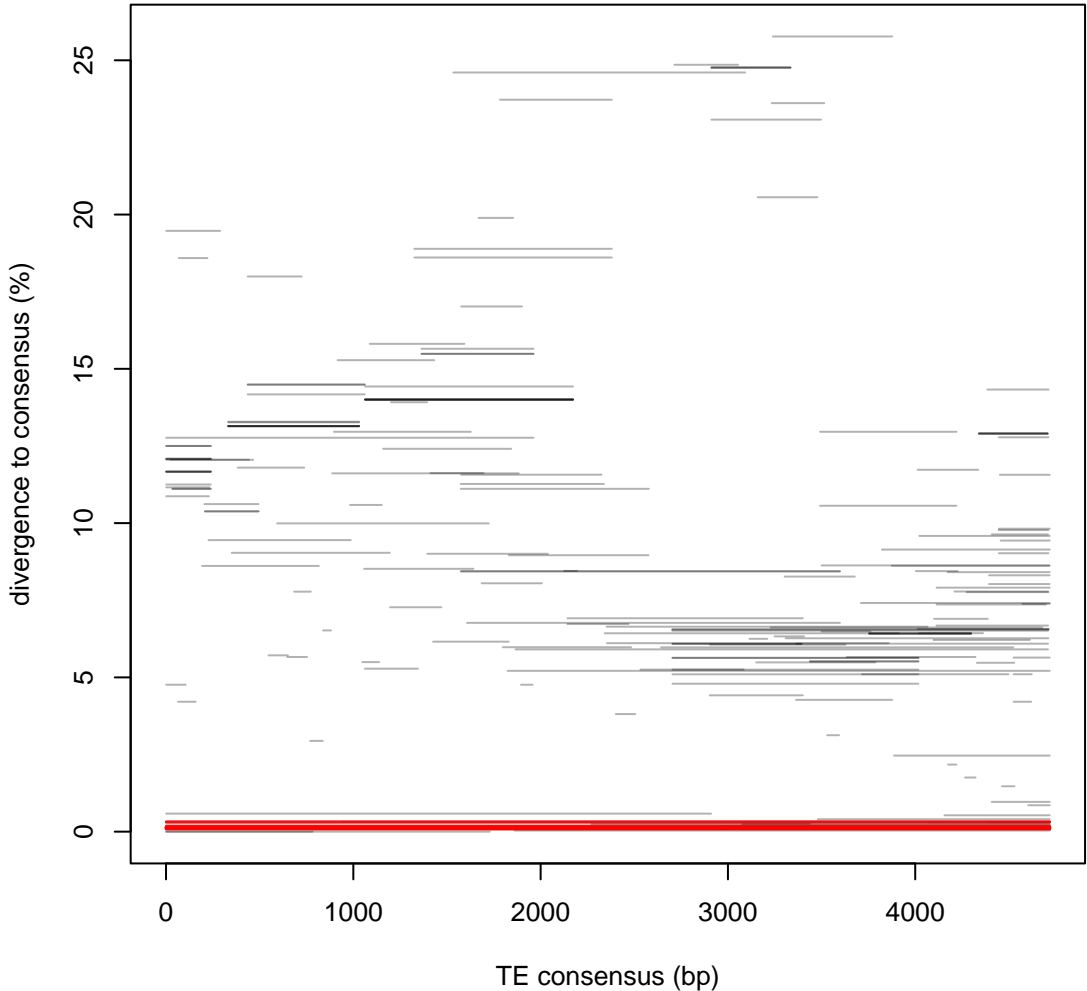
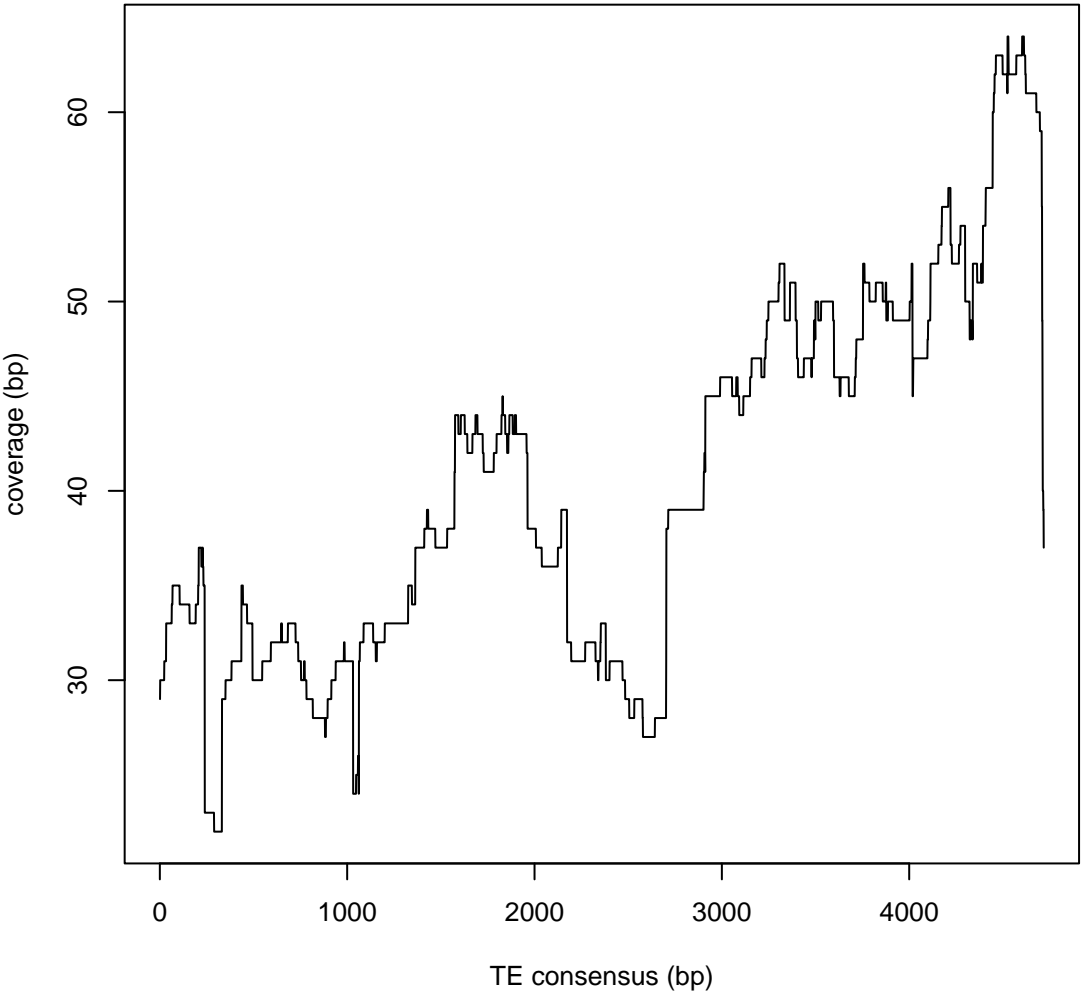


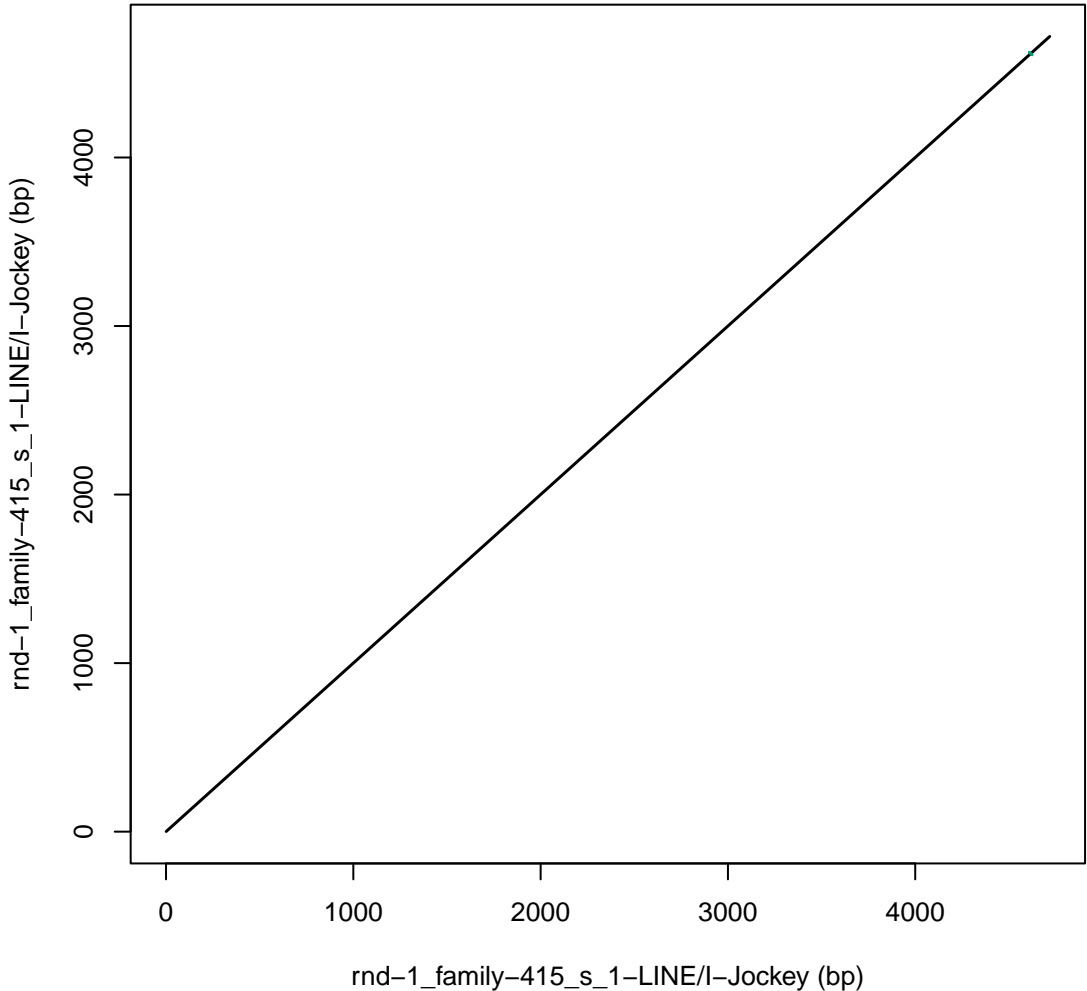
TE: rnd-1\_family-415\_s\_1-LINE/I-Jockey  
consensus size: 4718bp; fragments: 204; full length: 10 (>=4246.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

