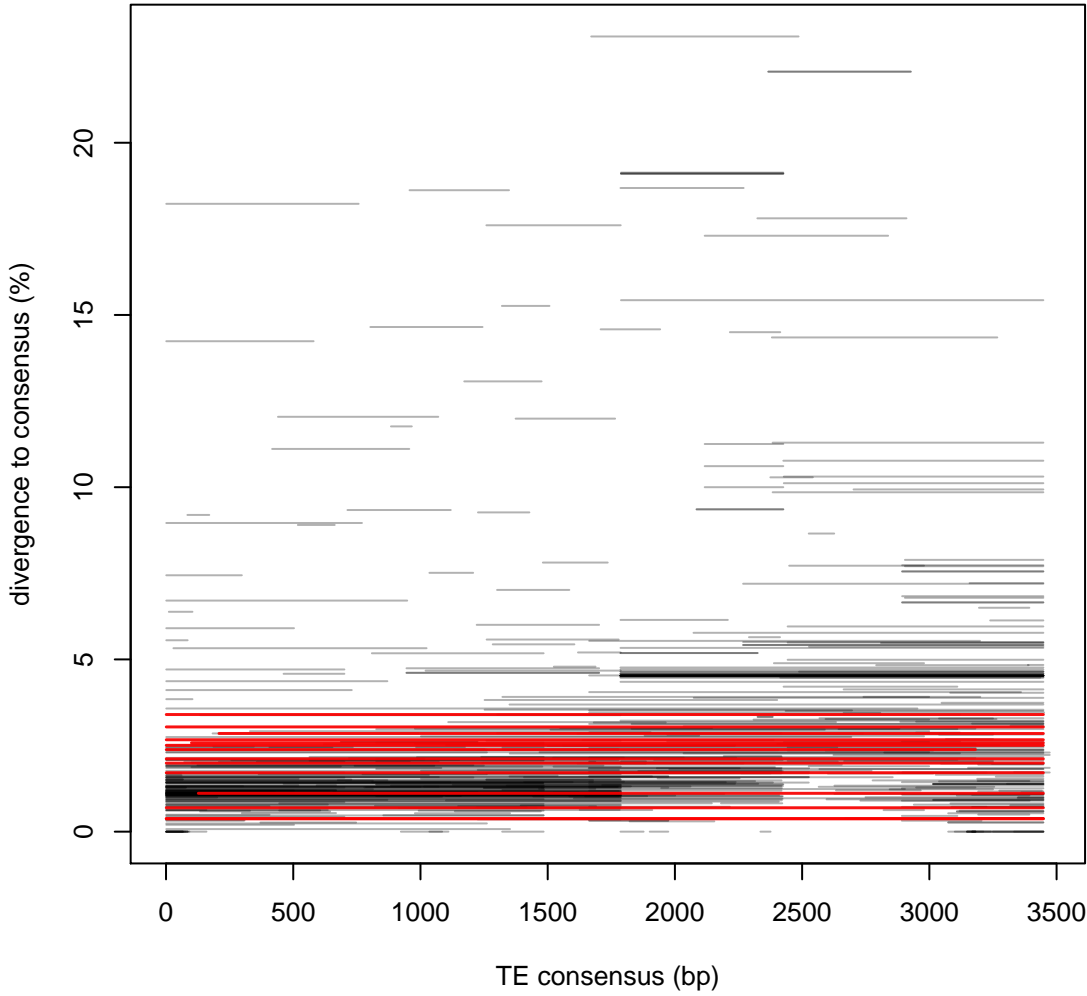
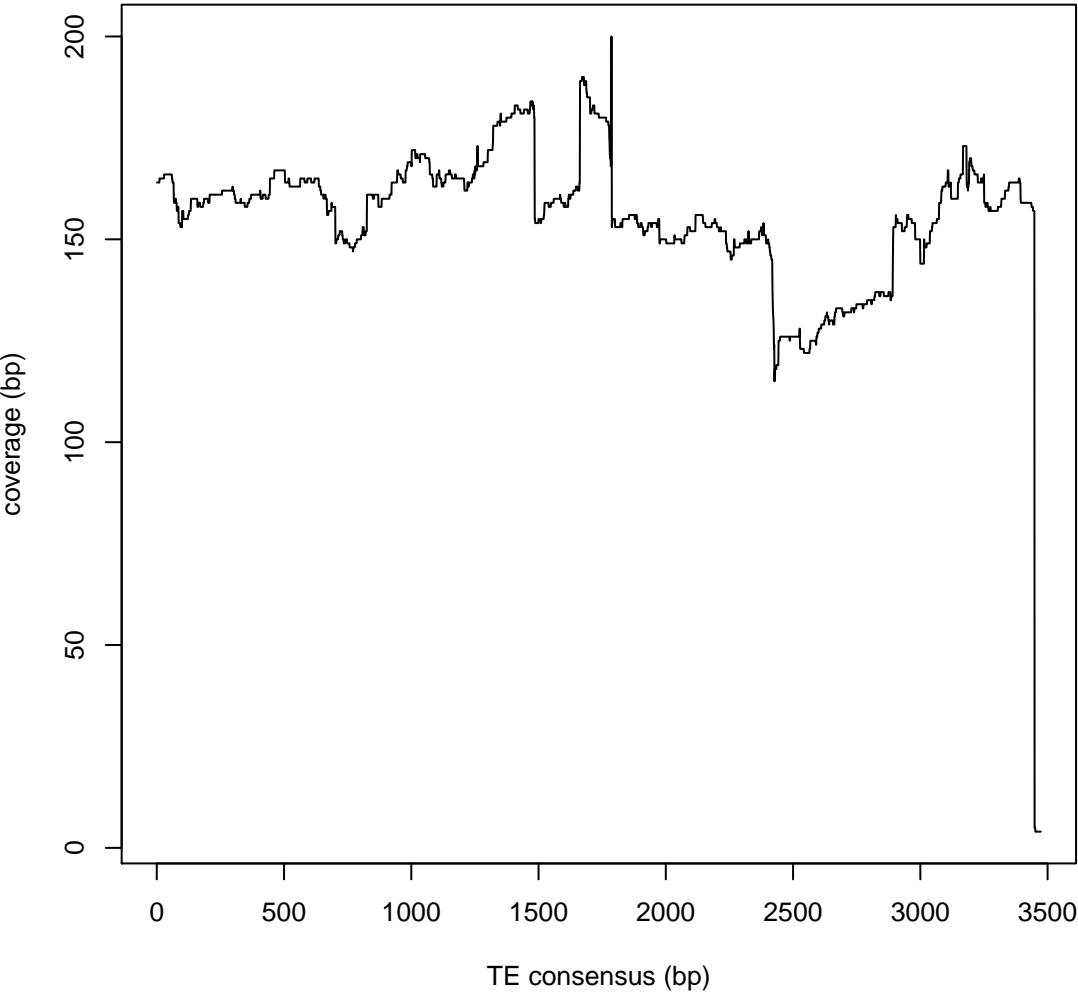


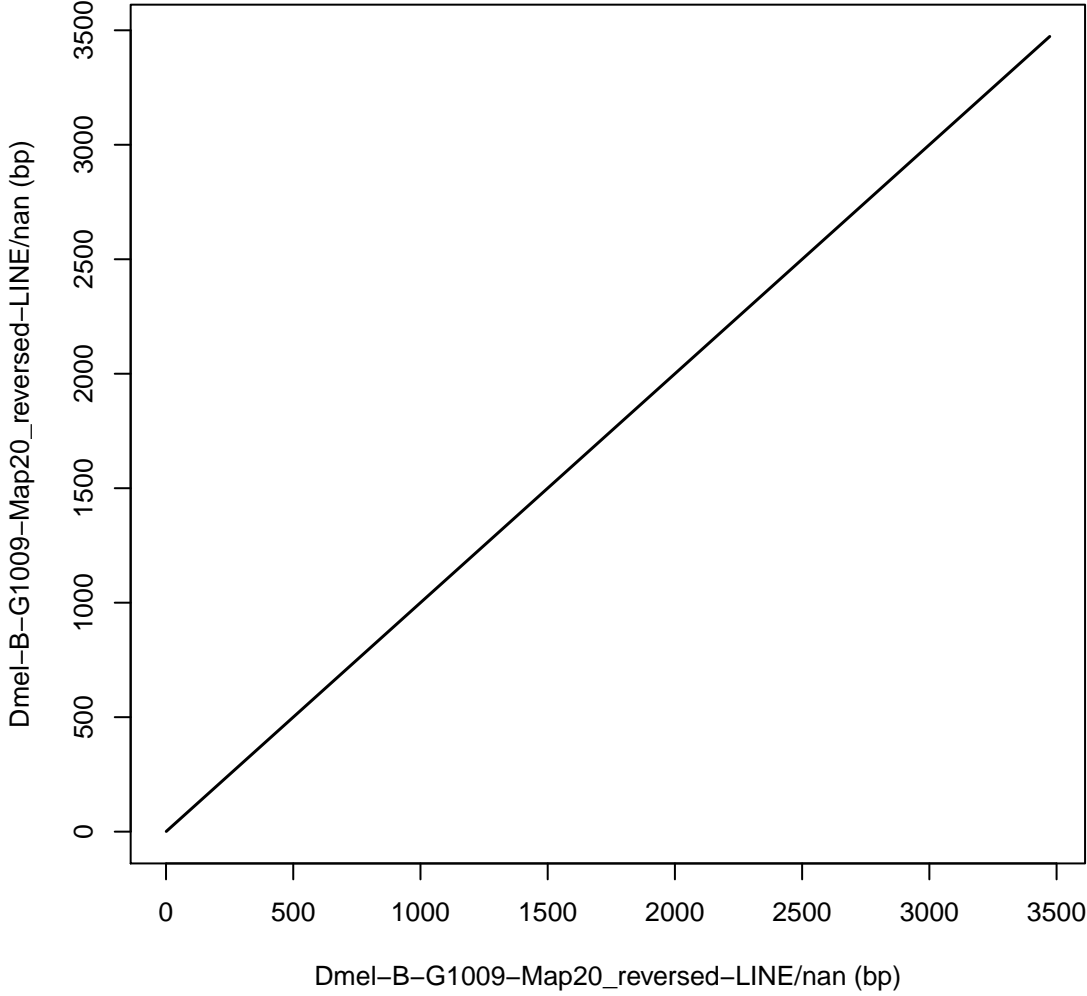
TE: Dmel-B-G1009-Map20_reversed-LINE/nan
consensus size: 3473bp; fragments: 564; full length: 14 (>=3125.7bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

