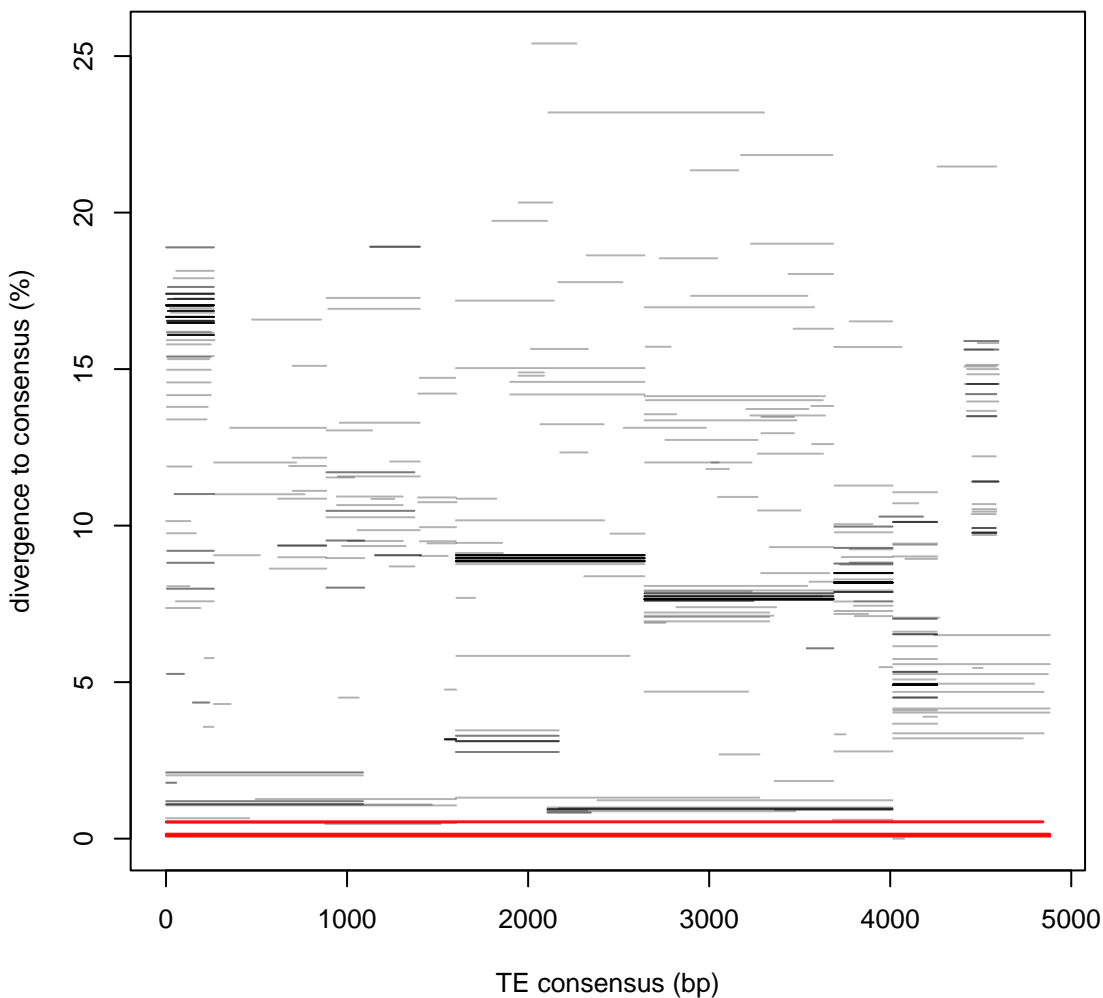
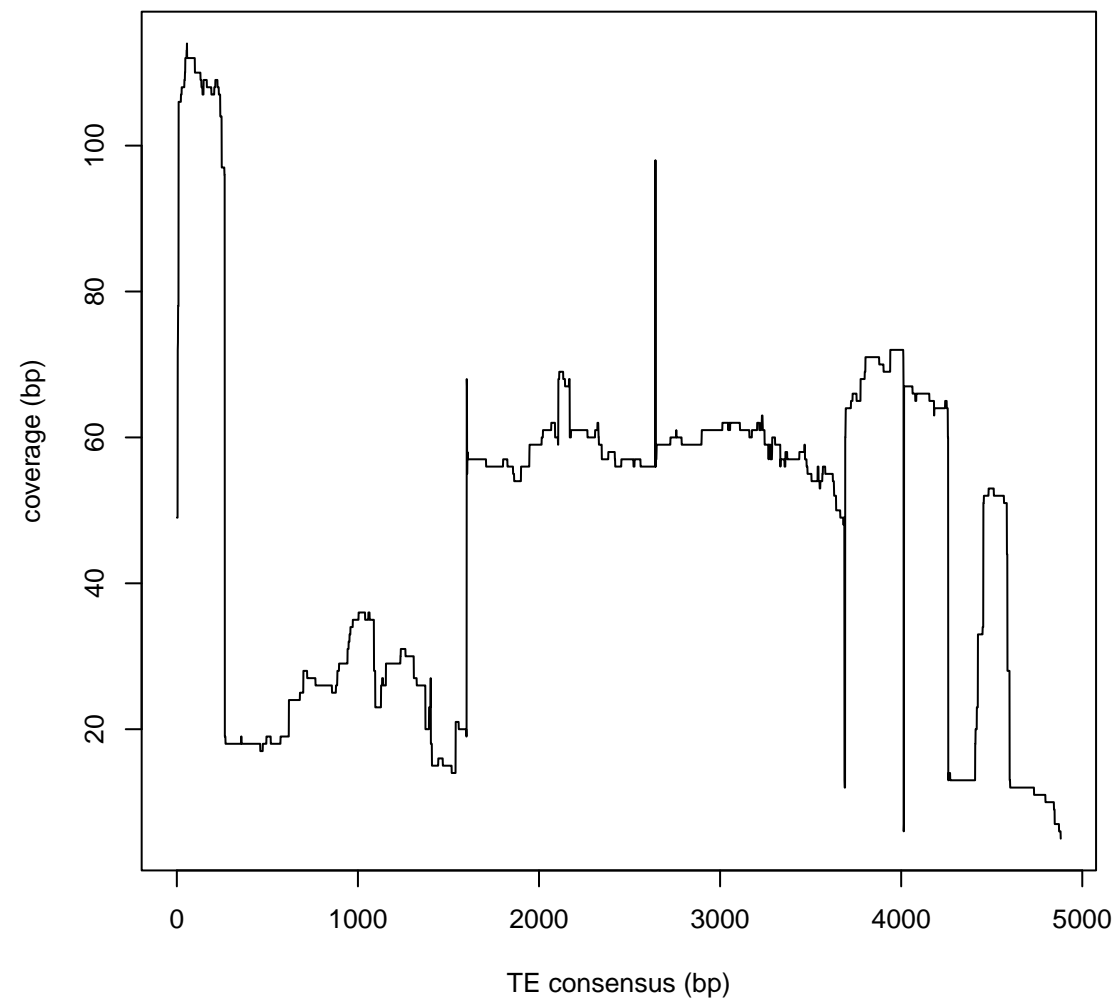


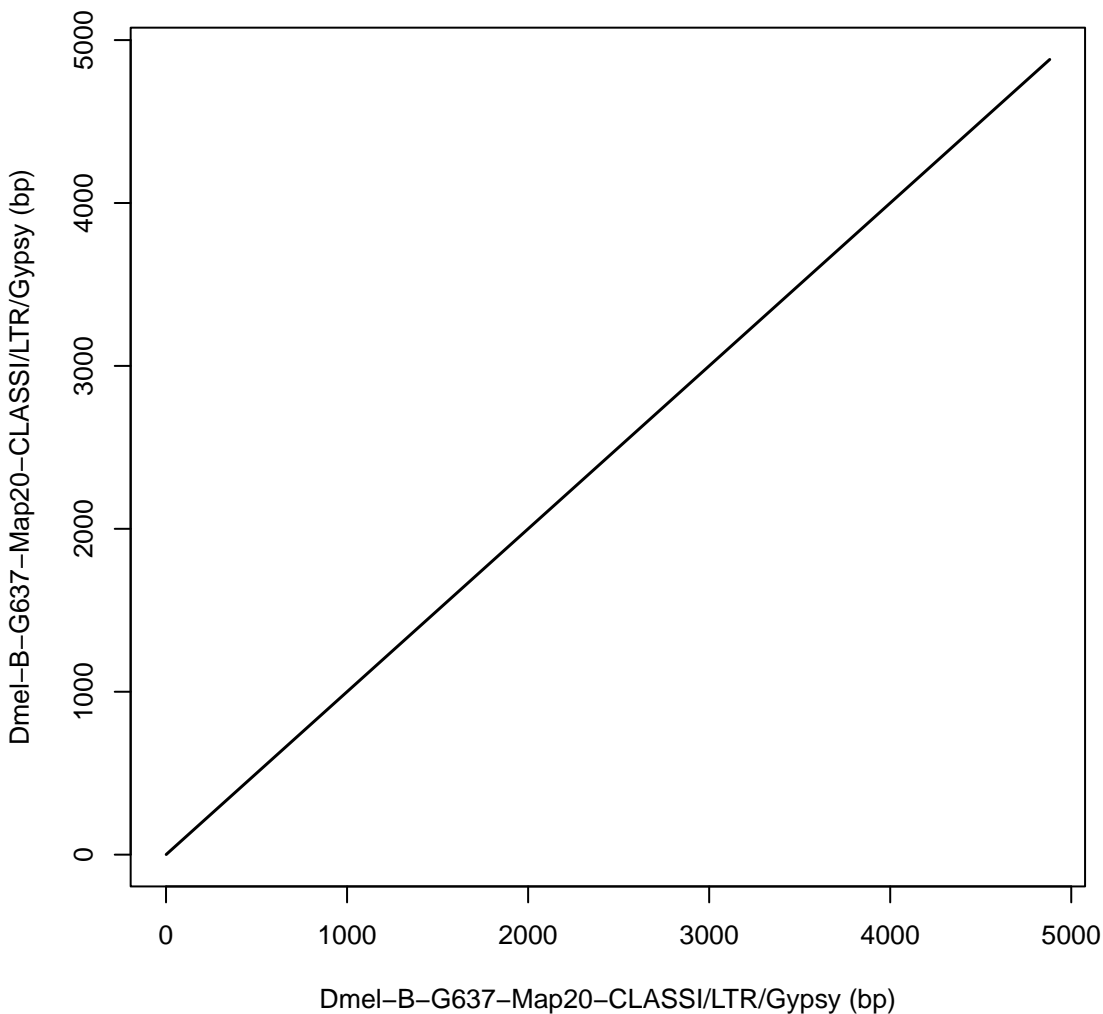
TE: Dmel-B-G637-Map20-CLASSI/LTR/Gypsy  
consensus size: 4881bp; fragments: 505; full length: 3 (>=4392.9bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

