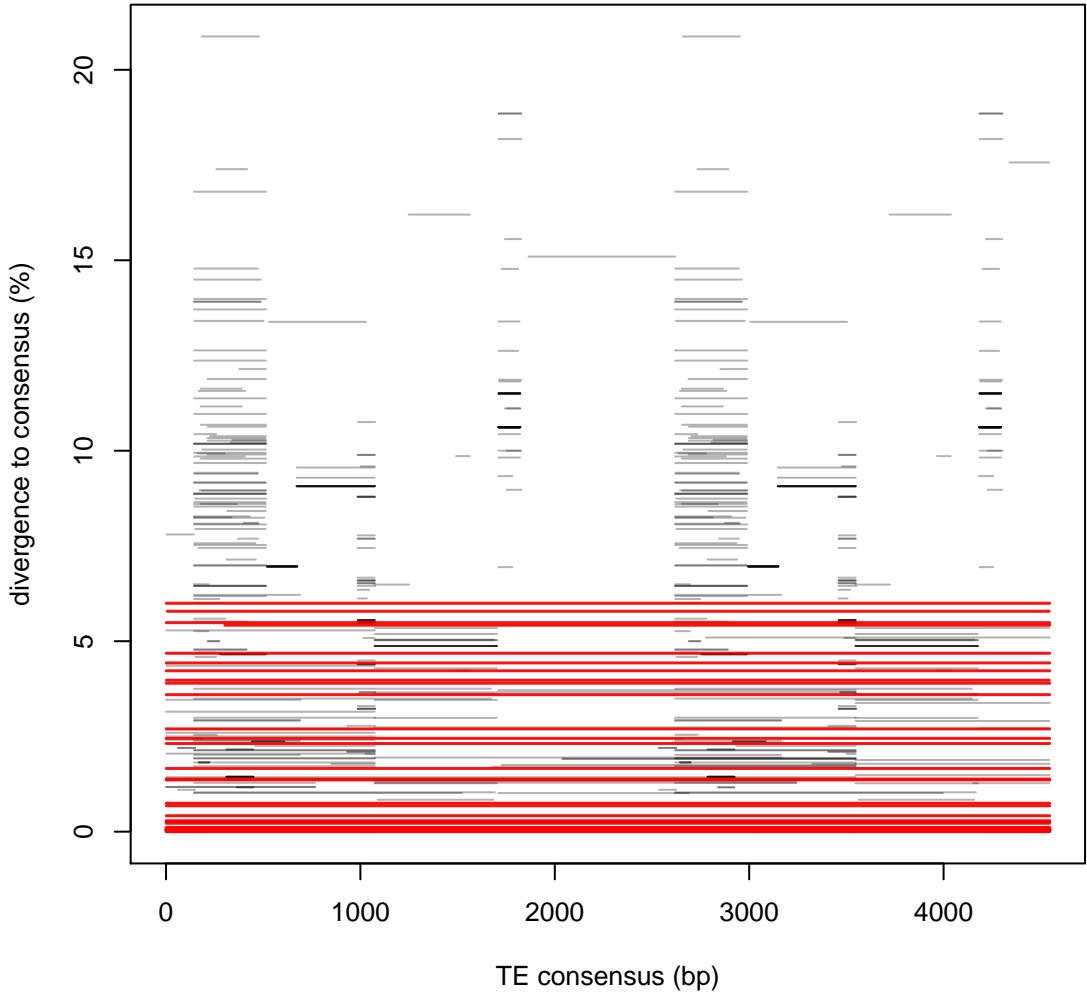
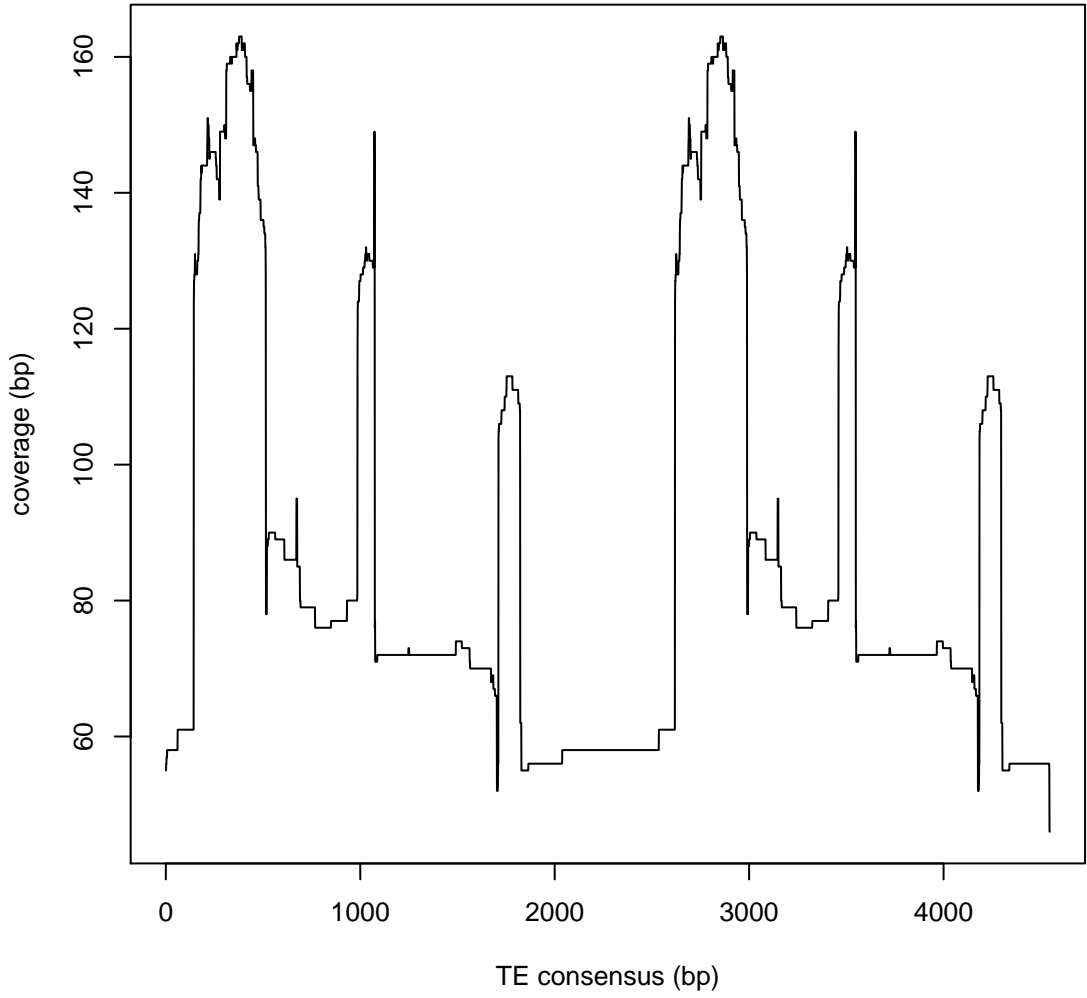


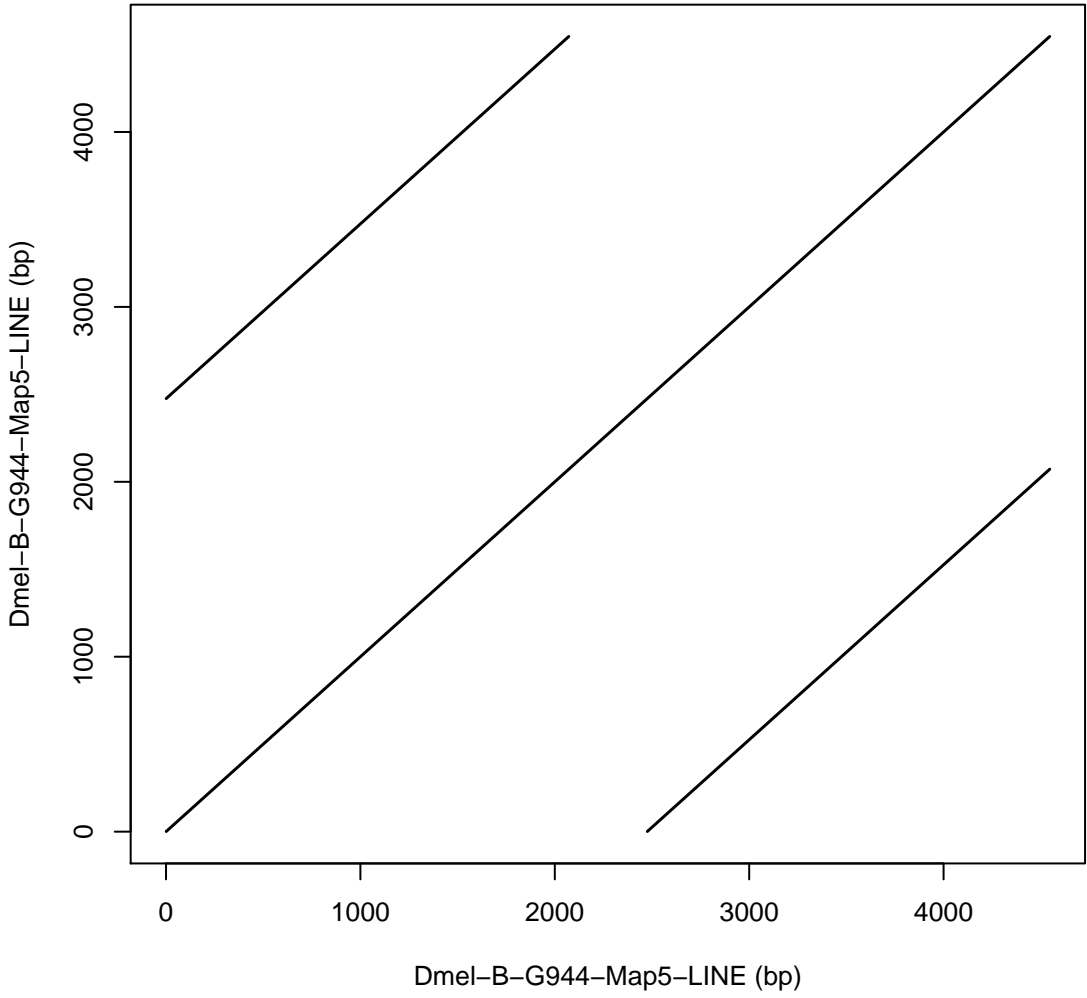
TE: Dmel-B-G944-Map5-LINE  
consensus size: 4546bp; fragments: 658; full length: 45 (>=4091.4bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

