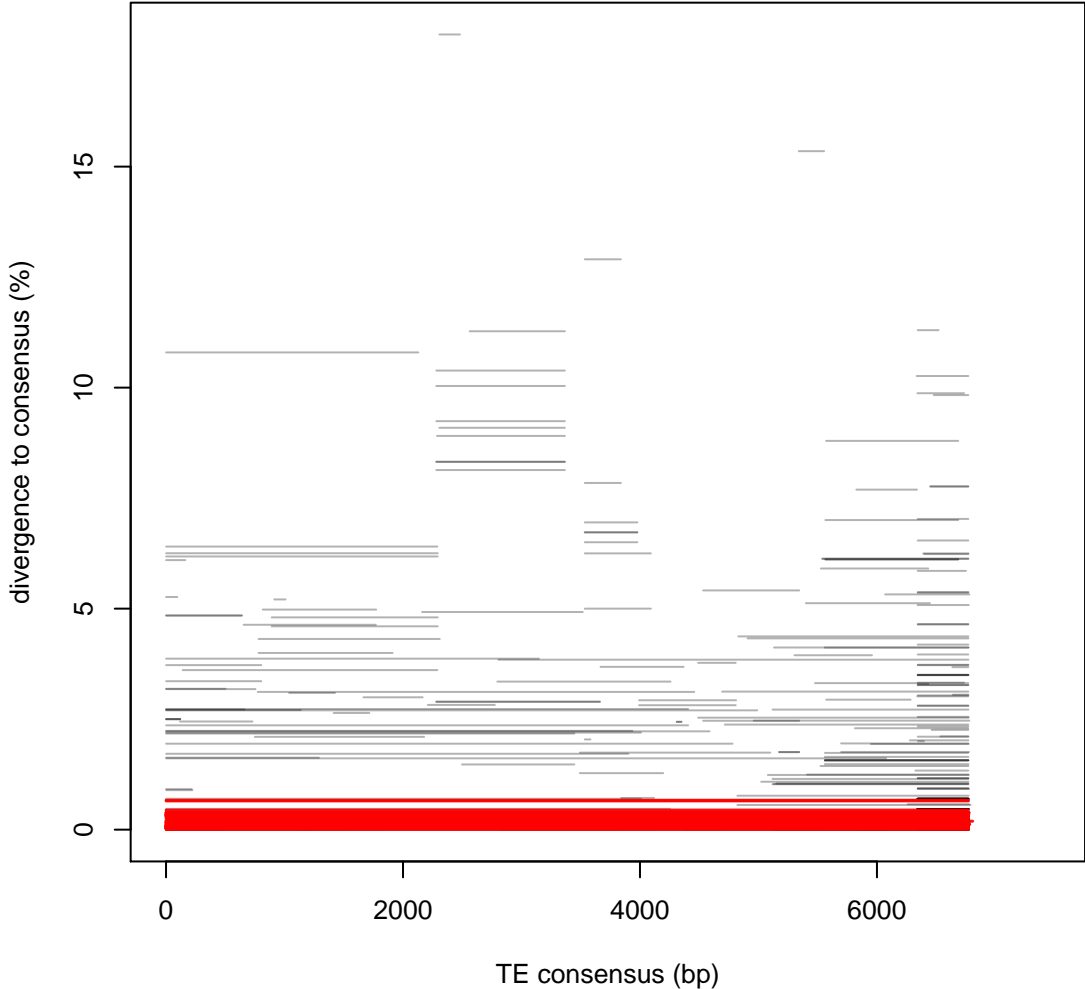
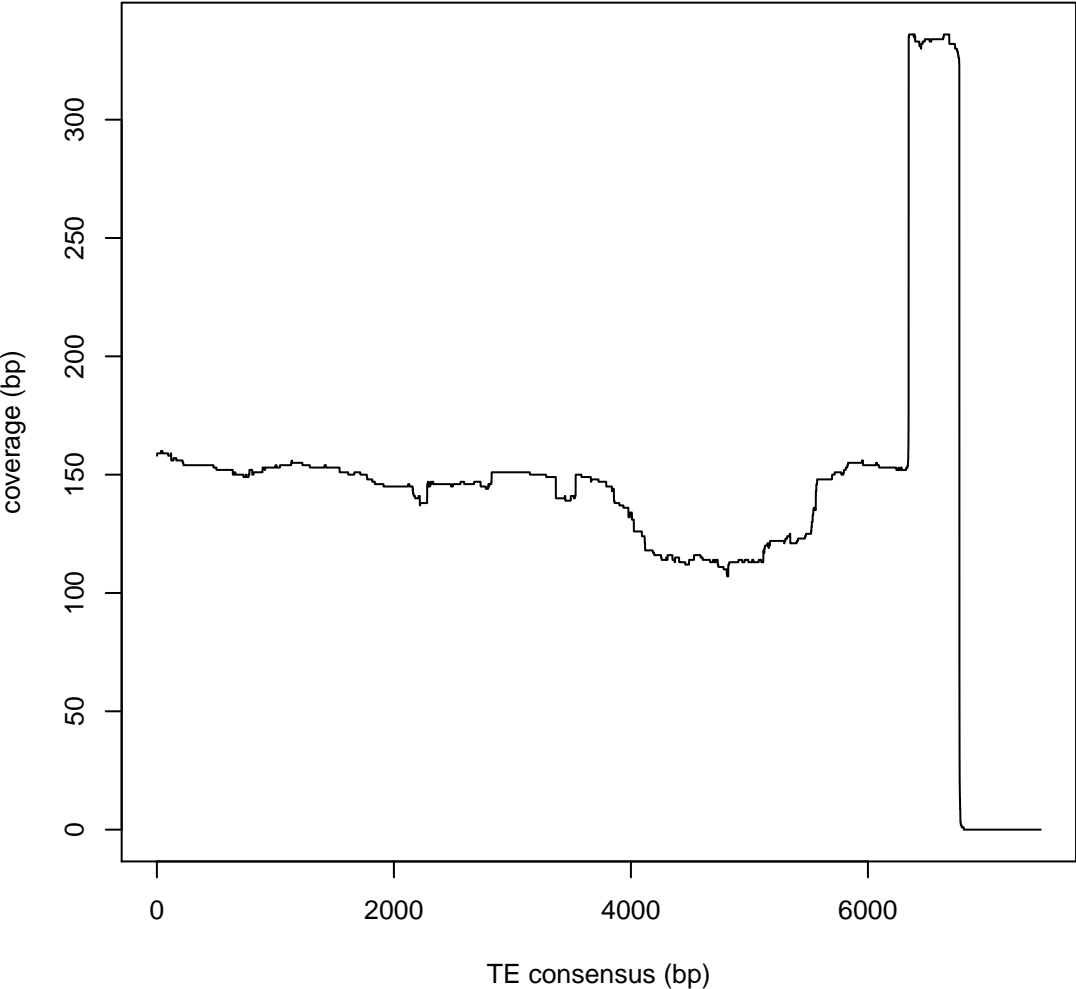


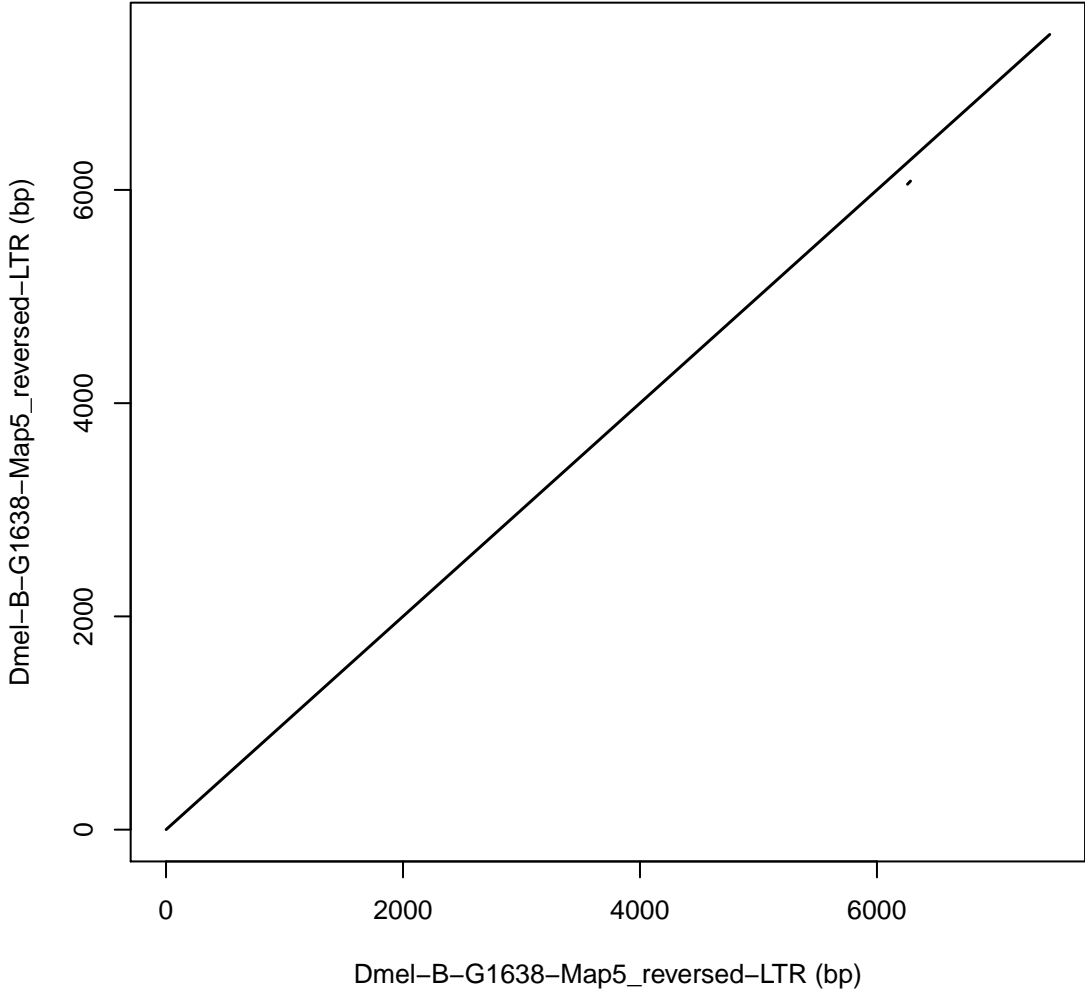
TE: Dmel-B-G1638-Map5_reversed-LTR
consensus size: 7458bp; fragments: 496; full length: 78 (>=6712.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

