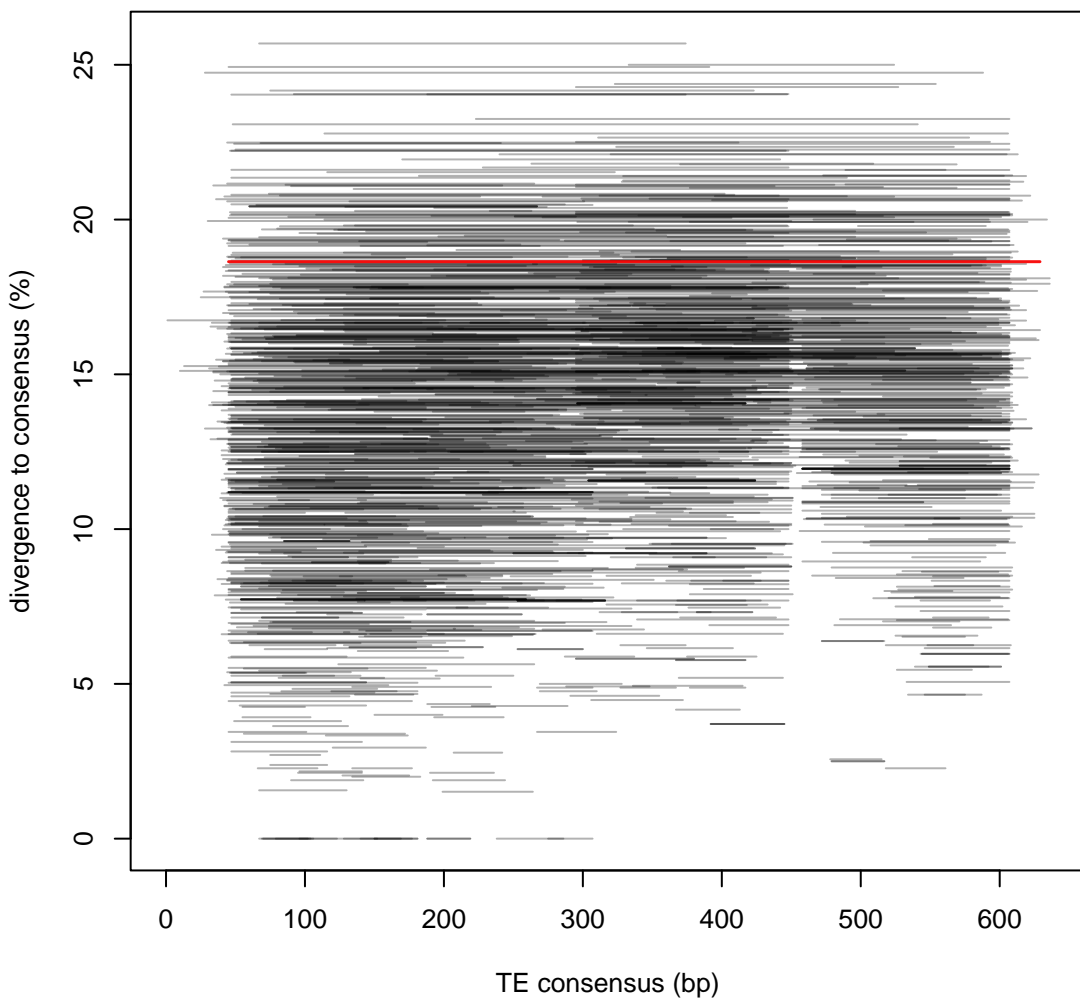
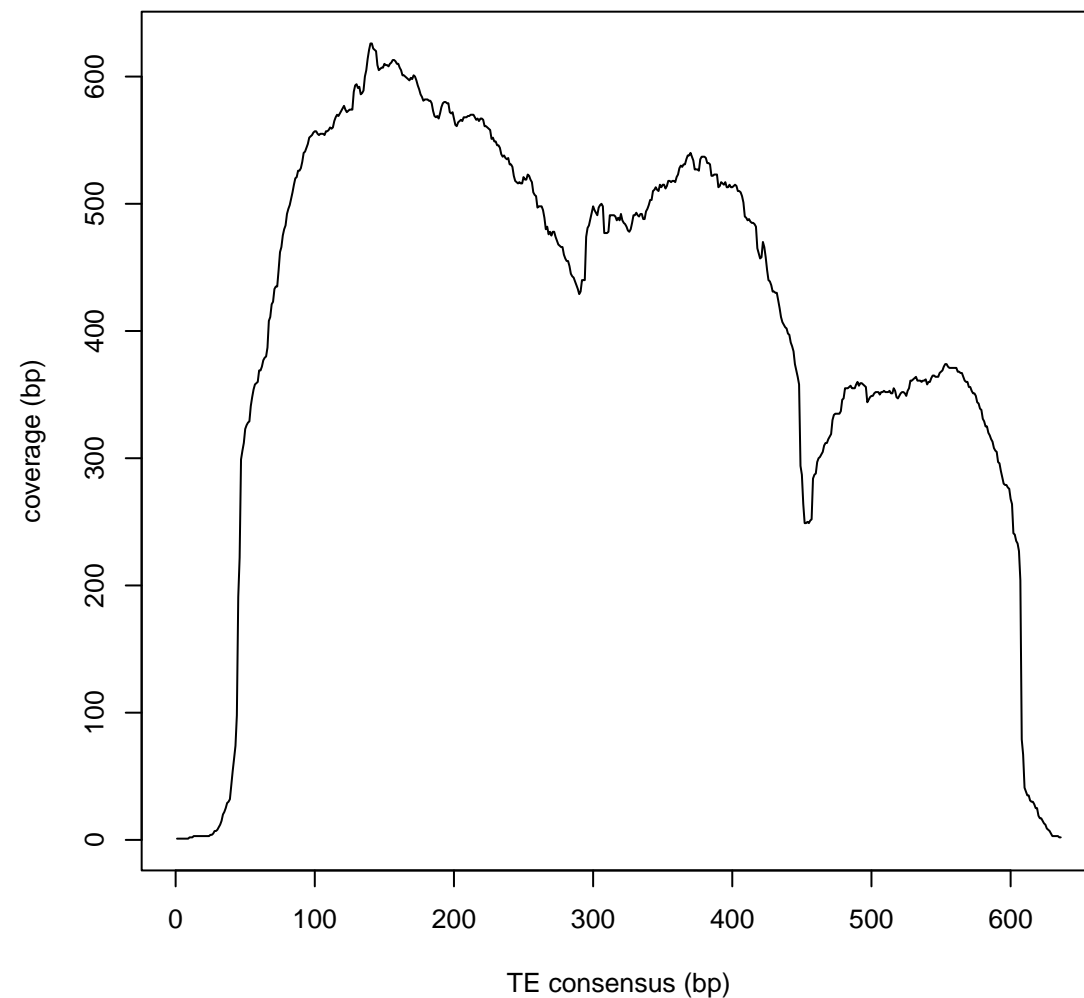


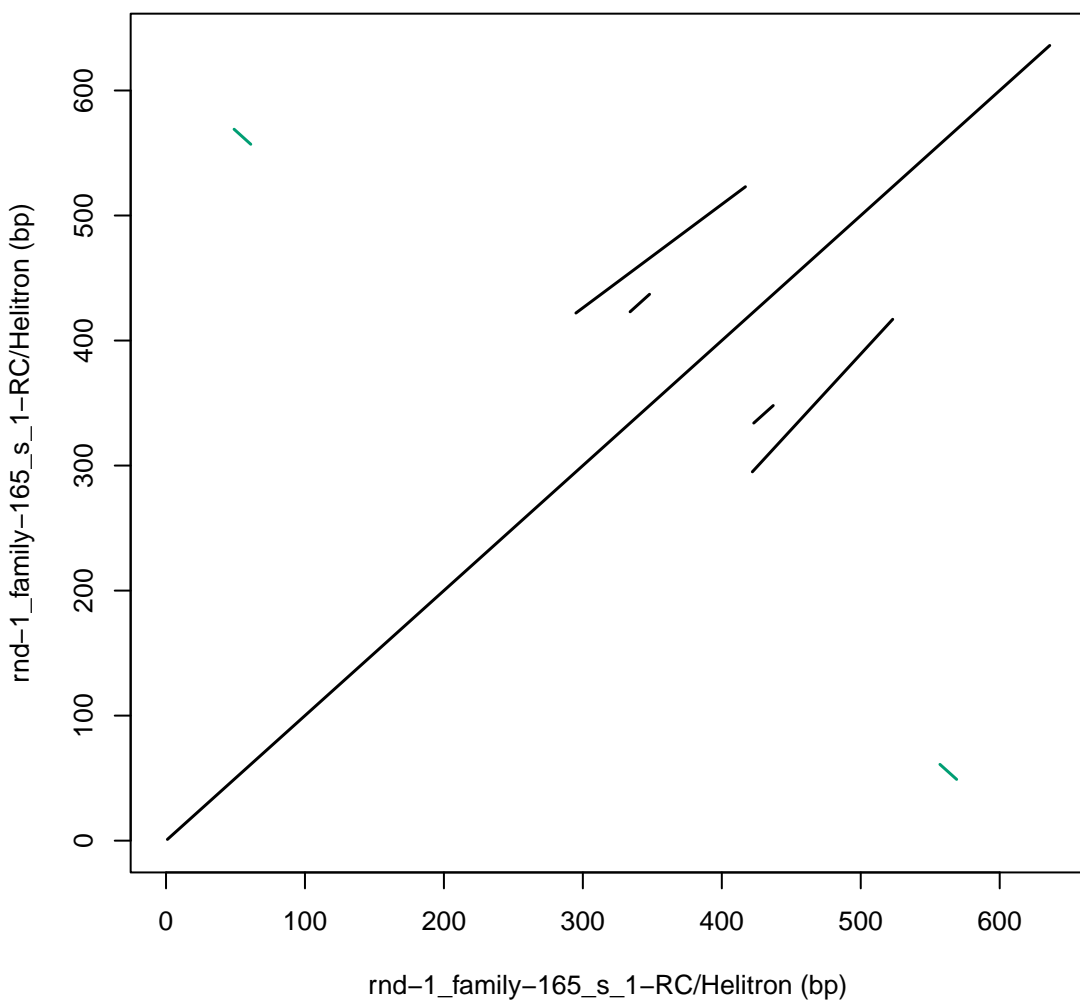
TE: rnd-1\_family-165\_s\_1-RC/Helitron  
consensus size: 636bp; fragments: 1411; full length: 1 (>=572.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

