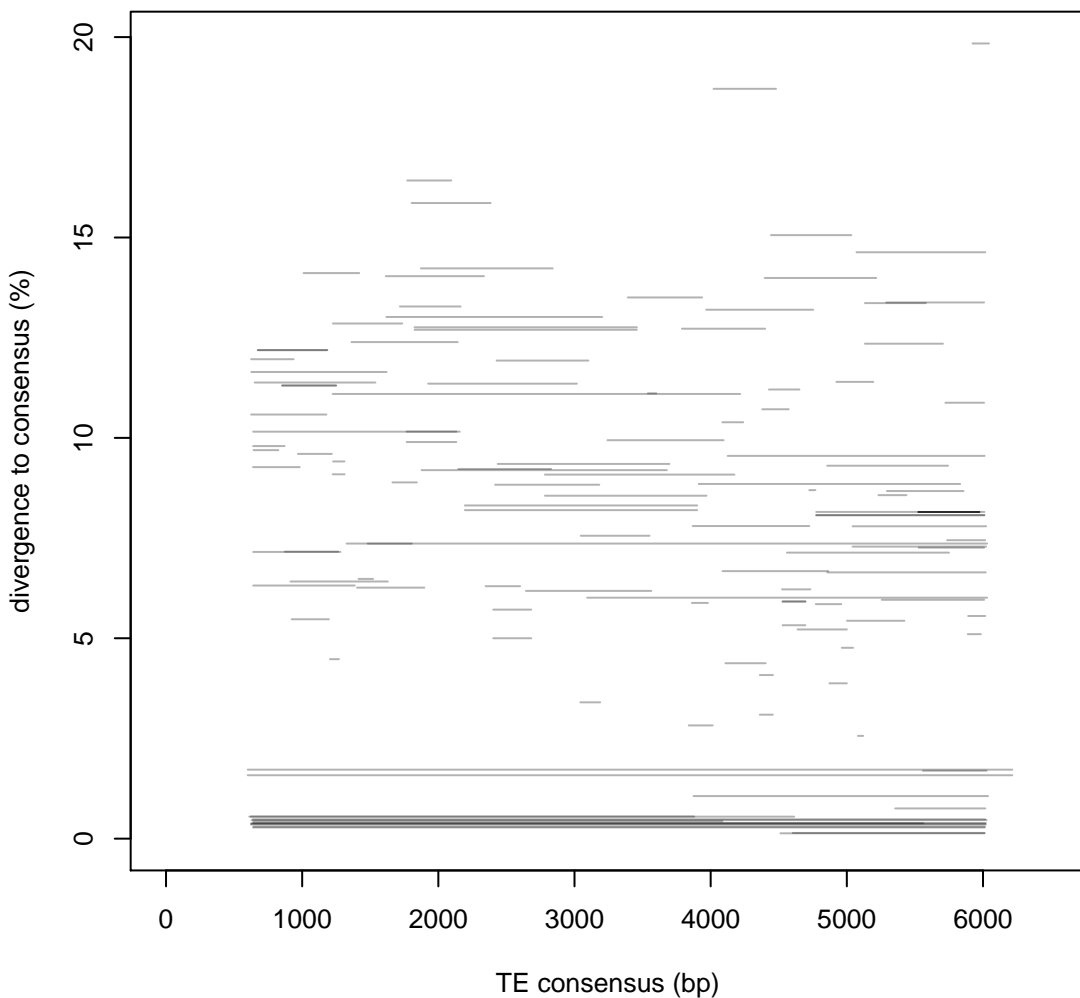
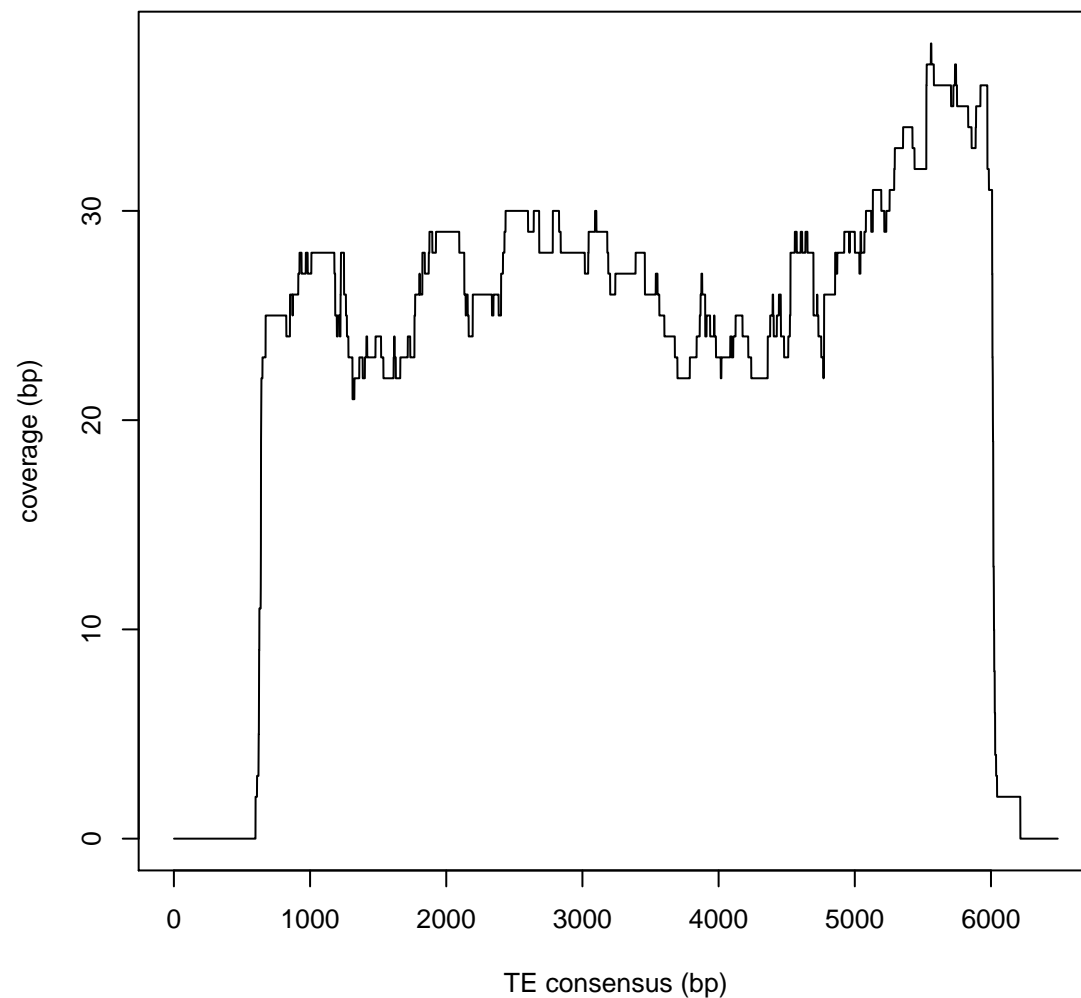


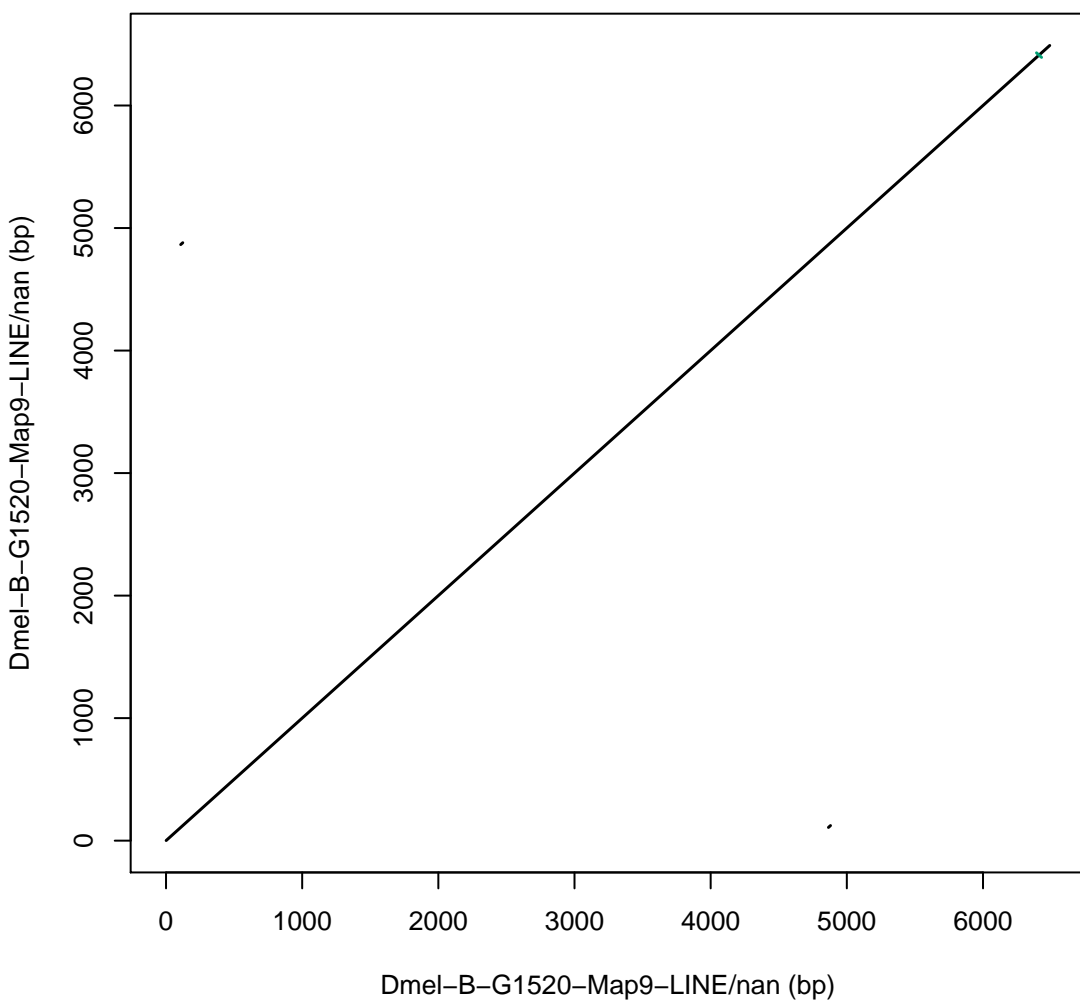
TE: Dmel-B-G1520-Map9-LINE/nan
consensus size: 6490bp; fragments: 132; full length: 0 (>=5841bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

