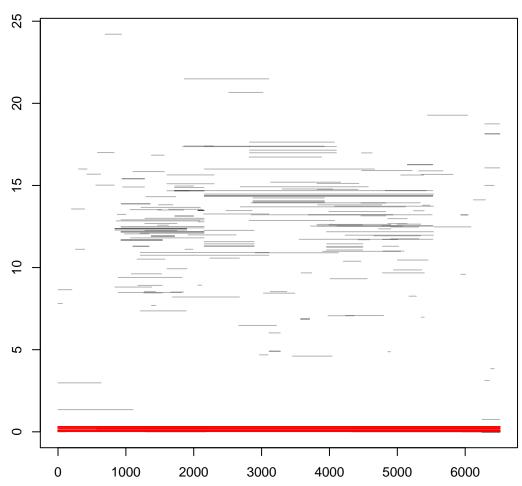
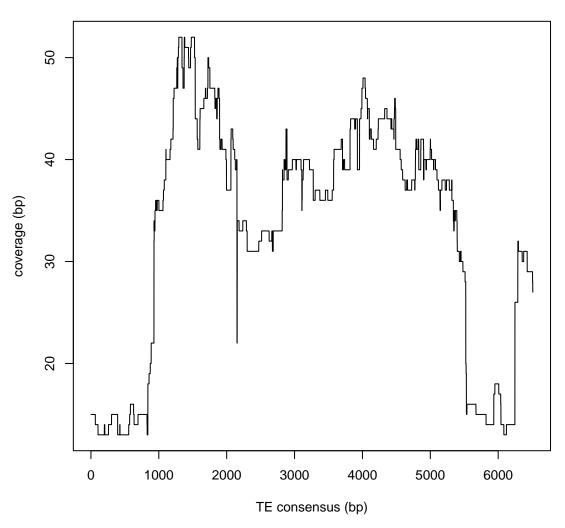
TE: rnd-1_family-73-LTR/Gypsy consensus size: 6511bp; fragments: 227; full length: 10 (>=5859.9bp)



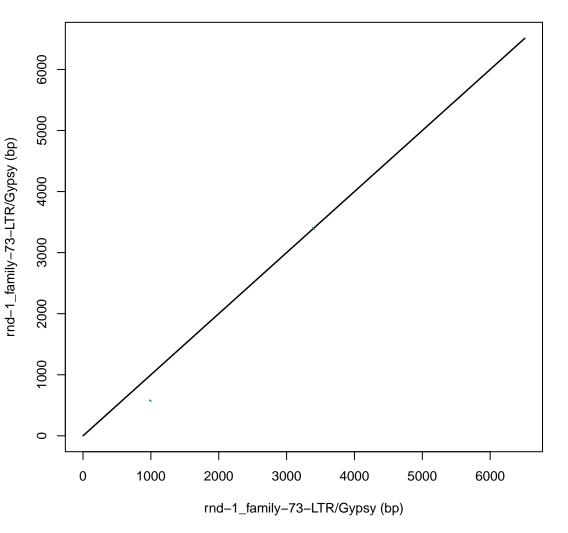
divergence to consensus (%)

TE consensus genomic coverage



TE consensus self dotplot (blastn)

TE consensus (bp)



TE consensus structure and protein hits

