

SUPPLEMENTARY INFORMATION

Global analysis of protein degradation in prion infected cells

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Provided separately as Excel spreadsheets:

Table S1 – Peptide level search data

Table S2 – Protein level search data

Table S3 – Protein degradation rates

Table S4 – Optimized gradients for individual fractions during QExactive LC-MS/MS

Table S5 – Inclusion list for SILAC labeled PrP peptides generated using Skyline software program

Table S6 – MaxQuant software parameters for data analysis

Full-length western blots for the corresponding to the main figures in the text are provided on the following pages.

Figure 2 - A (left)

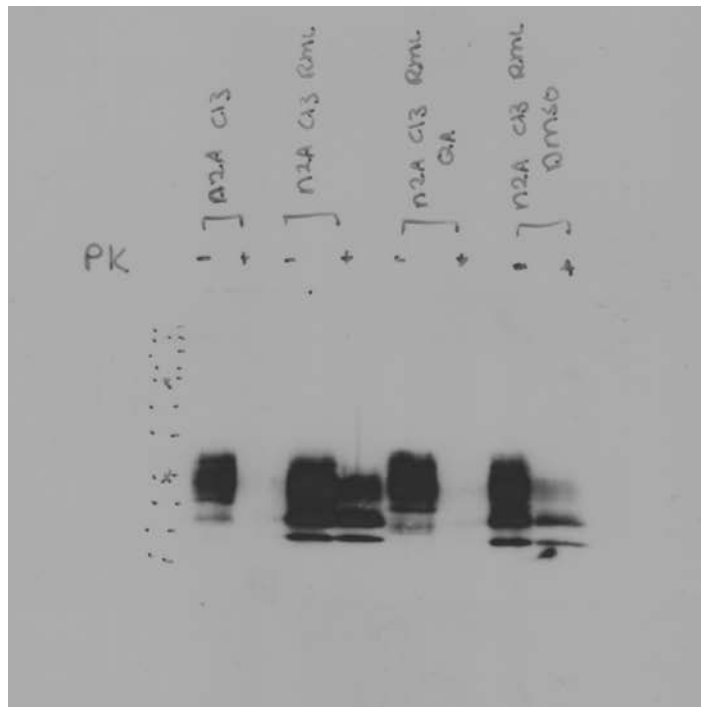


Figure 2 - A (right)

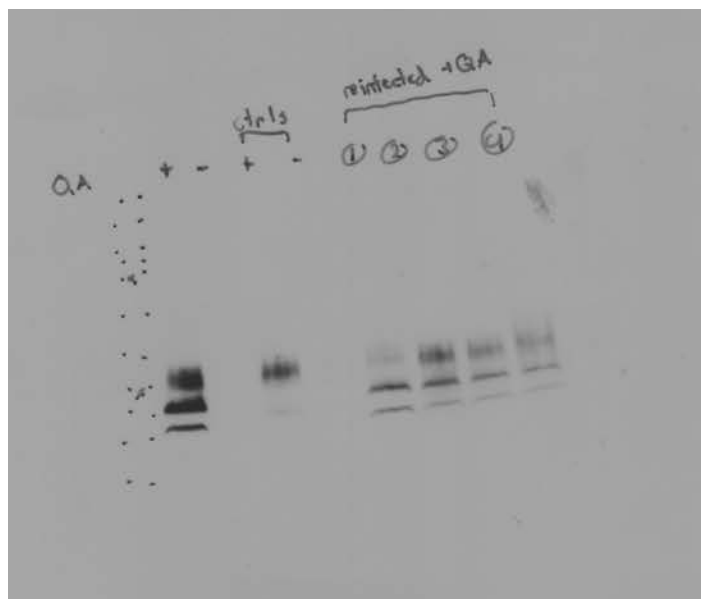


Figure 3 - D (left)

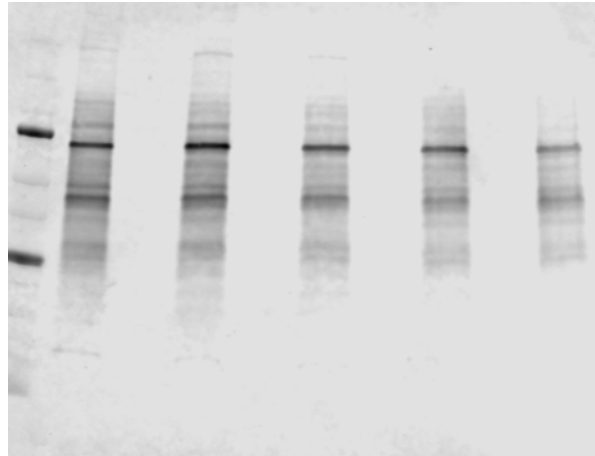


Figure 3 - D (right)

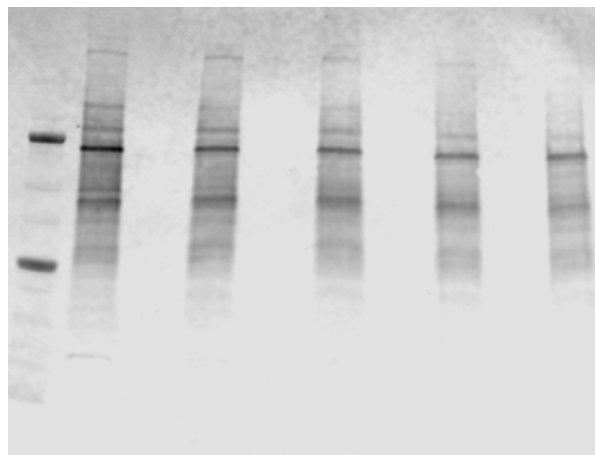


Figure 4 - C (upper left)

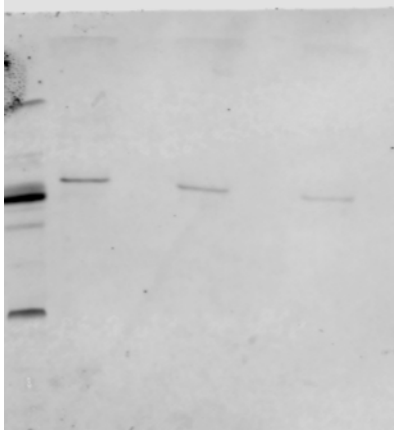


Figure 4 - C (upper right)

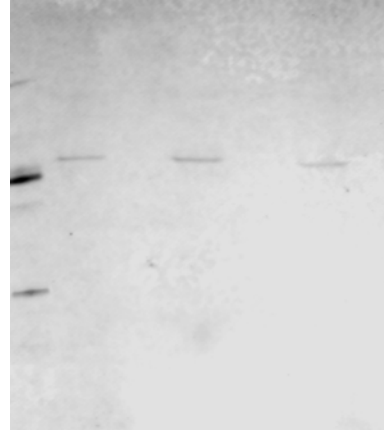


Figure 4 - C (lower left)

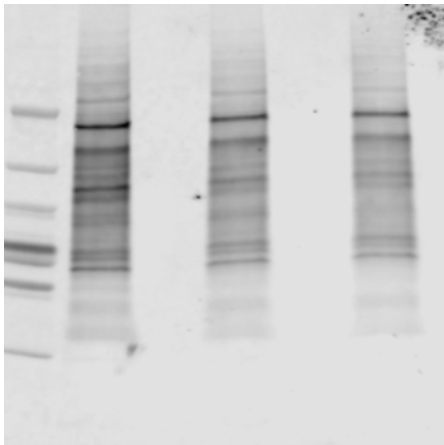


Figure 4 - C (lower right)

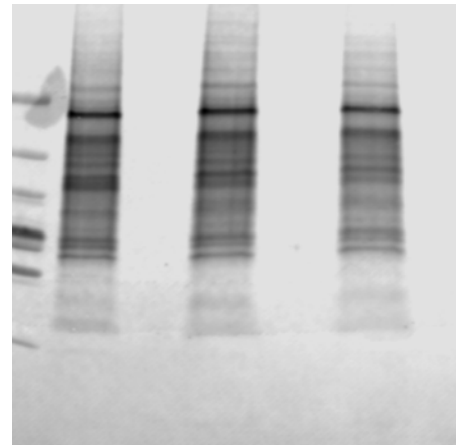


Figure 5 - A (cathepsin D)

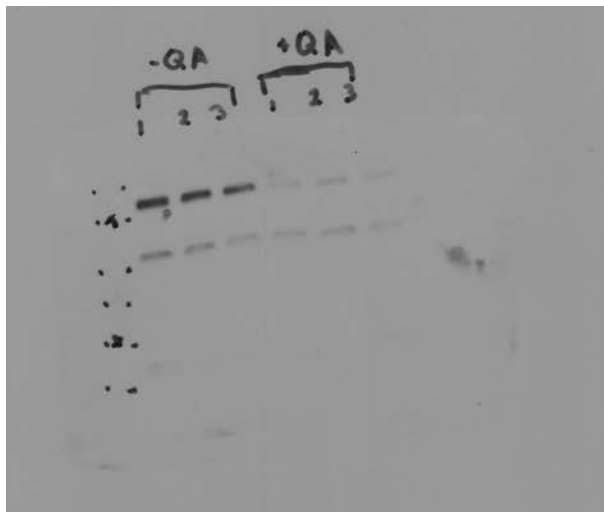


Figure 5 - A (cathepsin A)

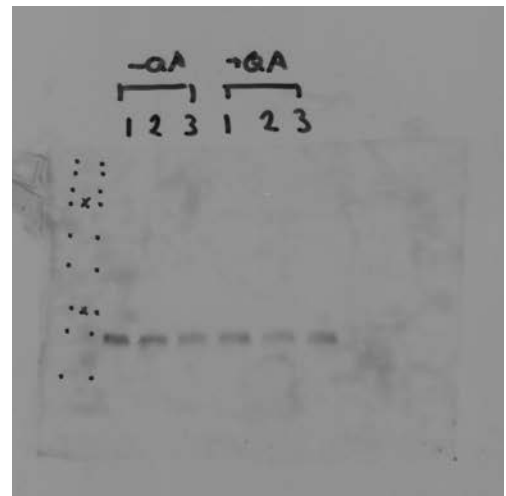


Figure 5 - A (cathepsin L)

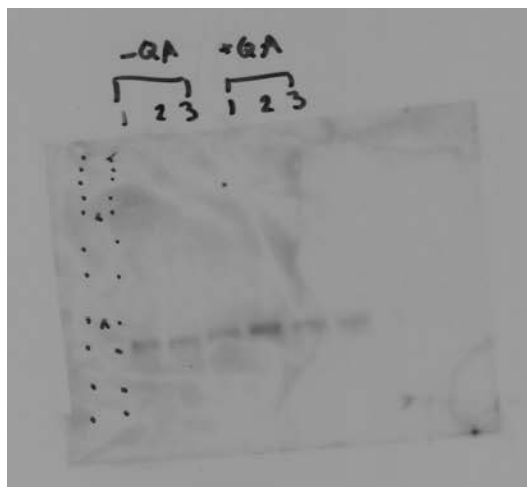


Figure 5 - A (p62/SQSTRM1)

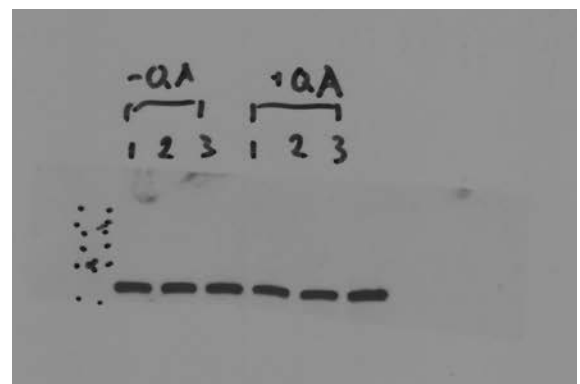


Figure 5 - A (LC3)



Figure 5 - A (Actin)

