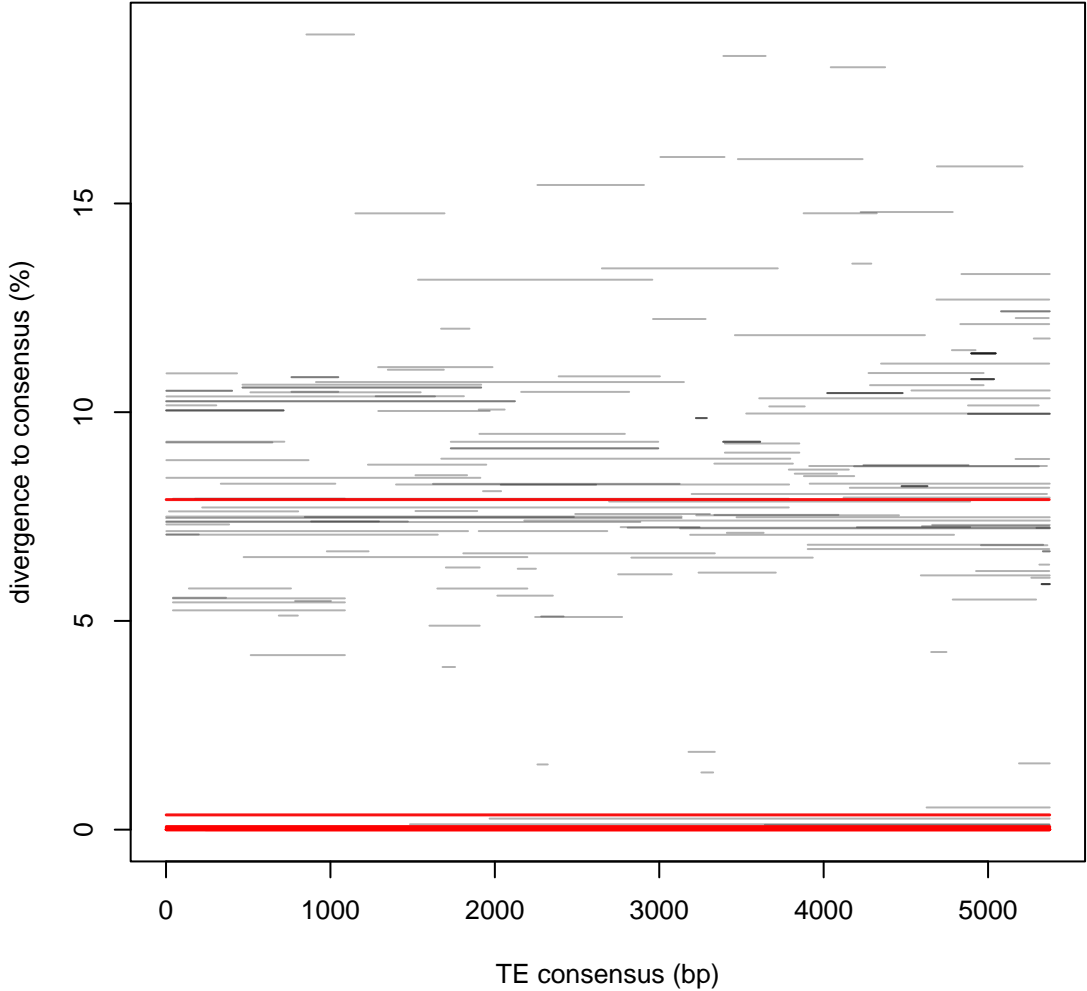
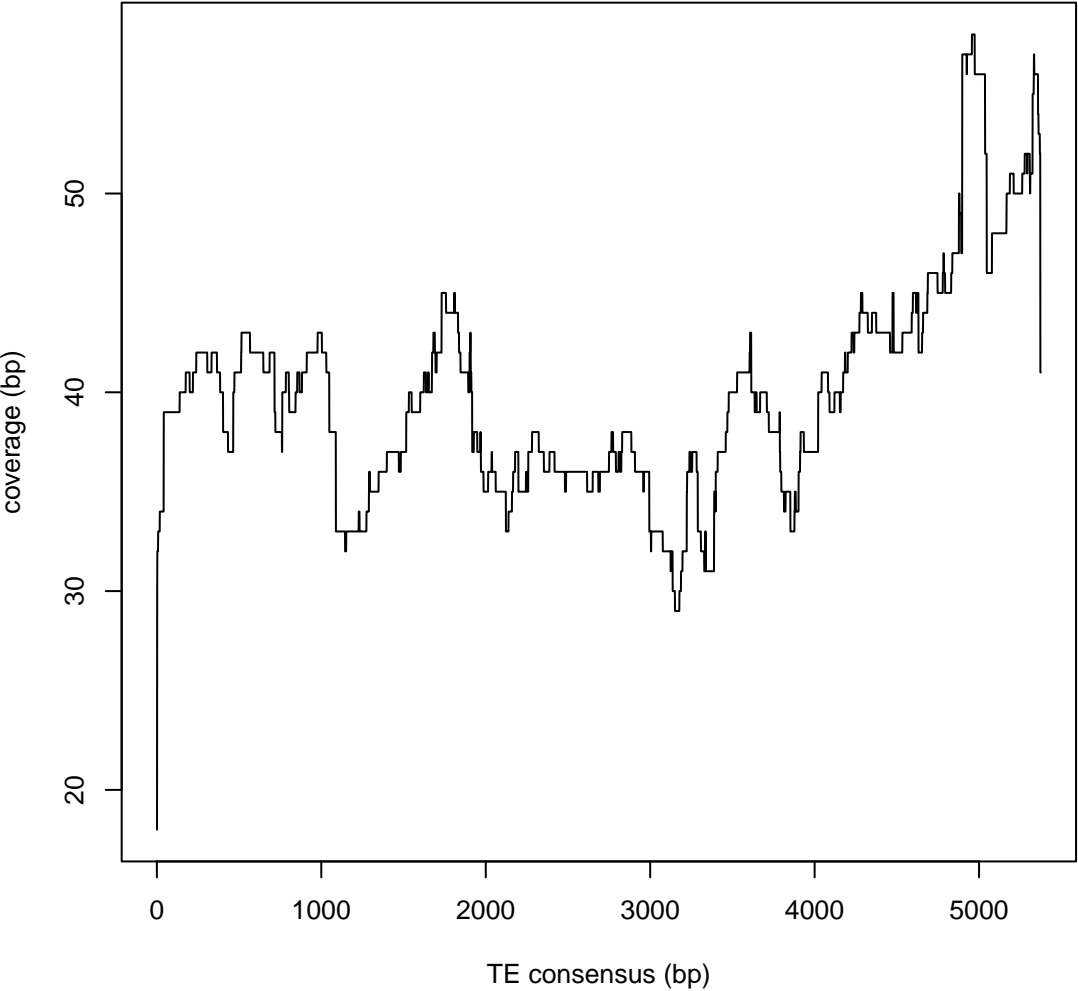


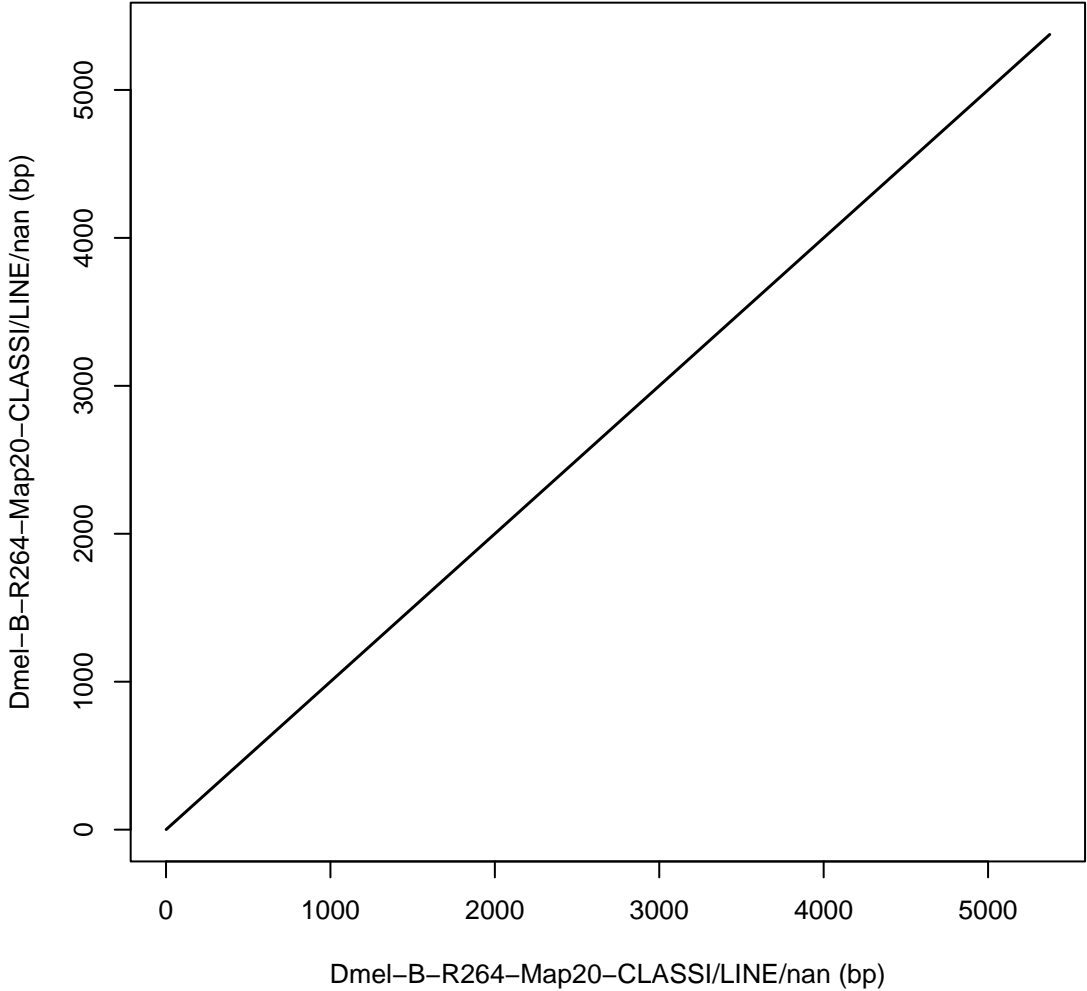
TE: Dmel-B-R264-Map20-CLASSI/LINE/nan
consensus size: 5375bp; fragments: 202; full length: 10 (>=4837.5bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

