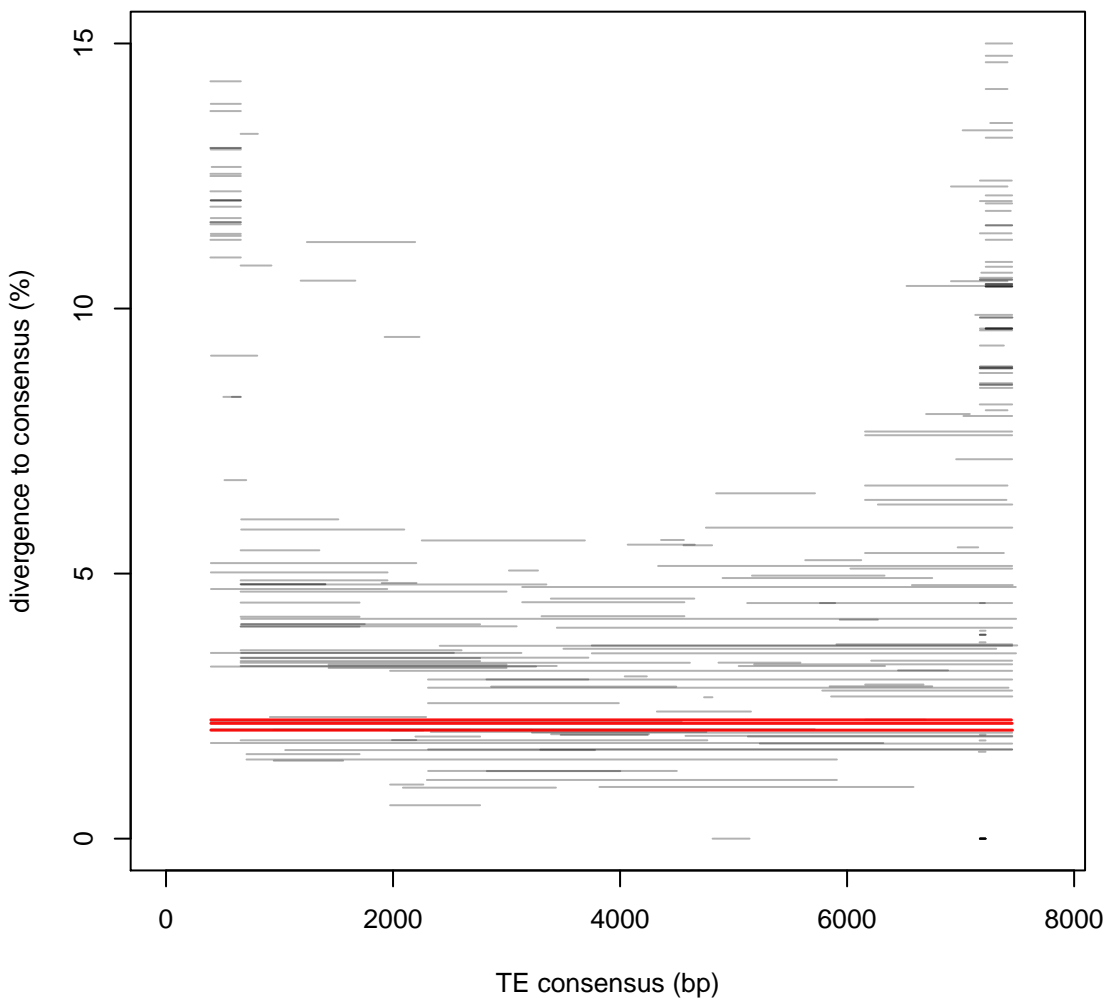
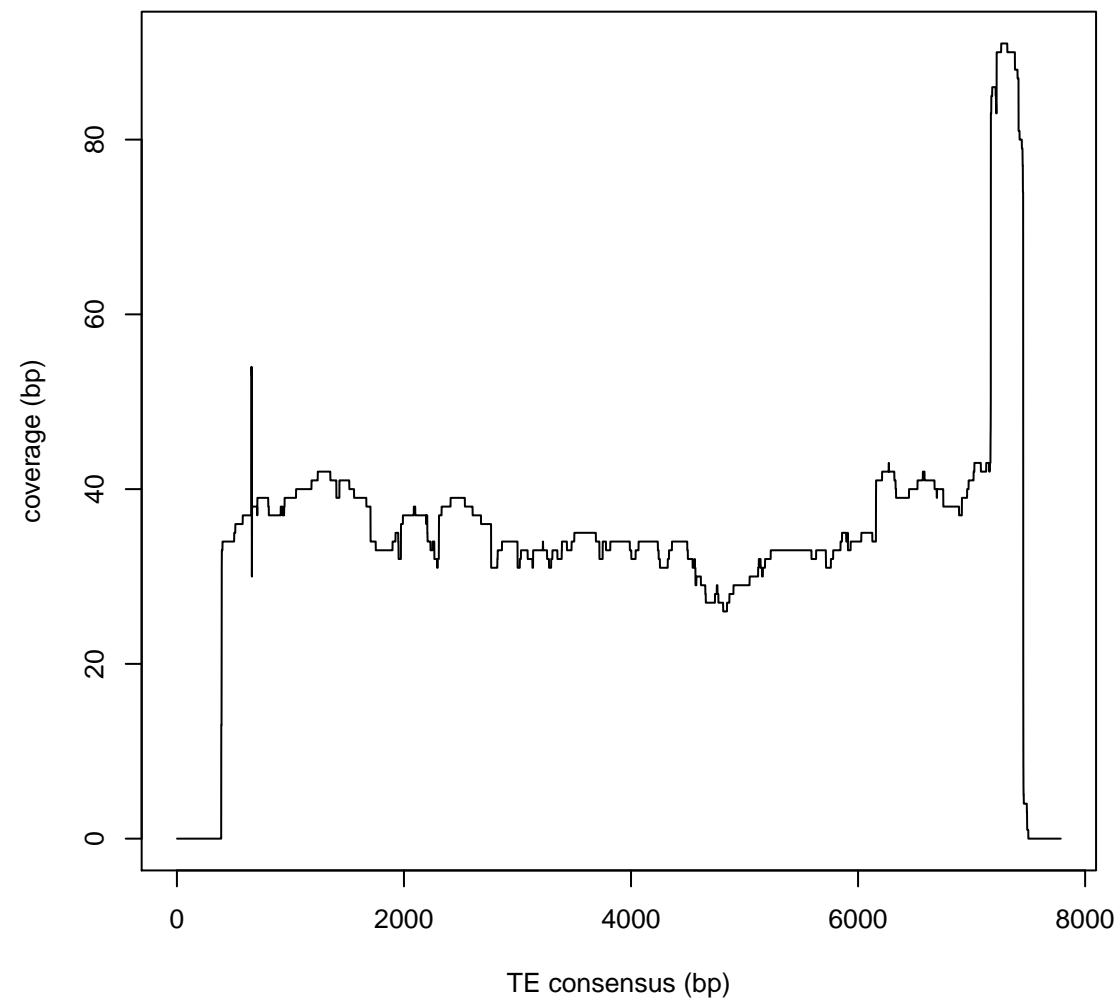


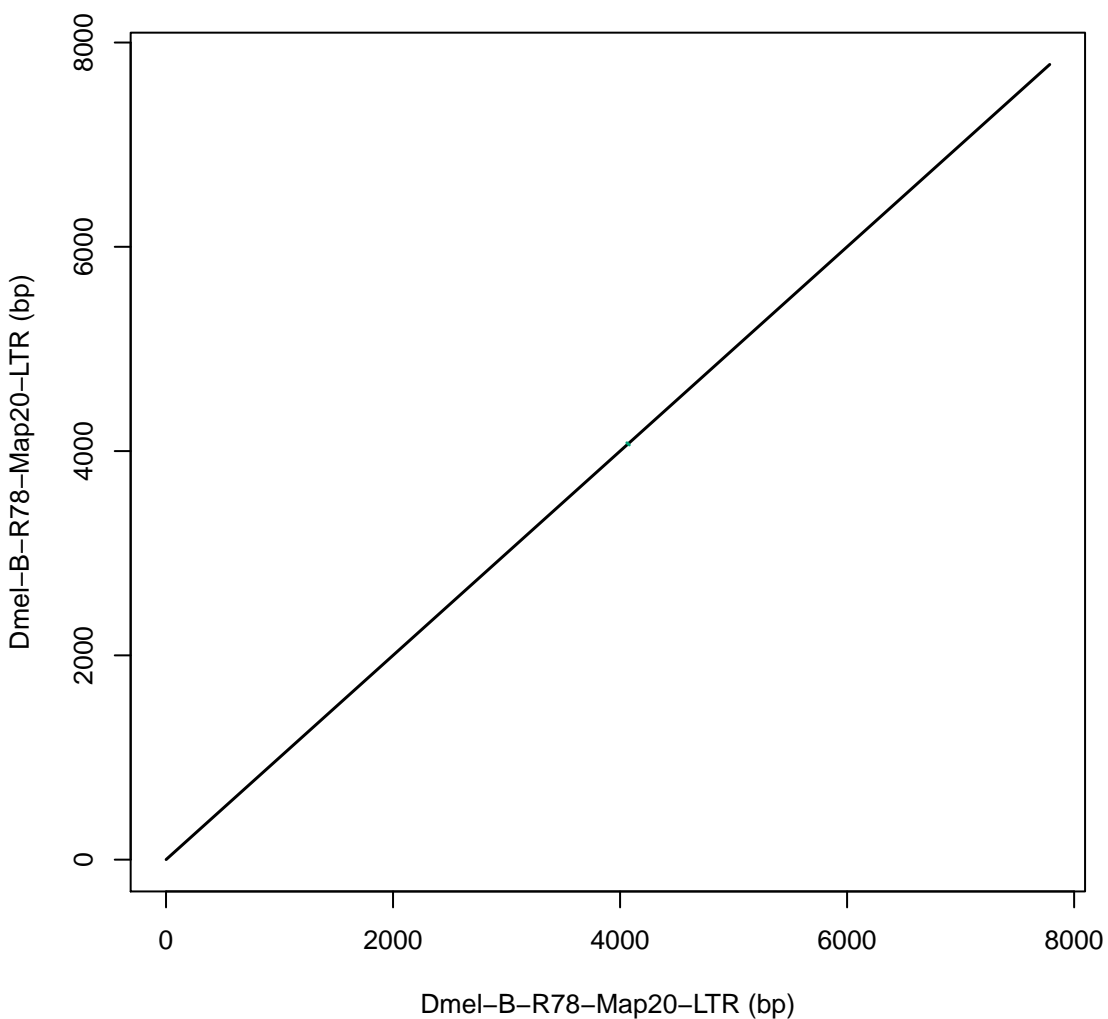
TE: Dmel-B-R78-Map20-LTR
consensus size: 7785bp; fragments: 225; full length: 3 (>=7006.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

