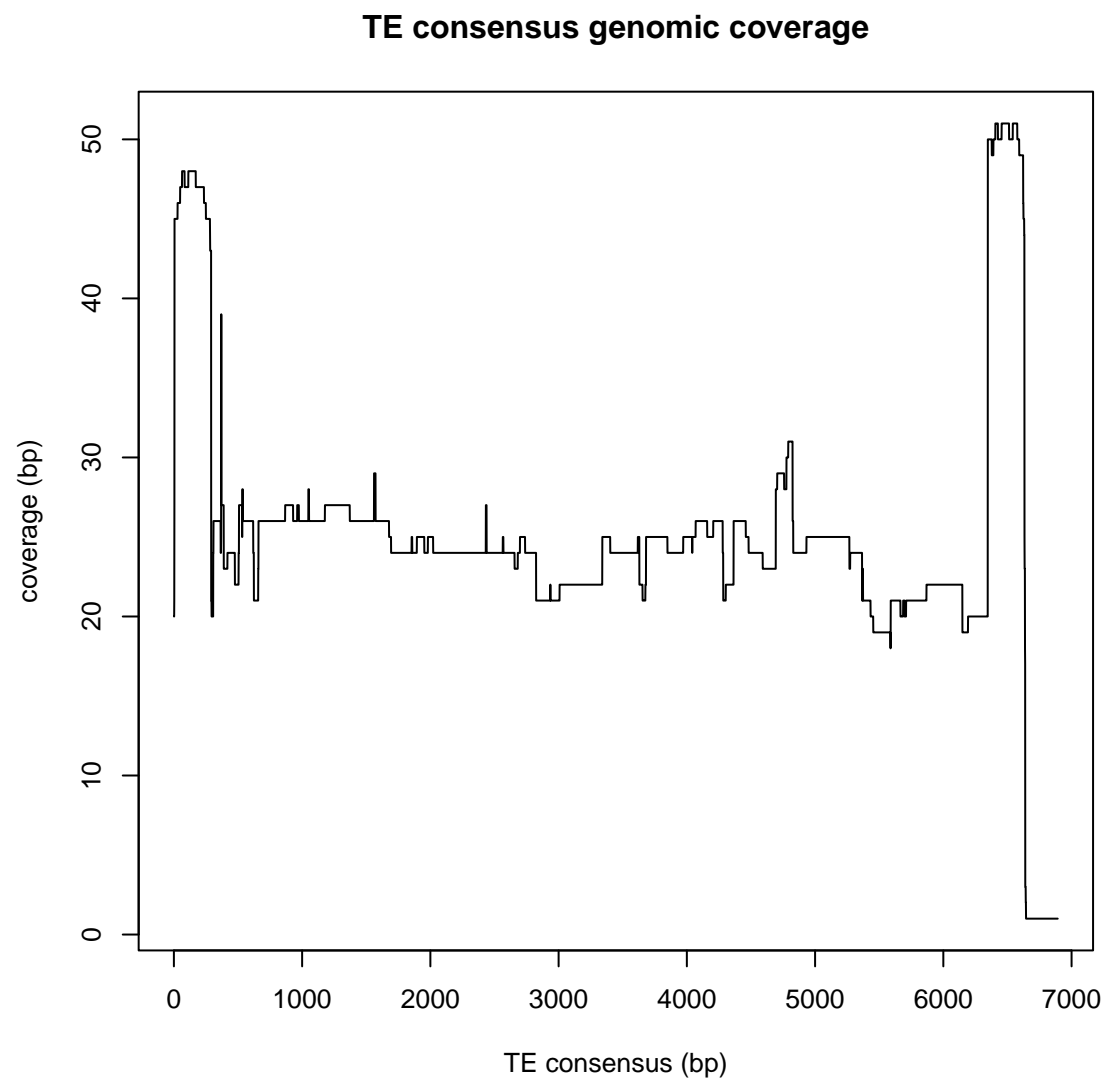
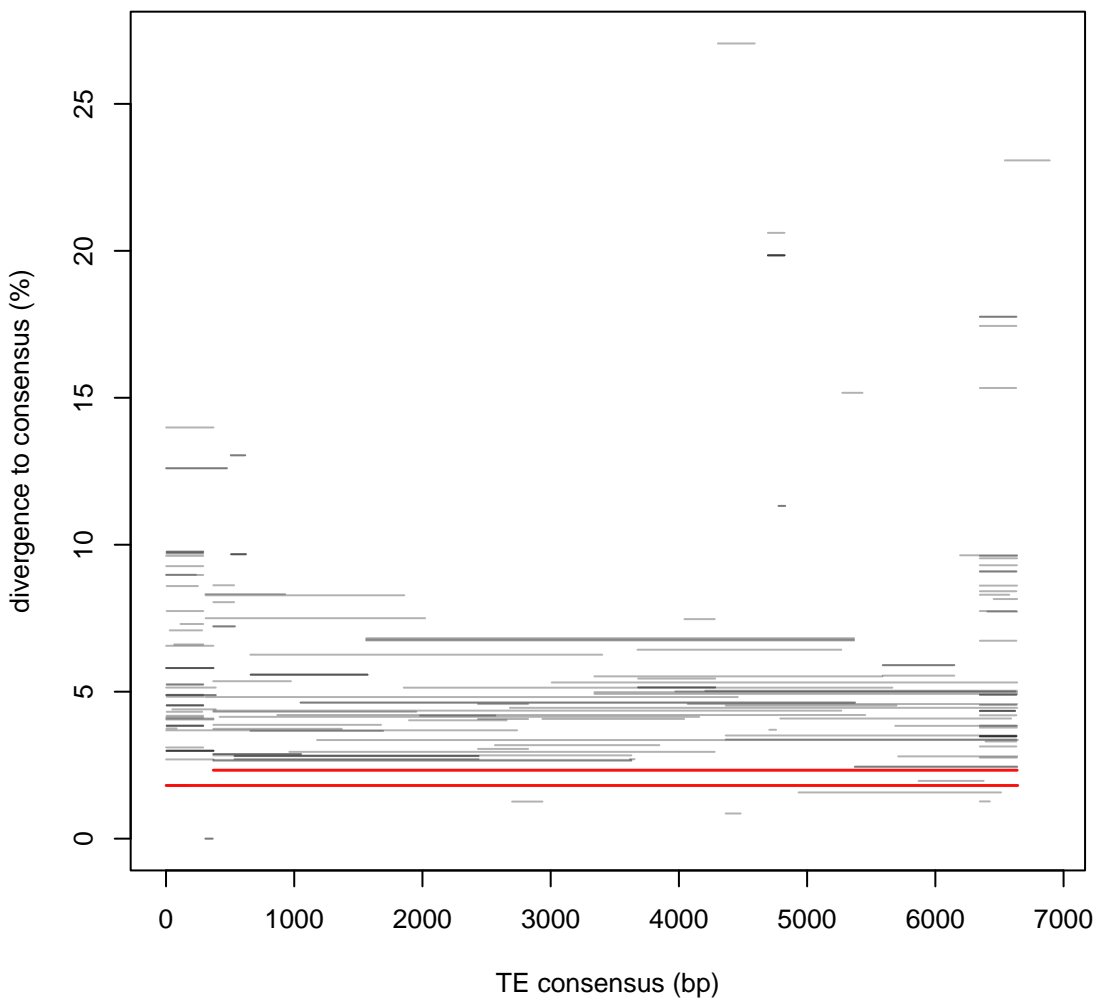
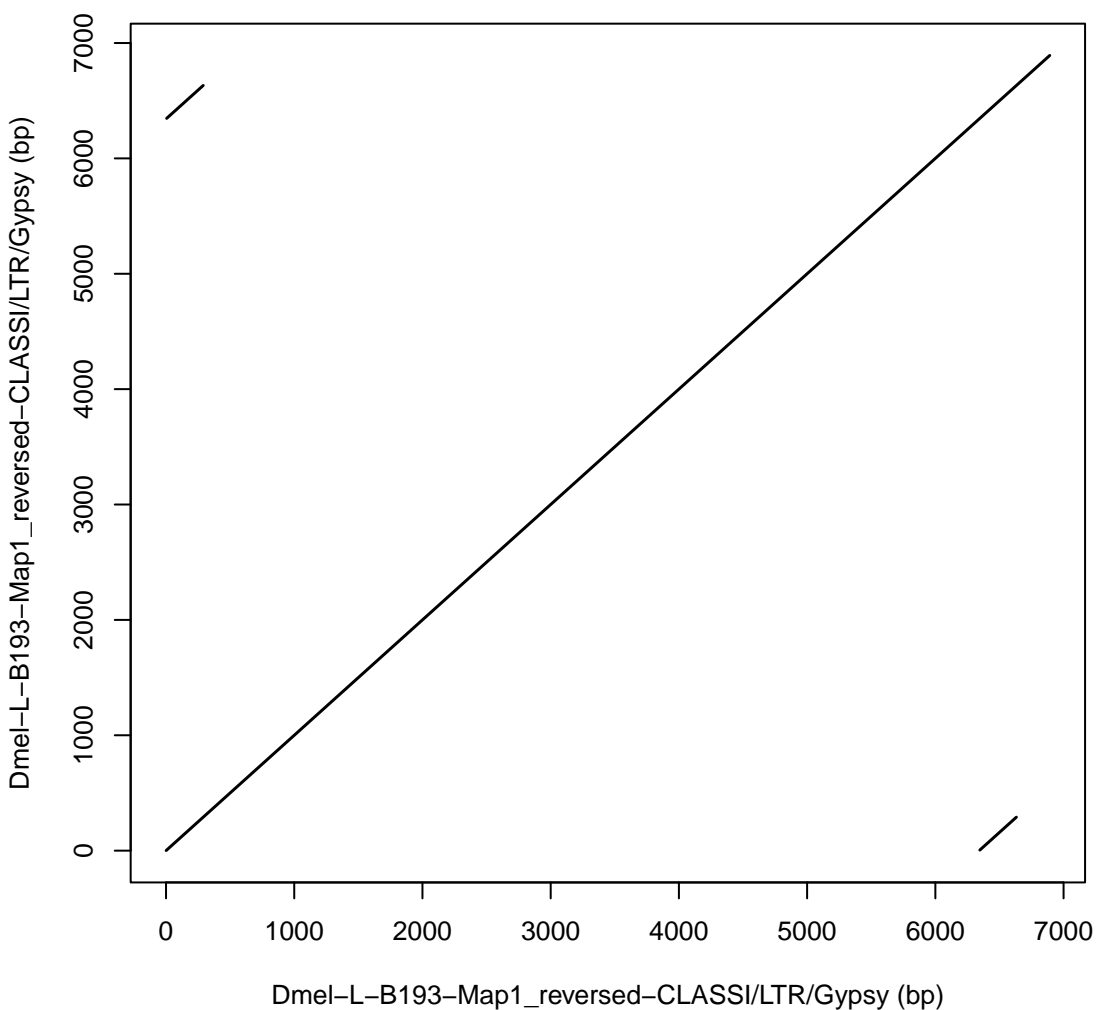


TE: Dmel-L-B193-Map1\_reversed-CLASSI/LTR/Gypsy  
consensus size: 6892bp; fragments: 174; full length: 2 (>=6202.8bp)



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

