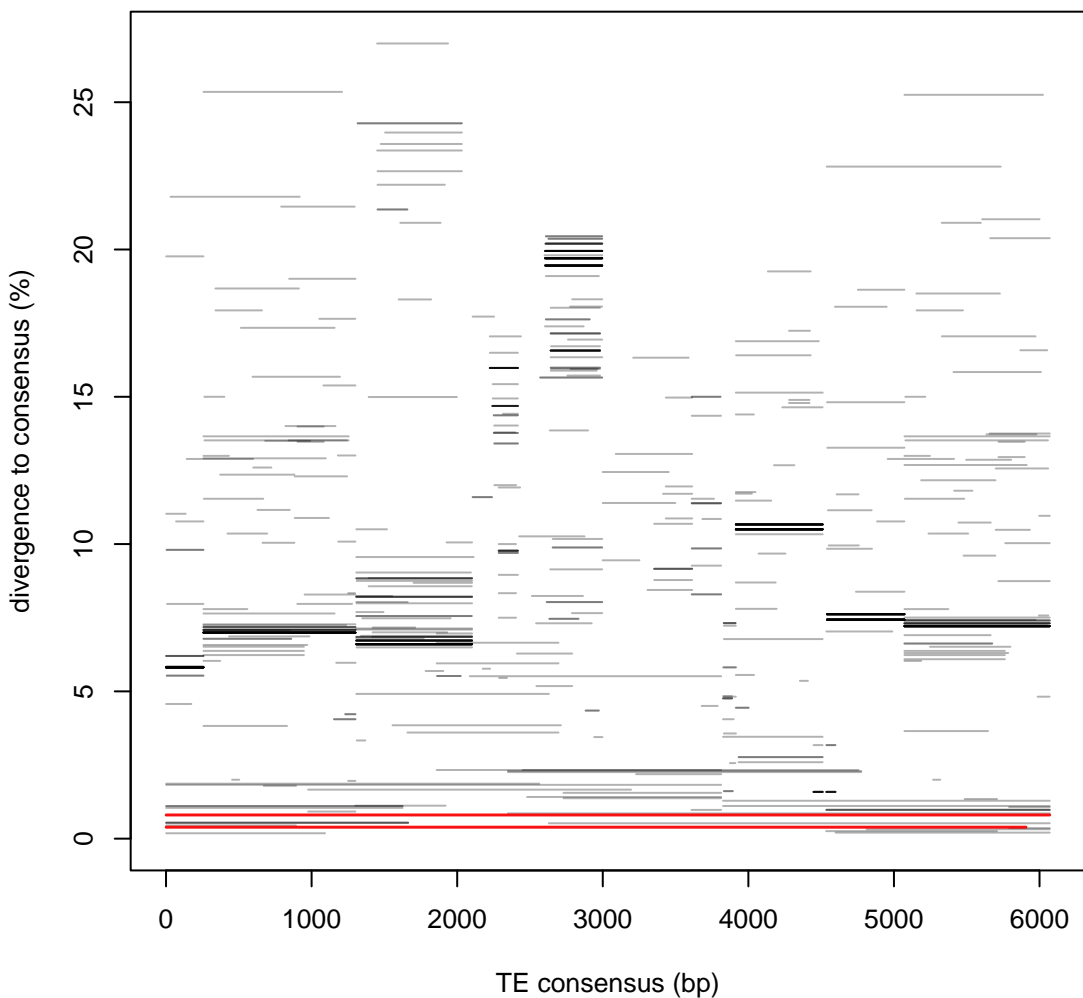
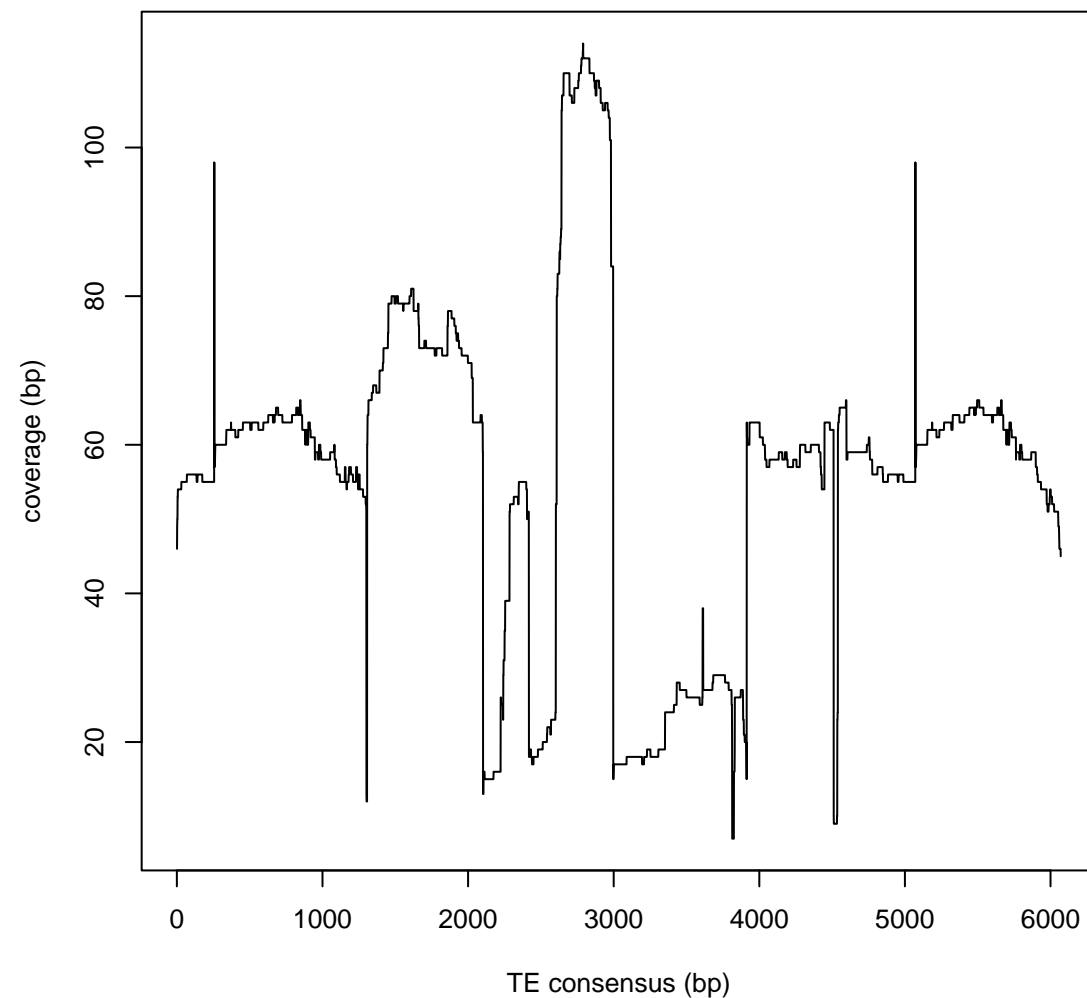


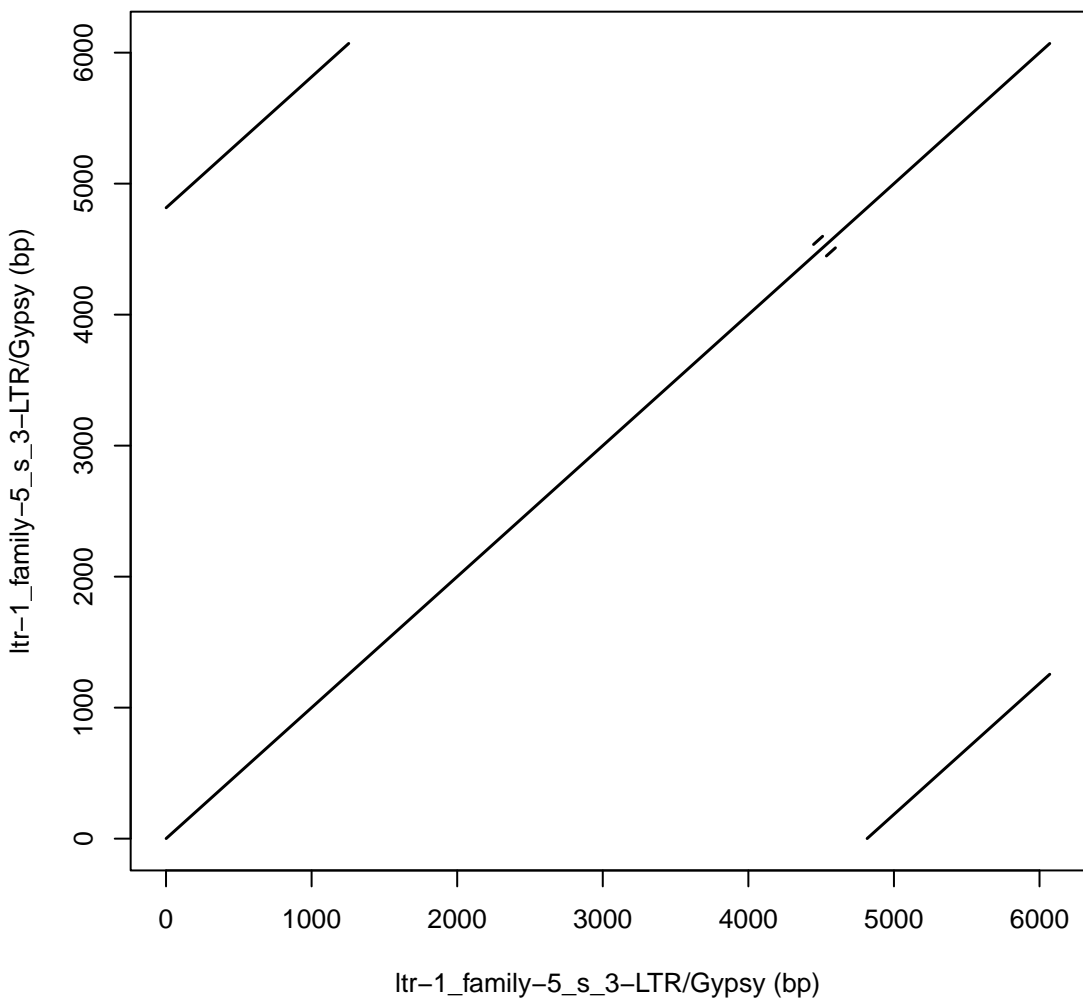
TE: ltr-1_family-5_s_3-LTR/Gypsy
consensus size: 6070bp; fragments: 640; full length: 2 (>=5463bp)



TE consensus genomic coverage



TE consensus self dotplot (blastn)



TE consensus structure and protein hits

