

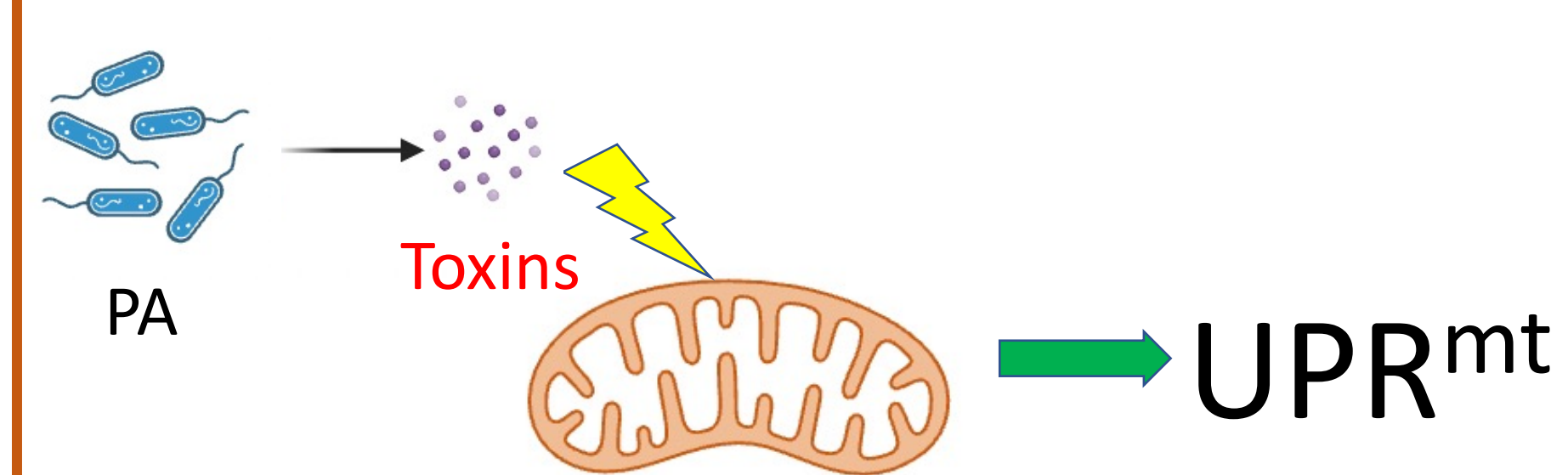


# The HMX/NKX homeodomain transcription factor MLS-2 mediates a cell non-autonomous UPR<sup>mt</sup> that promotes host resistance during infection.

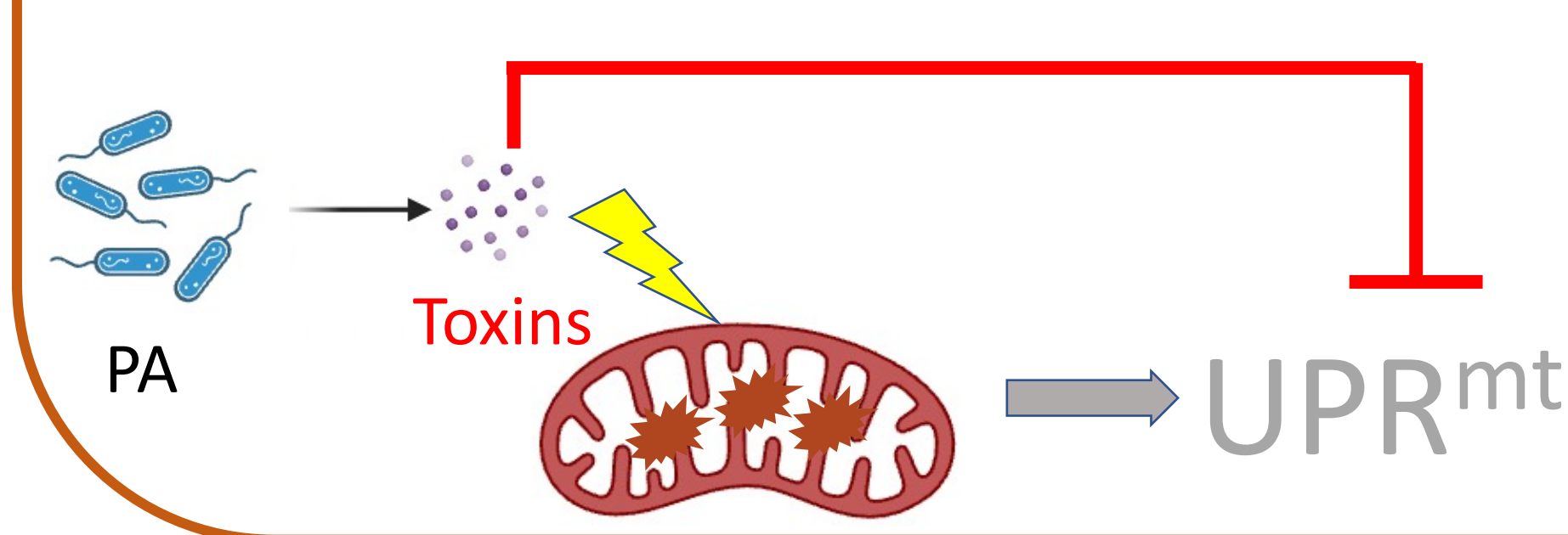
Yiting Xu and Mark W. Pellegrino

## BACKGROUND

24 hrs post-infection:

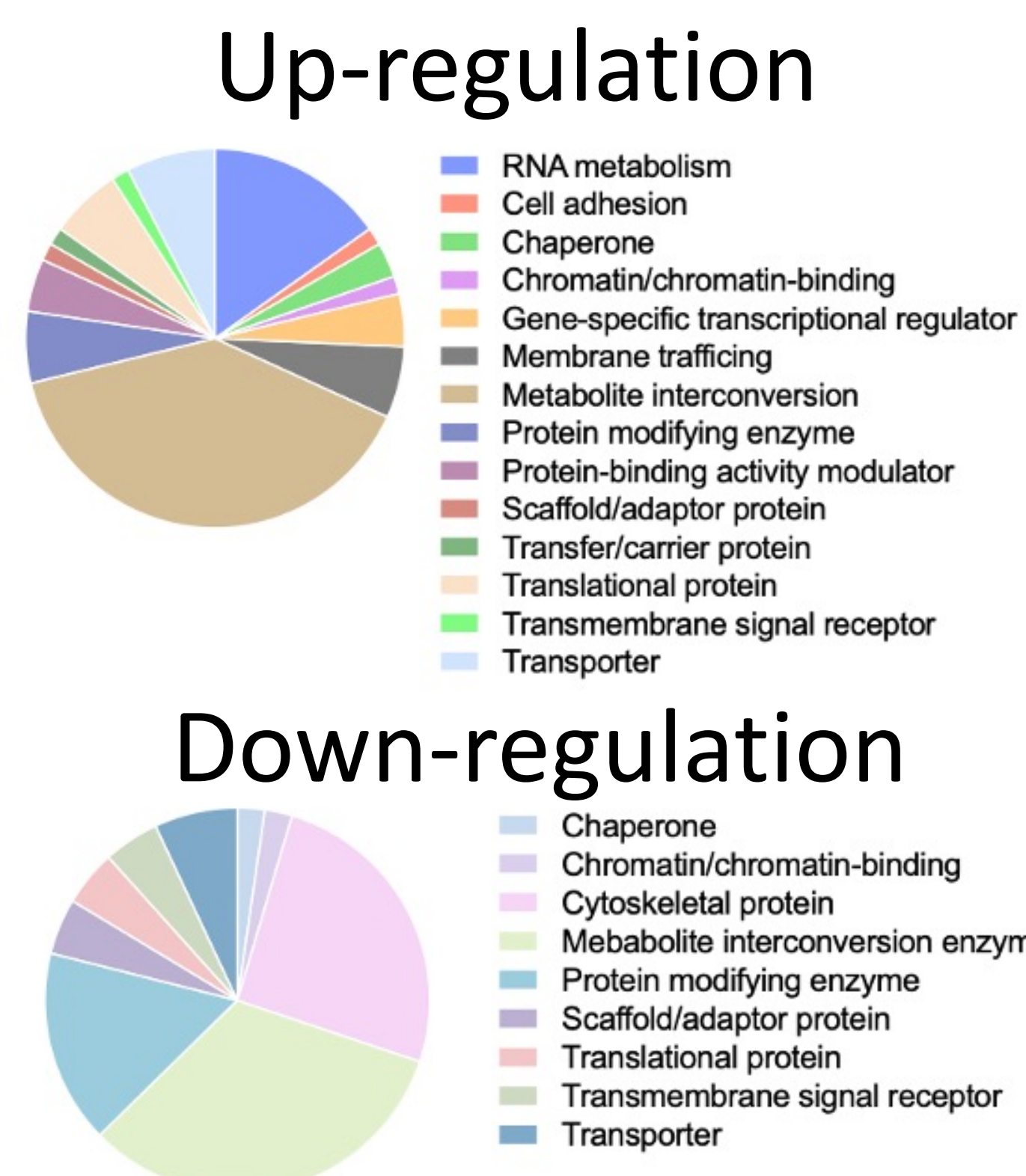
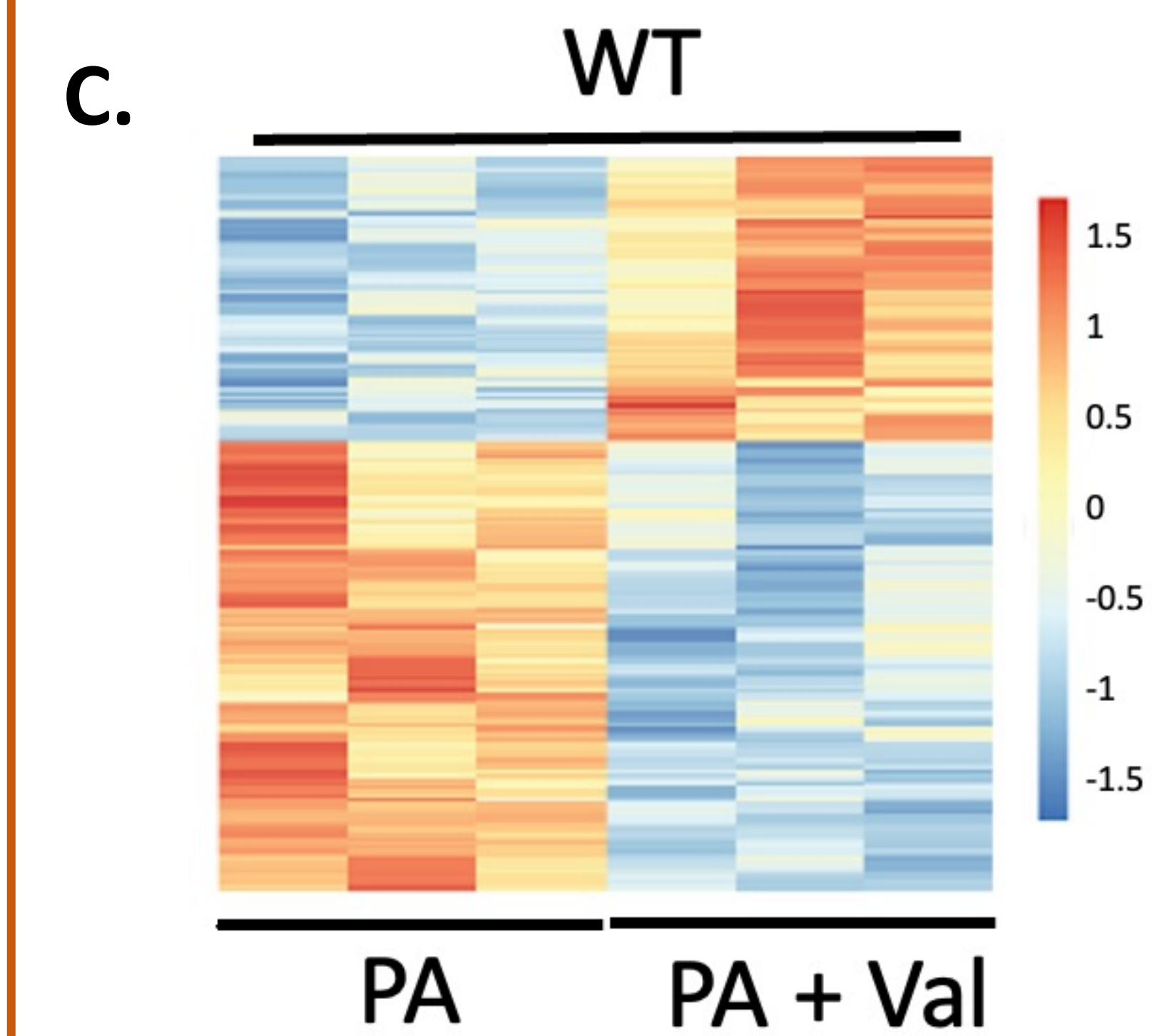
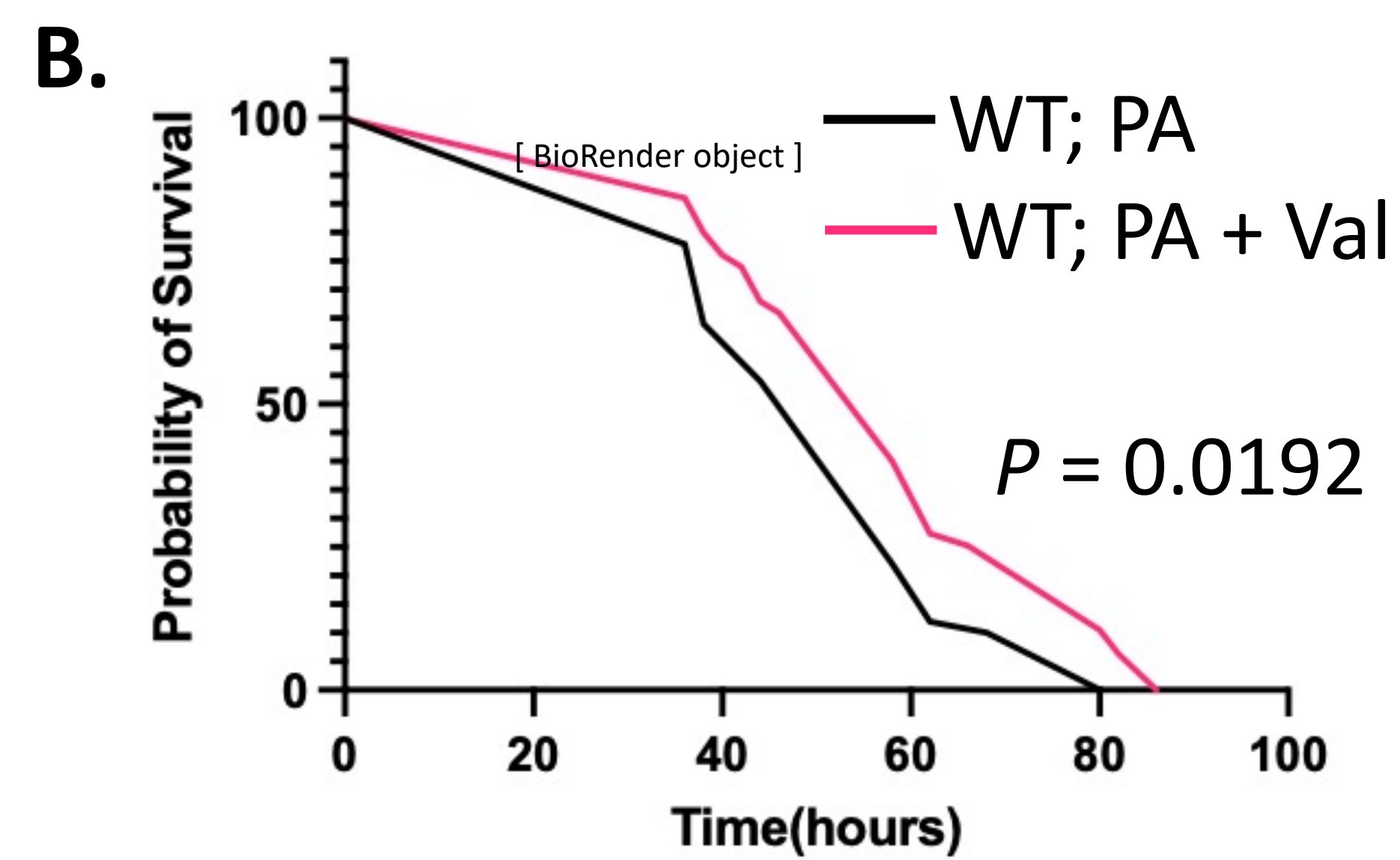
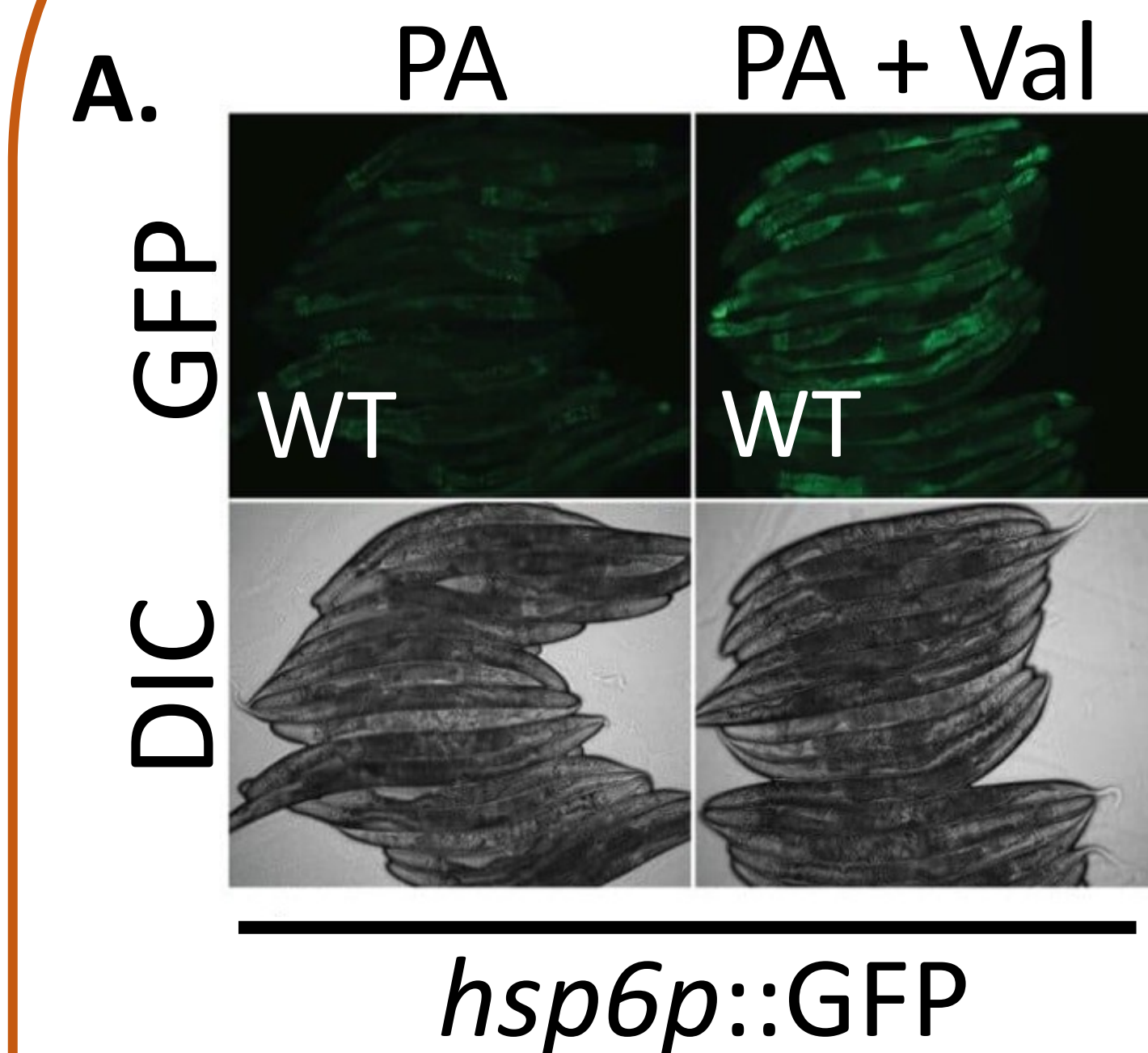


48 hrs post-infection:



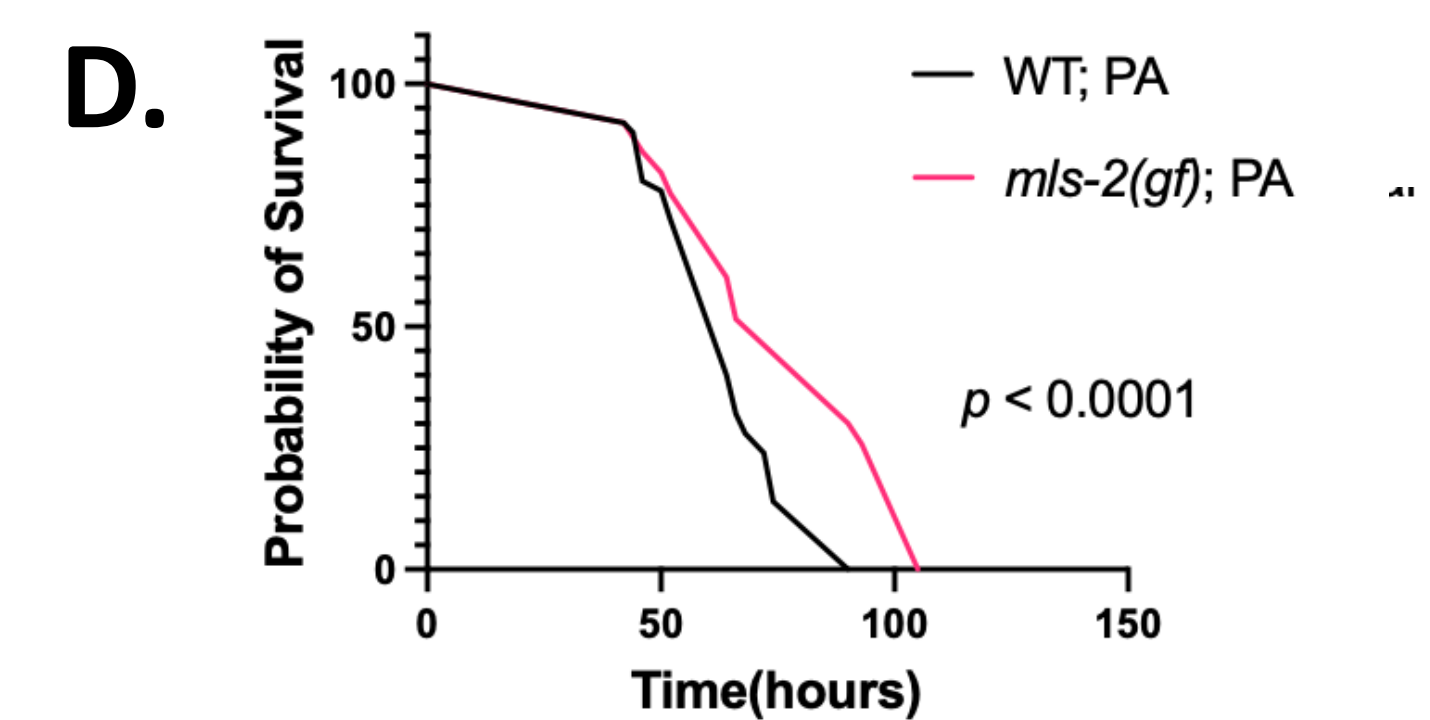
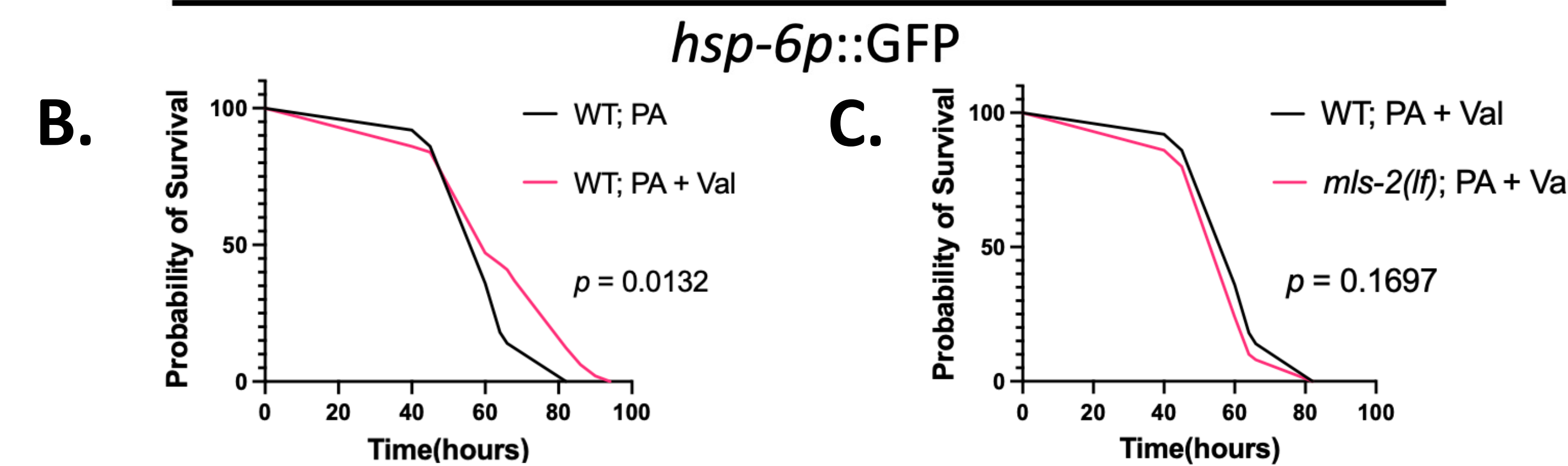
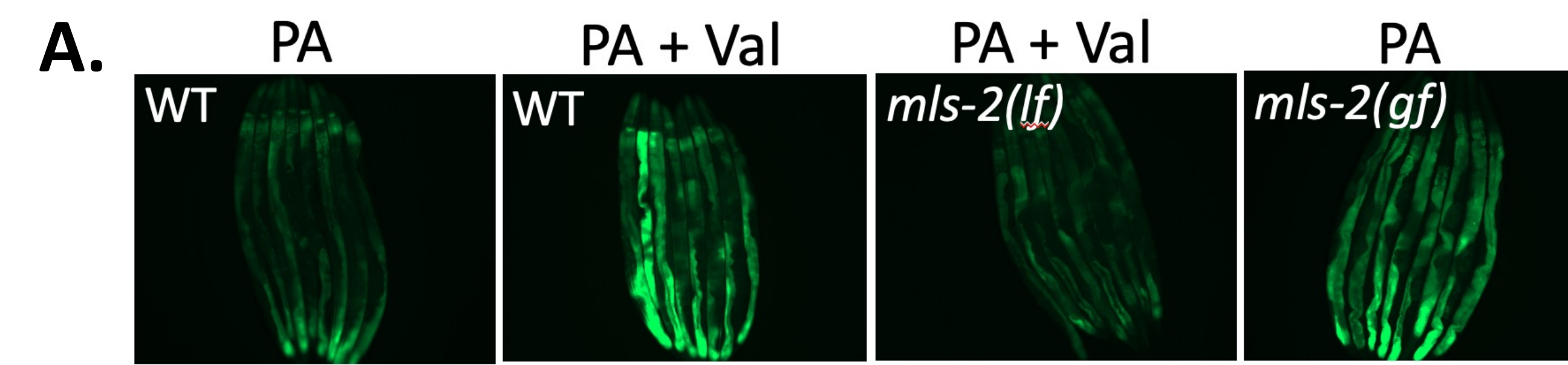
The UPR<sup>mt</sup> is a stress response pathway that transcriptionally reprograms the cell to promote mitochondrial recovery during stress (Qureshi, Haynes, and Pellegrino 2017). The bacterial pathogen *P. aeruginosa* activates the UPR<sup>mt</sup> but only transiently, suggesting a repressive force that is acting on this stress response during infection (Mahmud et al. 2020).

## Valine enhances UPR<sup>mt</sup> during infection



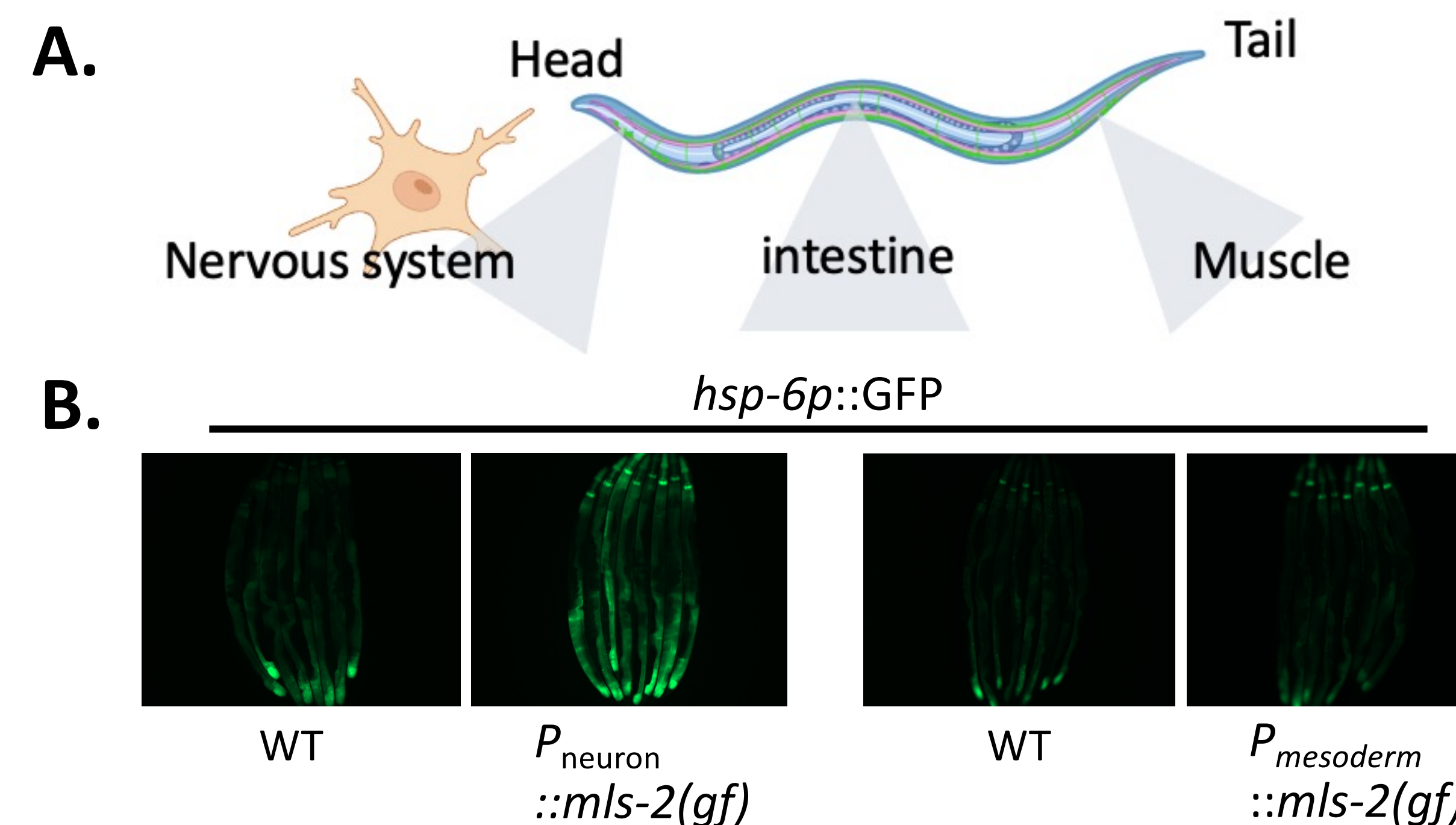
Chronic *P. aeruginosa* infection represses UPR<sup>mt</sup> due to deficiencies in branched-chain amino acids such as valine. A) Supplementation with valine restores UPR<sup>mt</sup> during chronic *P. aeruginosa* infection (visualized by the UPR<sup>mt</sup> reporter transgene *hsp-6p::GFP*), and B) promotes the survival of the *C. elegans* host. C) Valine supplementation transcriptionally reprograms *C. elegans*.

## MLS-2 HMX/NKX transcription factor mediates valine-induced UPR<sup>mt</sup>



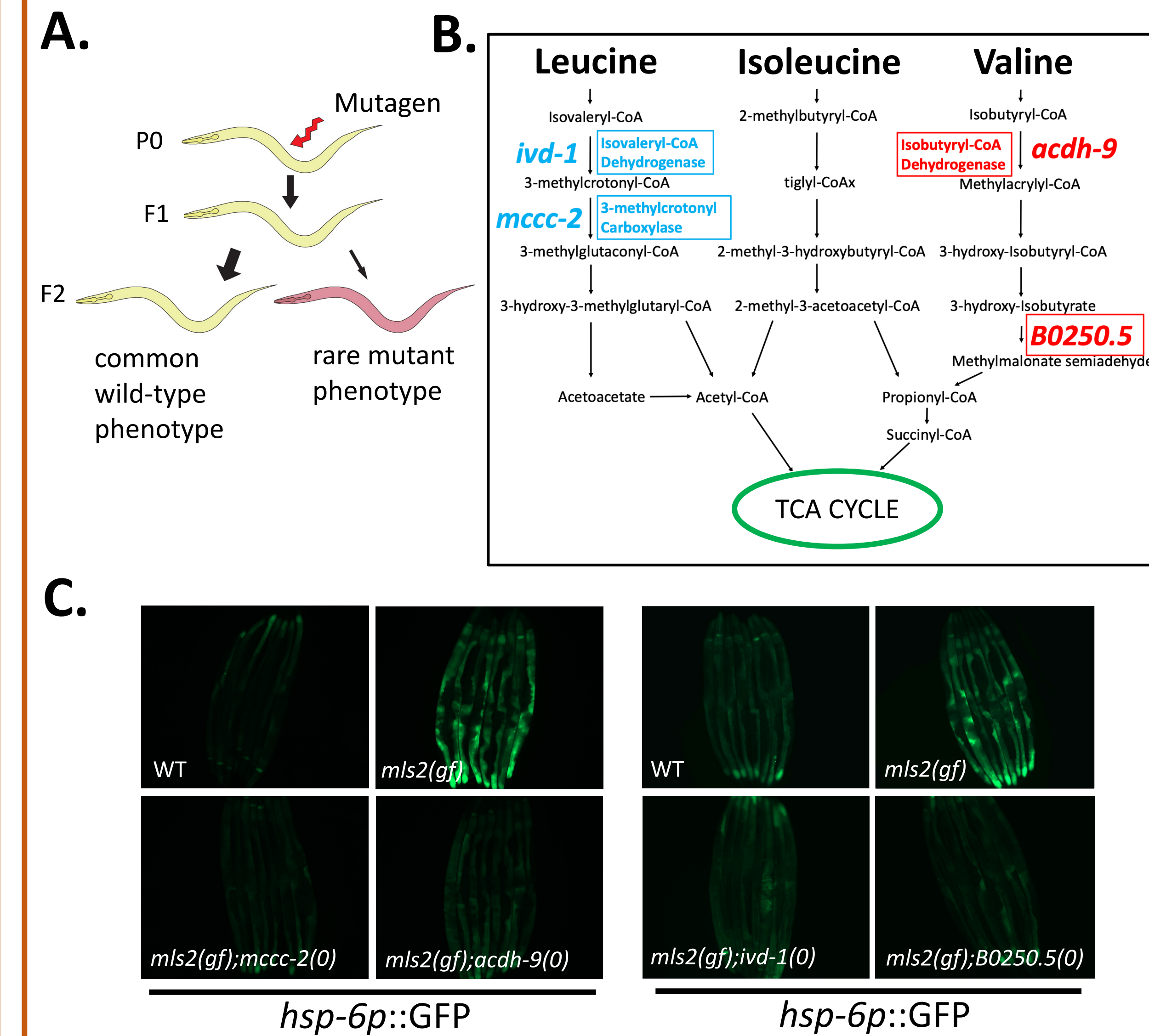
MLS-2 is required for valine-mediated UPR<sup>mt</sup> phenotypes. A) *mls-2* loss of function mutant suppresses UPR<sup>mt</sup> activation by valine during *P. aeruginosa* infection, B, C) as well as the survival benefit of this amino acid. D) *mls-2* gain of function is sufficient to promote host survival during infection.

## MLS-2 regulates UPR<sup>mt</sup> cell non-autonomously



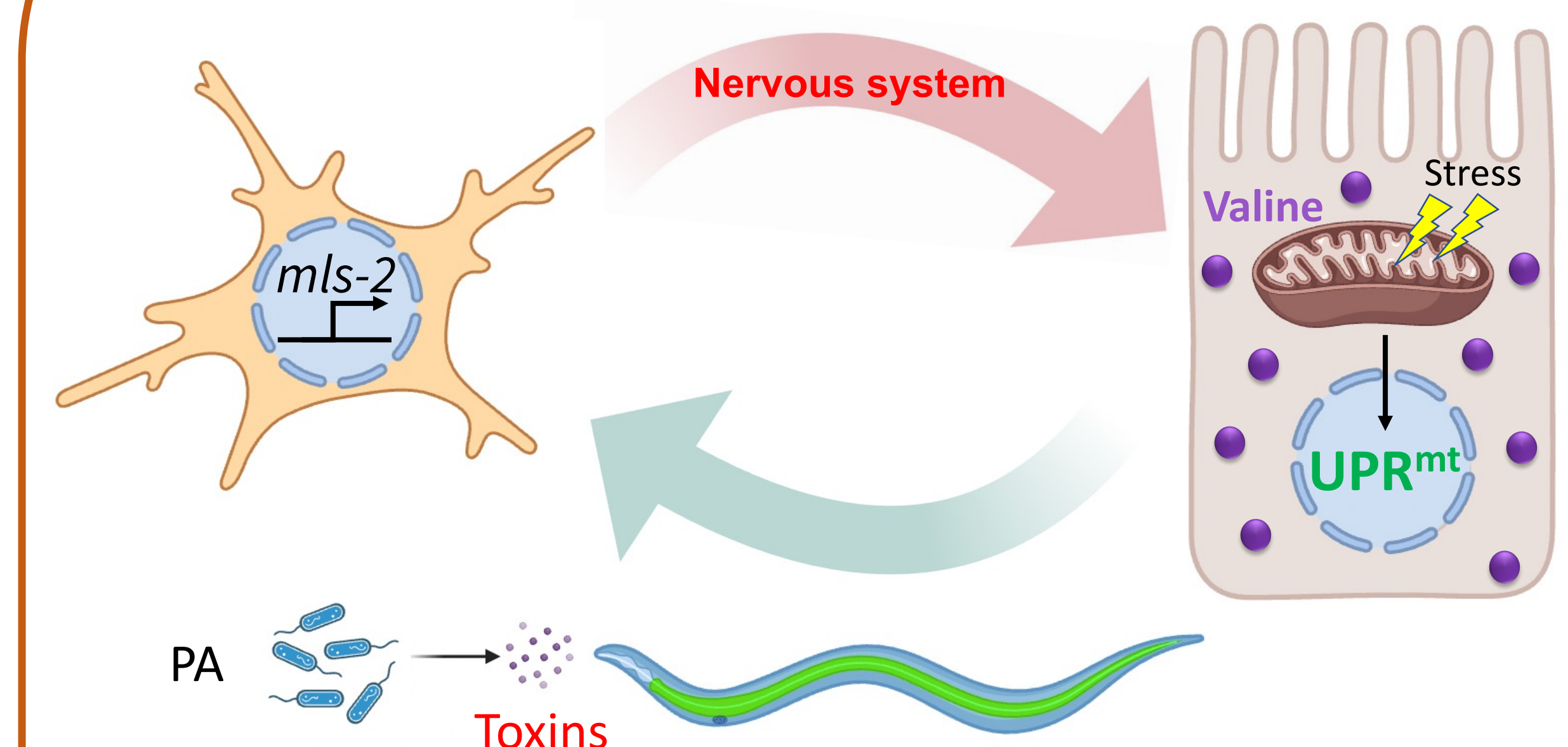
MLS-2 activates UPR<sup>mt</sup> cell non-autonomously. A) Simplified scheme of *C. elegans*. B) Expression of *mls-2(gf)* cDNA in neurons activates UPR<sup>mt</sup> in the intestine.

## MLS-2 regulates UPR<sup>mt</sup> via metabolic rewiring



BCAA catabolism is required for MLS-2-mediated UPR<sup>mt</sup>. A) Forward genetics mutagenesis scheme. B,C) Loss of function mutations in leucine and valine catabolic genes suppress *mls-2(gf)*-UPR<sup>mt</sup>.

## Model



We find that MLS-2 acts cell non-autonomously in the nervous system to regulate the UPR<sup>mt</sup> in the intestine, promoting the survival of the *C. elegans* host. Curiously, our genetic approaches suggest that MLS-2 stimulates the UPR<sup>mt</sup> by modulating energy metabolism.