

Preclinical modeling of chronic inhibition of the Parkinson's disease associated kinase LRRK2 reveals altered function of the endolysosomal system in vivo

Article

Supplemental Material

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These membranes were cut according to the molecular weight markers before blotting so that multiple proteins in different molecular-weight ranges could be probed using different strips of the membrane.

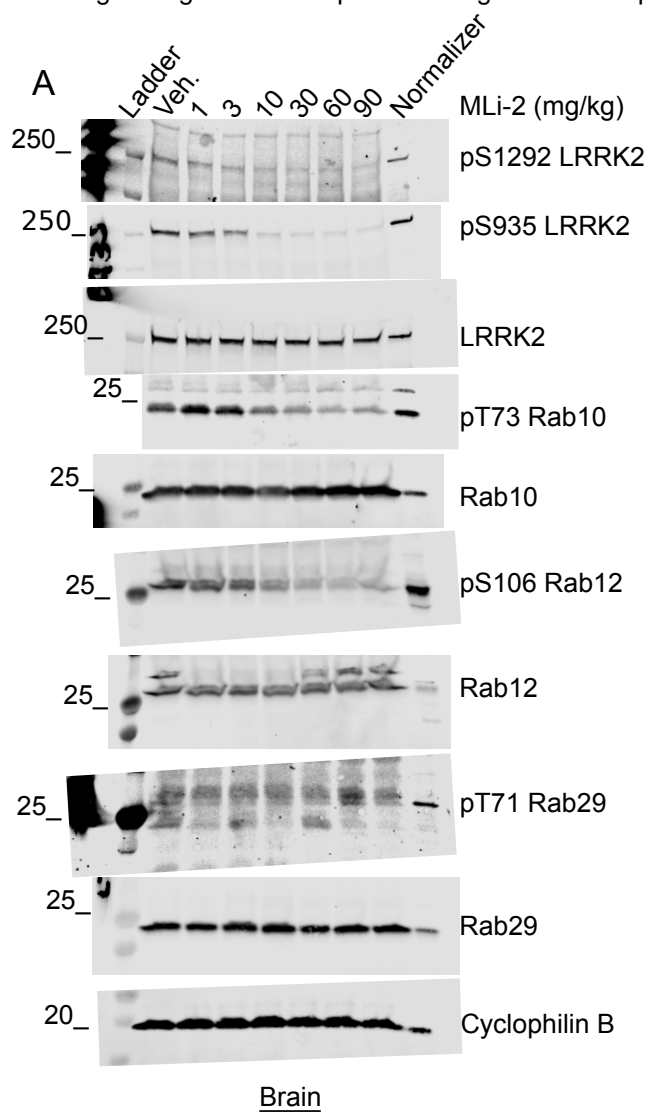


Fig. 1A Brain

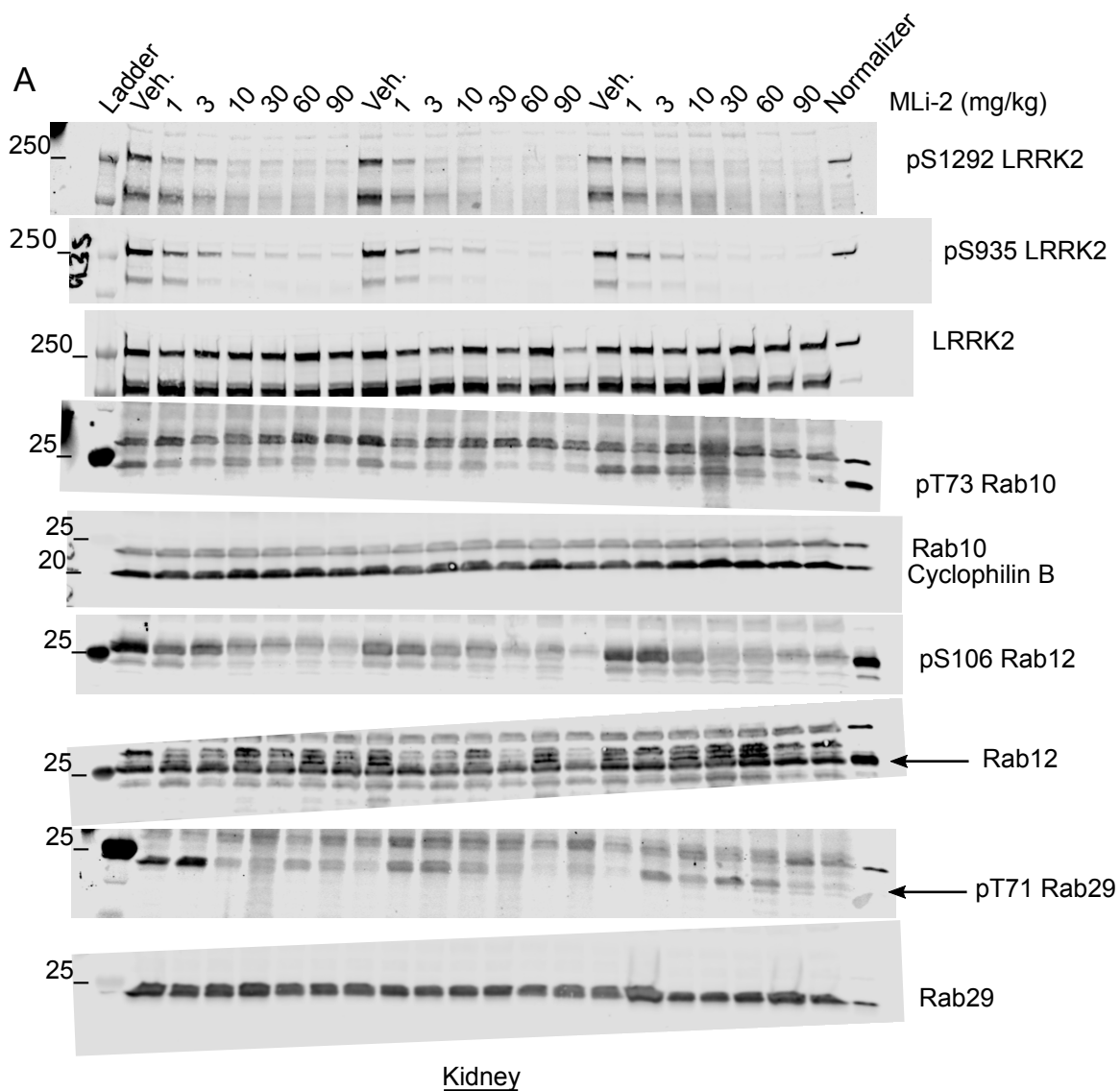


Fig. 1A Kidney

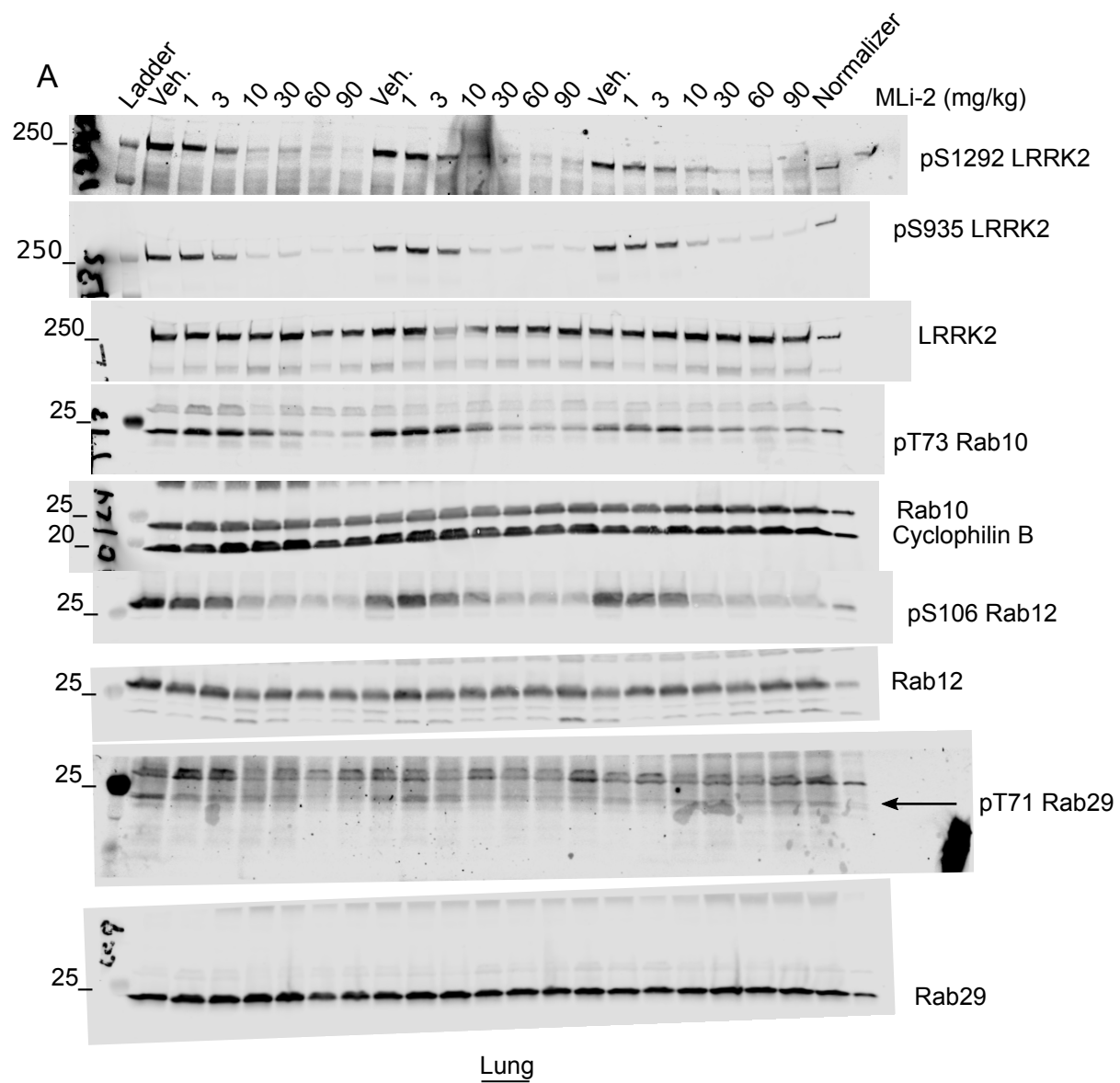


Fig. 1A Lung

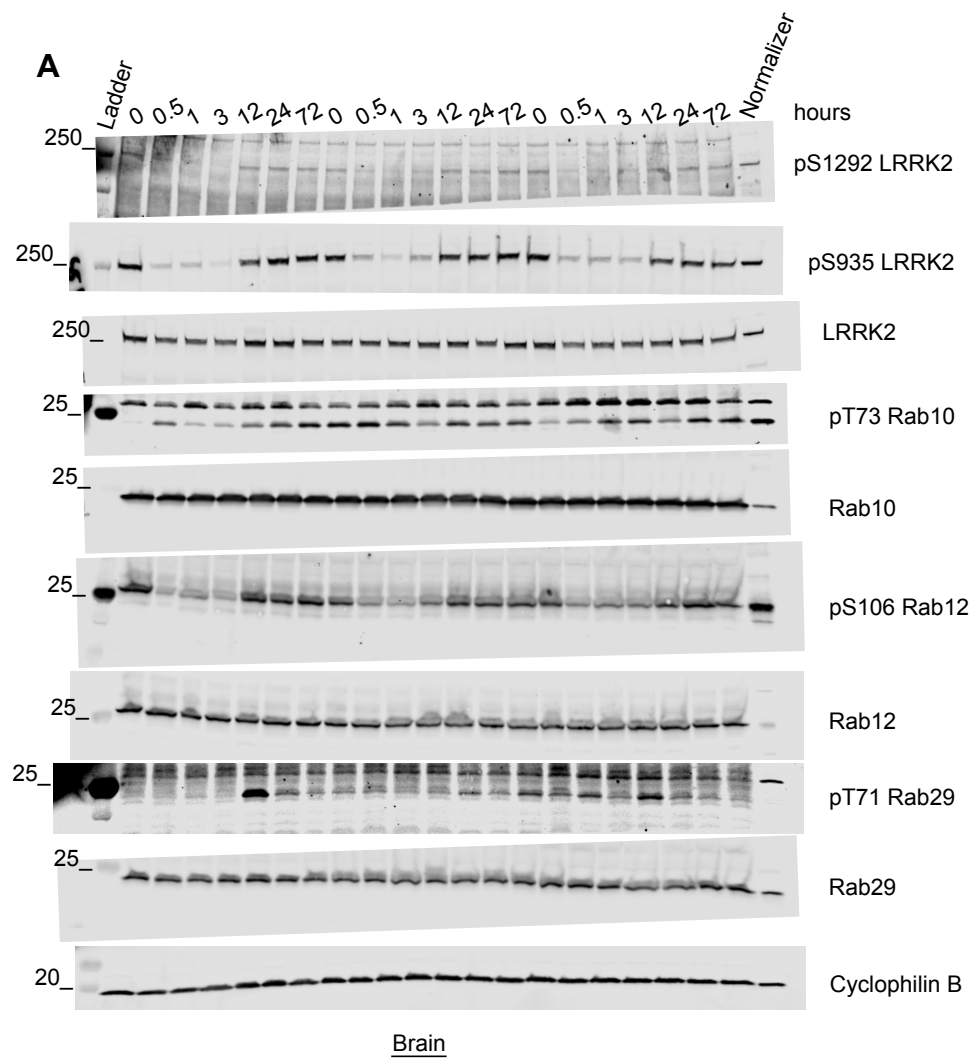


Fig. 2A Brain

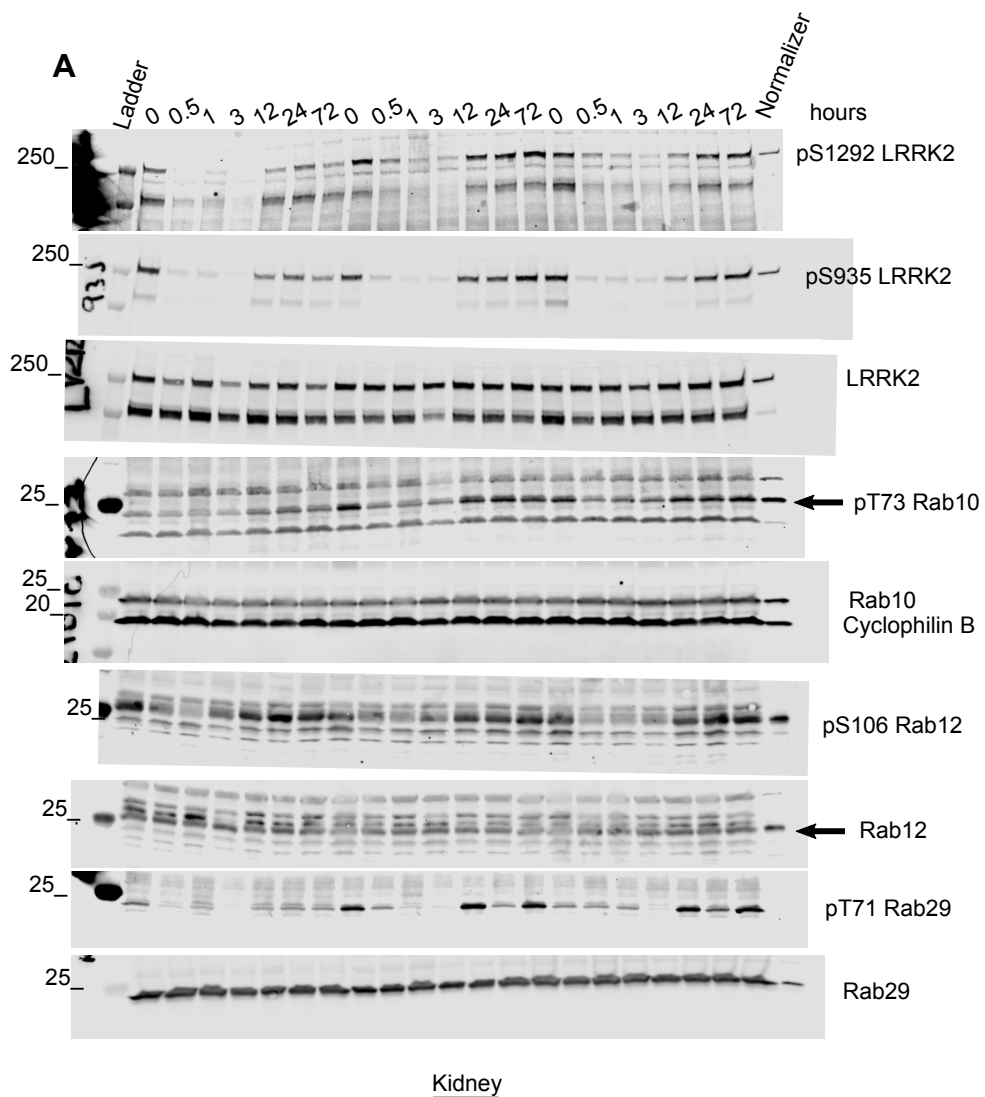


Fig. 2A Kidney

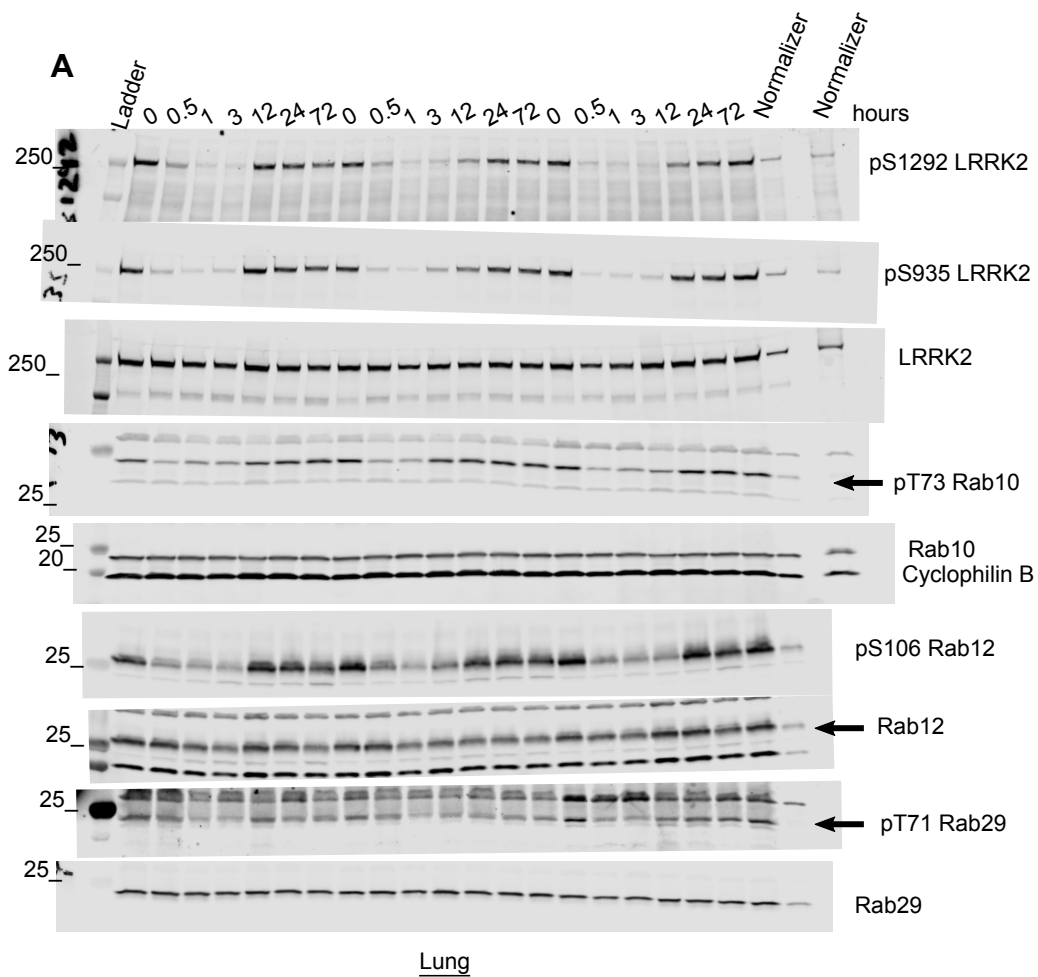


Fig. 2A Lung

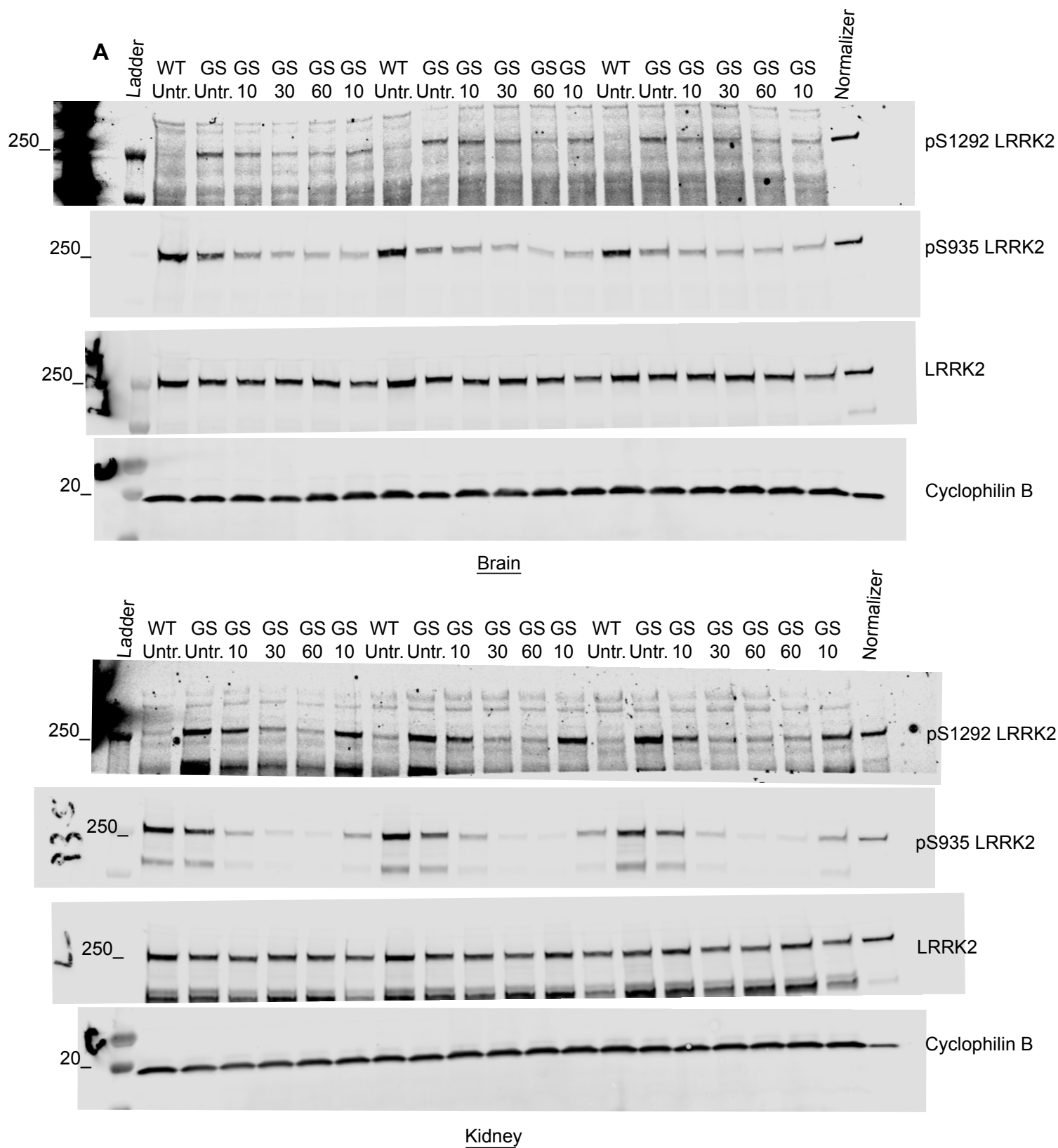


Fig. 3A Brain and Kidney

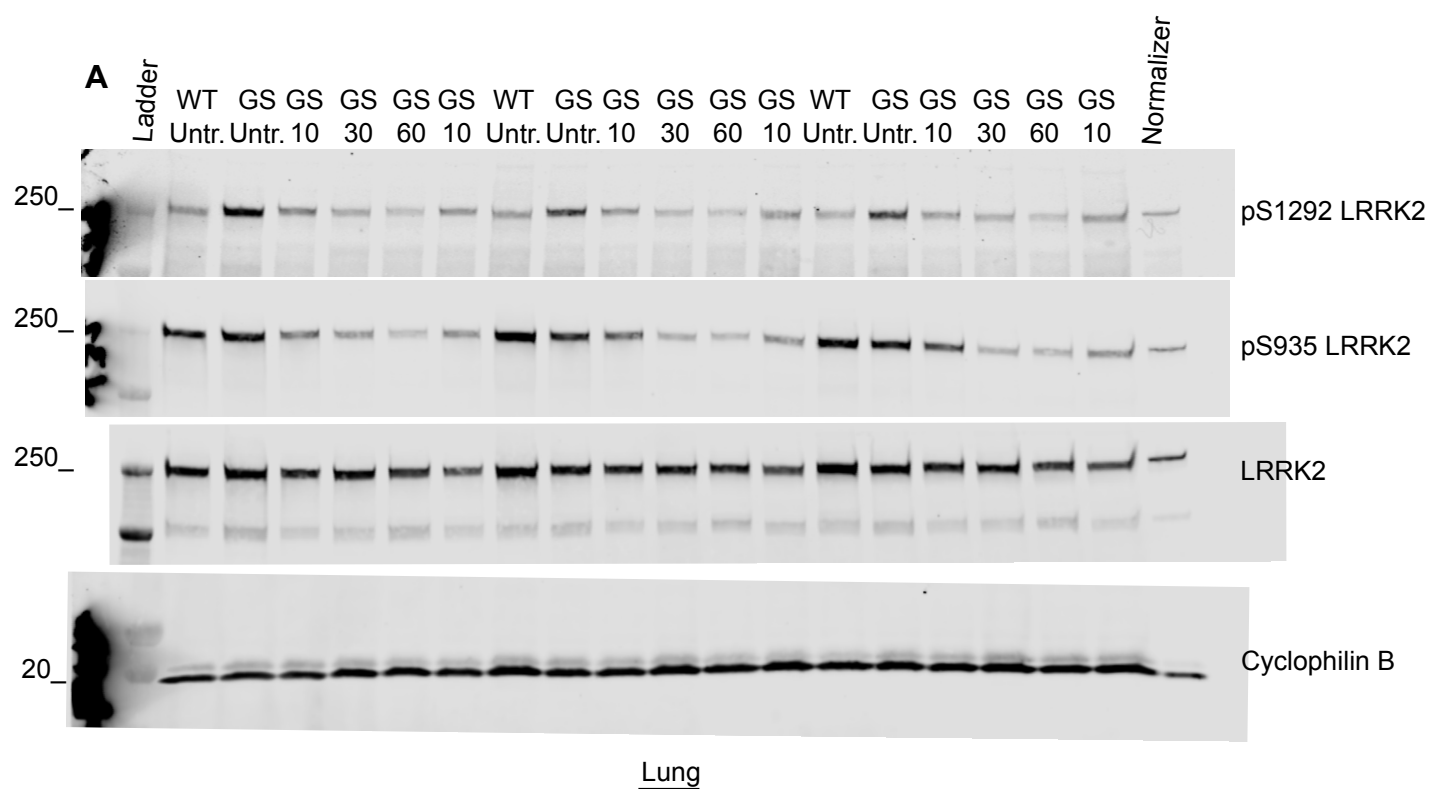


Fig. 3A Lung

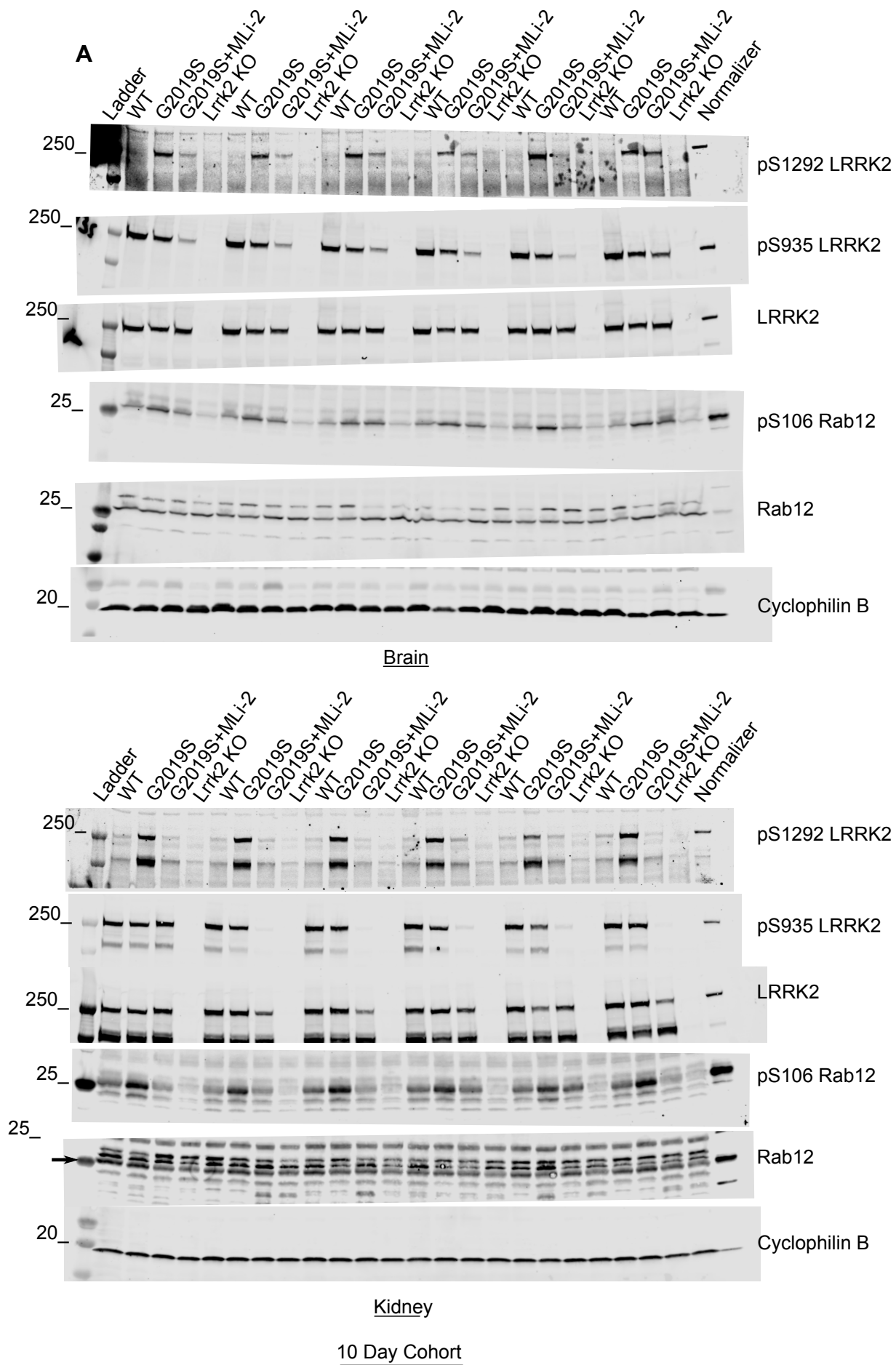


Fig. 5A Brain and Kidney 10 days

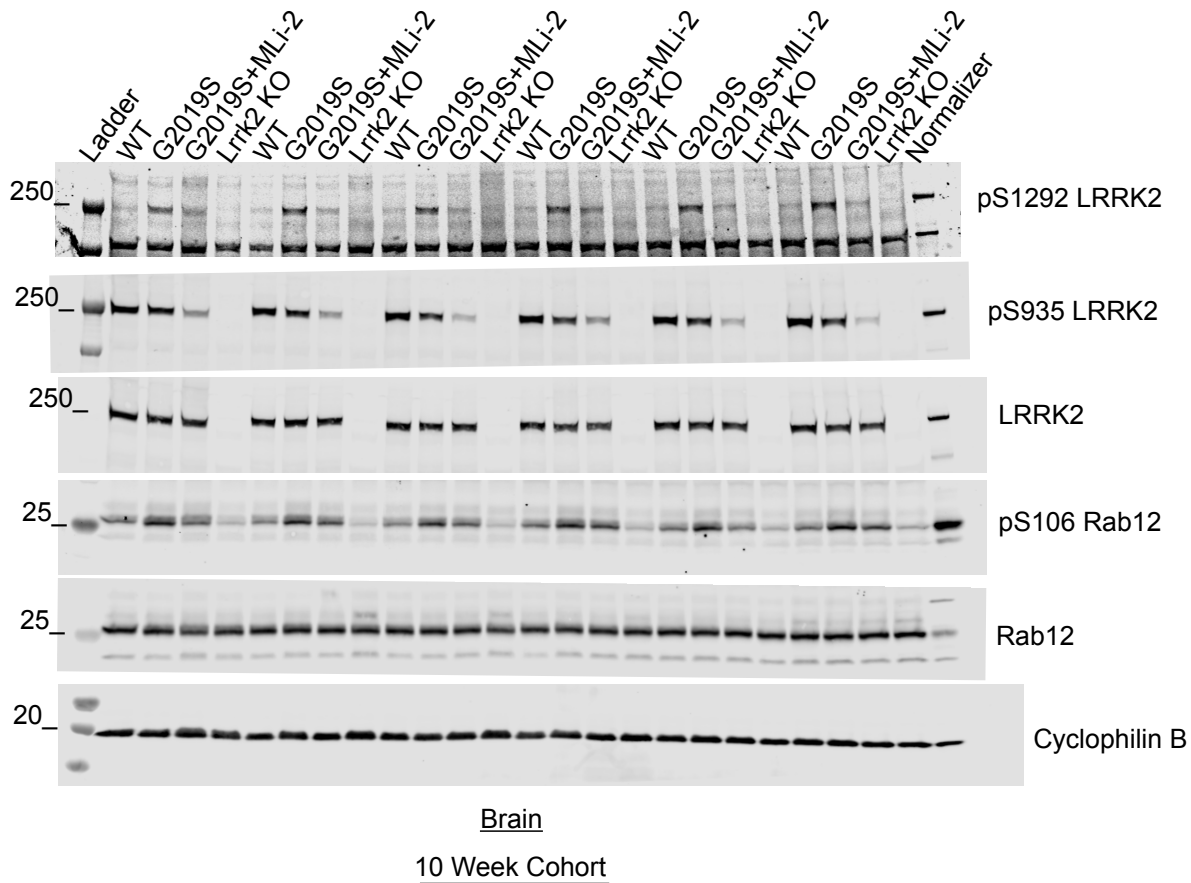
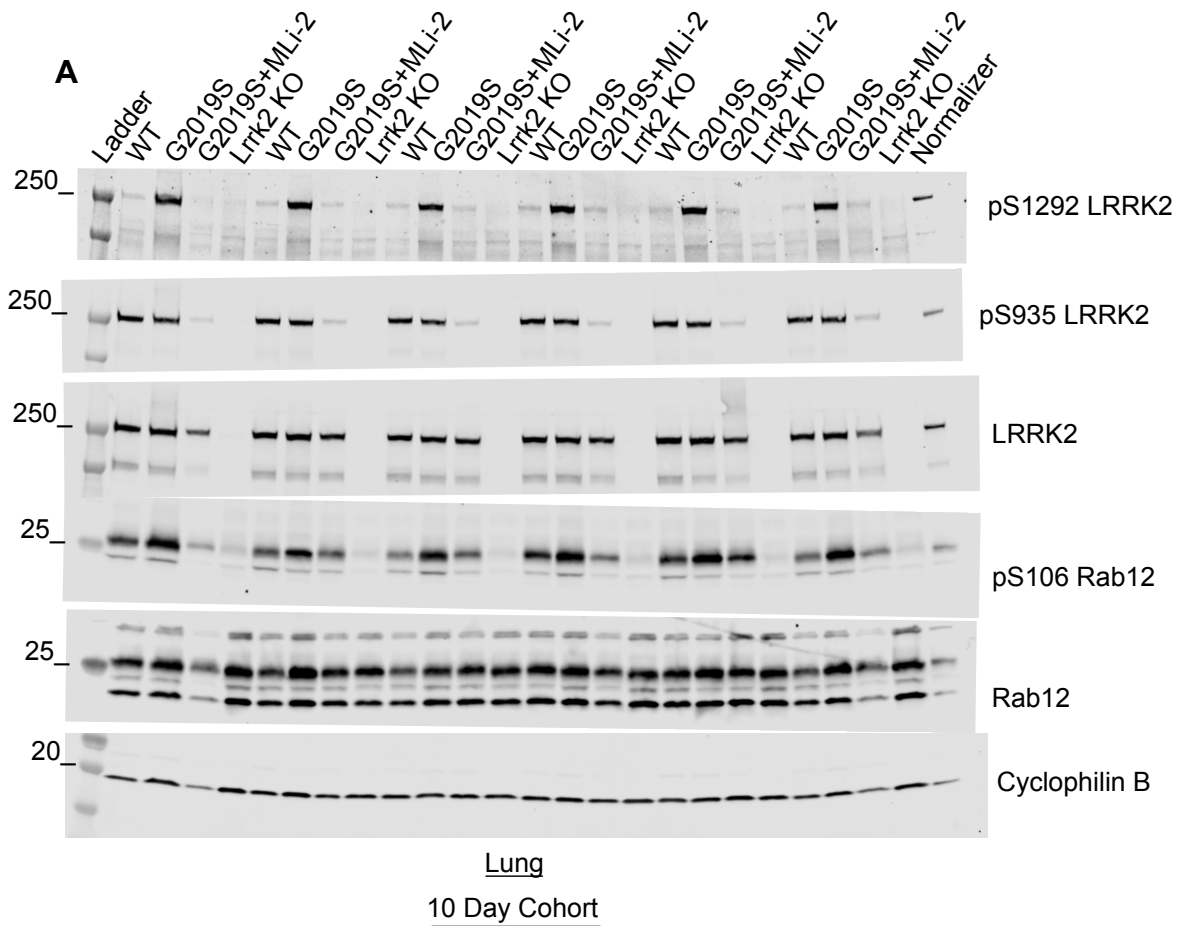


Fig. 5A Lung days and Brain weeks

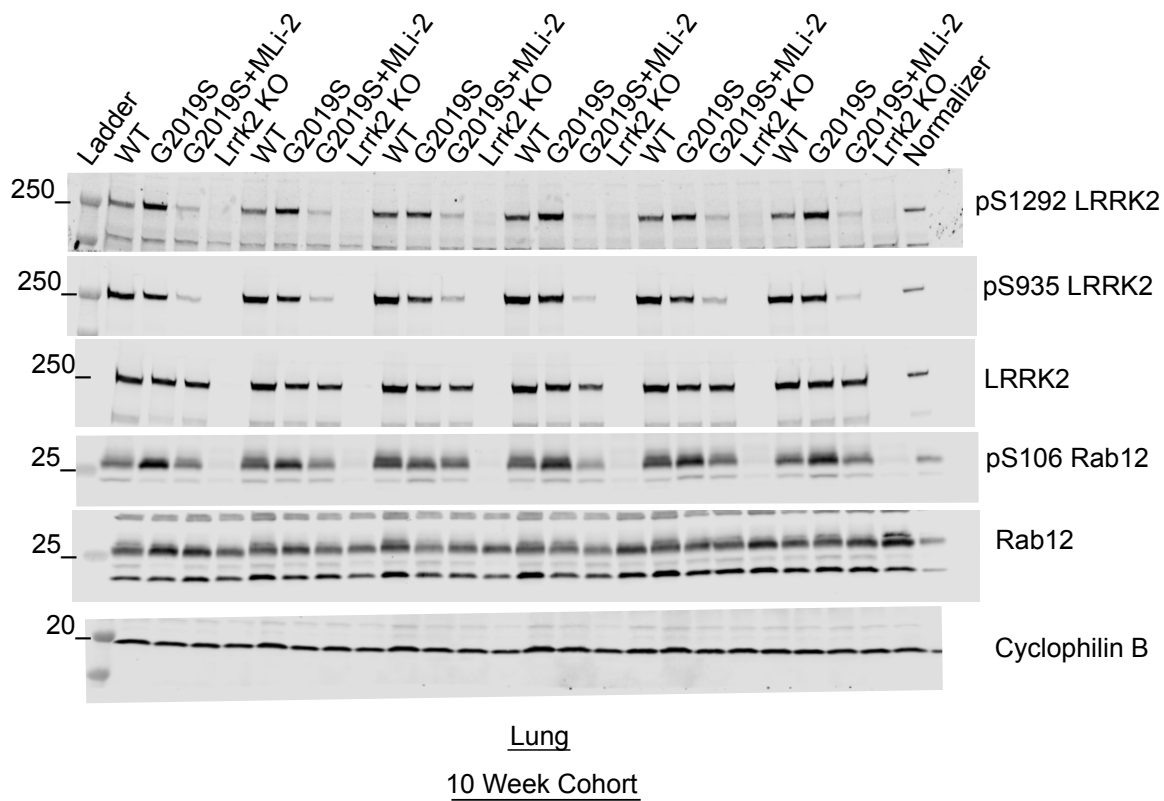
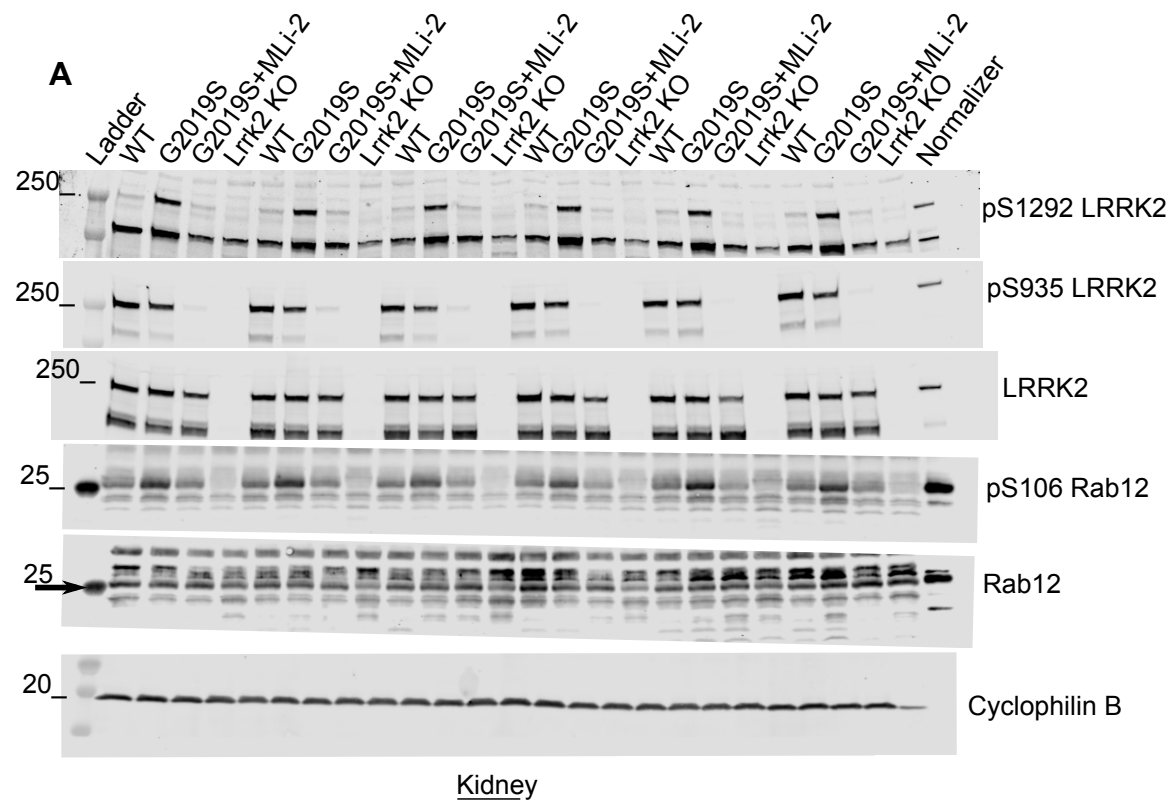


Fig. 5A Kidney and Lung weeks

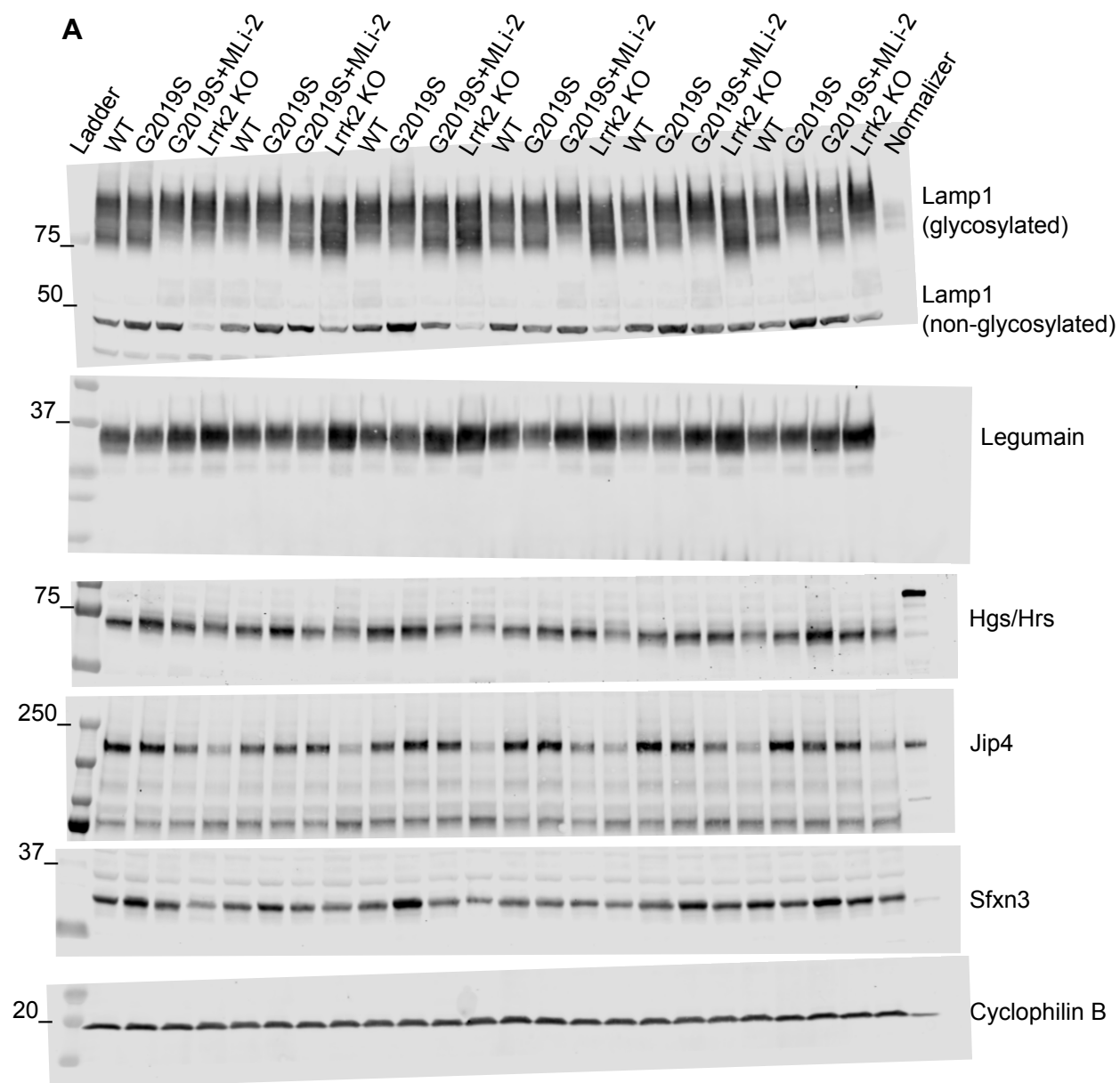


Fig. 8A

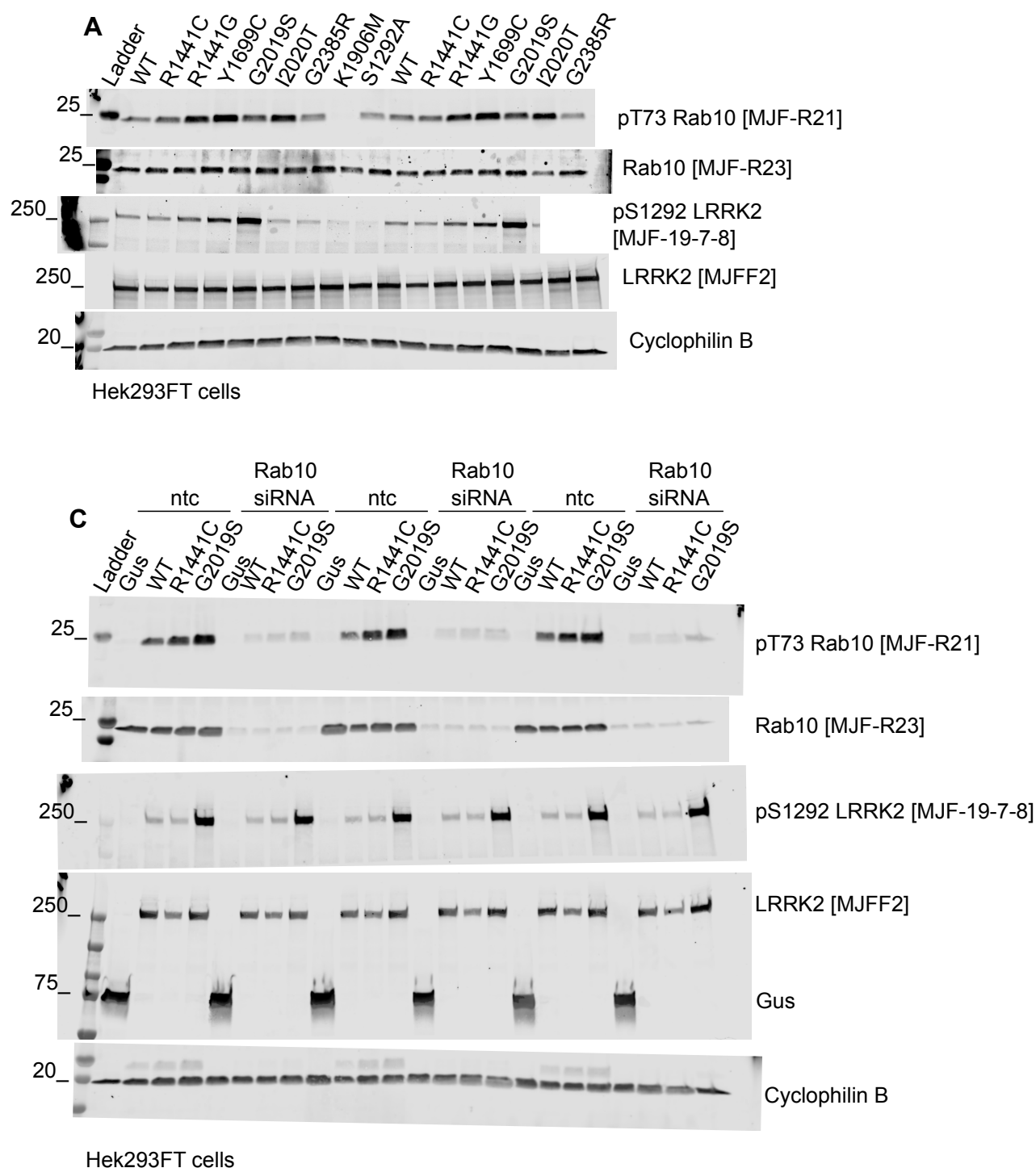


Fig. S2A, C

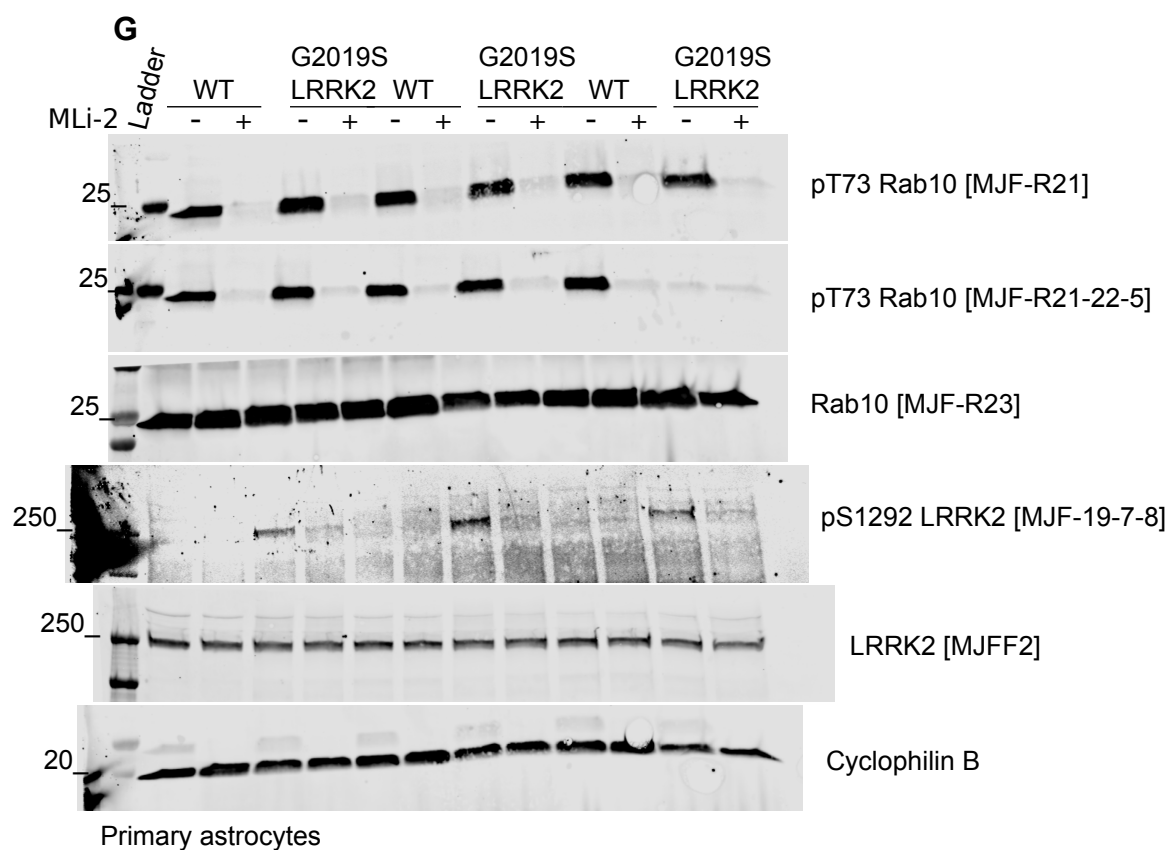
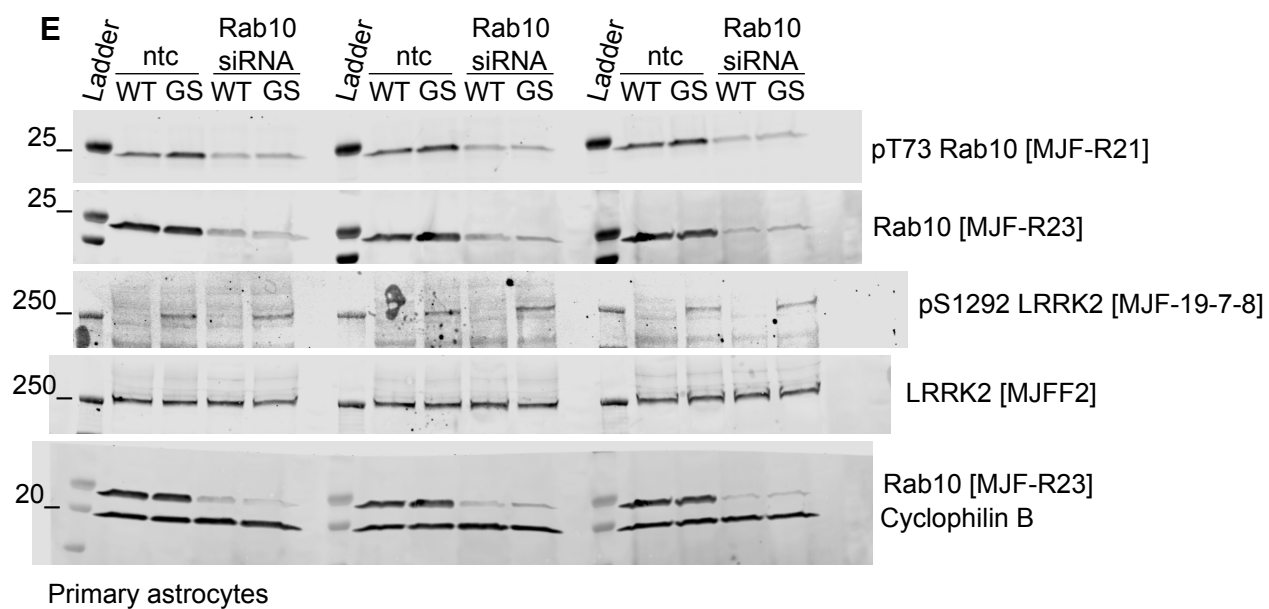


Fig. S2E, G

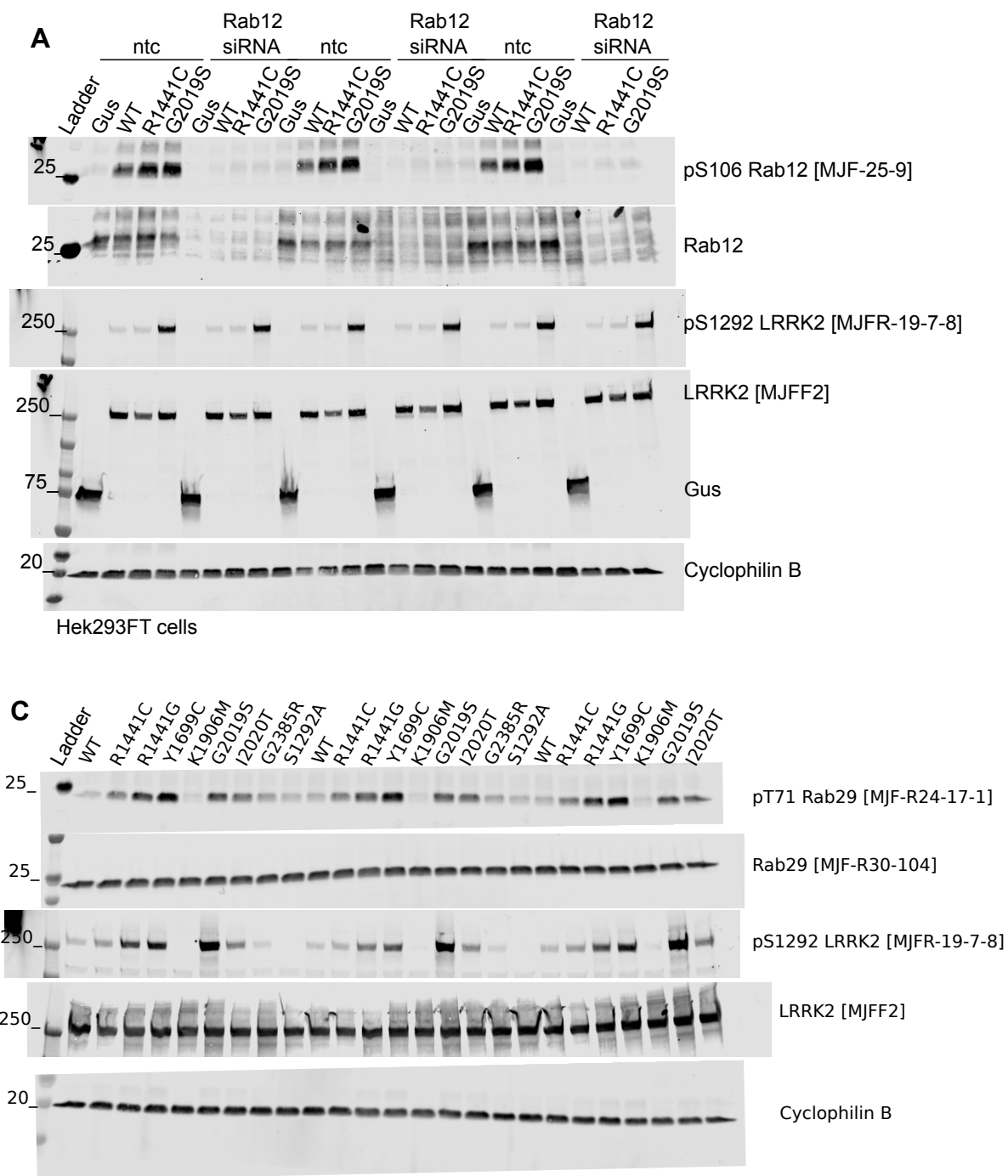


Fig. S3A, C

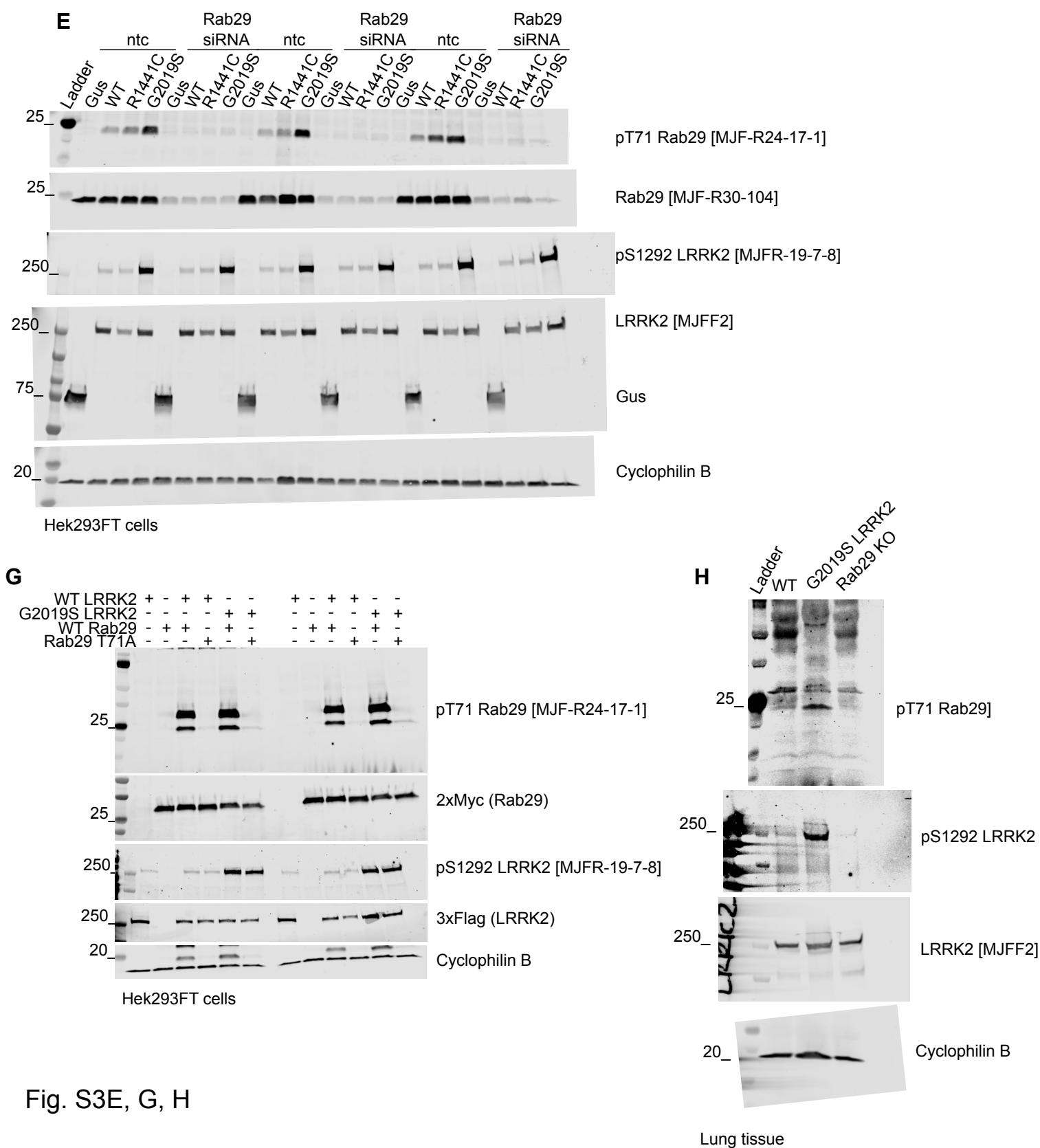


Fig. S3E, G, H

10 Day Cohort

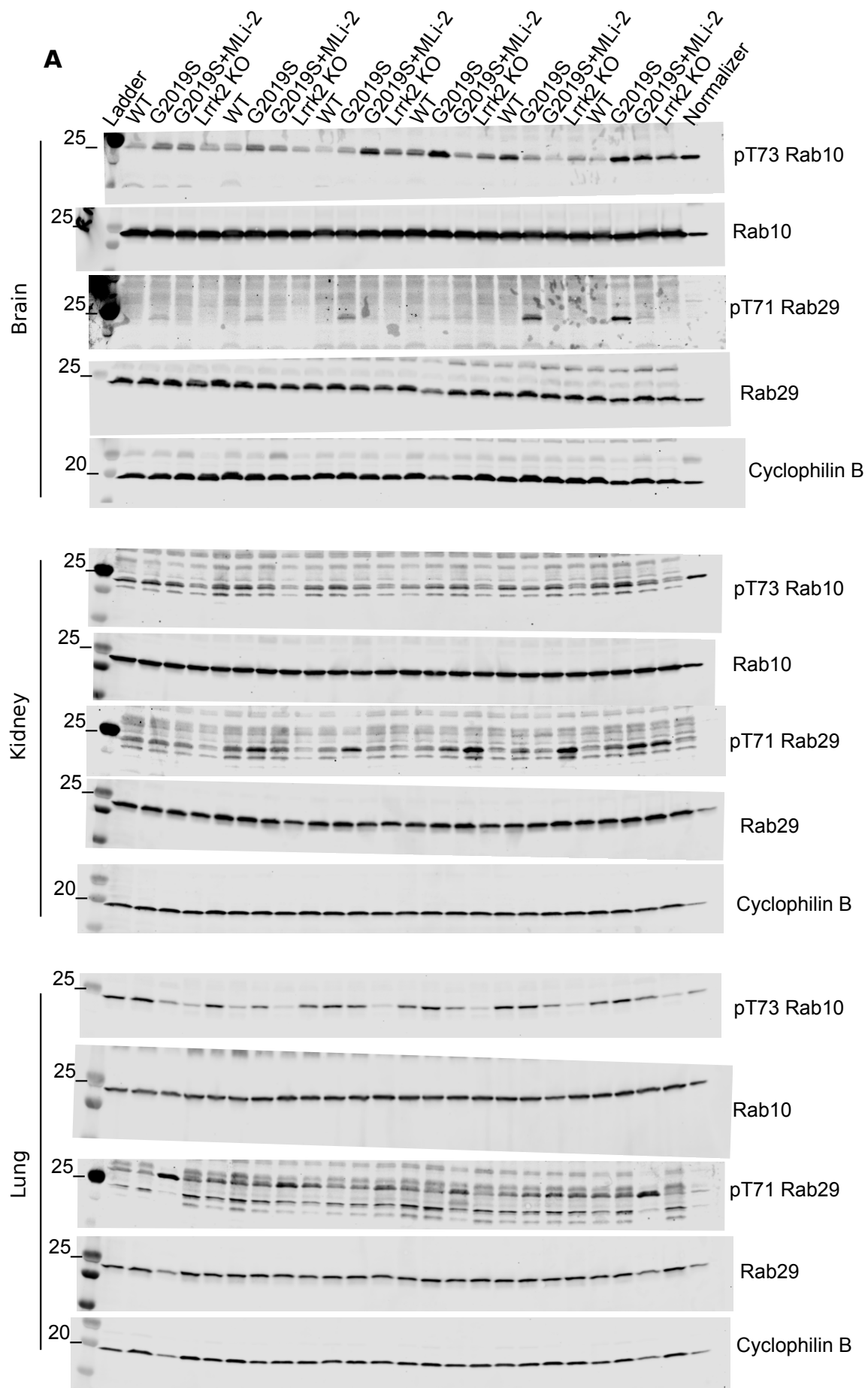


Fig. S4A Days

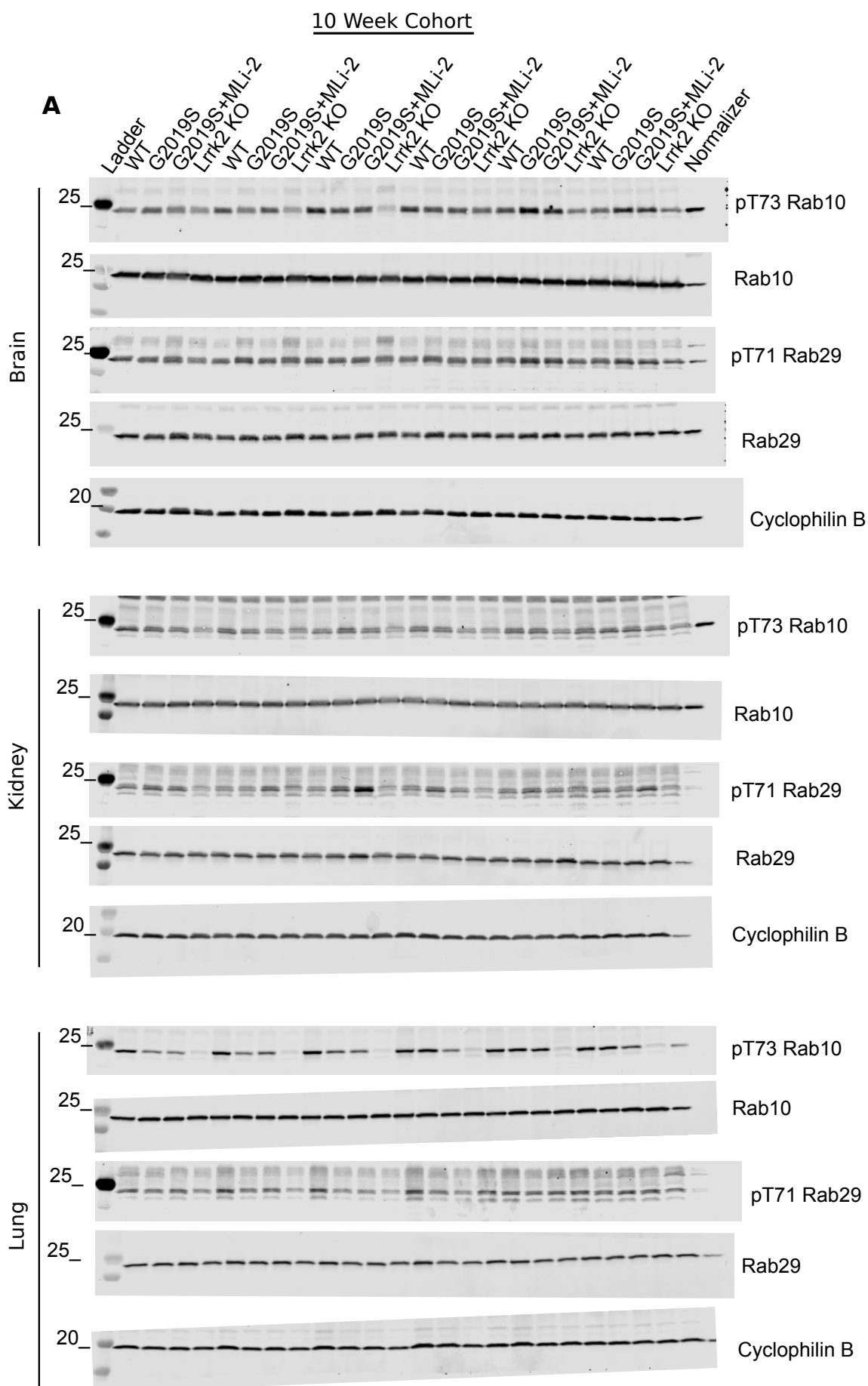


Fig. S4A Weeks

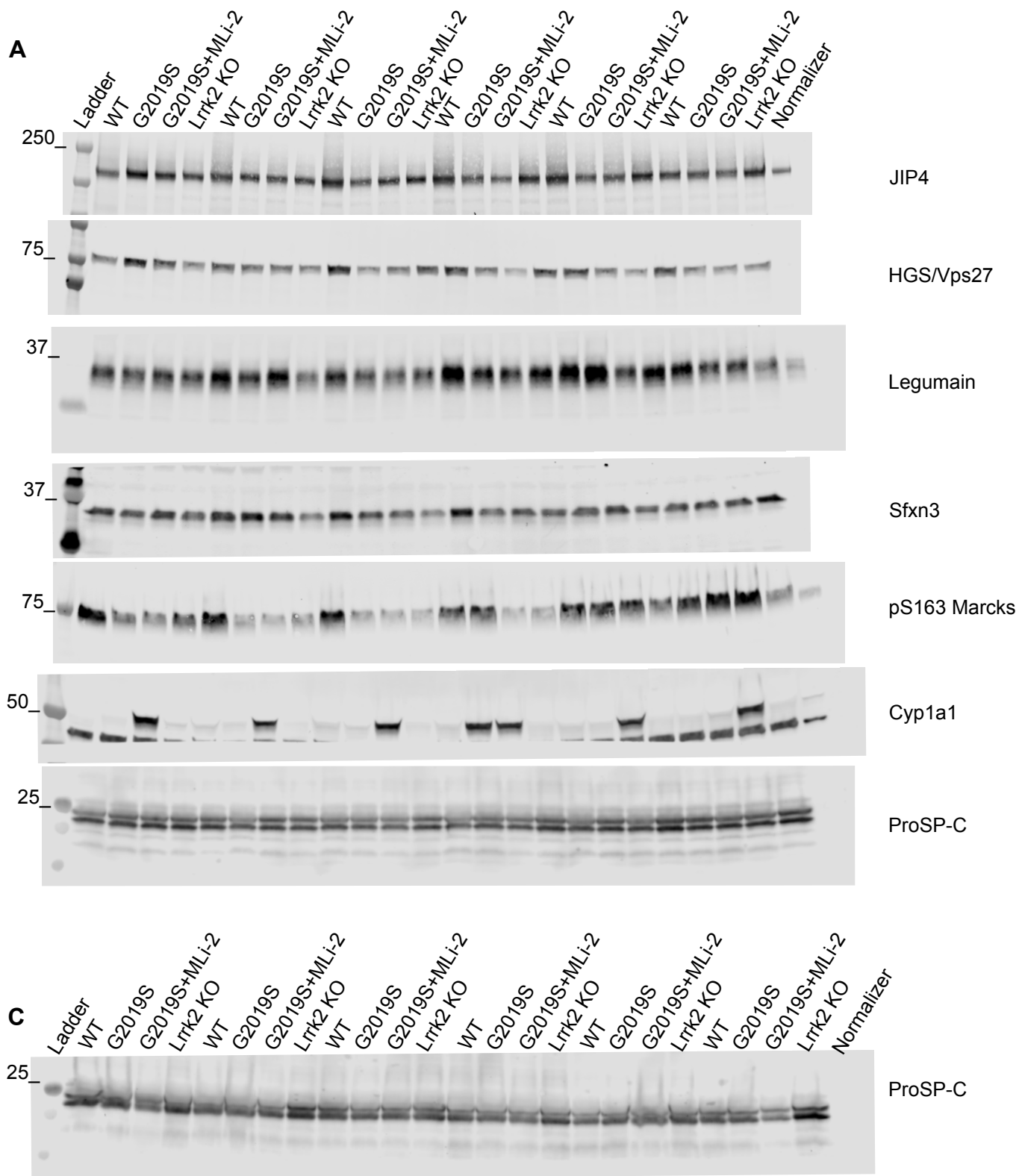


Fig. S6A, C

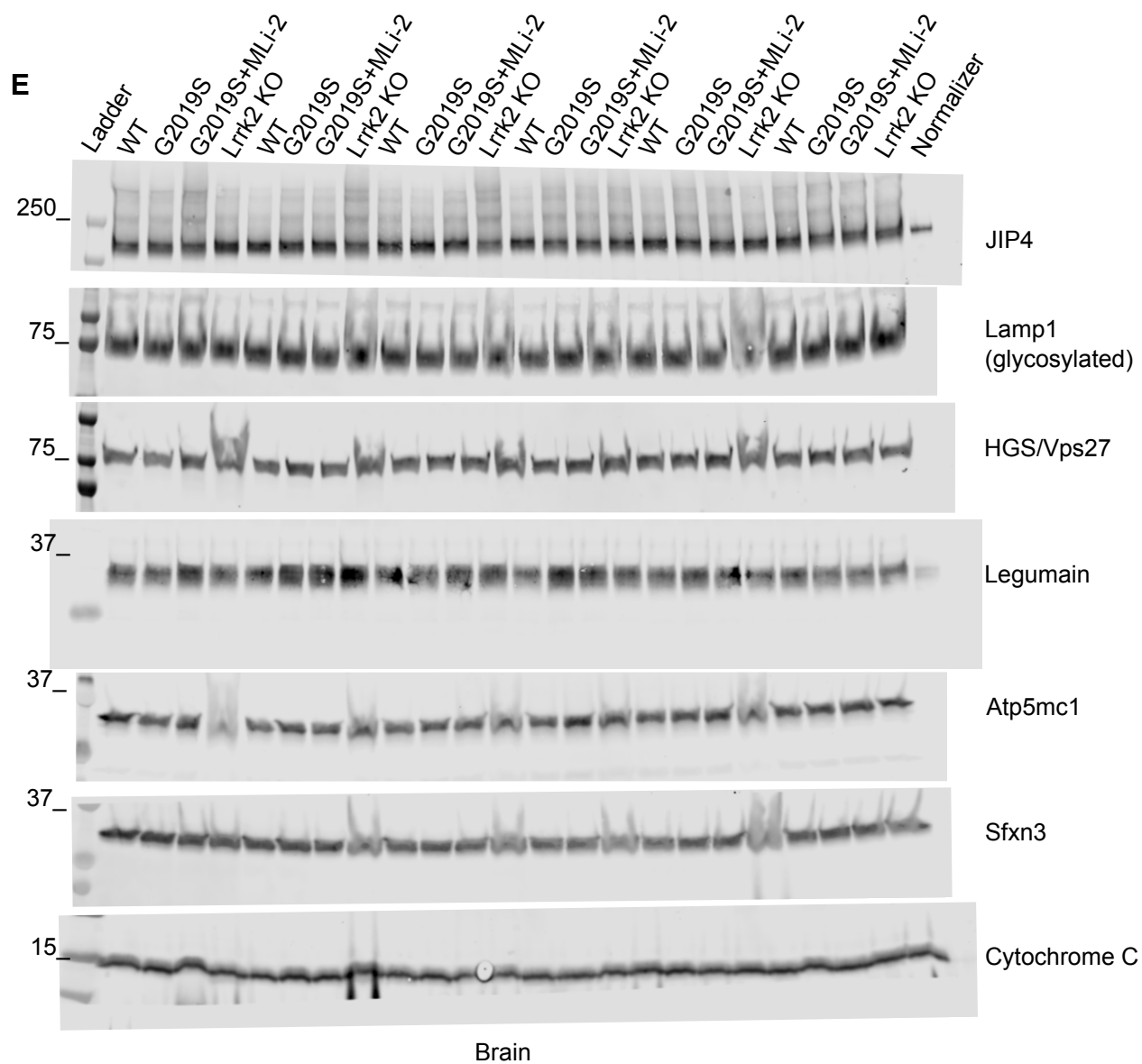


Fig. S6E

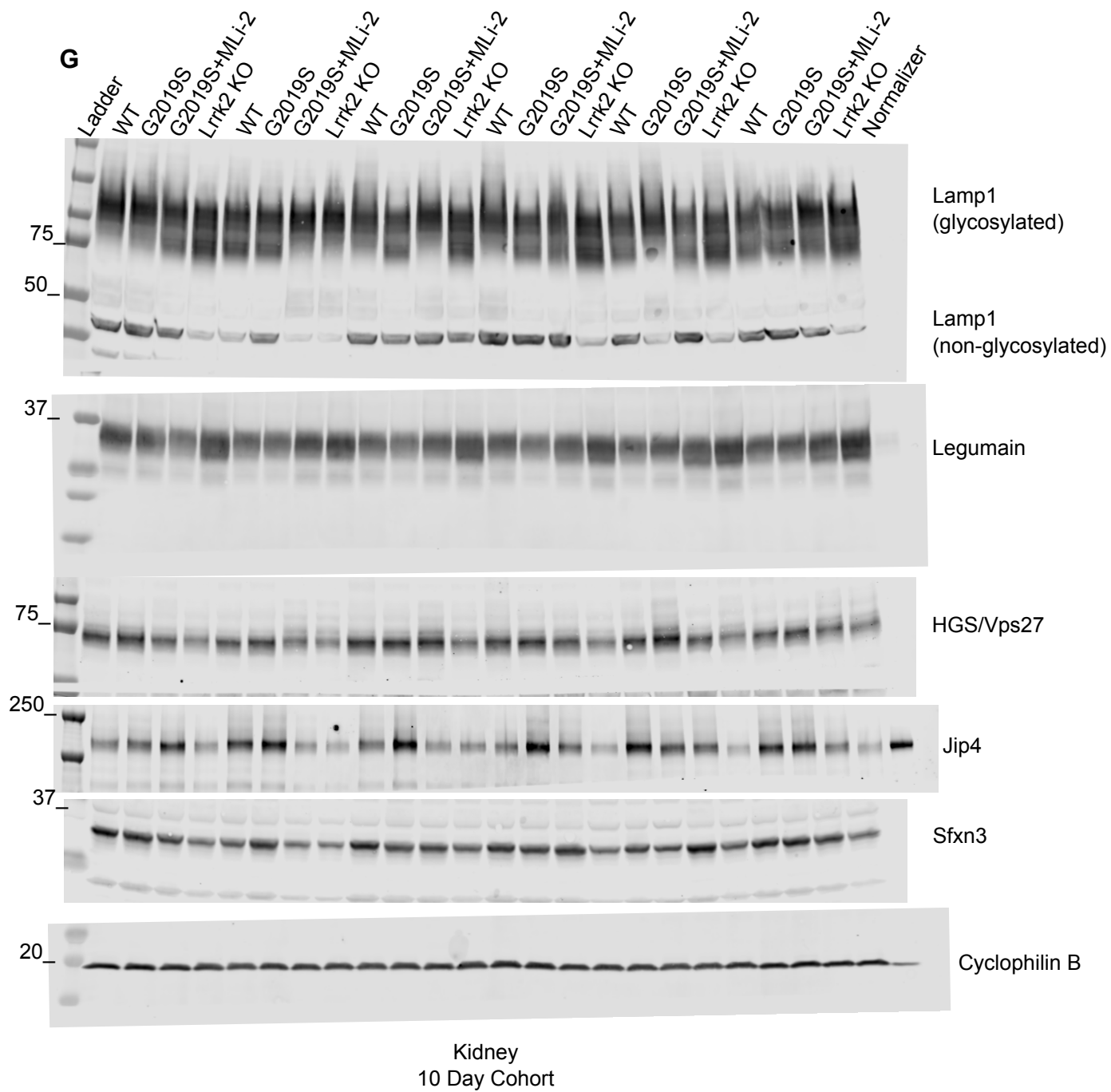


Fig. S6G