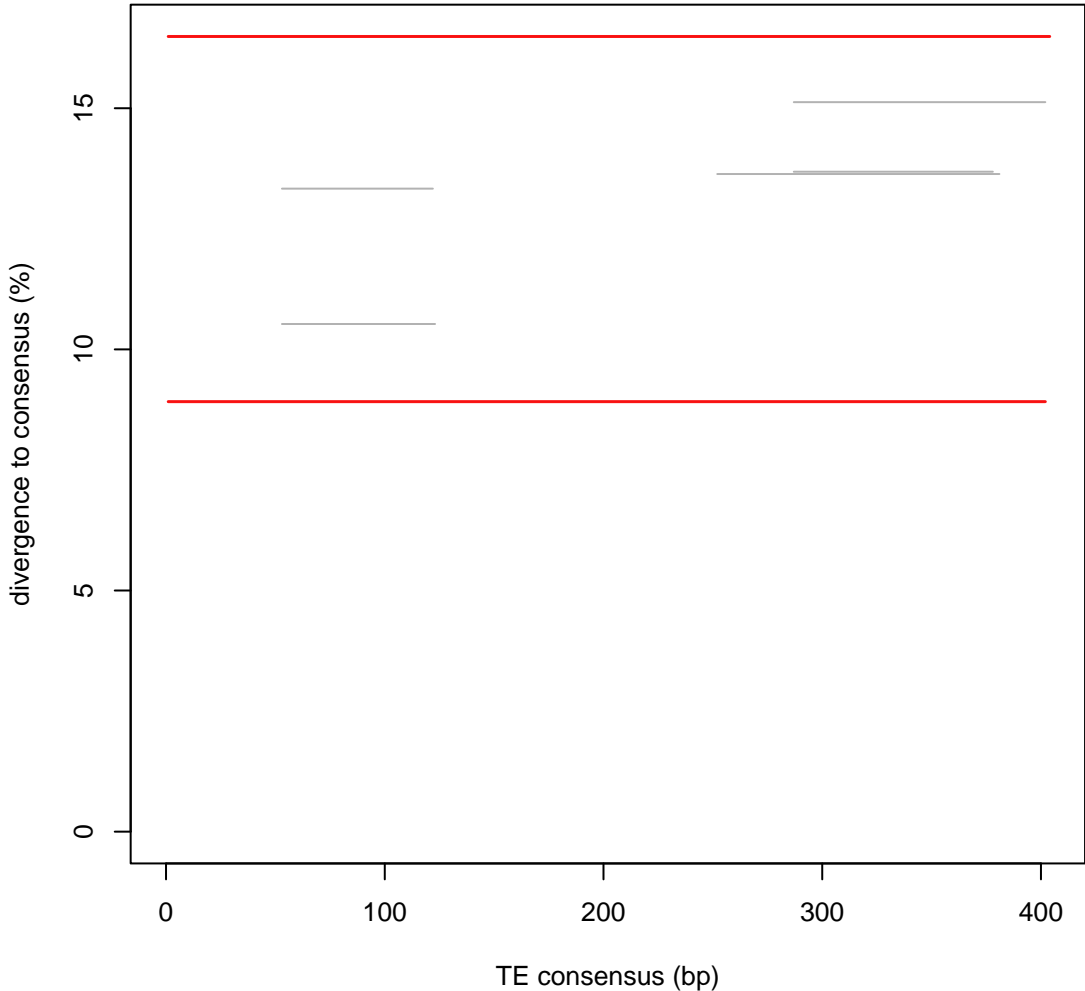
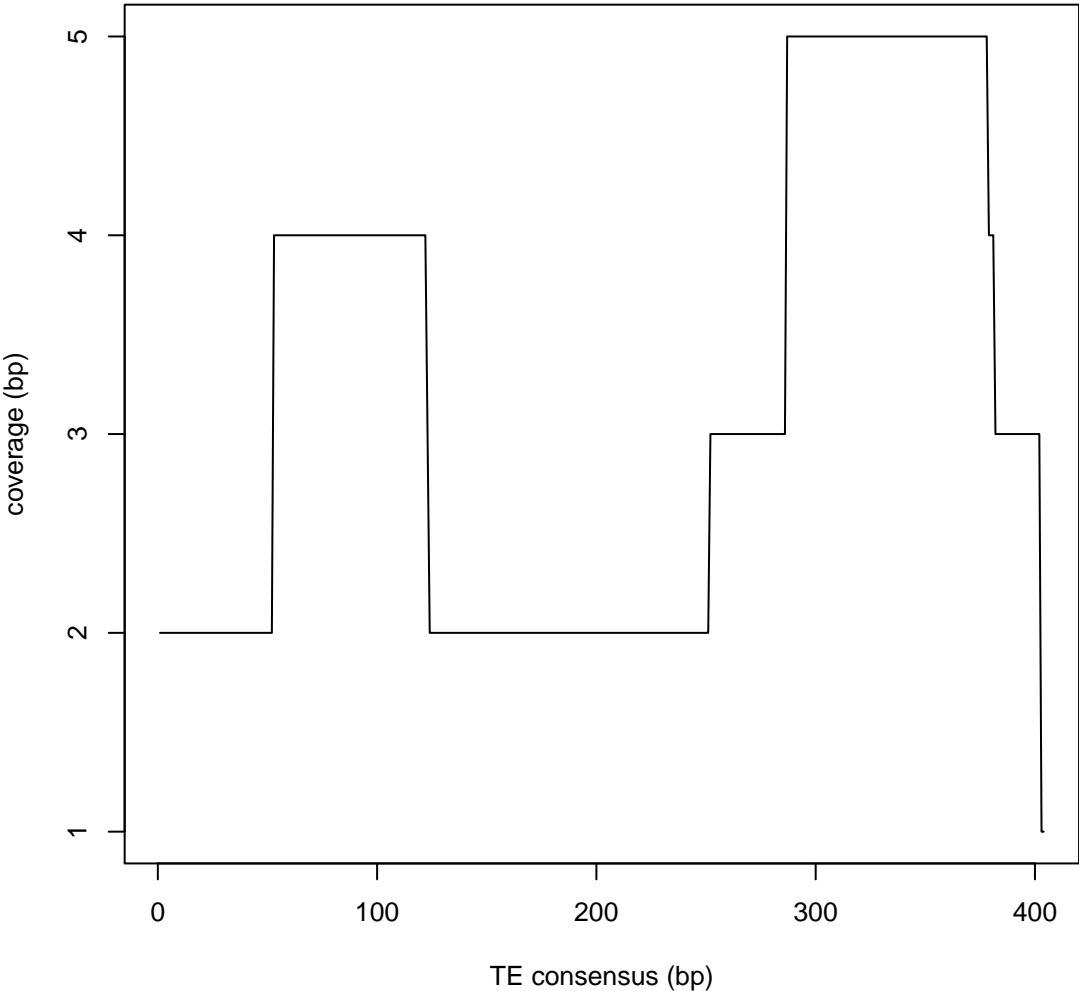


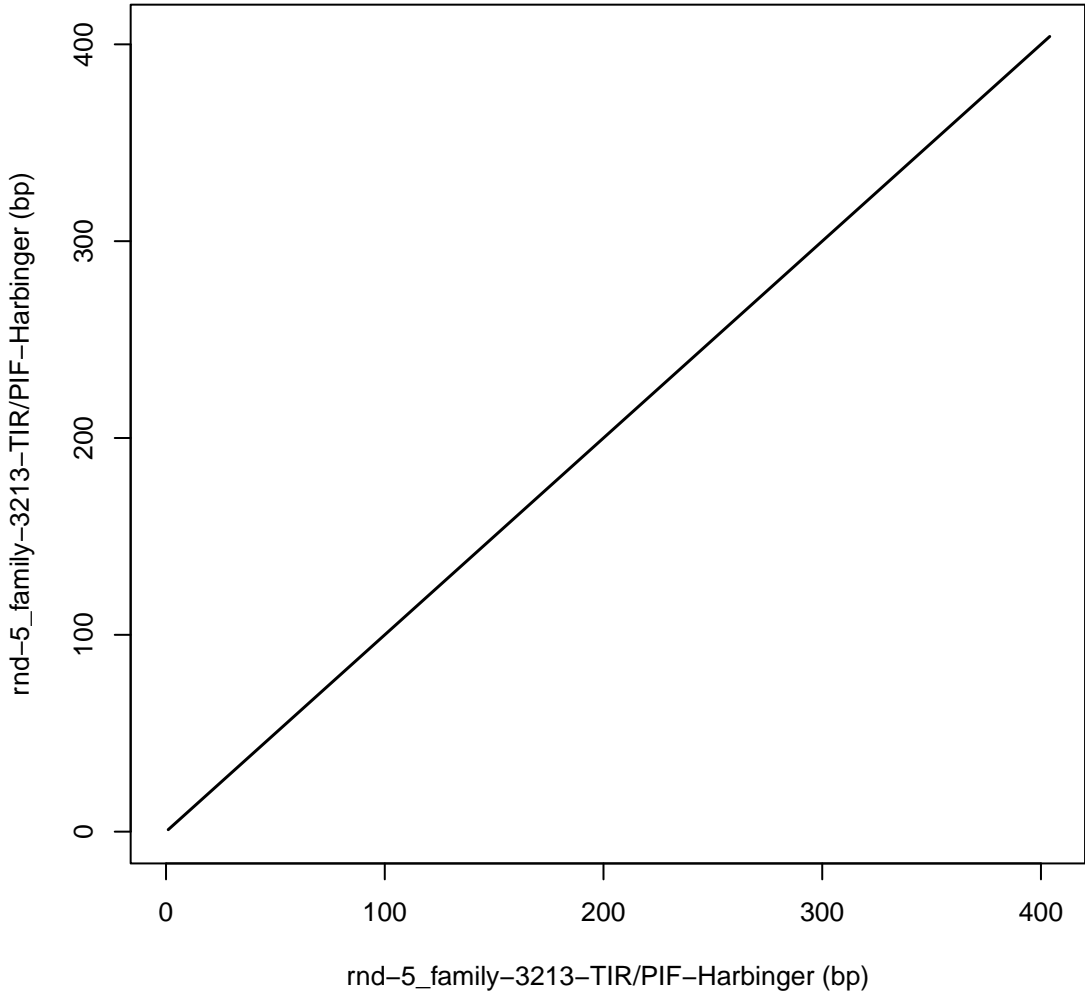
TE: rnd-5_family-3213-TIR/PIF-Harbinger
consensus size: 404bp; fragments: 7; full length: 2 (>=363.6bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

