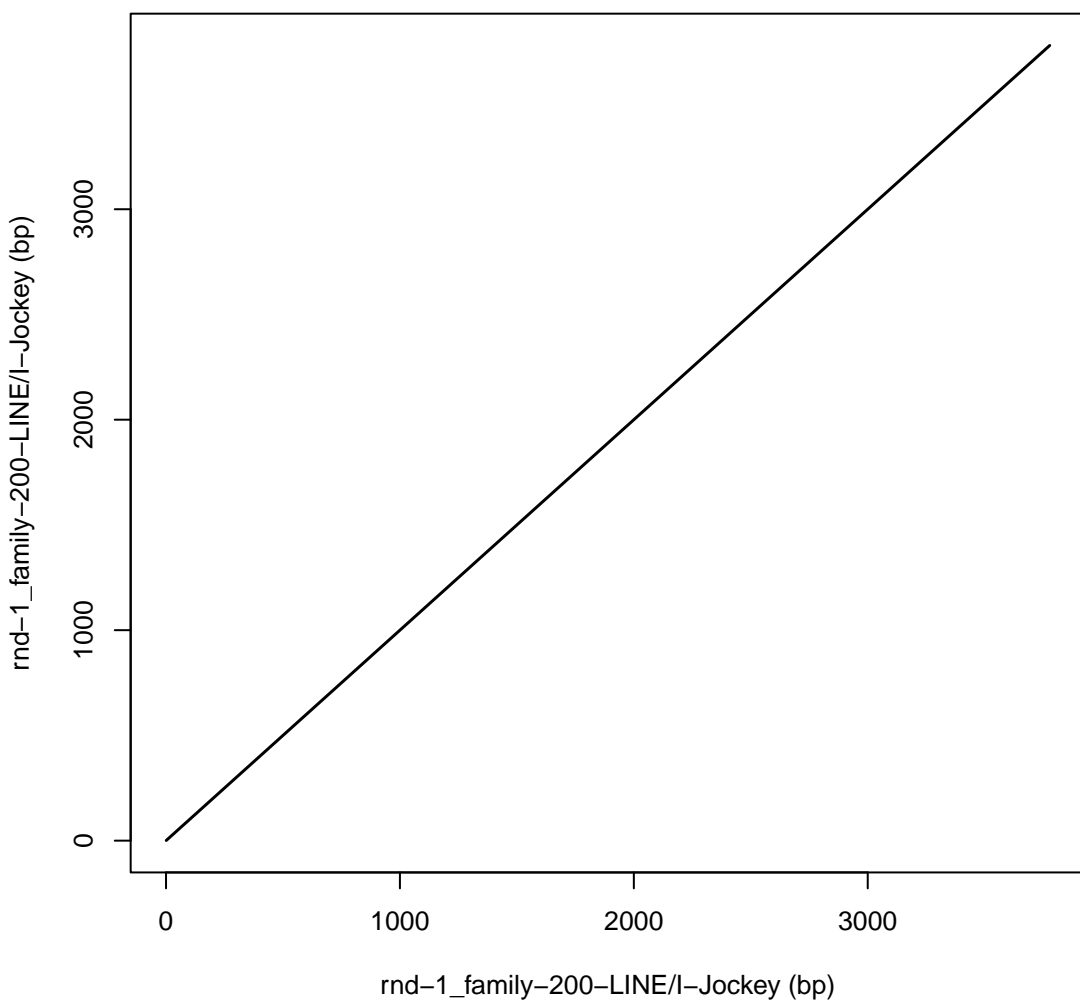
TE: rnd-1\_family-200-LINE/l-Jockey  
consensus size: 3778bp; fragments: 141; full length: 3 (>=3400.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

