Figure S3.

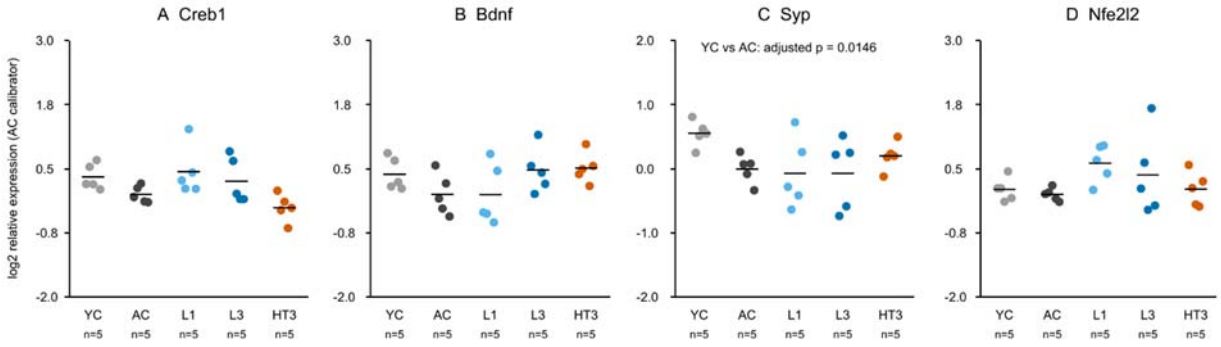


Figure S3. Brain transcripts. Exploratory whole-brain qPCR reanalysis for (A) Creb1, (B) Bdnf, (C) Syp, and (D) Nfe2l2. Each point represents one mouse (n = 5/group). ΔCt uses Actb as the reference gene and the AC mean as calibrator; plots show log2 relative expression. Horizontal lines indicate the mean. Two-sided Welch tests on ΔCt were adjusted with Benjamini–Hochberg correction separately across four age and twelve treatment contrasts. Syp differed between YC and AC (adjusted $p = 0.0146$). No treatment contrast met adjusted $p < 0.05$.