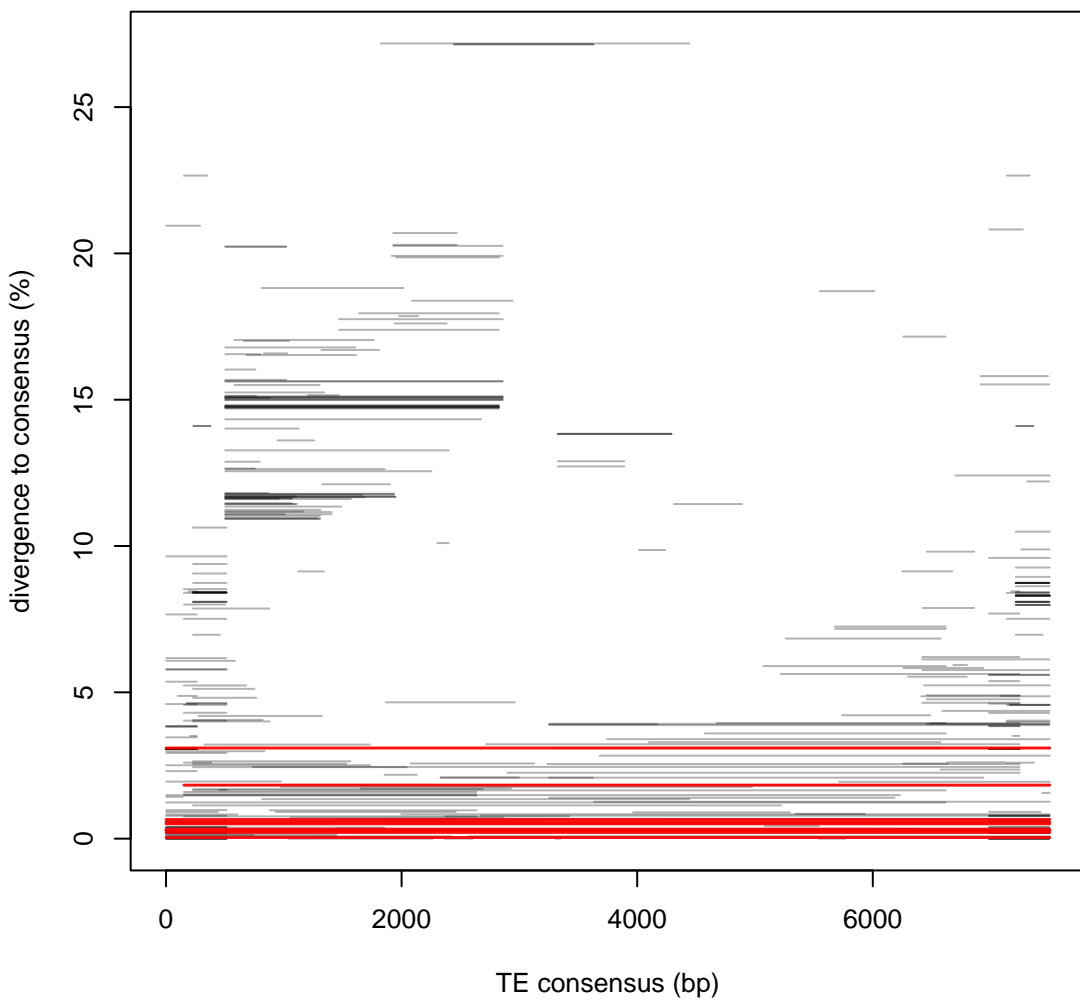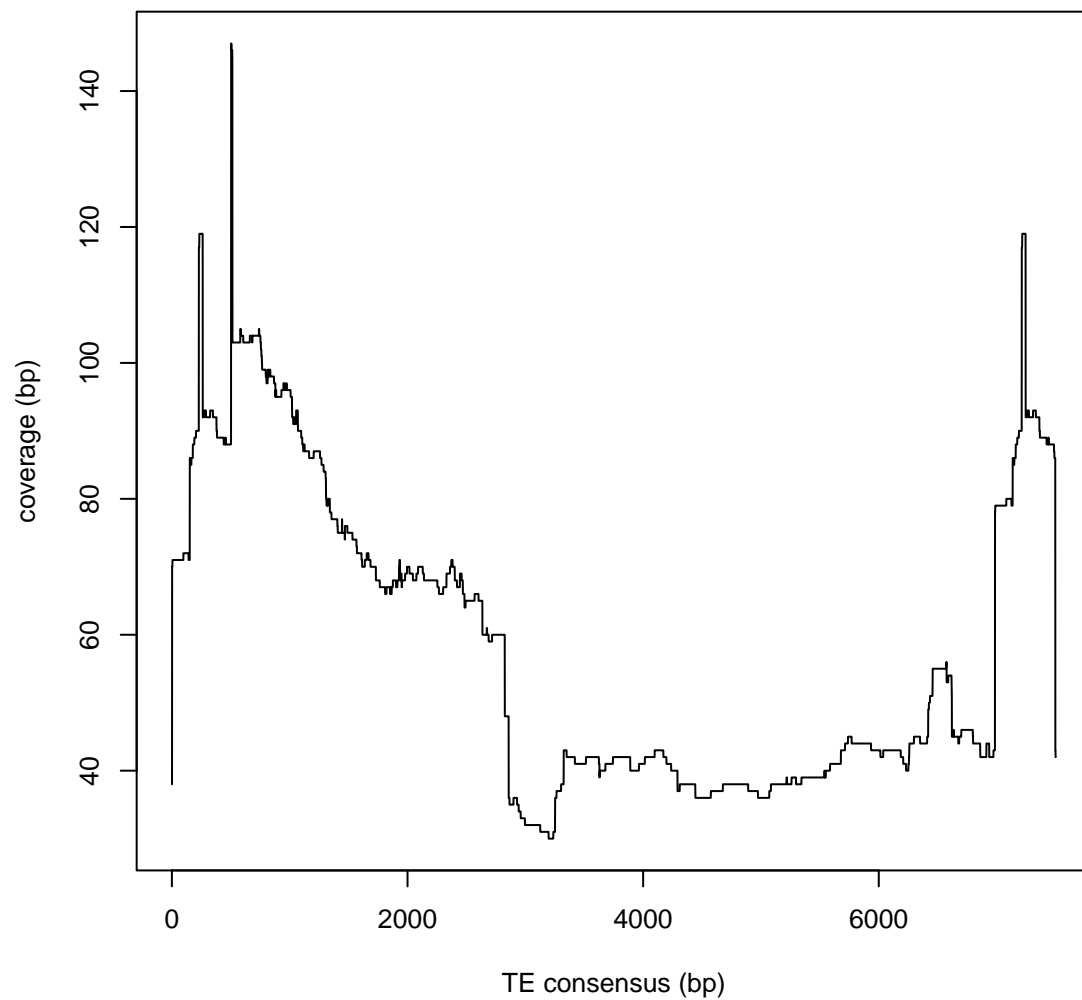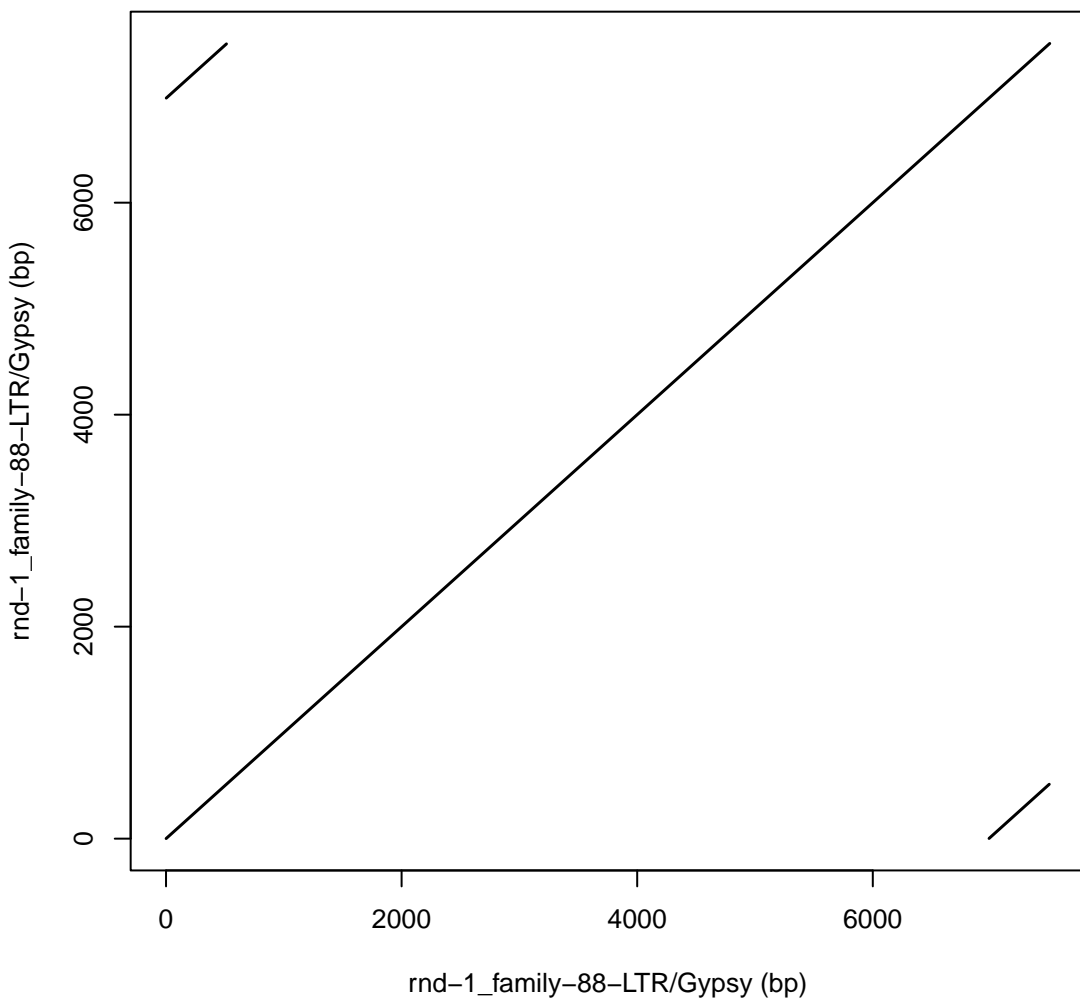**TE: rnd−1_family−88−LTR/Gypsy**
consensus size: 7502bp; fragments: 377; full length: 10 (>=6751.8bp)

TE consensus genomic coverage

TE consensus self dotplot (blastn)

TE consensus structure and protein hits

DM176_env LTR/Gypsy
DM176_pol LTR/Gypsy
Gypsy−15_DEl_gag LTR/Gypsy
Invader2 LTR/Gypsy