



Supplementary Figure S1. Semi-quantitative RT-PCR for seven mouse transcripts that contain CUG or CAG repeats.

The length of the repeat tract for each transcript is indicated. Assays were performed using RNA isolated from saline (sal) or siCAG (si) electroporated TA muscle of HSA^{LR} , and compared to untreated TA muscle from wild type mice (WT). The bar graph shows mean \pm SD from three treated mice and two independent RT-PCR assays, normalized to *Gapdh*. ** $P < 0.05$ for siCAG-tail versus saline treated muscles (Student's *t*-test).