

## **SUPPLEMENTARY DATA**

### **Emerging role of kinin B1 receptor in persistent neuroinflammation and neuropsychiatric symptoms in mice following recovery from SARS-CoV-2 infection**

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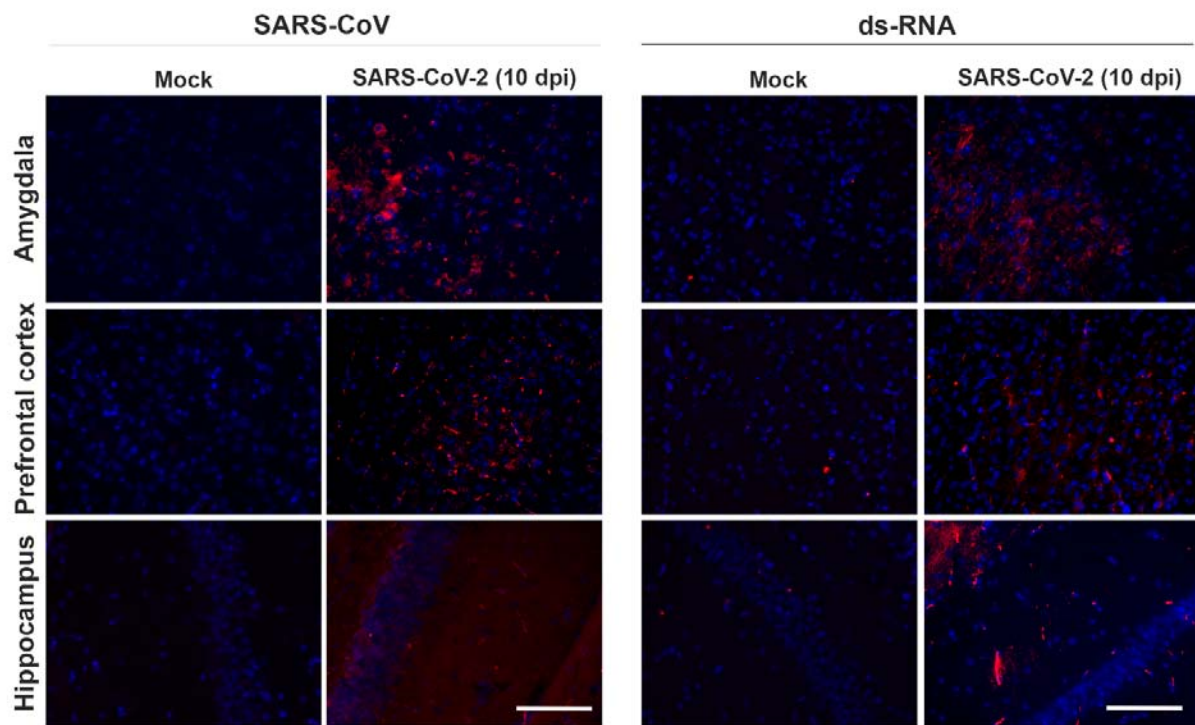
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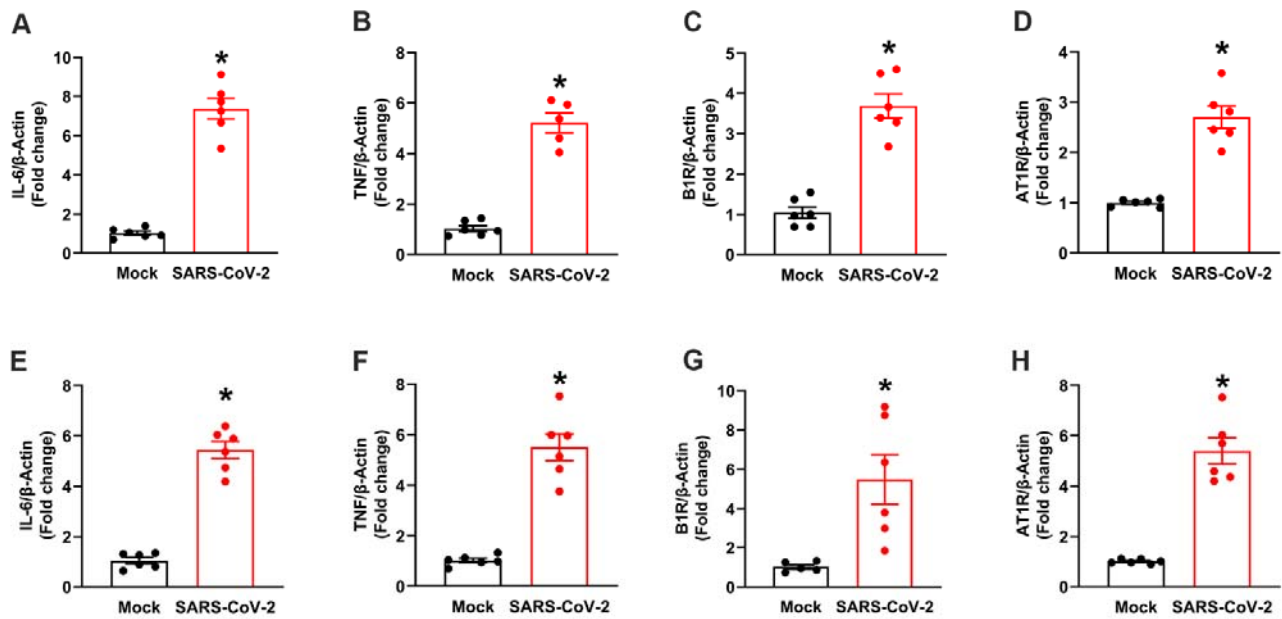
**Table S1. Primer sequences used for real-time quantitative reverse transcriptase polymerase chain reaction (qRT-PCR).**

<b>Gene</b>	<b>Forward primer (5'-3')</b>	<b>Reverse primer (5'-3')</b>
IL-6	TCC TTA GCC ACT CCT TCT GT	AGC CAG AGT CCT TCA GAG A
TNF	TCT TTG AGA TCC ATG CCG TTG	AGA CCC TCA CAC TCA GAT CA
B1R	AGA TCA GAA GCT GCC AAG TTA G	TCC TGT CCT TCT TCC TTT TGC
AT1R	GGC TGG CAT TTT GTC TGG ATA	CAG TAG AAG AGT TAA GGG CCA T
$\beta$ -Actin	GAC TCA TCG TAC TCC TGC TTG	GAT TAC TGC TCT GGC TCC TAG



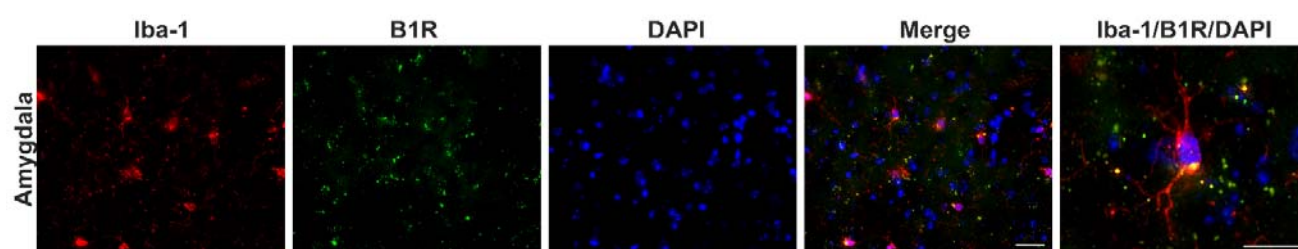
**Figure S1. SARS-CoV-2 infected K18-hACE2 mice show presence of virus in the brain.**

Representative images showing immunofluorescence staining of SARS-CoV virus and double stranded RNA (dsRNA) in amygdala, prefrontal cortex, and hippocampus of mock or virus infected mice. Staining reveals that SARS-CoV virus is detectable at 10 dpi in all 3 regions of virus infected mice compared to mock infected. Detection of dsRNA in 10 dpi mice indicates active viral replication compared to mock infected mice. Scale bar: 100  $\mu$ M.



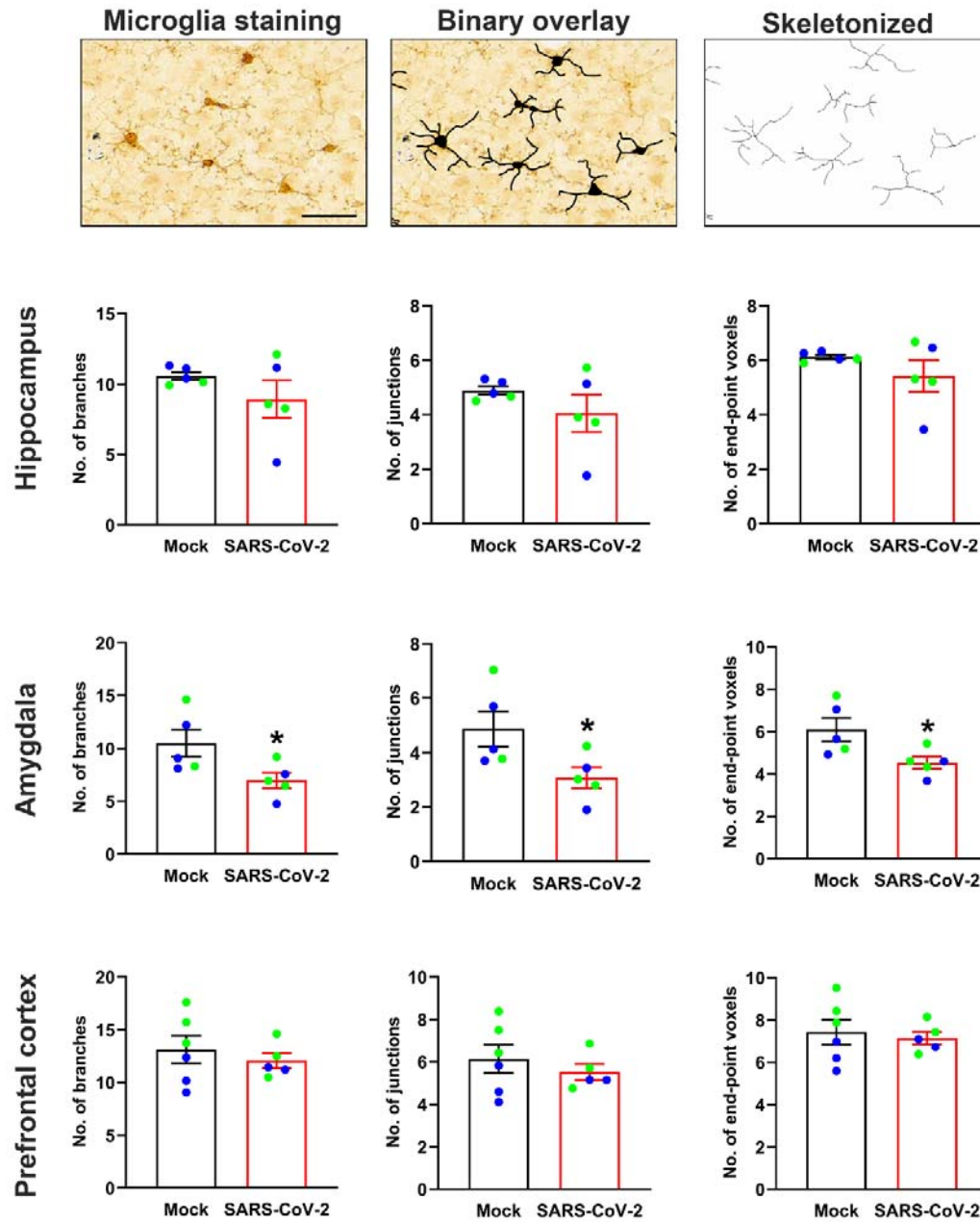
**Figure S2. SARS-CoV-2 infection increases inflammatory markers in the brain.**

Real time RT-PCR was used to measure mRNA expression of proinflammatory mediators in hippocampus and prefrontal cortex punches from mock and SARS-CoV-2 infected mice 45dpi. mRNA expression of IL-6, TNF, B1R, and AT1R is upregulated within the hippocampus (**A-D**) and prefrontal cortex (**E-H**) of SARS-CoV-2 infected mice compared to mock infected mice, indicating increased neuroinflammation.  $n=6$  per group. Data are mean  $\pm$  s.e.m. \* $p < 0.05$  (Student's t-test).



**Figure S3. Co-localized expression of B1R with microglia.**

Representative double immunofluorescence staining for microglia stained with ionized calcium-binding adaptor molecule 1 (Iba-1, red) and B1R expression (green) showing co-localization of B1R with microglia within the amygdala region of brain in SARS-CoV-2 infected mice. Scale bar: 100  $\mu$ M.



**Figure S4. Mild infection with SARS-CoV-2 and microglia ramifications in different brain regions.** Microglia morphology was quantified using photomicrographs of tissue in three different regions of the brain. The images were made binary, skeletonized, and analyzed using ImageJ plugins AnalyzeSkeleton (2D/3D) to determine the complexity of process branching (number of branches, number of junctions, and number of endpoints). Within the amygdala, there was a significant difference among the mock and SARS-CoV-2 infected groups for microglia in no. of branches, no. of junctions, and no. of end-point voxels. Blue data points represent male mice, green data points represent female mice. Data are mean  $\pm$  s.e.m. \* $p < 0.05$  (Student's t-test).