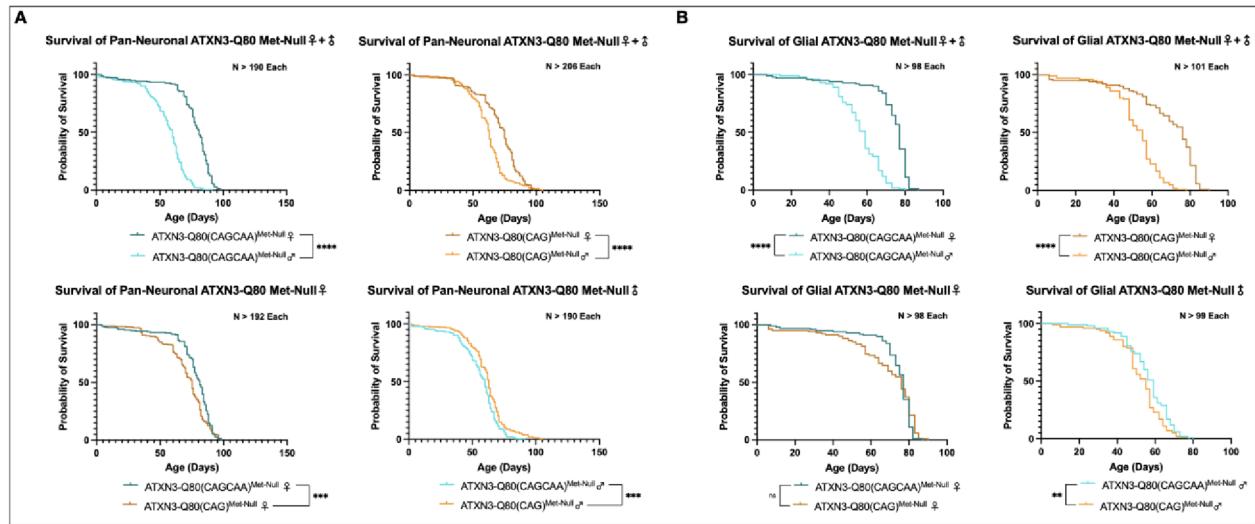


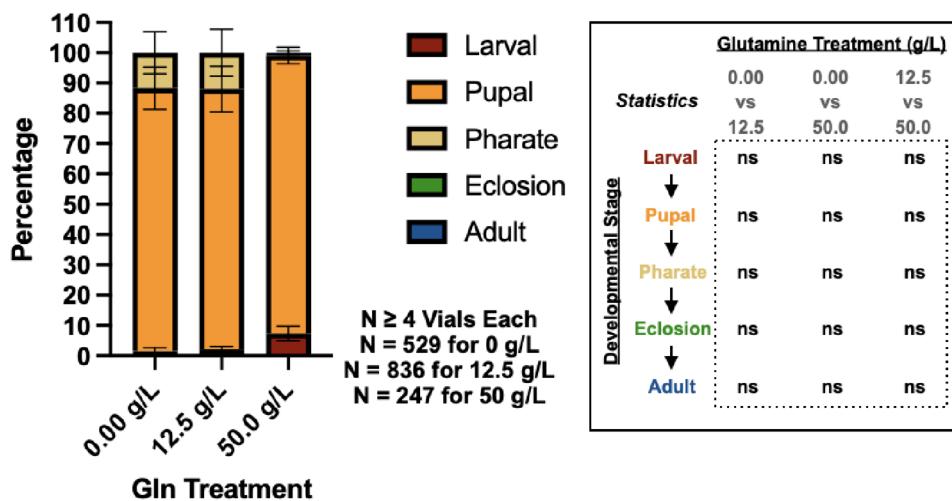
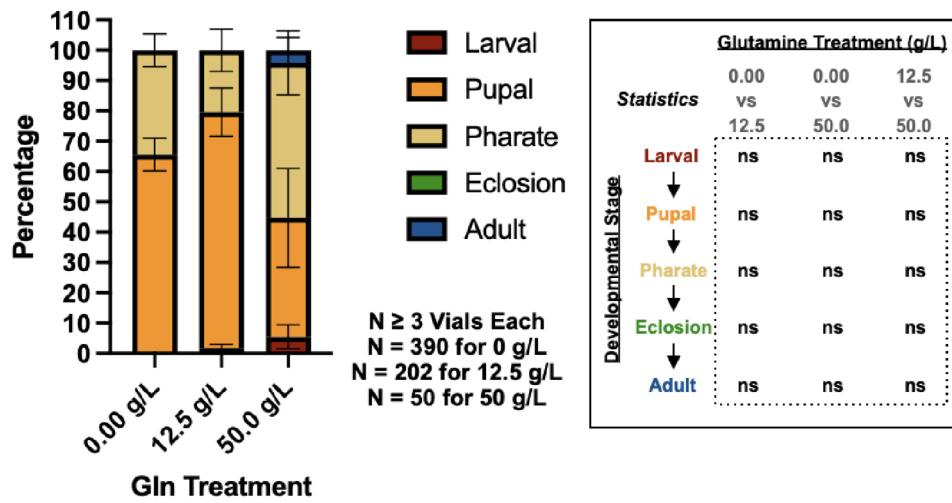
Figure S1. Ubiquitous expression longevity data and individual comparisons from survival analysis in figure 2

(A) Ubiquitous expression of ATXN3-Q80(CAG) and ATXN3-Q80(CAGCAA) individually was developmentally lethal in flies. Observations of developmental deaths in both lines based on the fly life cycle (shown at top of figure) showed ATXN3-Q80(CAG) flies dying primarily at earlier developmental stages than flies ubiquitously expressing ATXN3Q80(CAGCAA). These observations were confirmed with survival analysis in tissue-specific expression models shown in figure 2. (B) Individual comparisons of all pan-neuronal expressing fly lines shown in figure 2E. “\*\*”: p<0.0332; “\*\*\*”: p<0.0021; “\*\*\*\*”: p<0.0002; “\*\*\*\*\*”: p<0.0001. p-values from log-rank test. No adults eclosed in the glial expression analysis in figure 2F, thus no individual comparisons were made.



**Figure S2.** Individual comparisons from survival analysis in figure 6

(A) Individual comparisons of all pan-neuronal expressing fly lines shown in figure 6A. “\*\*”: p<0.032; “\*\*\*”: p<0.0021; “\*\*\*\*”: p<0.0002; “\*\*\*\*\*”: p<0.0001. p-values from log-rank test. (B) Individual comparisons of all glial expressing fly lines shown in figure 6C. “ns”: non-statistically significant; “\*\*”: p<0.032; “\*\*\*”: p<0.0021; “\*\*\*\*”: p<0.0002; “\*\*\*\*\*”: p<0.0001. pvalues from log-rank test.

**A****Ubiquitous ATXN3-Q80(CAG)****B****Ubiquitous ATXN3-Q80(CAGCAA)****Figure S3. Developmental deaths of ATXN3-Q80-expressing flies with glutamine supplementation**

(A) Developmental death tracking based on the percentage of fly deaths at each developmental stage in flies ubiquitously expressing ATXN3-Q80(CAG) in food with 0.00, 12.5, or 50.0 g/L glutamine supplementation. Driver was sqh-Gal4. Each liter of food made accounted for 100 fly vials, with each vial then receiving an additional 0.000, 0.125, or 0.500 g/L of glutamine respectively. Means of percentages at each stage +/- SEM. "ns": non-statistically significant. p-values of all comparisons shown to the right of graph from RM two-way ANOVA with Geisser-Greenhouse correction and Tukey's multiple comparison test. n ≥ 3 vials each. (B) Developmental death tracking based on percentage of fly deaths at each developmental stage in flies ubiquitously expressing ATXN3-Q80(CAGCAA) in food with 0.00, 12.5, or 50.0 g/L glutamine supplementation in food. Means of percentages at each stage +/- SEM. "ns": nonstatistically significant. p-values of all comparisons shown to the right of graph and from RM two-way ANOVA with Geisser-Greenhouse correction and Tukey's multiple comparison test. n ≥ 3 vials each.

Figure S4: Nucleotide sequences for Panel 5A

**Met-Null CAGCAA Sequence**

GAATTCACTTACCGAAATAGTAGGAGTCATCTCCACGAGAA  
ACAAGAAGGCTCACTTGCTCAACATTGCCGTGAATAACT  
TATTGCAAGGAGAATATTTAGCCCTGTGGAATTATCCTCAA  
TTGCACATCAGCTGGATGAGGAGGGAGAGGTAGAGATAGGC  
AGAAGGAGGAGTTACTAGTGAAGATTATCGCACGTTTAC  
AGCAGCCTCTGGAAATTAGGATGACAGTGGTTTCTCT  
ATTCAAGGTTATAAGCAATGCCTGAAAGTTGGGGTTAGAA  
CTAACCTGTTAACAGTCCAGAGTATCAGAGGCTCAGGAT  
CGATCCTATAAATGAAAGATCATTATGCAATTATAAGGAA  
CACTGGTTACAGTTAGAAAATTAGGAAAACAGTGGTTAA  
CTTGAATTCTCTCTGACGGGTCCAGAATTATCAGATAC  
ATATCTGCACCTTCTGGCTCAATTACAACAGGAAGGTTA  
TTCTATATTGCGTTAAGGGTGTCTGCCAGATTGCGAAGC  
TGACCAACTCCTGCAGTAGATTAGGGTCCAACAGTAGCAT  
CGACCAAAACTTATTGGAGAAGAATTAGCACAACAAAAAGA  
GCAAAGAGTCCATAAACAGACCTGGAACCGAGTGTAGAA  
GCAAATGATGGCTCAGGATAGTTAGACGAAGATGAGGAGG  
ATTGCAAGAGGGCTCTGGCACTAAGTCGCCAACAAATTGA  
CTAGGAAGATGAGGAAGCAGATCTCCGCAGGGCTATTCA  
CTAAGTTAGCAAGGTAGTCCAGAAACATATCTCAAGATTAG  
ACACAGACATCAGGTACAAATCTTACTTCAGAACAGCTCG  
GAAGAGACGAGAAGCCTACTTGAAAAACAGCAACAGAAA  
CAACAGCAACAGCAACAGCAACAGCAACAGCAACAGCAA  
CAGCAACAGCAACAGCAACAGCAACAGCAACAGCAACAG  
CAACAGCAACAGCAACAGCAACAGCAACAGCAACAGCAA  
CAGCAACAGCAACAGCAACAGCAACAGCAACAGCAACAA  
CAGCAAGGGGACCTATCAGGACAGAGTTCACATCCATGCG  
AAAGGCCAGCCACCAGTTCAGGAGCAGTCCAGGGAGCGATC  
TCGGCGATGCTTAGAGCGAAGAAGACTAGCTTCAGGCAG  
CTGTCACCTAGTCTTGGAAACTGTCAGAAACGATTGAAA  
ACAGAAGGAAAAAAATACCCATACGATGTTCCAGATTACGC  
TTAATAAAATGGAACAAAAACTTATTCTGAAGAAGATCTGT  
GGTAAGCCTATCCCTAACCCCTCCCTCGGTCTGATTCTAC  
GTAATAATCTAGA

**Met-Null CAGCAG Sequence**

GAATTCACTTACCGAAATAGTAGGAGTCATCTCCACGAGAA  
ACAAGAAGGCTCACTTGCTCAACATTGCCGTGAATAACT  
TATTGCAAGGAGAATATTTAGCCCTGTGGAATTATCCTCAA  
TTGCACATCAGCTGGATGAGGAGGGAGAGGTAGAGATAGGC  
AGAAGGAGGAGTTACTAGTGAAGATTATCGCACGTTTAC  
AGCAGCCTCTGGAAATTAGGATGACAGTGGTTTCTCT  
ATTCAAGGTTATAAGCAATGCCTGAAAGTTGGGGTTAGAA  
CTAACCTGTTAACAGTCCAGAGTATCAGAGGCTCAGGAT  
CGATCCTATAAATGAAAGATCATTATGCAATTATAAGGAA  
CACTGGTTACAGTTAGAAAATTAGGAAAACAGTGGTTAA  
CTTGAATTCTCTCTGACGGGTCCAGAATTATCAGATAC  
ATATCTGCACCTTCTGGCTCAATTACAACAGGAAGGTTA  
TTCTATATTGCGTTAAGGGTGTCTGCCAGATTGCGAAGC  
TGACCAACTCCTGCAGTAGATTAGGGTCCAACAGTAGCAT  
CGACCAAAACTTATTGGAGAAGAATTAGCACAACAAAAAGA  
GCAAAGAGTCCATAAACAGACCTGGAACCGAGTGTAGAA  
GCAAATGATGGCTCAGGATAGTTAGACGAAGATGAGGAGG  
ATTGCAAGAGGGCTCTGGCACTAAGTCGCCAACAAATTGA  
CTAGGAAGATGAGGAAGCAGATCTCCGCAGGGCTATTCA  
CTAAGTTAGCAAGGTAGTCCAGAAACATATCTCAAGATTAG  
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GATTCTACGTAATAATCTAGA