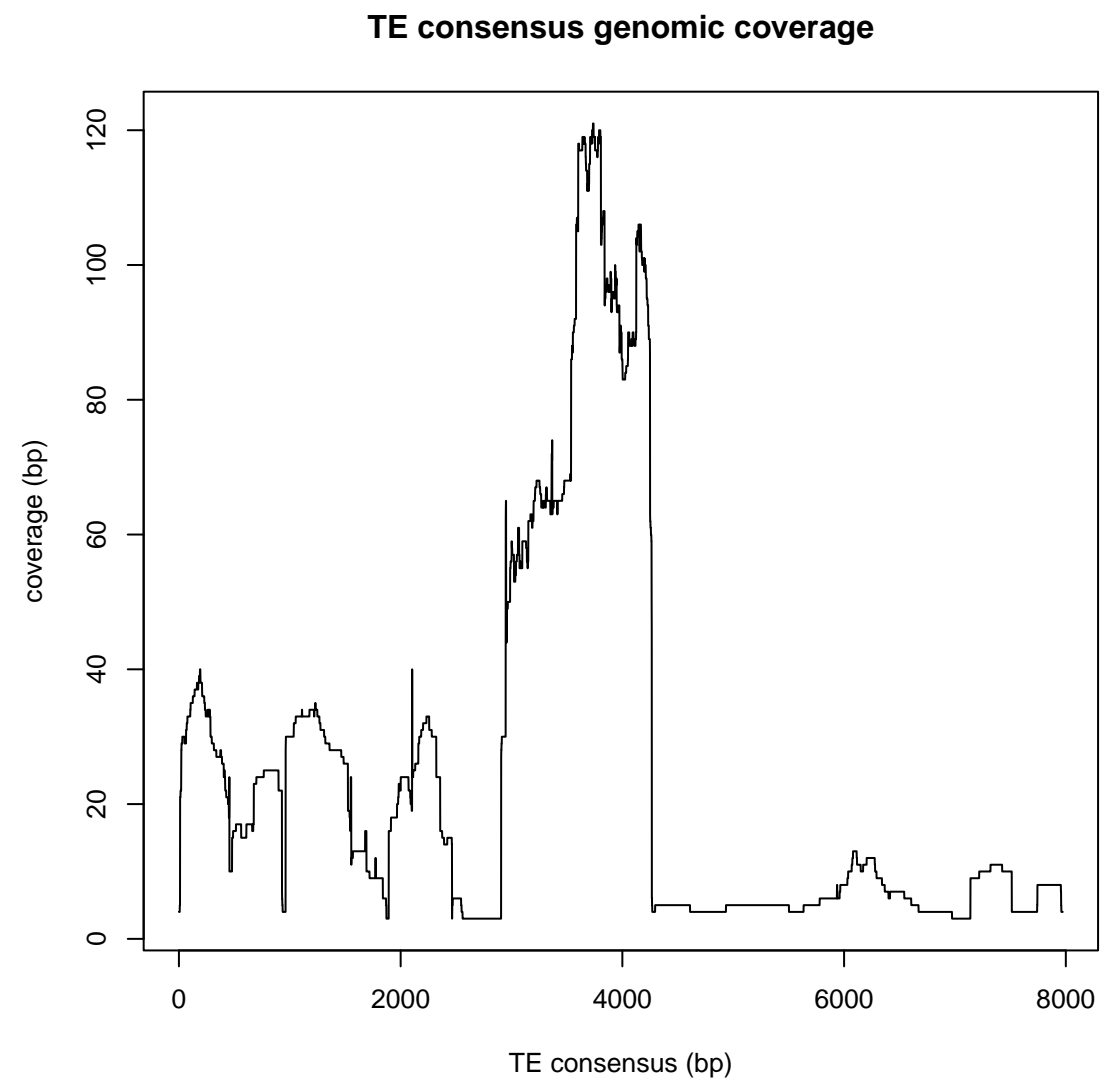
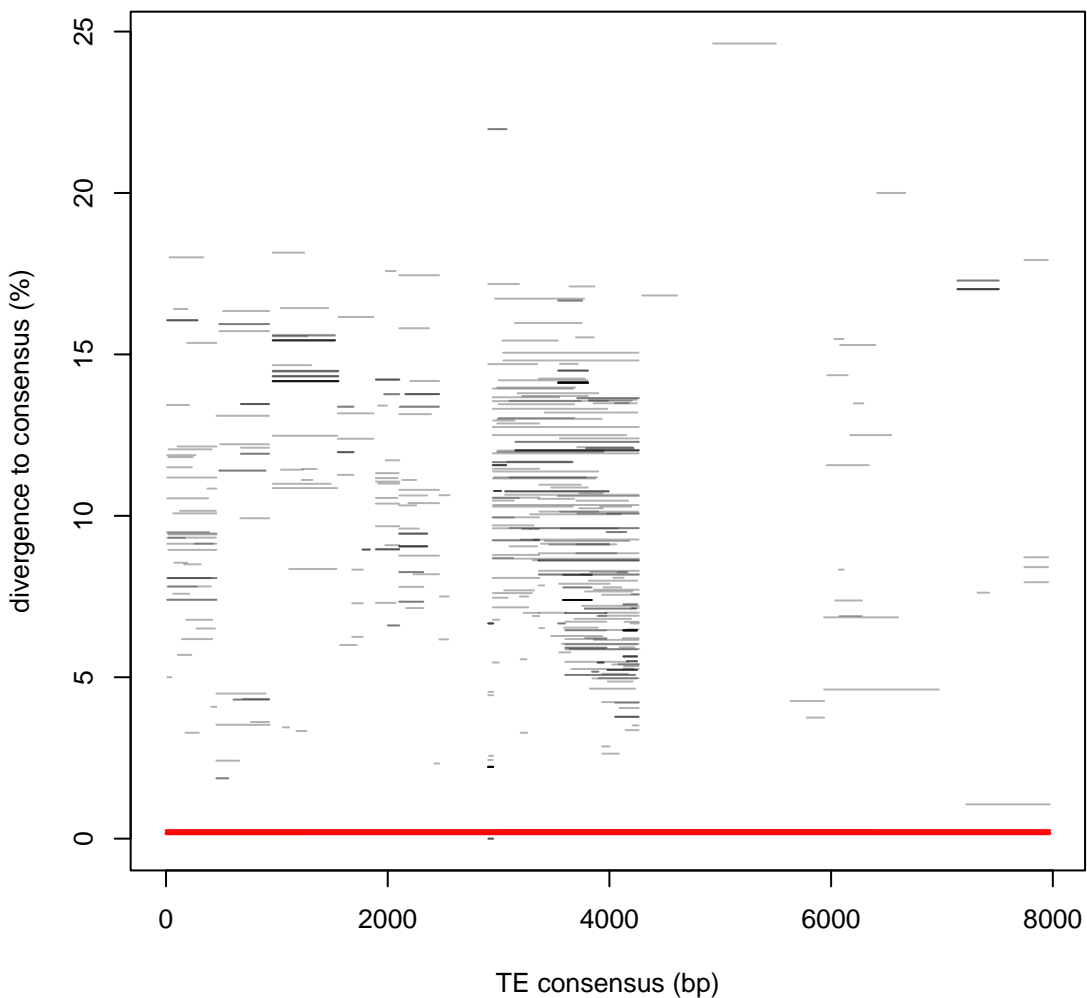
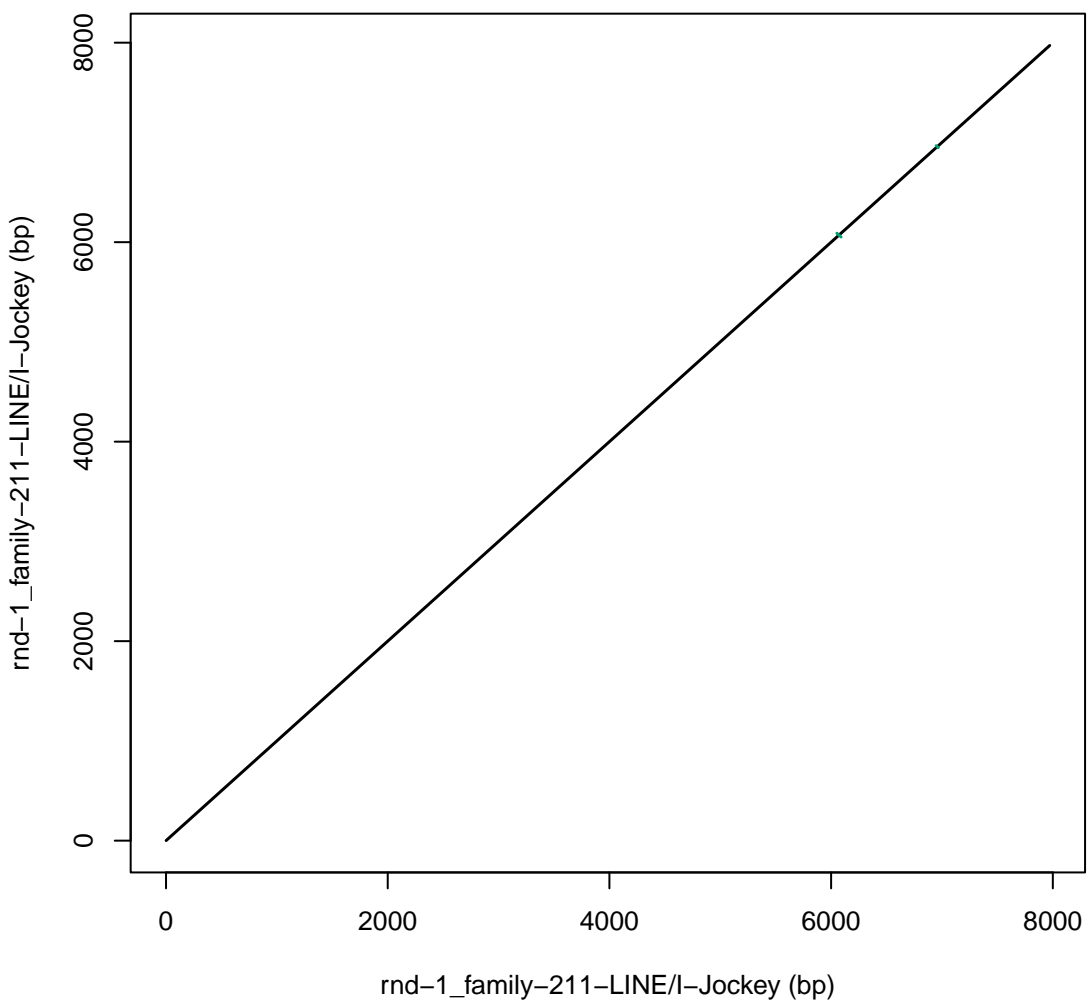


TE: rnd-1_family-211-LINE/I-Jockey
consensus size: 7972bp; fragments: 481; full length: 3 (>=7174.8bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

