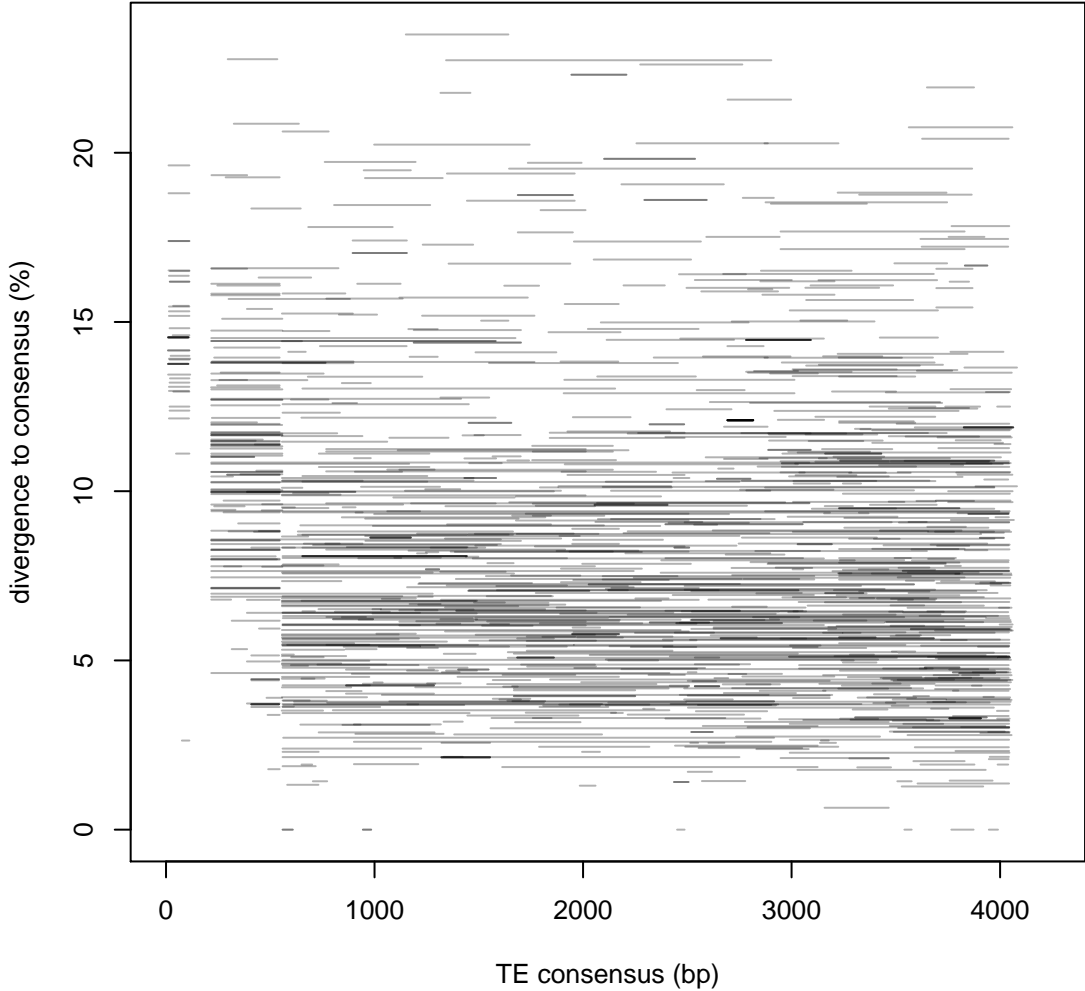
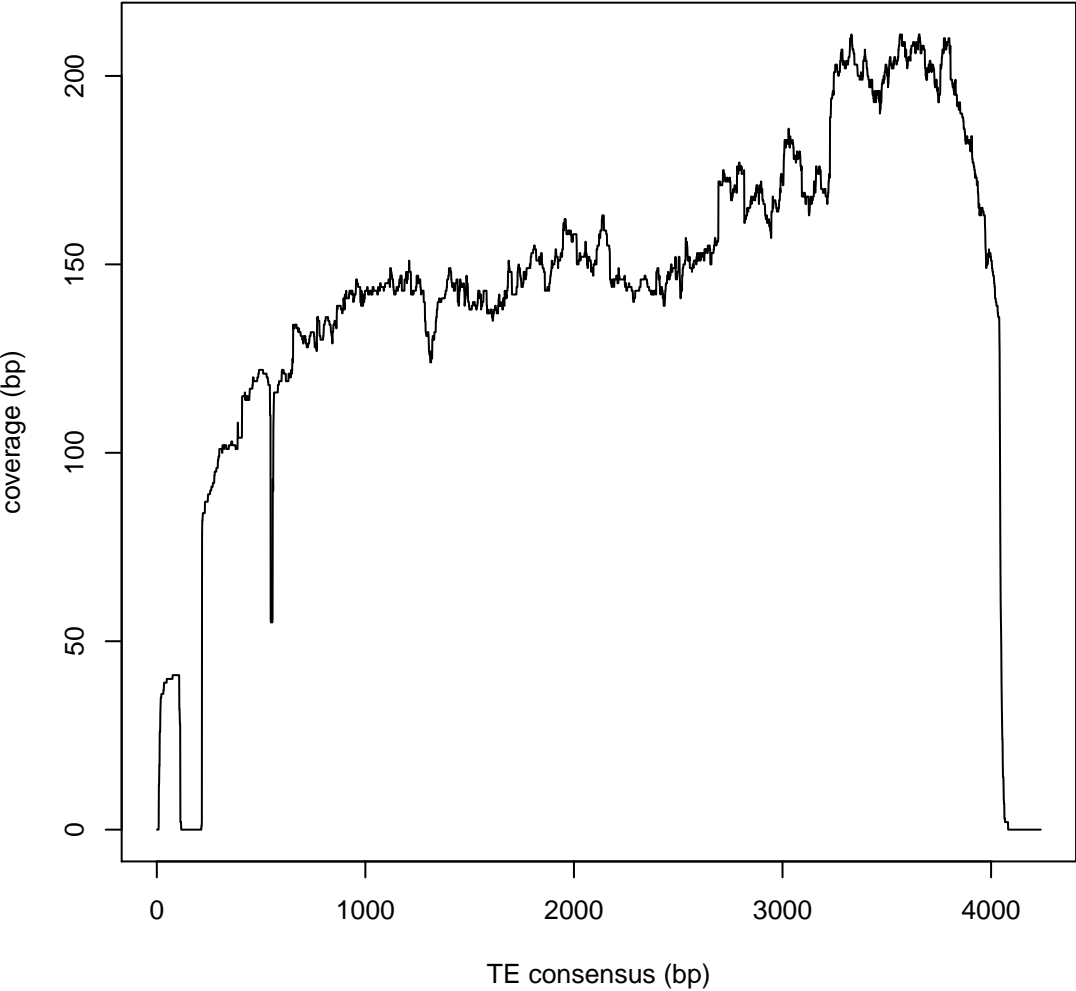


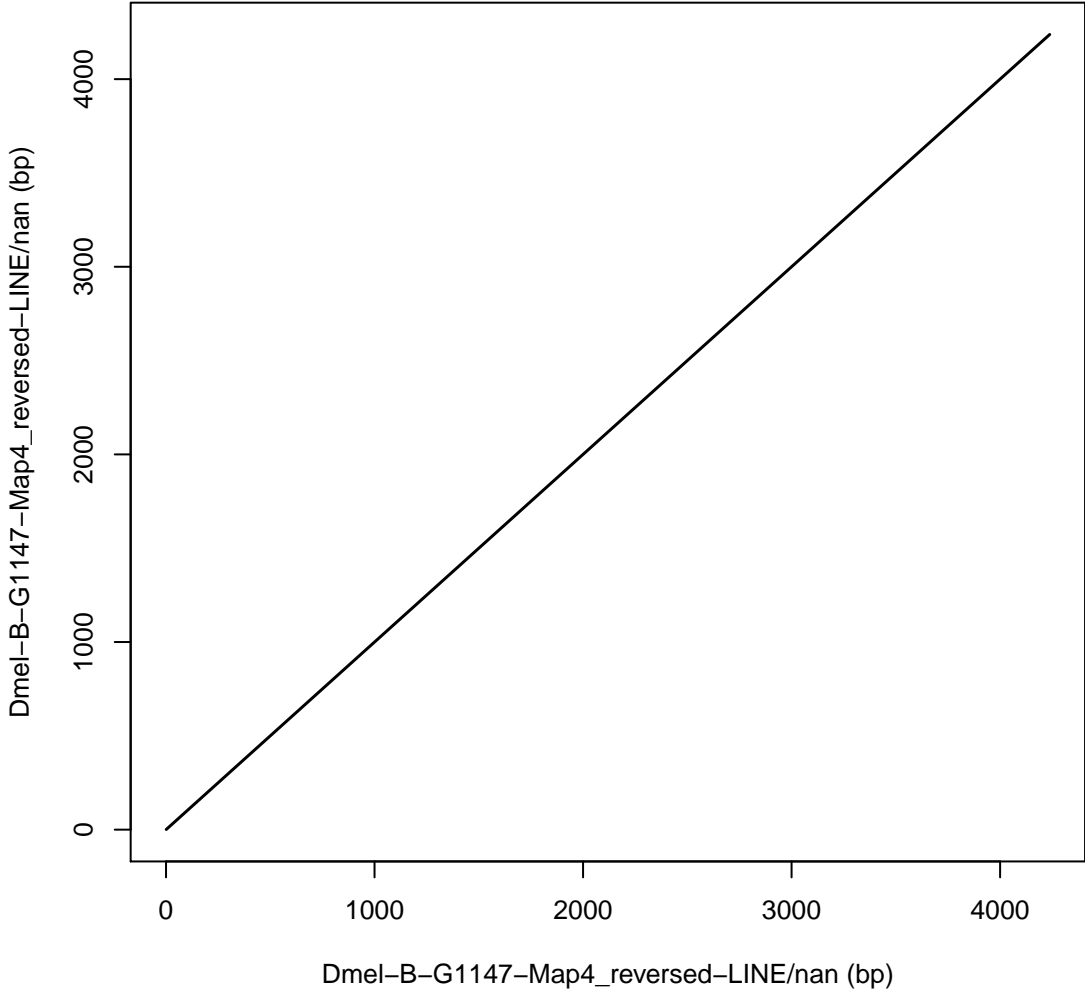
TE: Dmel-B-G1147-Map4\_reversed-LINE/nan  
consensus size: 4238bp; fragments: 1160; full length: 0 (>=3814.2bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

