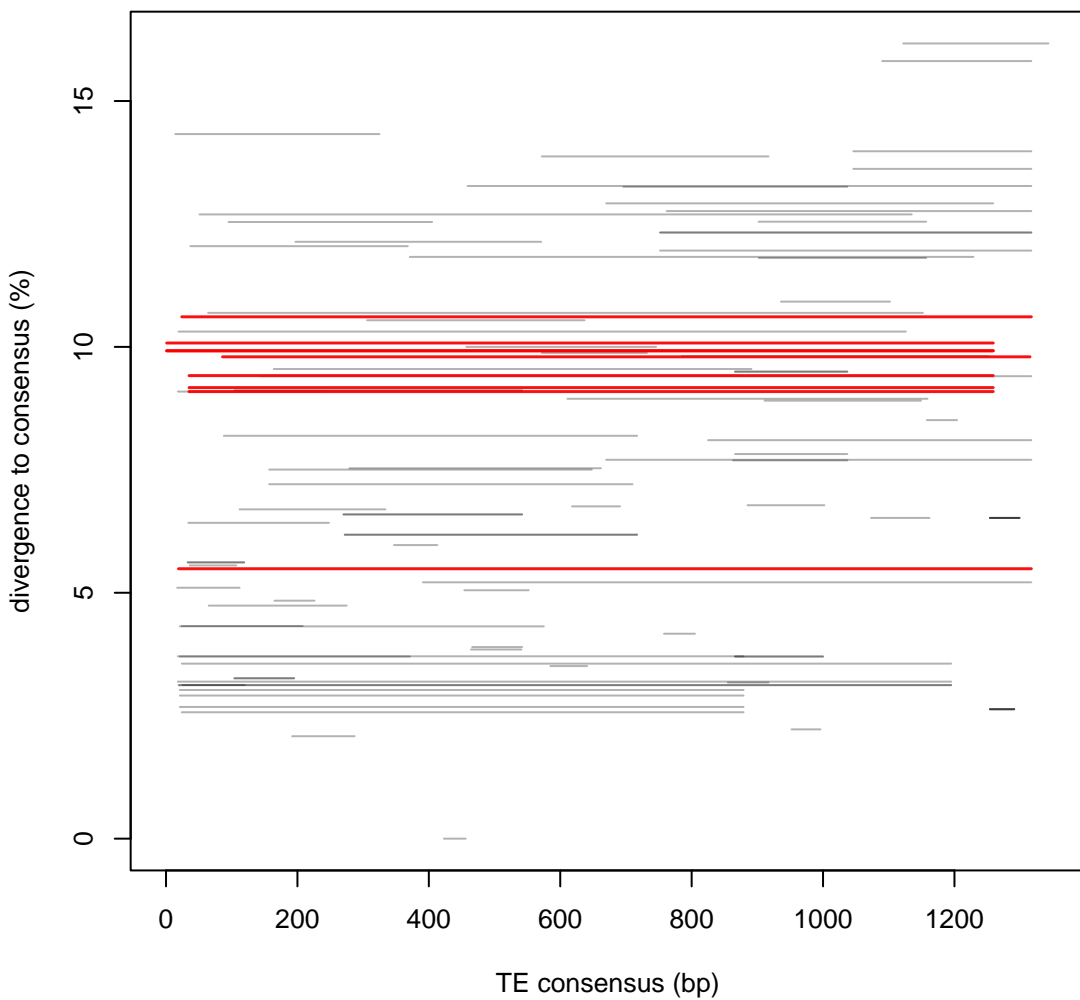
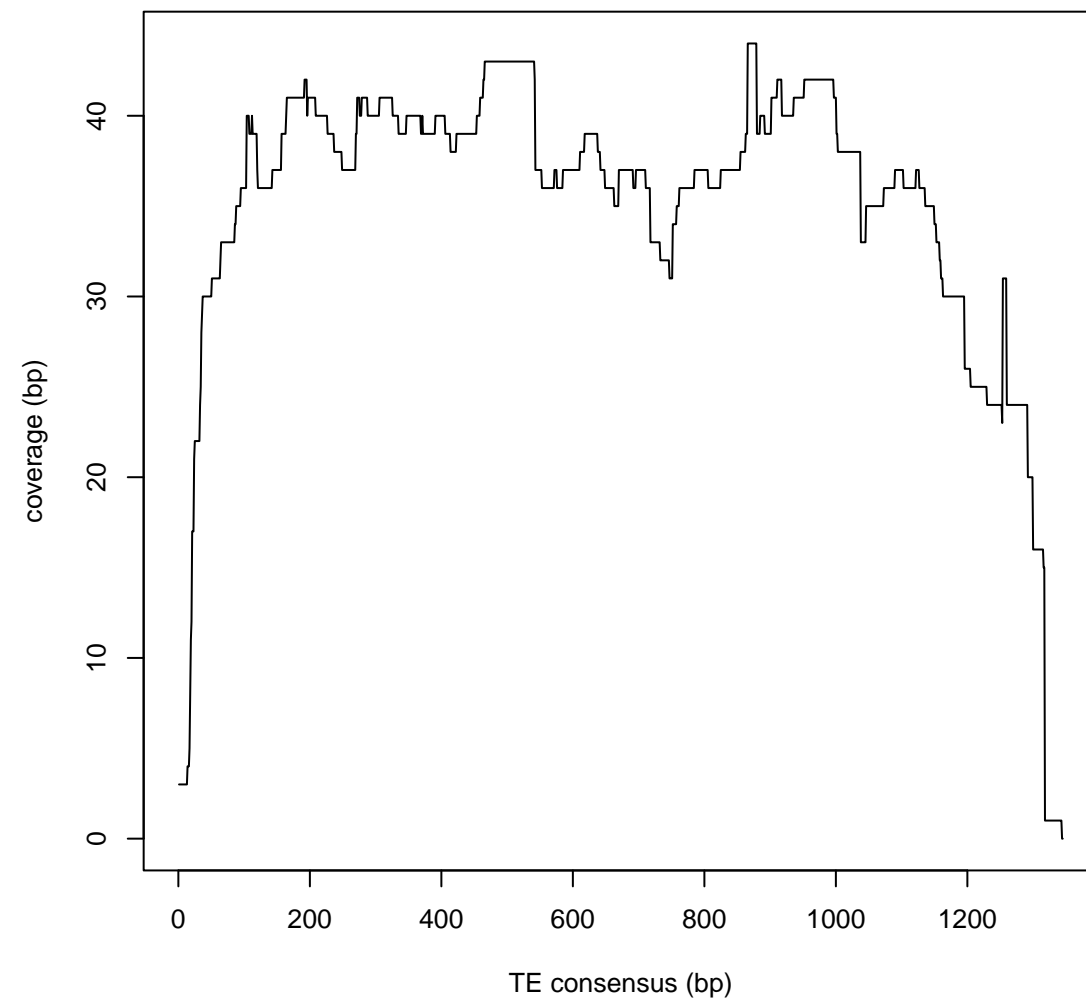


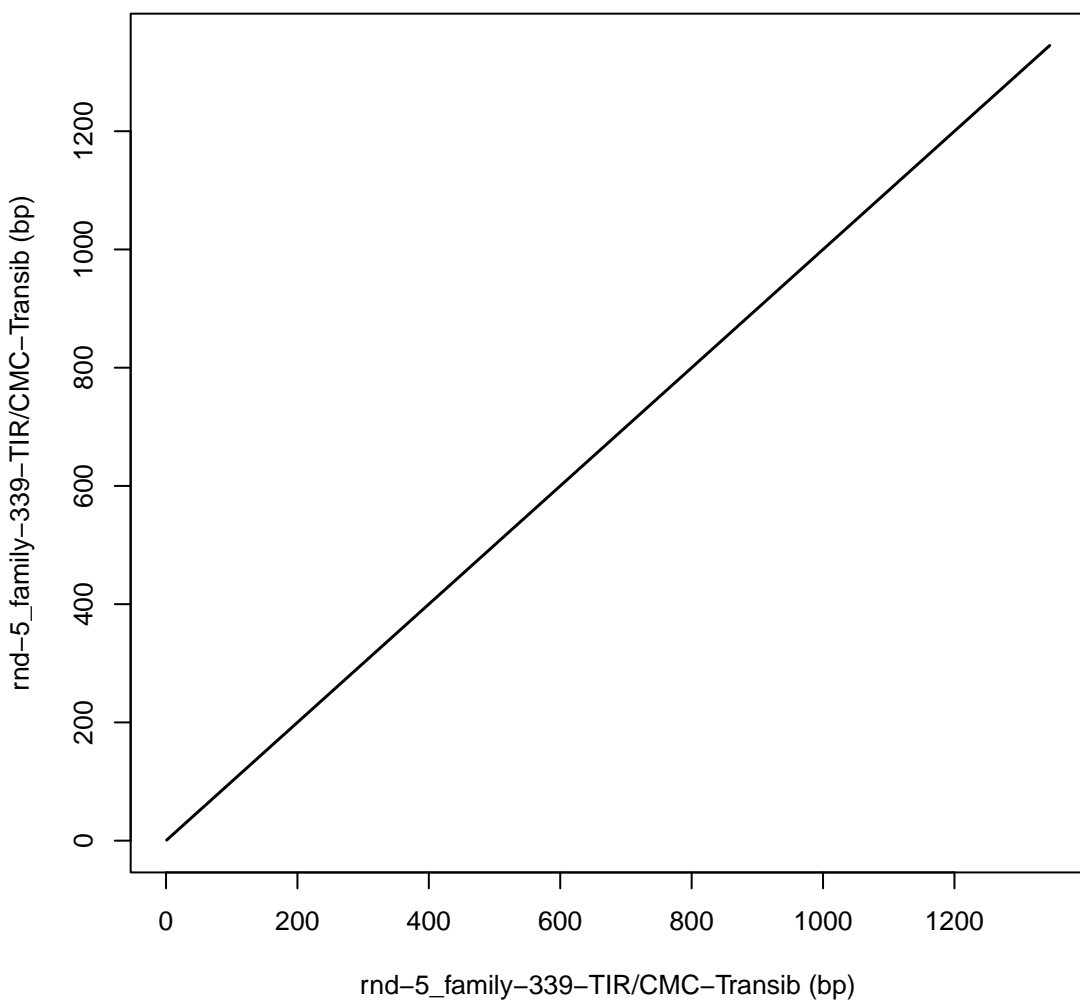
TE: rnd-5\_family-339-TIR/CMC-Transib  
consensus size: 1345bp; fragments: 105; full length: 9 (>=1210.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

