

Gene/ Target	GenBank	Primer Sequence (5'-3')	Ref.
<i>Arc</i>	NM_018790.3	Fwd: CAGATTGGTAAGTGCCGAGC Rev: GTGCAACCCTTTTCAGCTCTC	
<i>Arg-1</i>	NM_007482.3	Fwd: CGCCTTTCTCAAAGGACAG Rev: CCAGCTCTTCATTGGCTTTC	(Jang, Lee et al. 2013)
<i>Cd68</i>	NM_009853.1	Fwd: CCACAGGCAGCACAGTGGACA Rev: TCCACAGCAGAAGCTTTGGCCC	(Kobayashi, Imagama et al. 2013)
<i>Cd163</i>	NM_001170395.1	Fwd: CCTCCTCATTGTCTTCCTCCTGTG Rev: CATCCGCCTTTGAATCCATCTCTTG	(Ly, Baghat et al. 2010)
<i>Egr2</i>	NM_001373987.1	Fwd: CGAGGGGACACACTGACTGTT Rev: CTGACTCTCTCCTGCCTGTGA	
<i>Fos</i>	NM_010234.3	Fwd: GTTCCGGCATCATCTAGGC Rev: GACGTGTAAGTAGTGCAGCC	
<i>Gapdh</i>	NM_008084.2	Fwd: AGGTCGGTGTGAACGGATTTG Rev: TGTAGACCATGTAGTTGAGGTCA	(Maung, Hoefler et al. 2014)
<i>Grm5</i>	NM_001081414.2	Fwd: CTTGCCTGCTTCTCAGTTGTC Rev: CTCAGGAAGCACCCTAGGAC	
<i>Homer1</i>	NM_147176.4	Fwd: AGGGTTCTCAGCACCAATGG Rev: AGCTGCCGGGAGGTGTAAG	
<i>iNOS</i>	NM_010927.3	Fwd: TGACGCTCGGAACTGTAGCAC Rev: TGATGGCCGACCTGATGTT	(Crain, Nikodemova et al. 2013)
<i>Lcn2</i>	NM_008491.1	Fwd: CCAGTTCGCCATGGTATTTT Rev: GGTGGGGACAGAGAAGATGA	
<i>Mrc1</i>	NM_008625.2	Fwd: TGGCTACCAGGAAGTCCATC Rev: TGTAGCAGTGGCCTGCATAG	(Jang, Lee et al. 2013)
<i>Tnfa</i>	NM_013693.2	Fwd: CATCTTCTCAAATTCGAGTGACAA Rev: ACTTGGGCAGATTGACCTCAG	(Jang, Lee et al. 2013)

Supplementary Table 1. List of primers used for qRT-PCR.

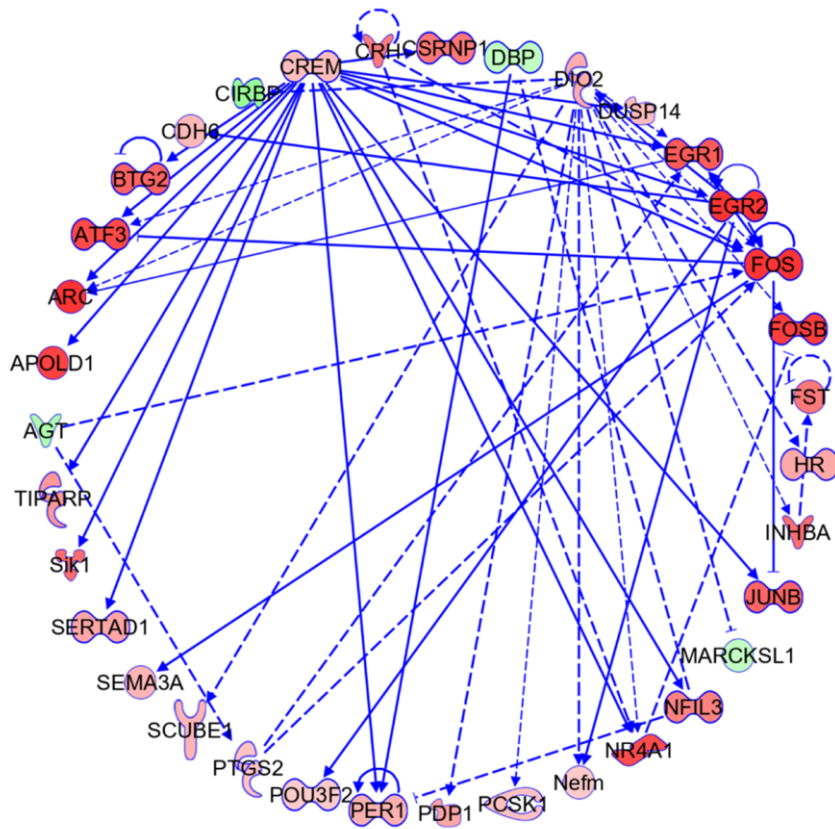


Figure S1. Top upregulated networks predicted by IPA. The top functions associated with the network of hDEGs in early prion disease were 'Neurological Disease' and 'Lipid Metabolism'. Color gradient is representative of fold change with highest up- or down-regulated genes having the darkest color. Red = genes significantly upregulated, and green = genes significantly downregulated.

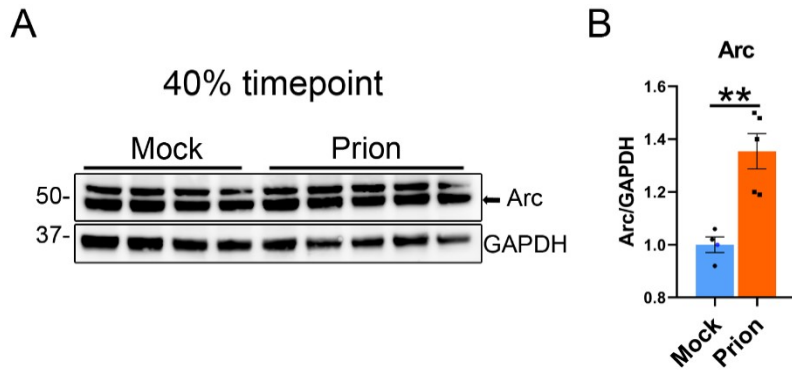


Figure S2. Arc is significantly increased in the hippocampus of 22L-infected mice versus mock-inoculated mice at the 40% timepoint (third male cohort). (A-B) Immunoblot of Arc at the 40% timepoint compared to age-matched, mock-inoculated controls. $n = 4 - 5$ male mice per group. Welch's t -test, $*P \leq 0.05$; $**P \leq 0.01$.

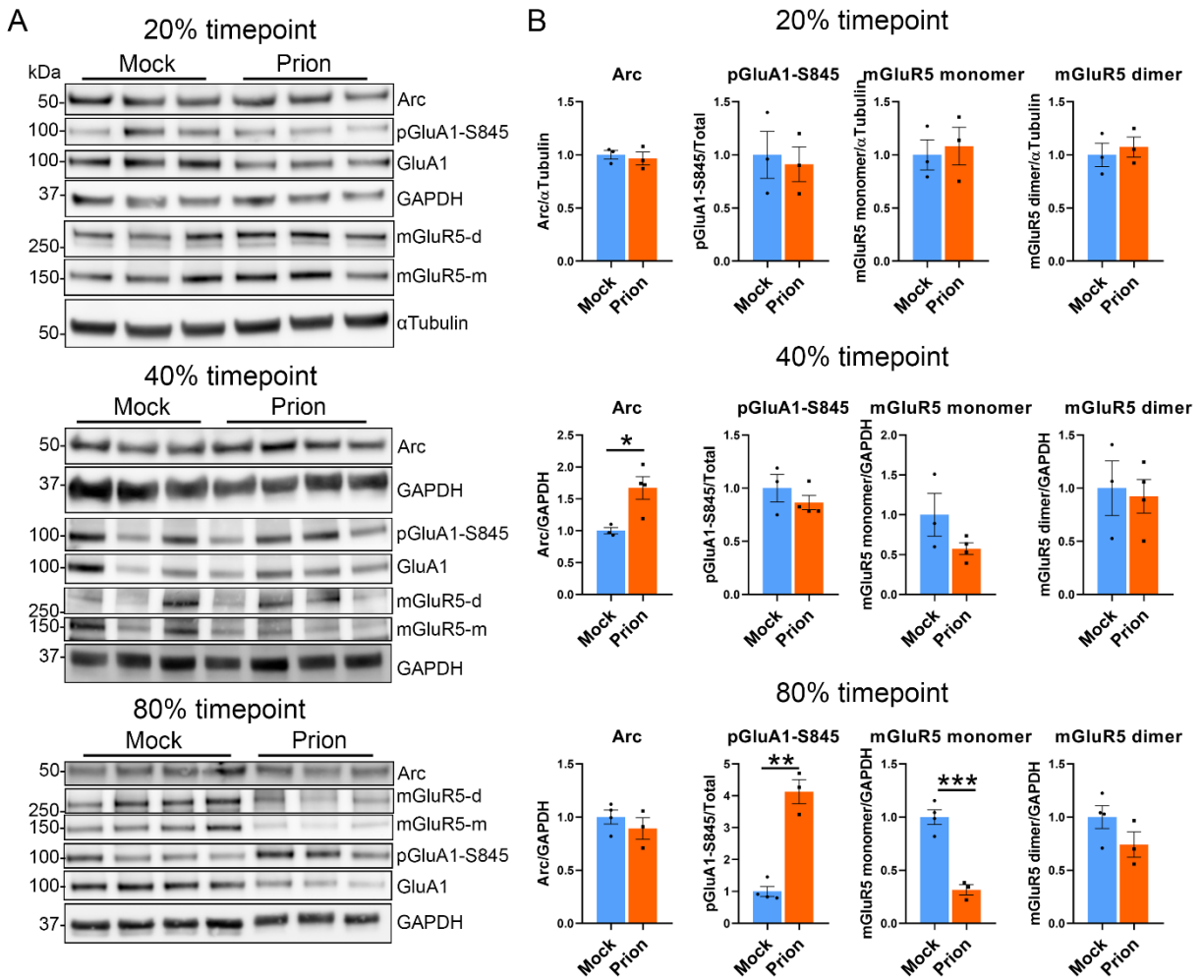


Figure S3. Comparison of Arc, mGluR5, and pGluA1-S845 in the hippocampus of mock- and 22L-inoculated mice collected at the 20%, 40%, and 80% timepoints. (A) Immunoblots of Arc, pGluA1-S845, and mGluR5 at the 20%, 40%, and 80% timepoints in prion-infected mice compared to age-matched, mock-inoculated controls. **(B)** Quantification. $n = 3 - 4$ male mice per group. Welch's t -test, $*P \leq 0.05$; $**P \leq 0.01$.

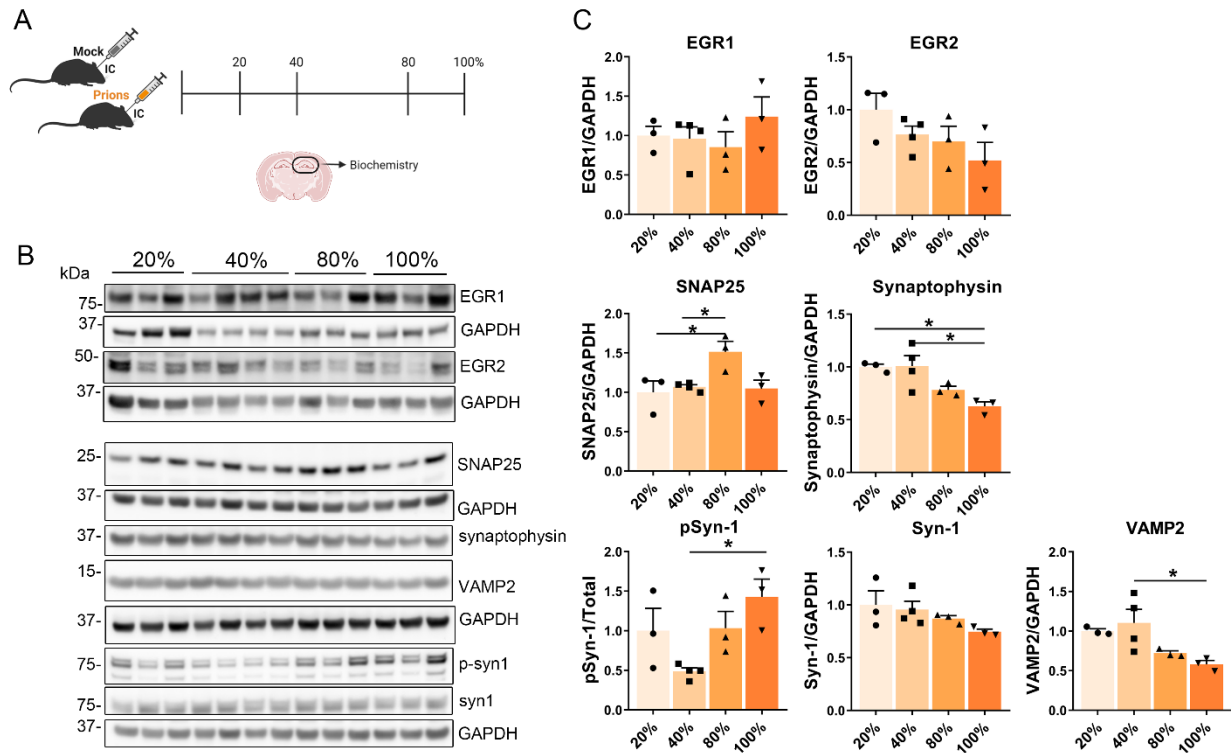


Figure S4. Additional IEGs and synaptic proteins evaluated in a longitudinal study of the prion-infected hippocampus *in vivo*. (A) A schematic summary shows the four timepoints used in this study. (B, C) Western blot and quantification of IEGs and pre-synaptic proteins. $n = 3 - 4$ mice per group. One-way ANOVA and Tukey HSD post-hoc test; $*P \leq 0.05$.

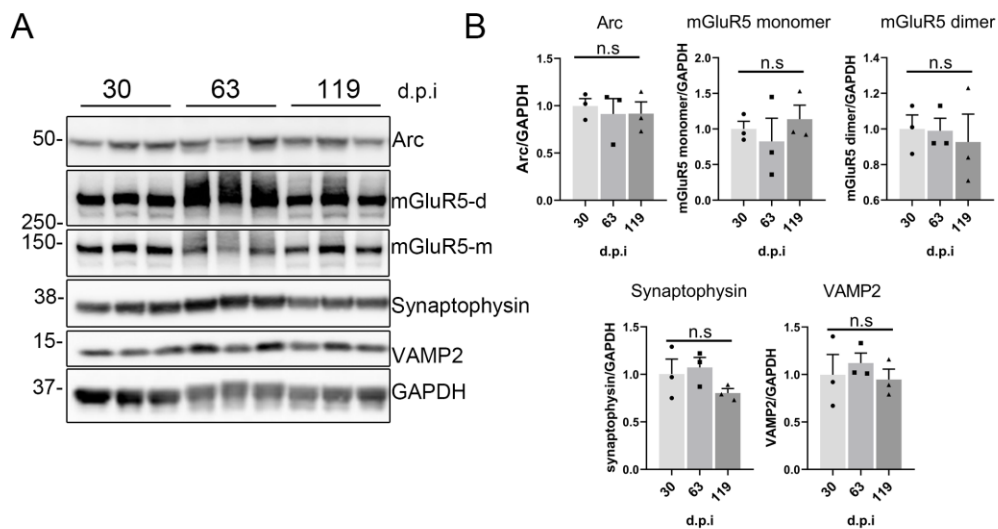


Figure S5. Hippocampal Arc and mGluR5 are not altered following inoculation of WT mice with uninfected brain (mock). (A, B) Western blot and quantification of Arc, mGluR5, VAMP2, and synaptophysin. $n = 3$ mice per group. One-way ANOVA and Tukey HSD post-hoc test; $*P \leq 0.05$.

References

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