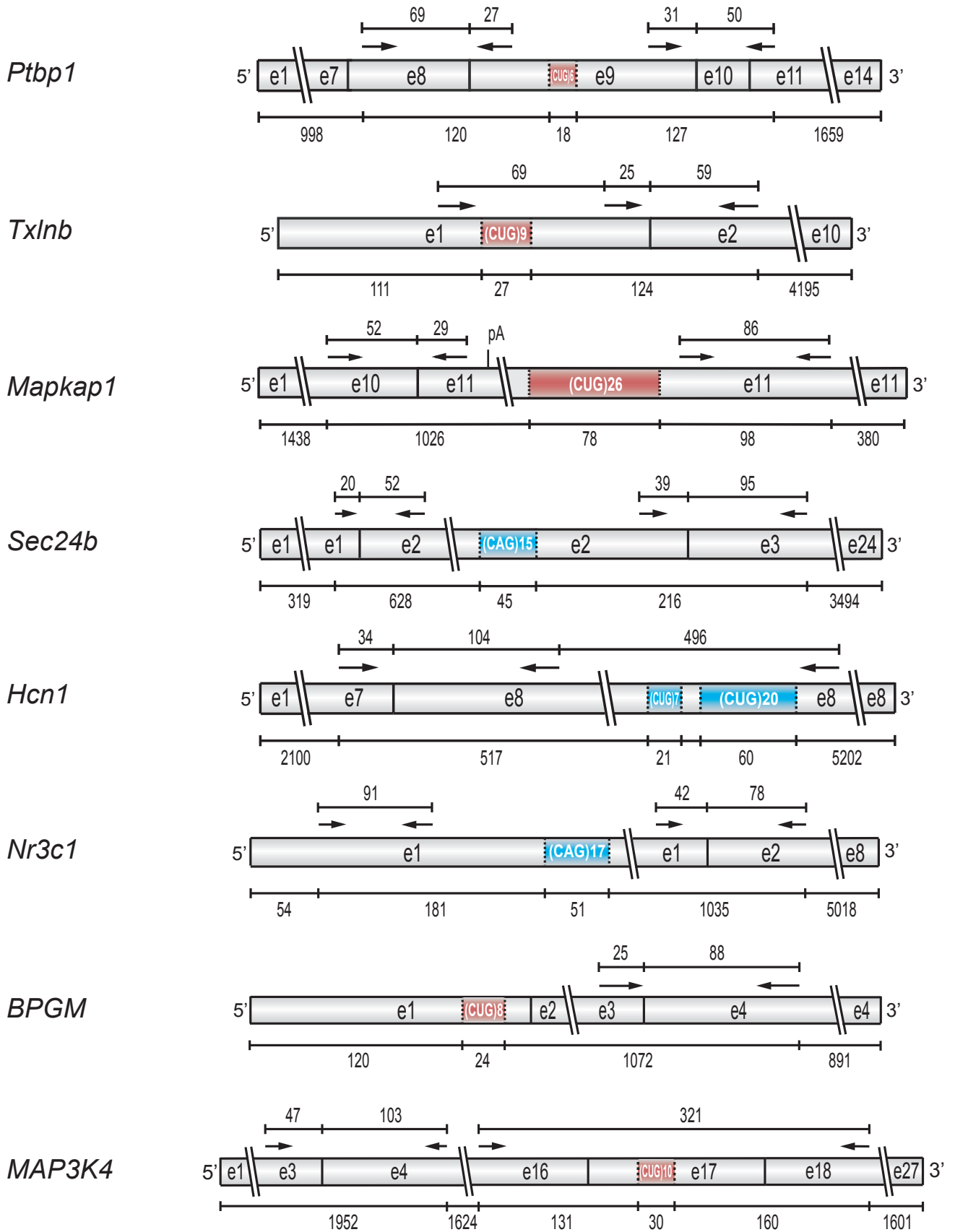


# FIGURE S4



## Supplementary Figure S4.

Schematic representation of triplet repeat transcripts and segments thereof, analyzed by RT-(q)PCR. Boxes indicate exons. (CUG)*n* and (CAG)*n* tracts are highlighted in red and blue, respectively. PCR primers are represented by arrows. Transcript lengths are based on NCBI Reference Sequences and indicate nucleotides. pA indicates alternative polyadenylation site in *Mapkap1*.