



Figure S7: Increased recruitment of TF2C-type complexes containing polyglutamine expanded ATXN7 at different regions of the *Rho* gene.

Chromatin immunoprecipitation assays (ChIP) were performed using formaldehyde-fixed chromatin extracts of retina from 2 month-old control (WT or R7N; dark grey) and R7E (light grey) mice. Specific regions within the *Rho* gene were analyzed by real-time PCR, after ChIP using an antibody against a TF2C-specific subunit (Spt3). Quantification of fold enrichment was calculated as indicated in Figure S9. Each bar represents the mean value \pm SEM ($n = 4-6$).