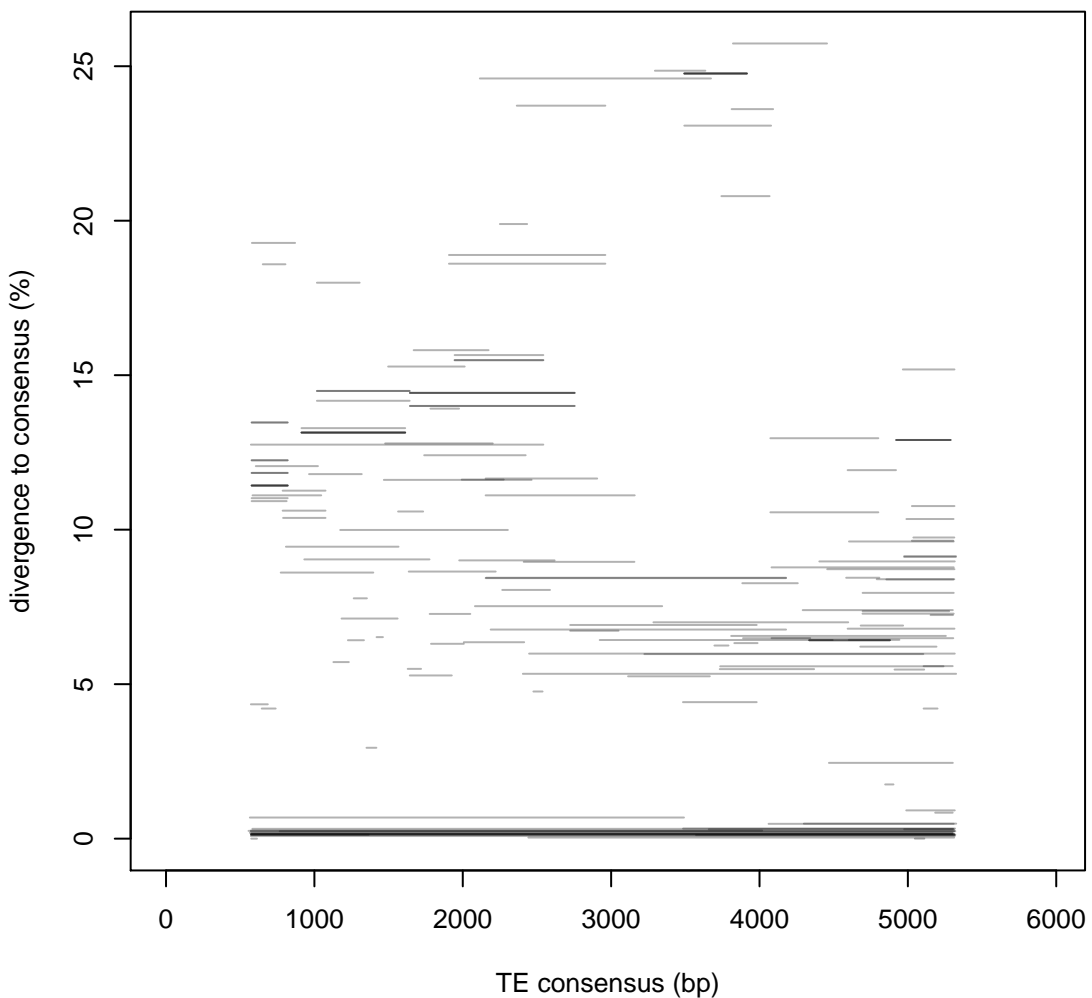
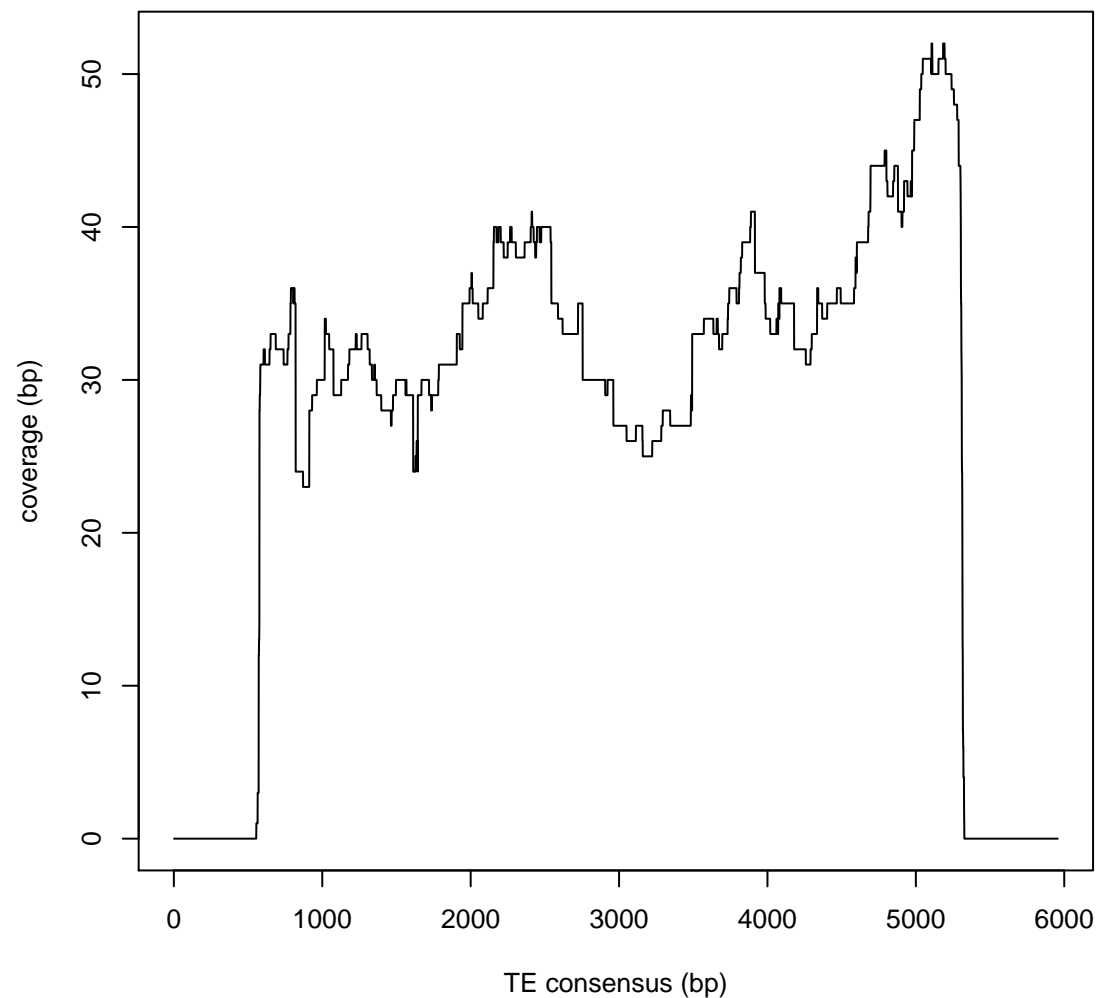


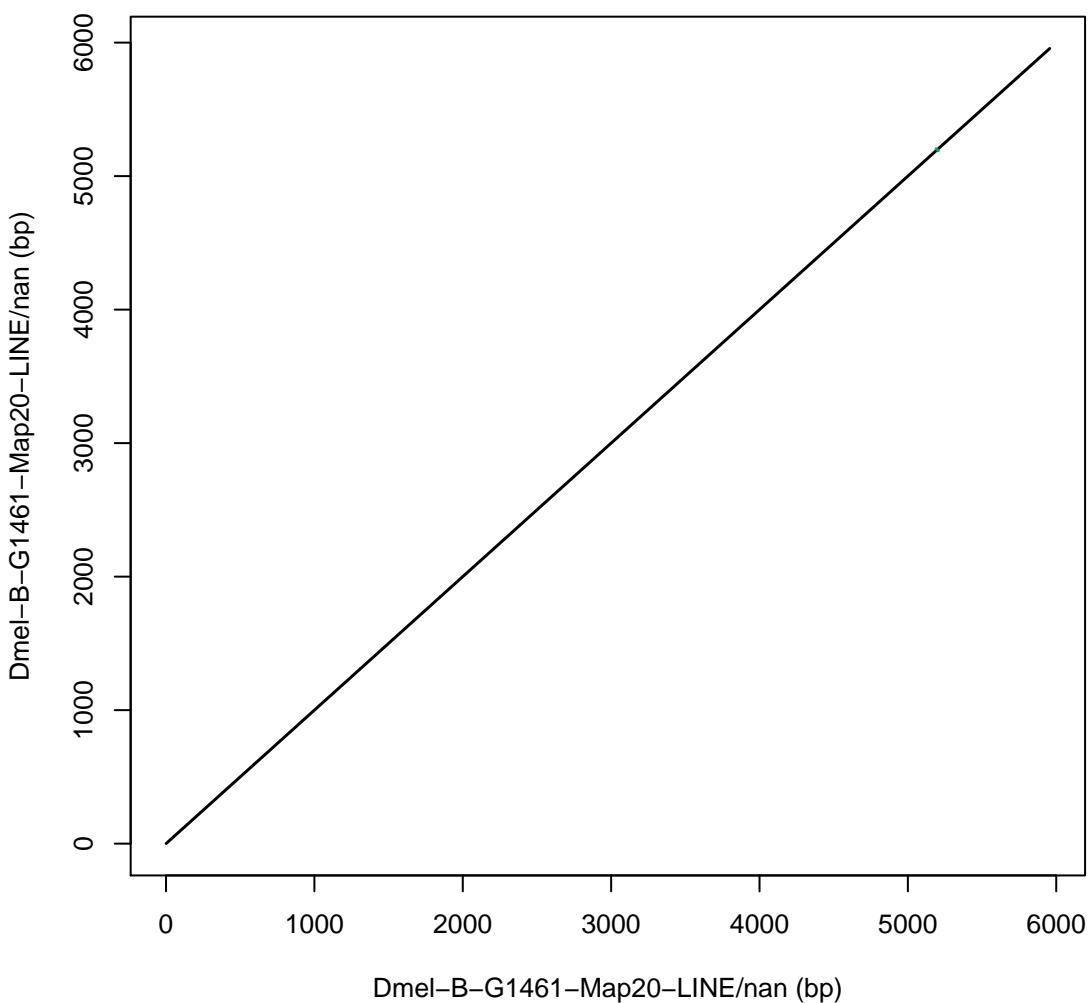
TE: Dmel-B-G1461-Map20-LINE/nan
consensus size: 5956bp; fragments: 168; full length: 0 (>=5360.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

