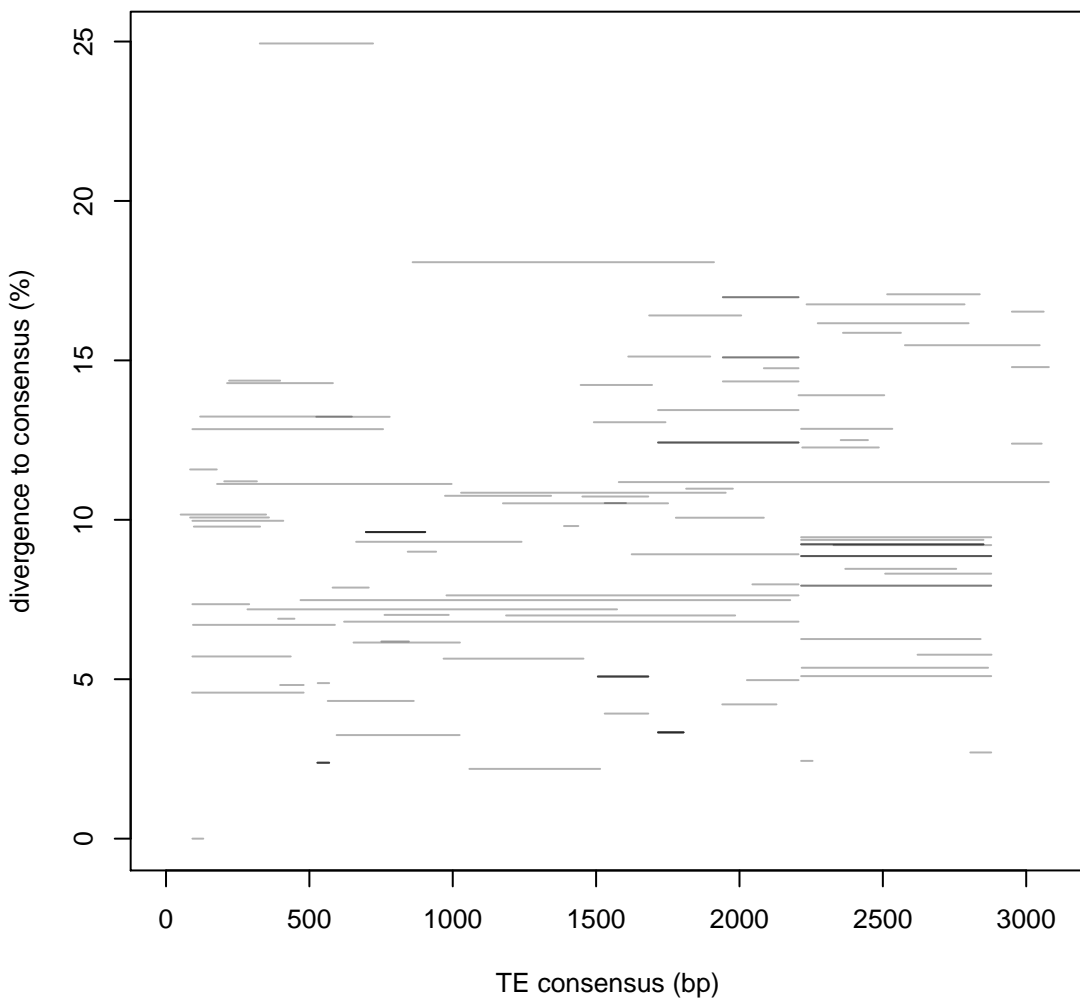
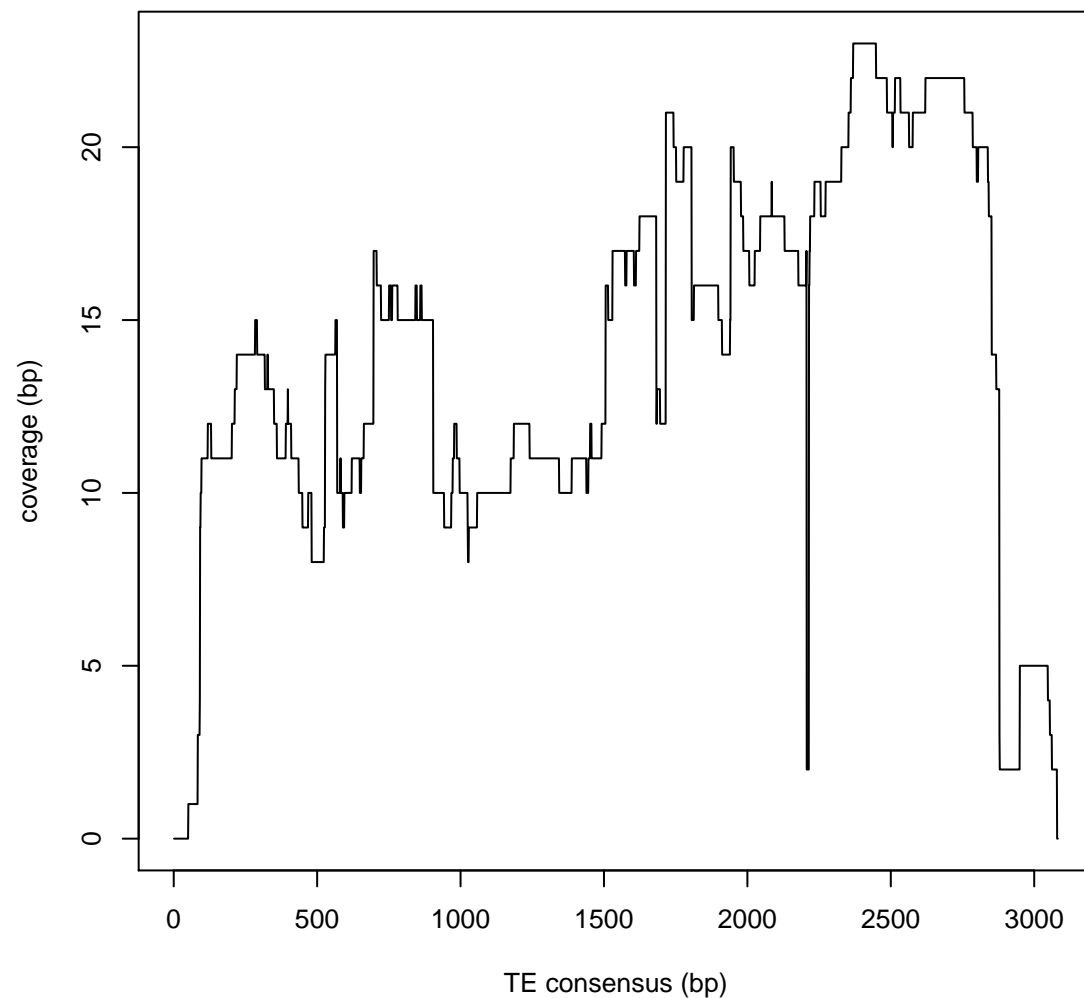


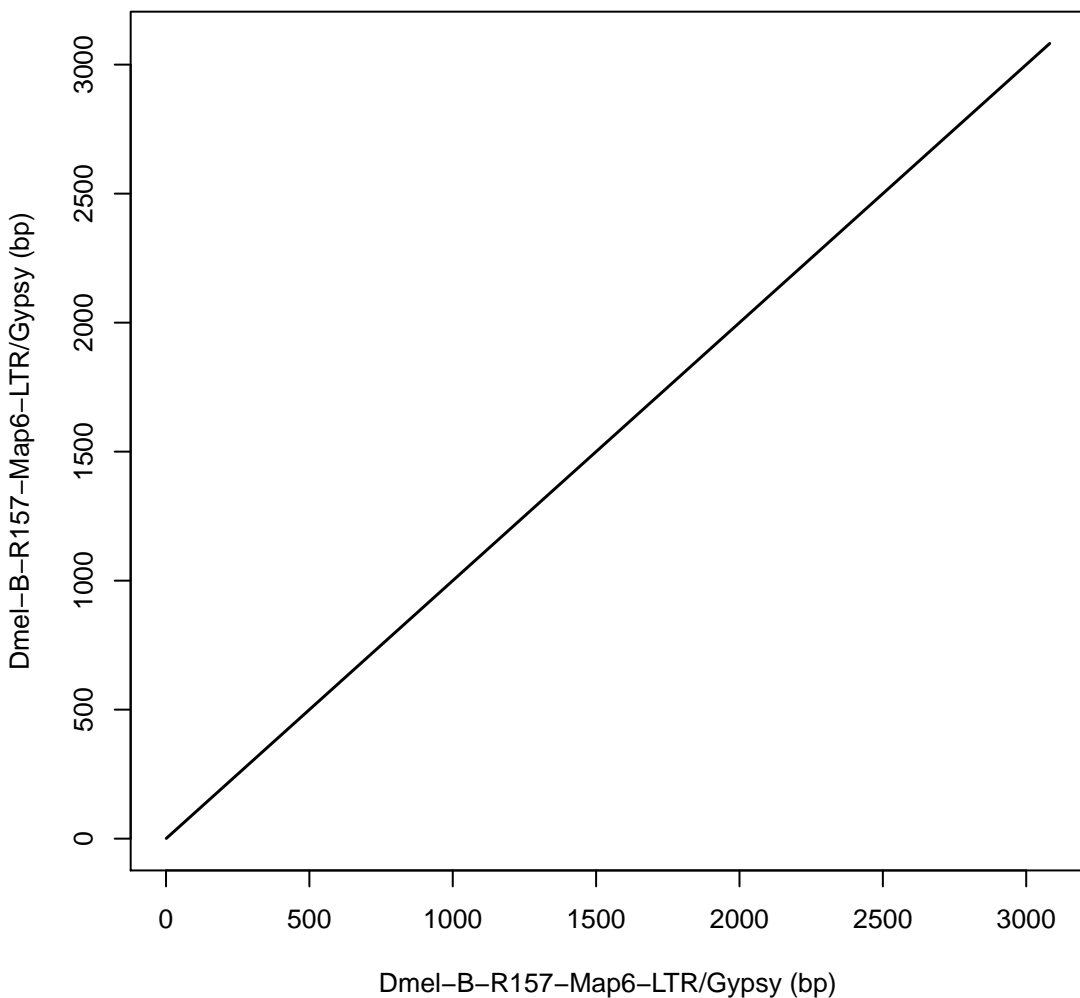
TE: Dmel-B-R157-Map6-LTR/Gypsy  
consensus size: 3082bp; fragments: 114; full length: 0 (>=2773.8bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

