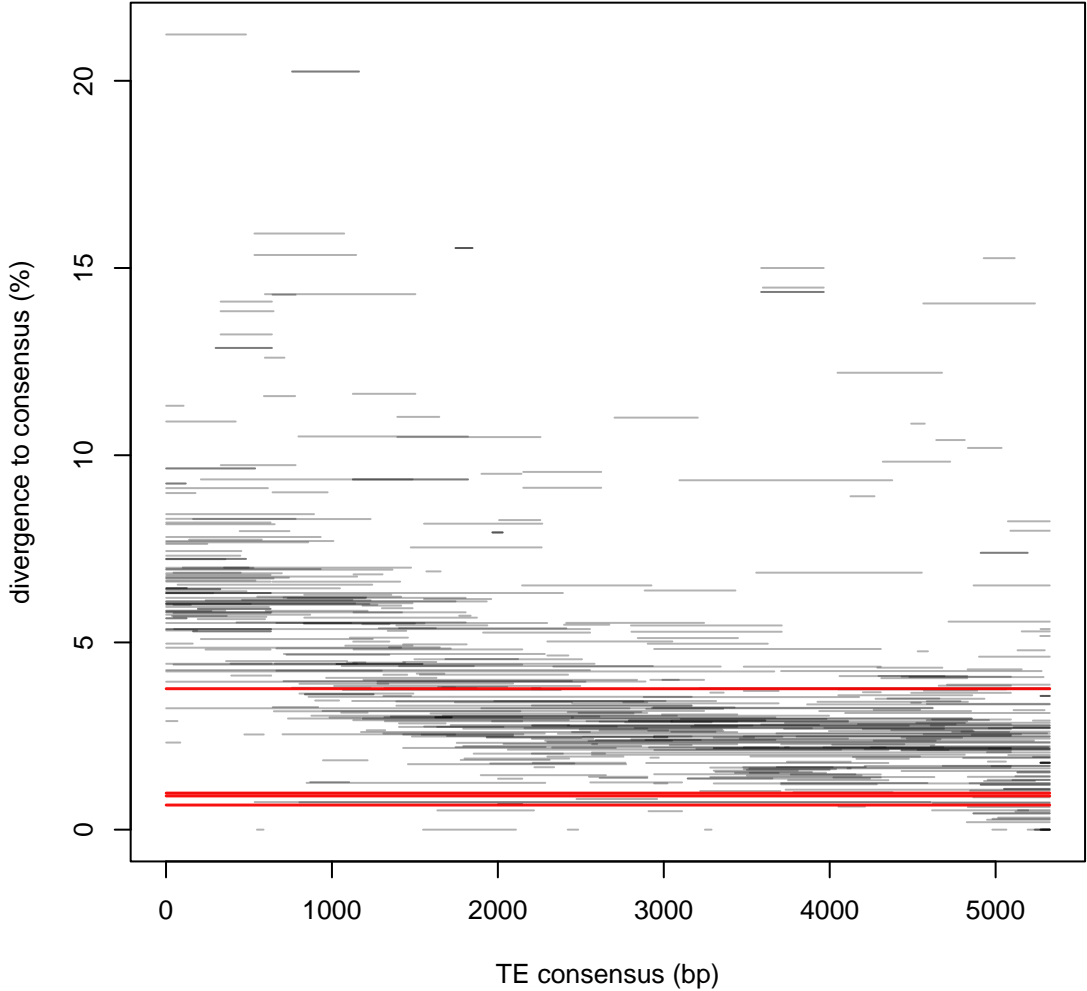
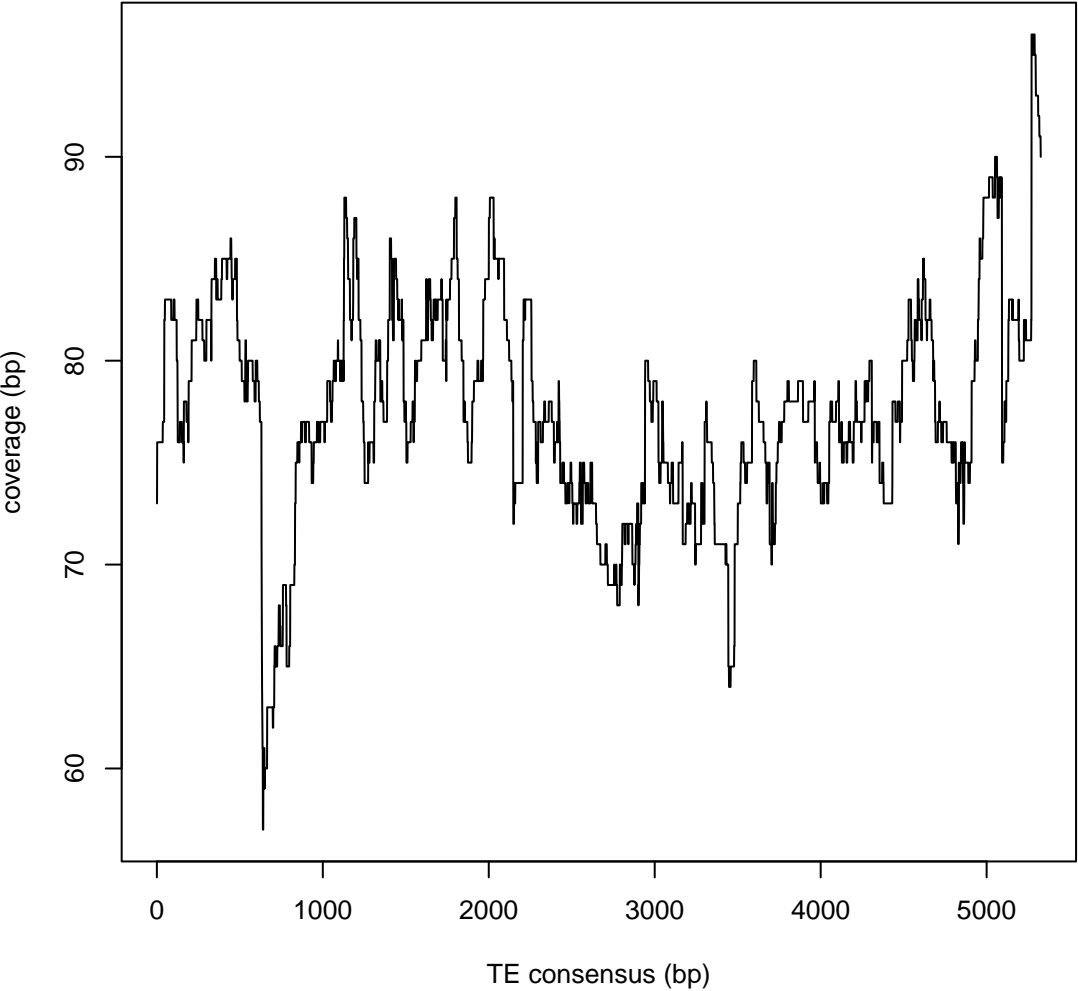


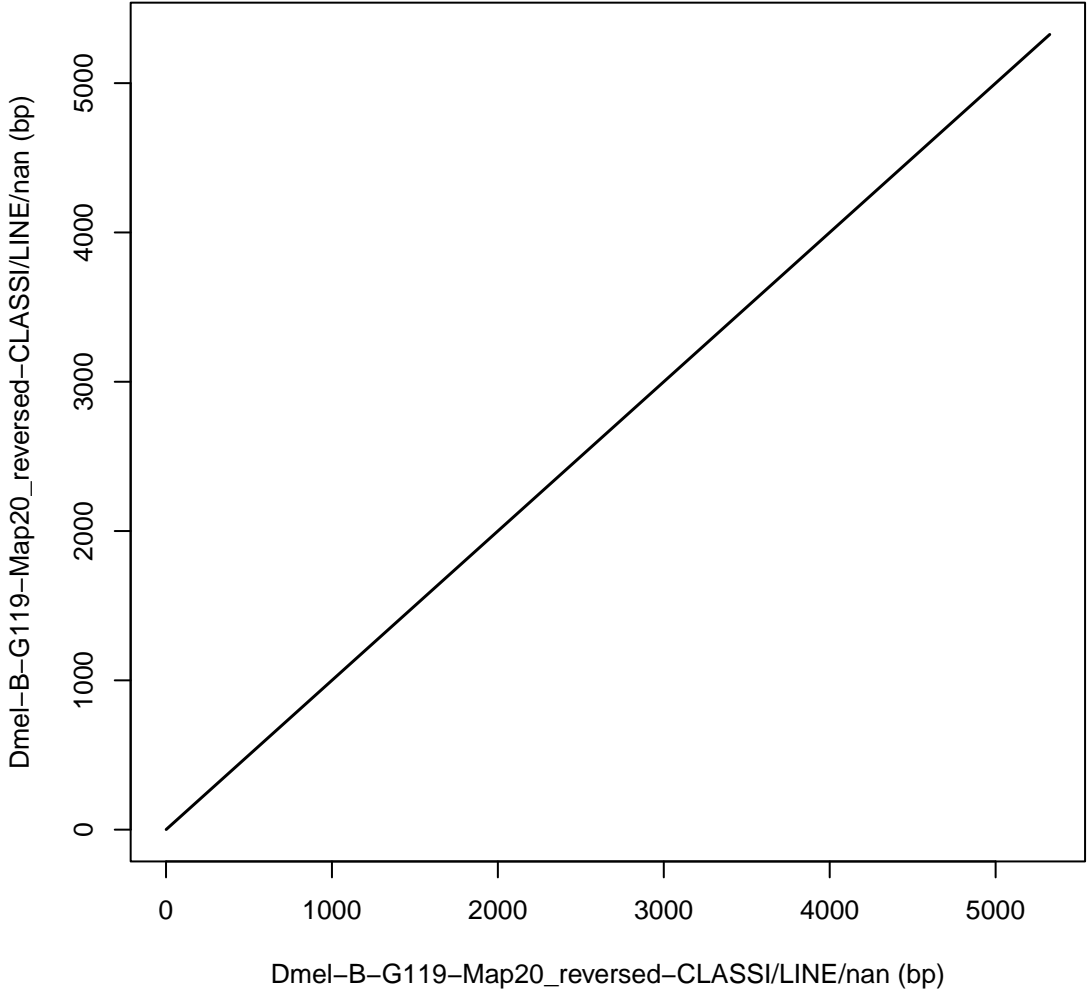
TE: Dmel-B-G119-Map20_reversed-CLASSI/LINE/nan
consensus size: 5326bp; fragments: 527; full length: 4 (>=4793.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

