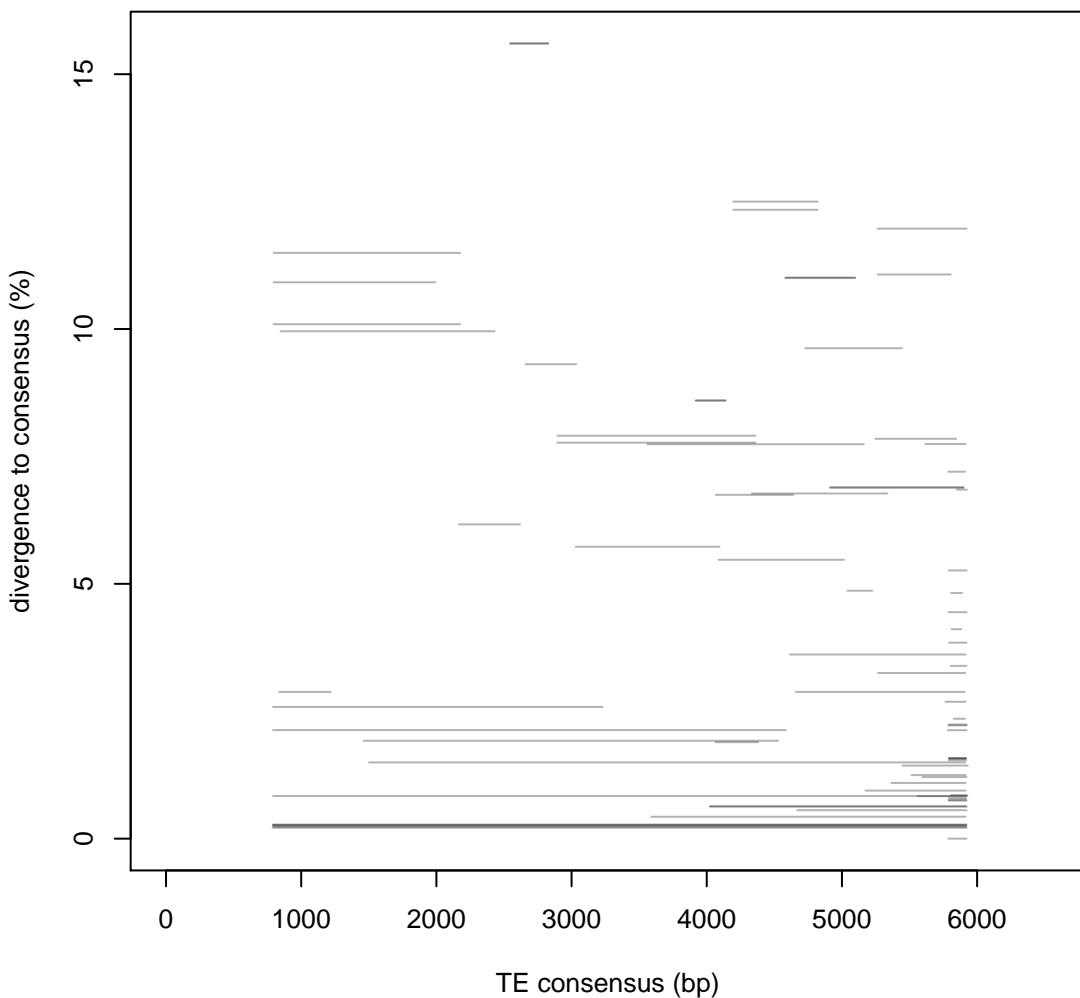
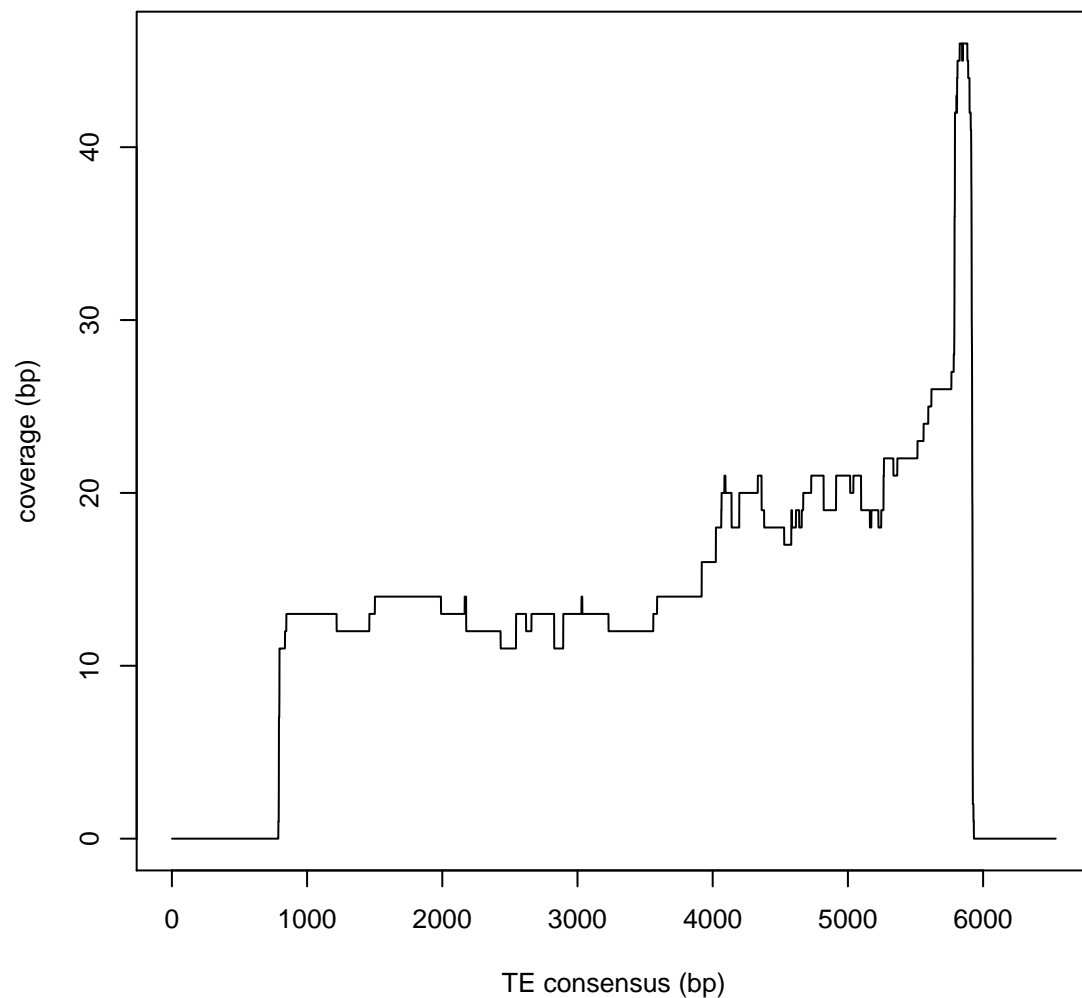


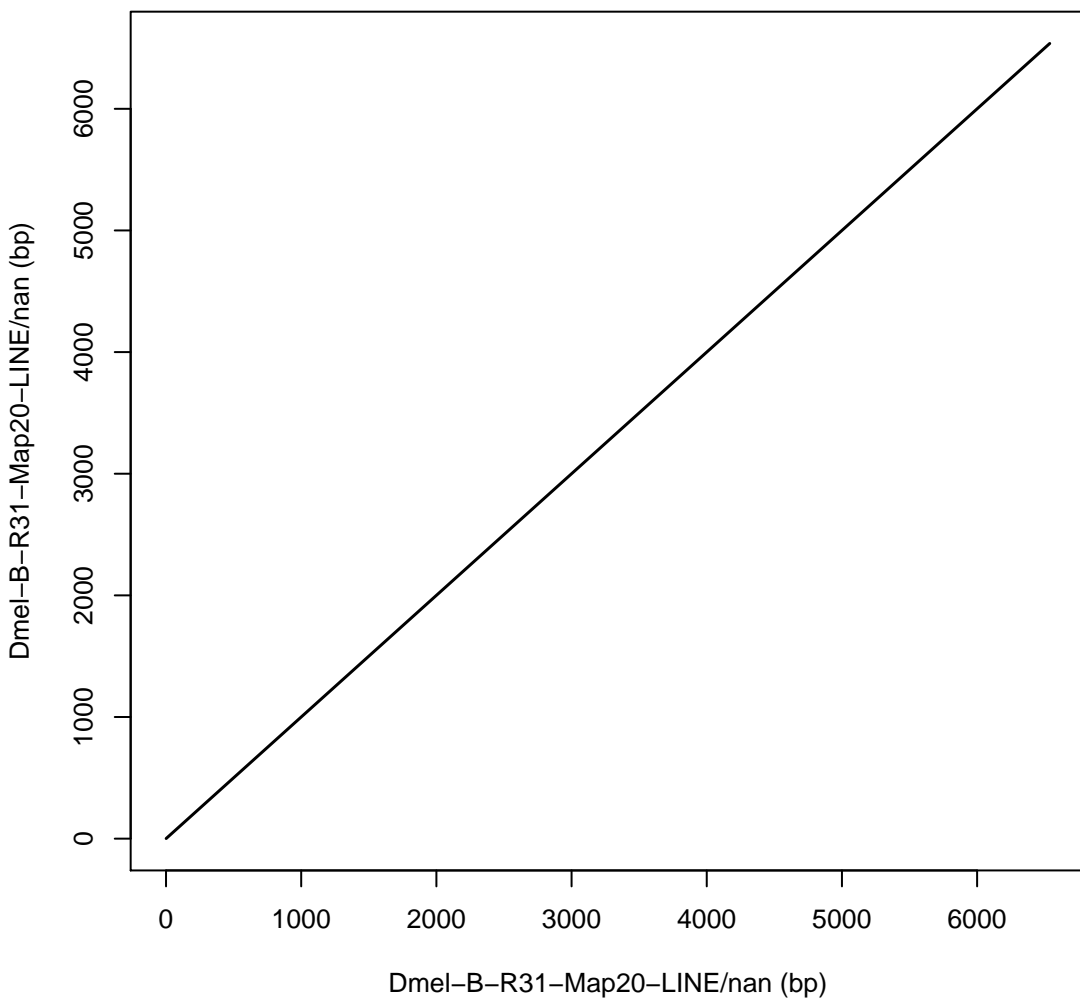
TE: Dmel-B-R31-Map20-LINE/nan
consensus size: 6537bp; fragments: 76; full length: 0 (>=5883.3bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

