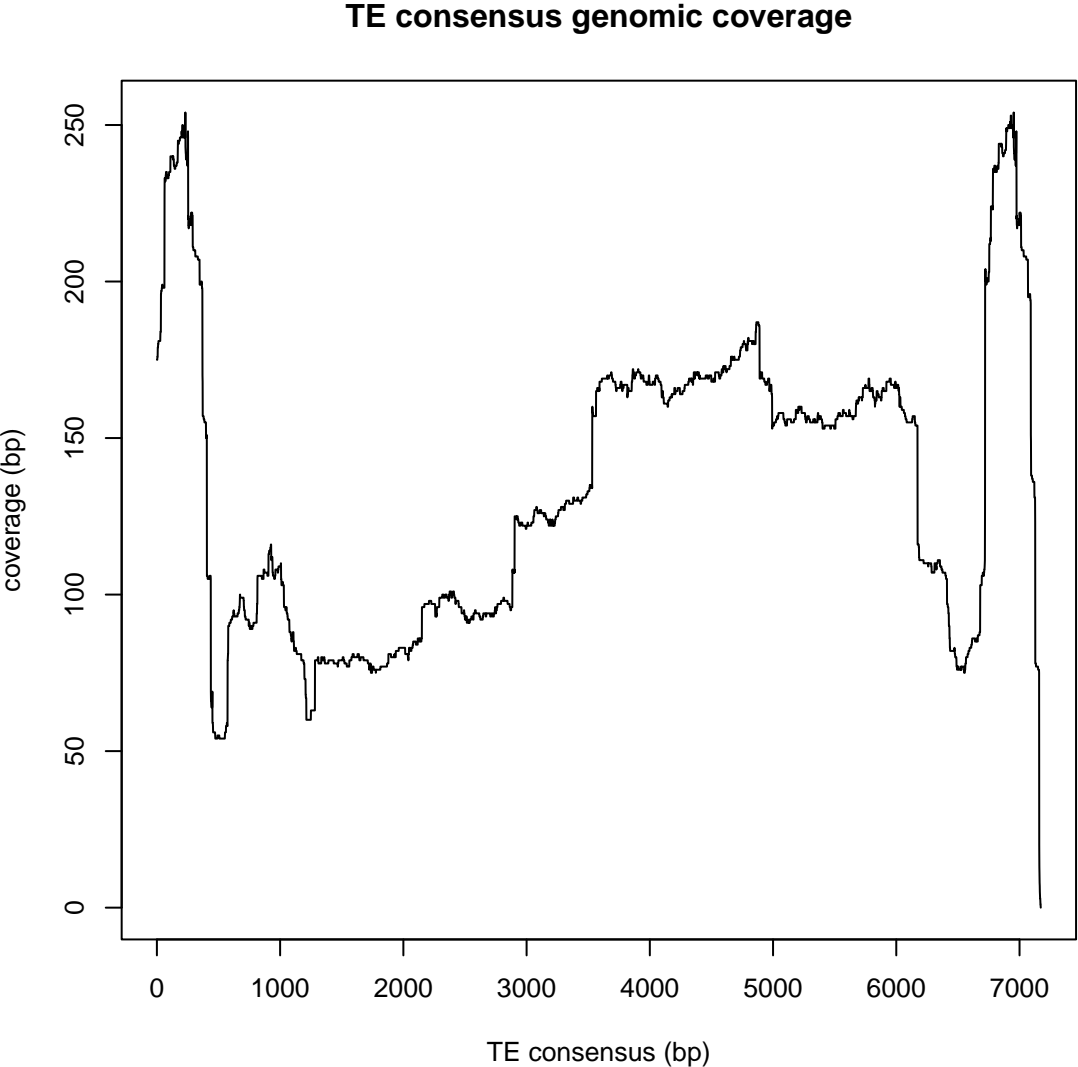
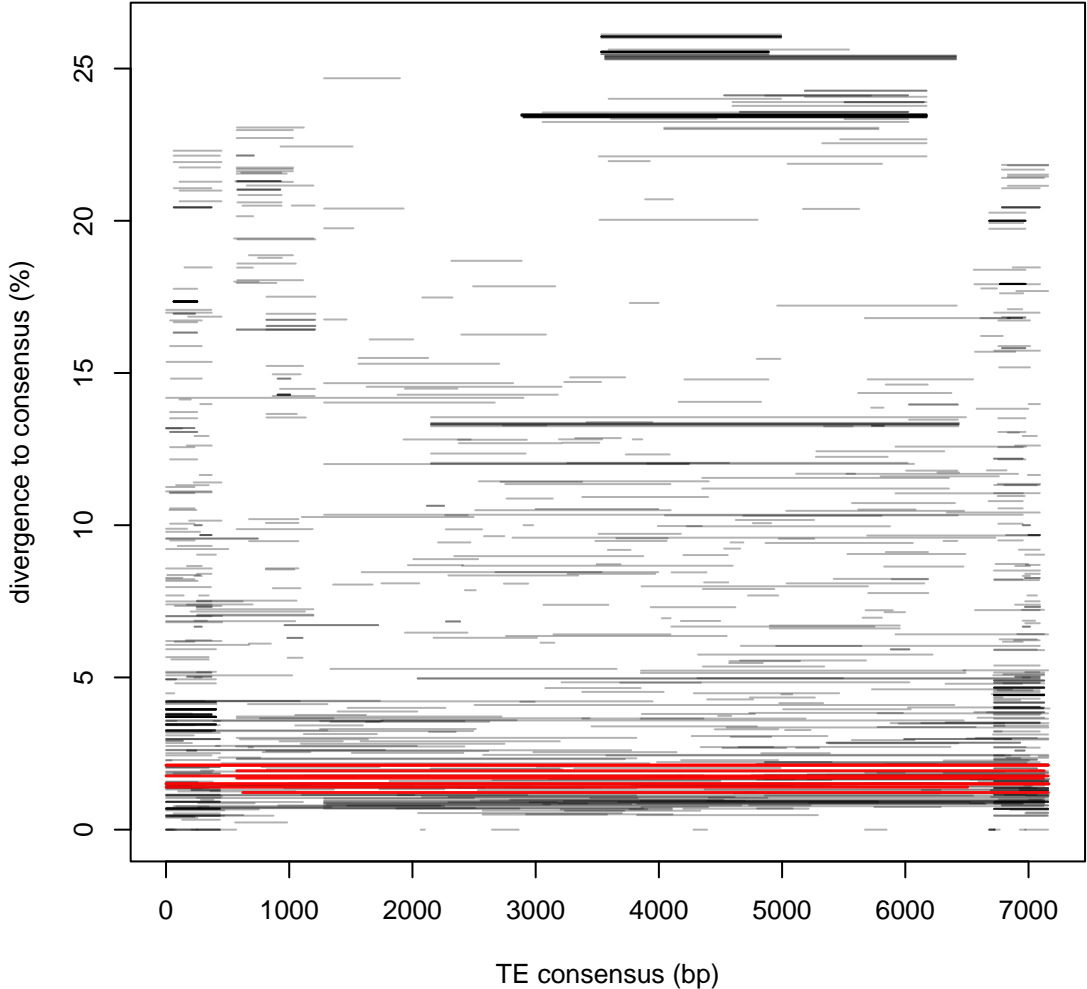
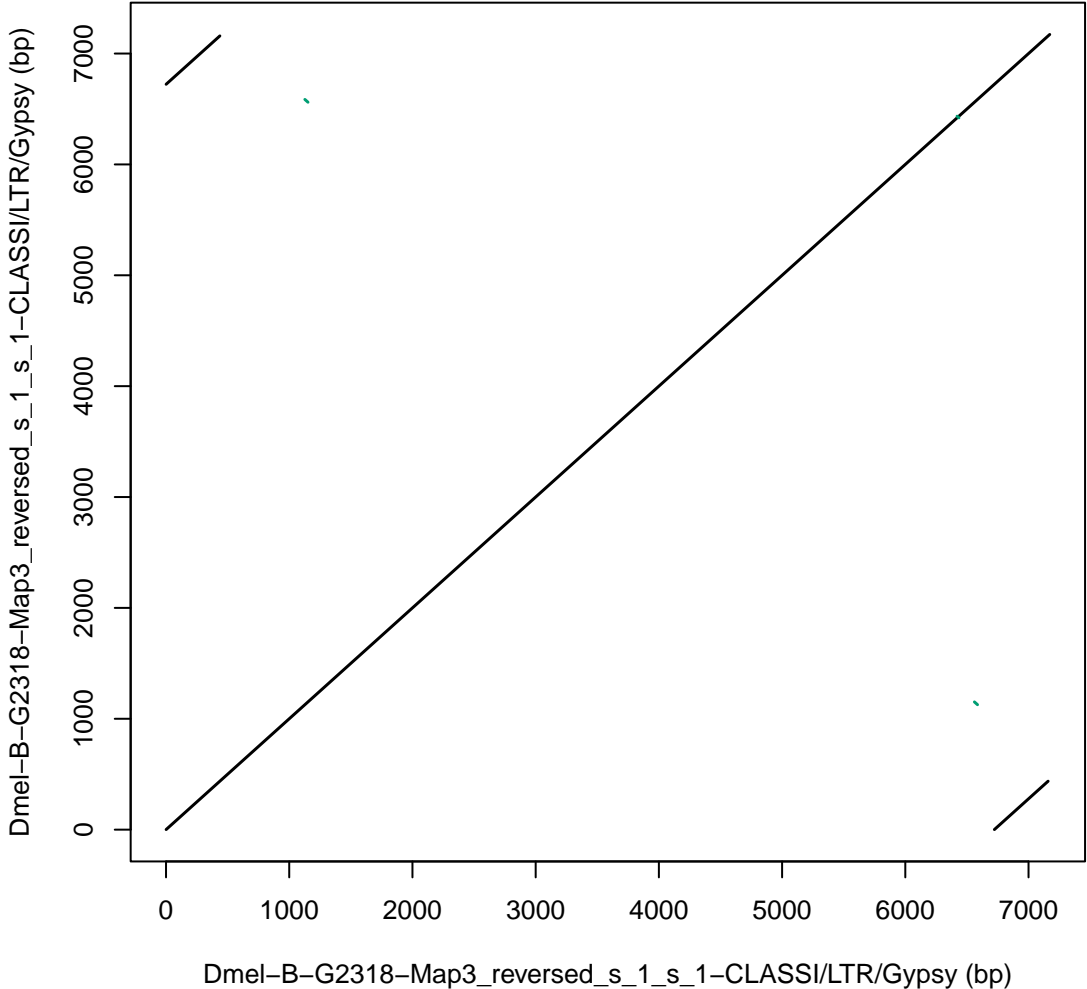


TE: Dmel-B-G2318-Map3\_reversed\_s\_1\_s\_1-CLASSI/LTR/Gypsy  
consensus size: 7172bp; fragments: 1051; full length: 9 (>=6454.8bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

