

Characterization of disease progression in a mouse model of COVID-19 Nikki Seilhamer, James Chung, Rachel Olson, James Amos-Landgraf, Craig Franklin Department of Veterinary Pathobiology, College of Veterinary Medicine, University of Missouri, Columbia, Missouri

Background

- SARS-CoV-2, the agent of COVID-19, remains a threat to public health Severe clinical cases can result in bilateral pneumonia and subsequent ARDS, driven by a proinflammatory cytokine storm
- The B6.Cg-Tg(K18-ACE2)2Prlmn/J mouse expressing human angiotensinconverting enzyme 2 (ACE2) is used to study SARS-CoV-2 pathogenesis
- Initial studies have shown that mice develop interstitial pneumonia during acute disease; however, the full scope of disease is not yet understood

Objective: Characterize disease progression in SARS-CoV-2 infected K18-hACE2 mice over an 8-week period, with a focus on lung and brain cytokines and histopathologic changes.



BAFF	IFN gamma	IL-6	IL-17A (CTLA-8)	IL-31	MCP-3 (CCL7)
Betacellulin (BTC)	IL-1 alpha	IL-7	IL-18	IL-33	MIP-1 alpha (CCL
ENA-78 (CXCL5)	IL-1 beta	IL-7R alpha	IL-19	IL-33R (ST2)	MIP-1 beta (CCL4
Eotaxin (CCL11)	IL-2	IL-9	IL-22	IP-10 (CXCL10)	MIP-2 alpha (CXCI
G-CSF (CSF-3)	IL-2R	IL-10	IL-23	LIF	RANKL
GM-CSF	IL-3	IL-12p70	IL-25 (IL-17E)	Leptin	RANTES (CCL5)
GRO alpha (CXCL1)	IL-4	IL-13	IL-27	M-CSF	TNF alpha
IFN alpha	IL-5	IL-15	IL-28	MCP-1 (CCL2)	VEGF-A

Sample collection (weeks post-inoculation)	Total Mice	Male / Fema
3 weeks	14	8M / 6F
4 weeks	13	7M / 6F
5 weeks	16	8M / 8F
6 weeks	11	6M / 5F
7 weeks	14	7M / 7F
8 weeks	10	5M / 5F

Figure 1. A) Schematic of methods for characterization of acute disease. B) Table illustrating cytokines that were tested in the 48-plex assay. C) Table illustrating mice used to study ongoing disease and resolution. Cohorts of mice were euthanized weekly for collection of brains and lungs used for histologic analysis.



Transition from lymphohistiocytic interstitial pneumonia to lymphoplamacytic aggregates



Figure 2. Representative images of mouse lung histology, stained with H&E and taken at 20x magnification. A) Normal mouse lung. B) Lung lesion from a mouse euthanized 6-days post-inoculation. Red arrows indicate alveolar macrophages, and the red dashed circle denotes an area with apoptotic debris. Typically, acute disease (6-8 days post-inoculation) was characterized by a multifocal mild to moderate lymphohistiocytic interstitial pneumonia. C) Lung from mouse euthanized 8-weeks post-inoculation. Orange arrows indicate plasma cells. Lung lesions at 8-weeks were primarily mild multifocal lymphoplasmacytic aggregates which was suggestive of ongoing disease resolution

Severity of pneumonia is similar between clinical and asymptomatic mice



Figure 4. Illustration of histopathologic scores of lungs from mice 6-days postinoculation. Bars represent mean and SEM. Independent t-test analysis revealed that the difference in lesion severity between mice with and without clinical signs was not statistically significant.

Meningoencephalitis is most evident in mice with clinical signs



Figure 5. Stacked bar chart showing the distribution of brain lesions between mice with and without clinical signs. Note that nearly all the mice with clinical signs had brain lesions. Also of note, mice with no clinical signs and evidence of meningoencephalitis had very mild to equivocal lesions (data not shown). Chi-square analysis revealed a statistically significant relationship between the presence of clinical signs and lesions (** indicates p < 0.001; $\chi 2 = 23.54$)



Meningoencephalitis during acute disease Figure 3. Representative images of mouse brain histopathology, stained with H&E and taken a 20x magnification. Both images are from a mouse euthanized 6-days post-inoculation. Typical brain lesions during acute disease were characterized by perivascular lymphoid cuffing (inset), meningeal lymphoid infiltration, and scattered apoptotic debris (arrow) most notably in the brainstem



resolution in mice that survived.

Collectively, these findings suggest a correlation between brain lesions and clinical disease.

This project is funded by the Mutant Mouse Research and Resource Center. The stipend for Nikki Seilhamer is supported by the Kent Tomazi Memorial Research Fund in Veterinary Medicine and an endowment established by IDEXX-BioAnalytics. We would also like to thank Alex DeWitt and Chris Johanning for their assistance with mice health checks and necropsies.



