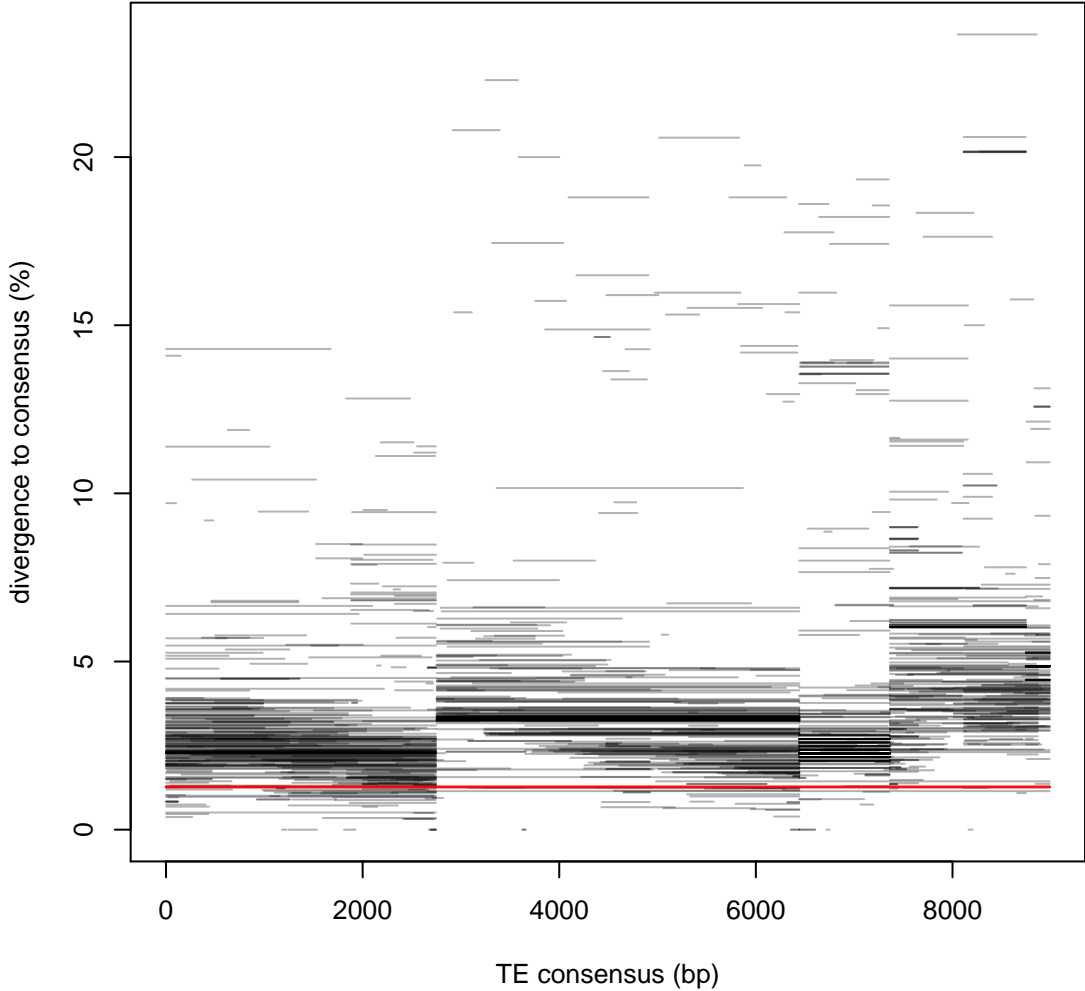
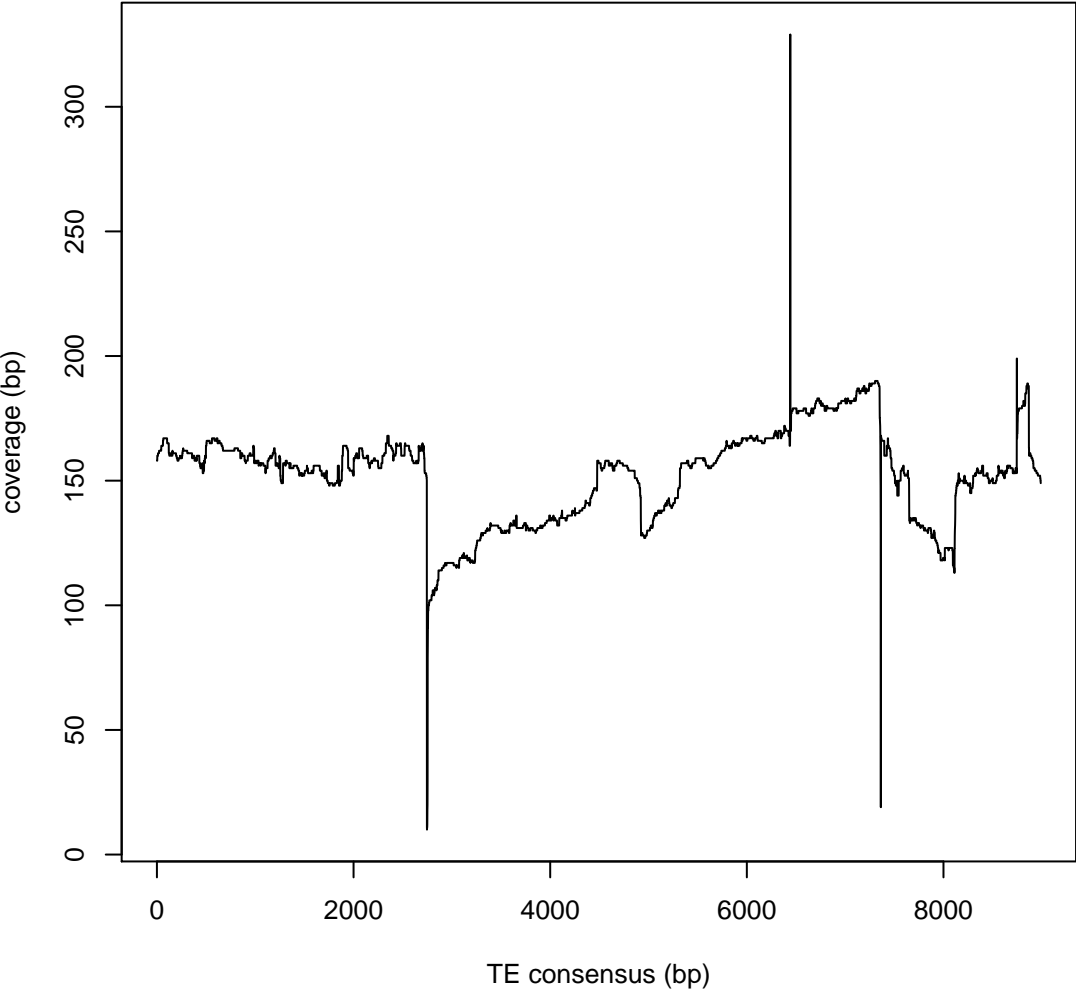


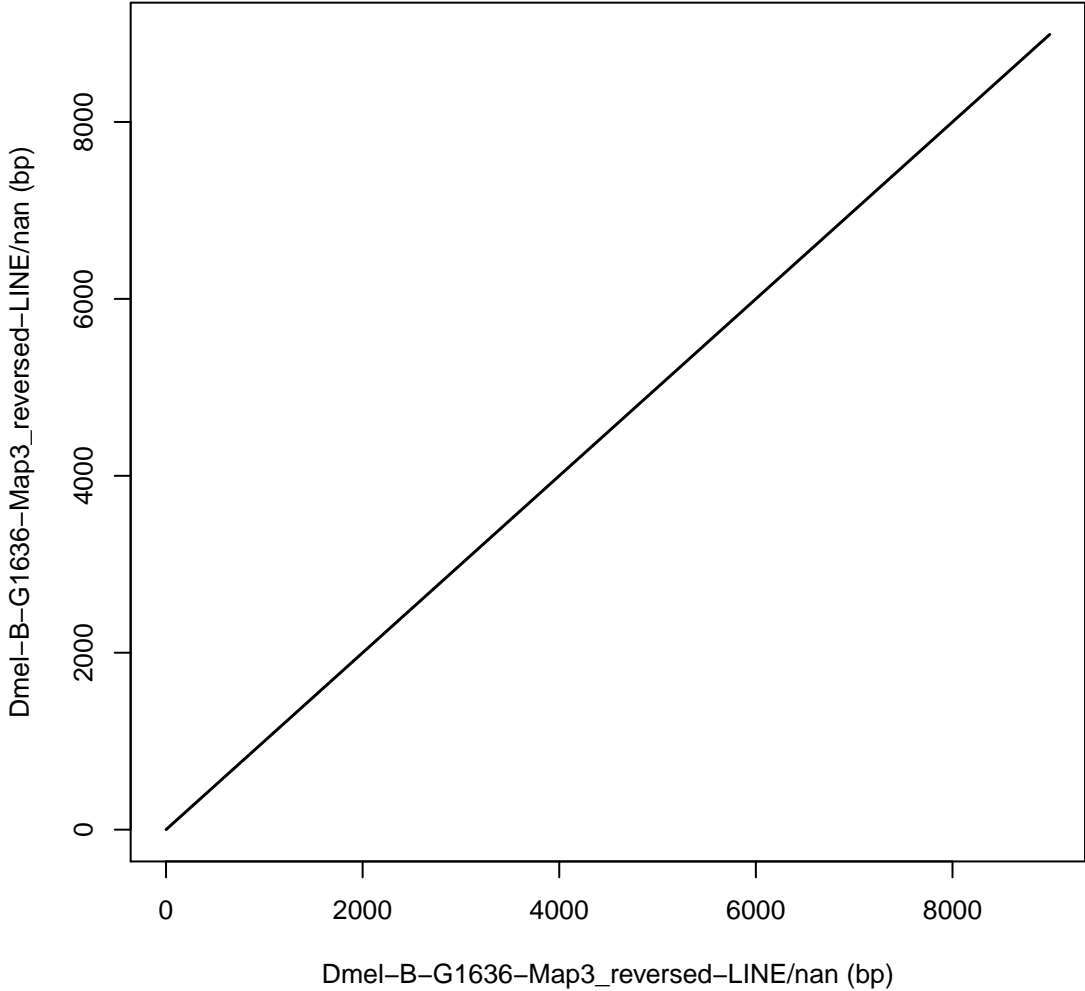
TE: Dmel-B-G1636-Map3\_reversed-LINE/nan  
consensus size: 8989bp; fragments: 1247; full length: 1 (>=8090.1bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

