

Examining the Effects Of SARS-CoV-2 on Metastatic Breast Cancer in a Murine Model

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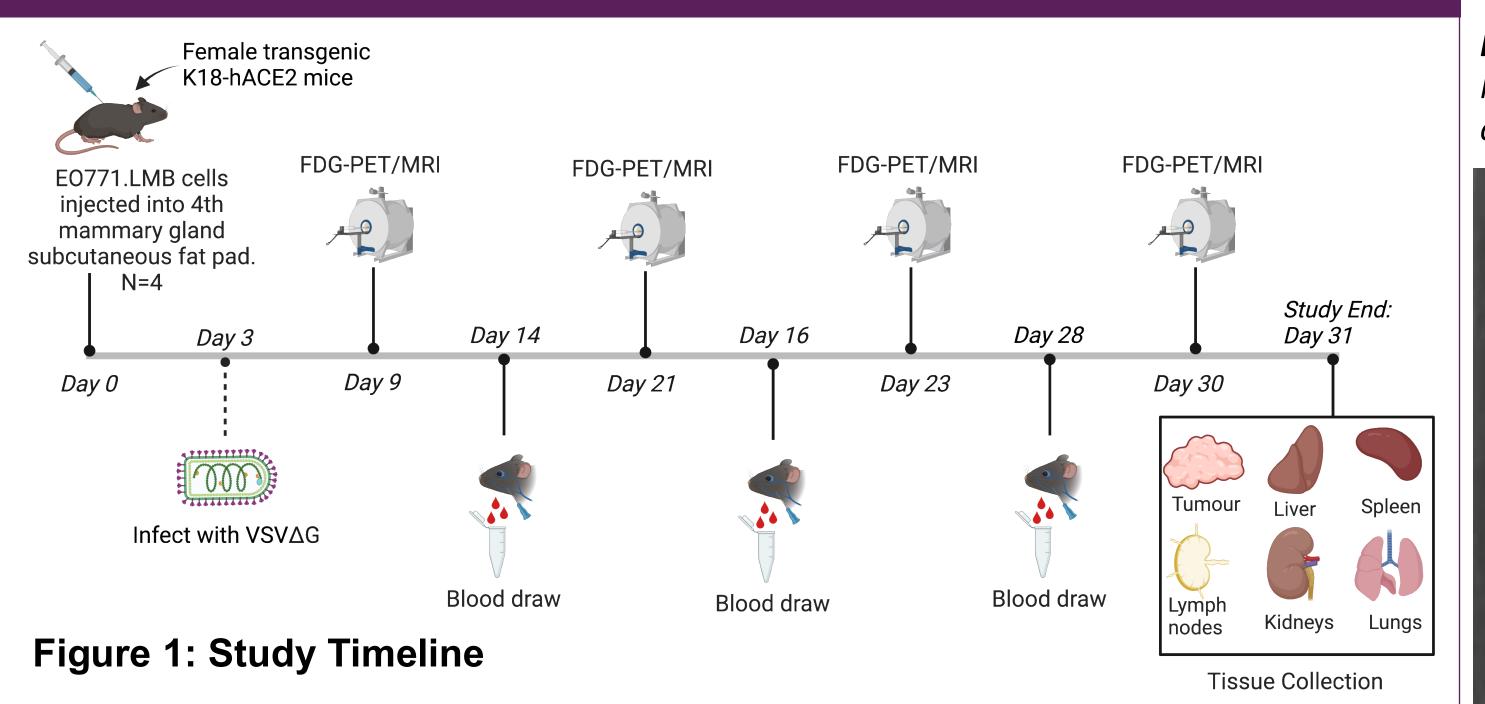
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Background and Aims

- SARS-CoV-2 has led to 771 million infections and ~7 million deaths (Nov 2, 2023)¹
- Estimated that at least 12% of Canadians and 39% of Americans have been infected (likely a significant underestimation)
- Infection and vaccination seen to interfere with clinical FDG-PET scans²
- Fluoro-2-deoxy-D-glucose (FDG) is preferentially taken up by cells with increased glucose metabolism^{3,4}
 - COVID-19 inflammation mimics cancer FDG uptake patterns
- COVID-19 is a CL3 pathogen → Expensive and difficult to work with
- VSVΔG S expresses the spike protein of COVID-19 to facilitate early research

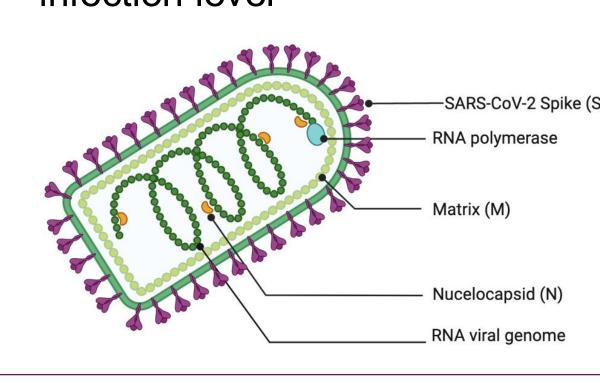
Goal: Characterize imaging and biological features of infection with VSV expressing SARS-CoV-2 spike in breast cancer model. Distinguish FDG uptake mediated by tumour and infection.

Methods



Experimental Design

- Mouse model:
- K-18-hACE2 mice (C57BL/6 background)
- Express human ACE2 under control of the K18 promotor
- Directed expression to epithelial tissues, mimics human ACE2 distribution
- Biological analysis:
- Samples: weekly and terminal blood, terminal organ collection
- Flow cytometry for immune phenotyping
- Level of GFP+ cells as marker of infection level



Groups:

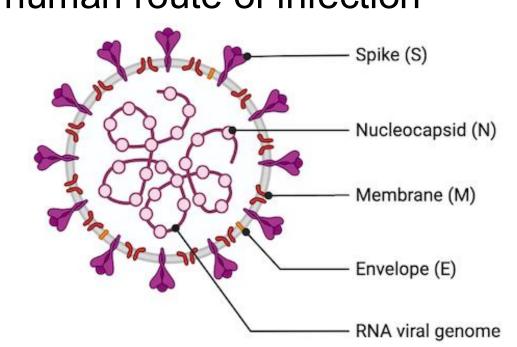
- E0771.lmb implant only 5X10⁴ cells in 100µl HBSS/Matrigel
- E0771.lmb implant + VSVΔG S (SARS-CoV-2 omicron) infection 1X10⁶ PFU/mL

• FDG-PET/MRI:

- ~350µCi injection 1 hour prior to scan
- Simultaneous PET/MRI

Virus model:

- VSVΔG S (GFP+)
- Pseudotype virus expressing the S protein of COVID-19 Omicron
- Intranasal administration to mimic human route of infection



Results

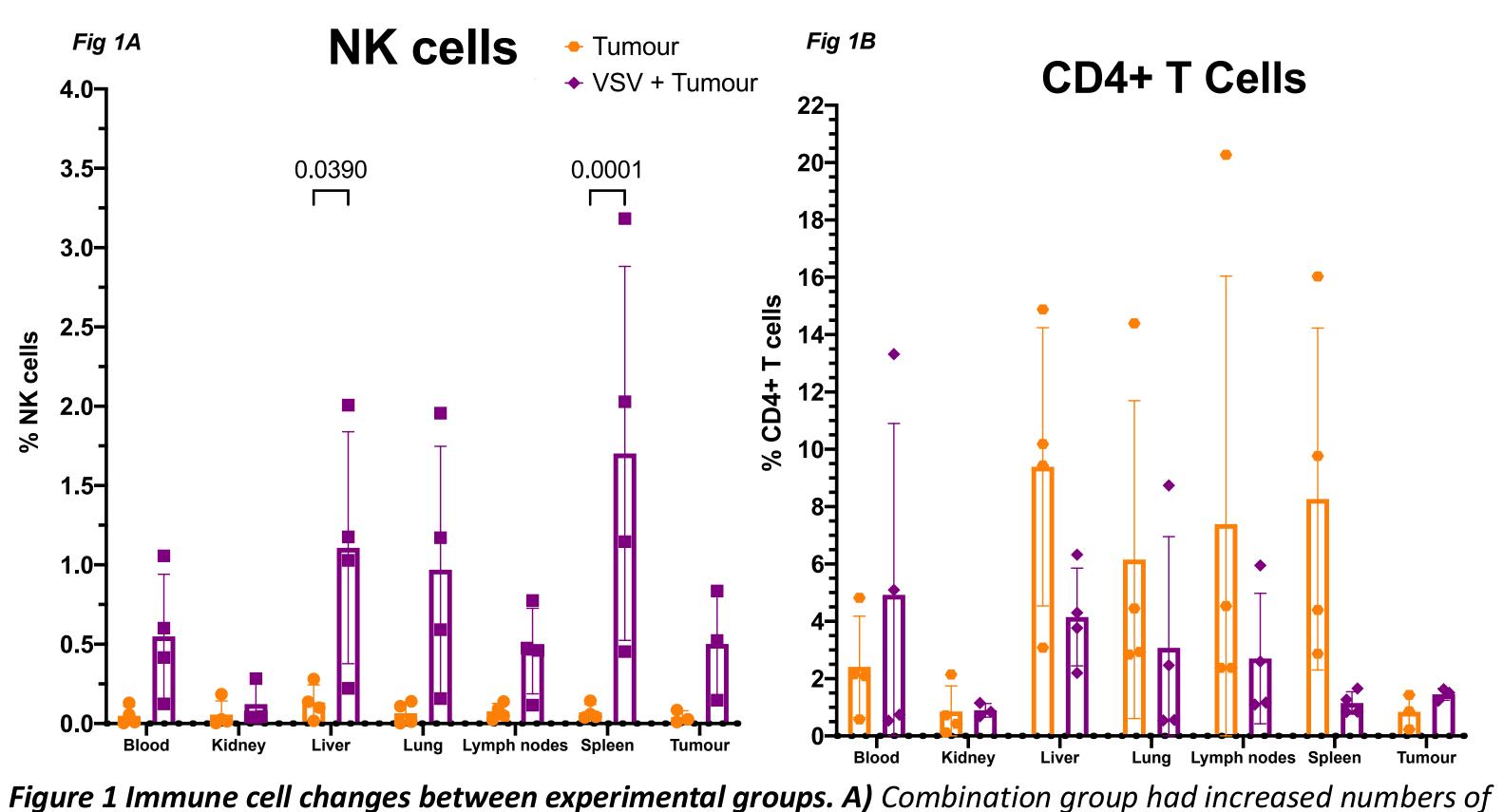


Figure 1 Immune cell changes between experimental groups. A) Combination group had increased numbers of NK cells in all organs compared to tumour alone. This was most pronounced in the spleen and liver. **B)** CD4+ T cell levels were increased overall for tumour only mice, particularly in the liver and spleen.

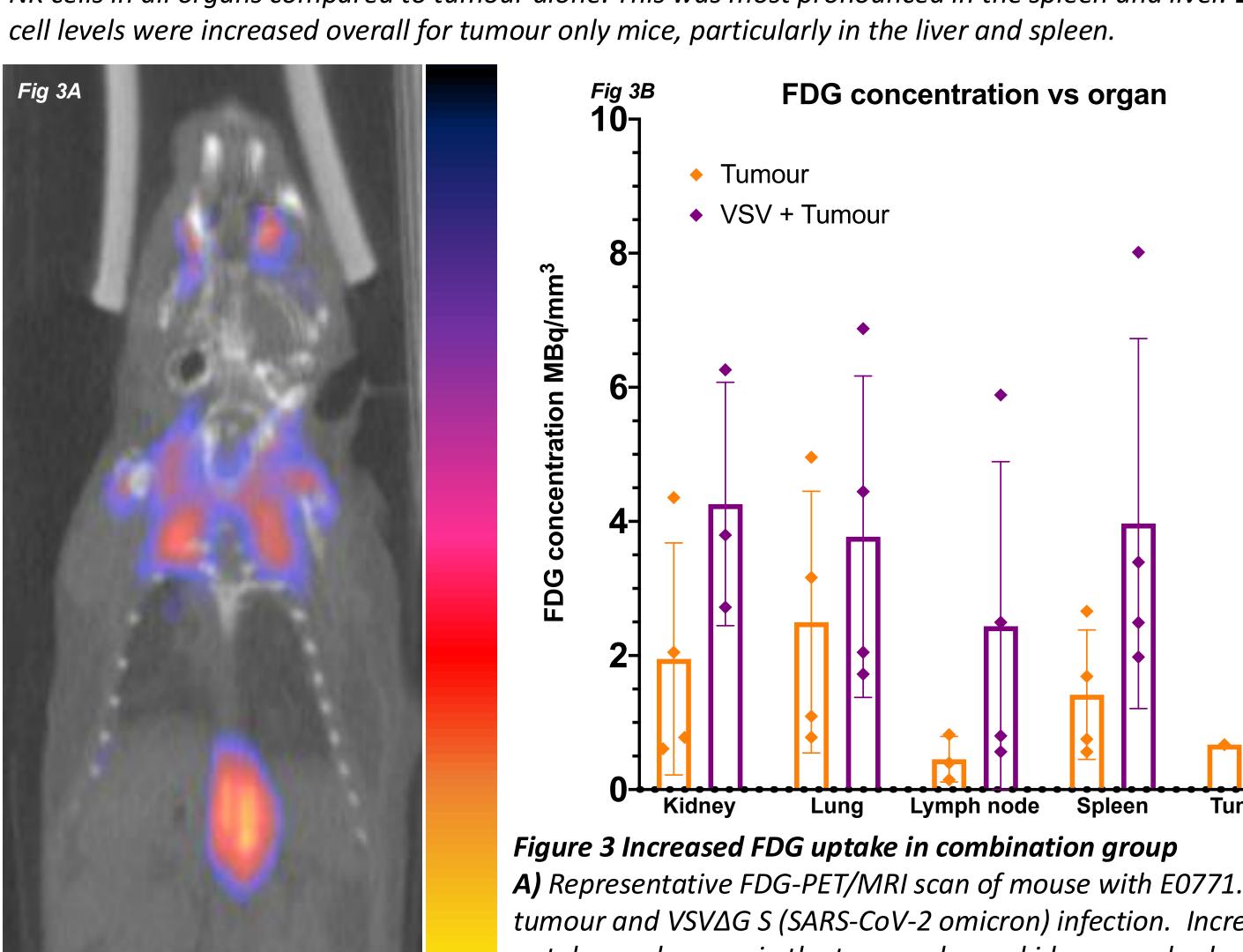


Figure 3 Increased FDG uptake in combination group

A) Representative FDG-PET/MRI scan of mouse with E0771.lmb

tumour and VSVΔG S (SARS-CoV-2 omicron) infection. Increased FDG

uptake can be seen in the tumour, lungs, kidneys, and spleen. B) FDG

concentration vs organ determined from FDG-PET/MRI imaging.

Values have been standardized against muscle tissue, FDG dose, and

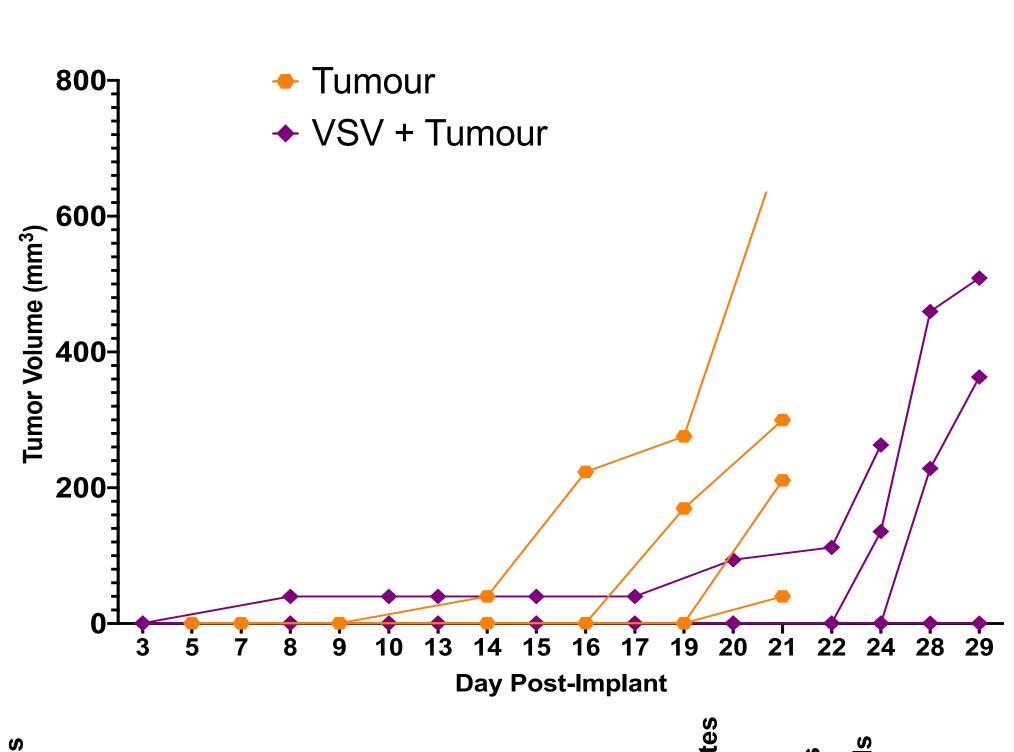
mouse weight. Overall, the combination group had increased FDG

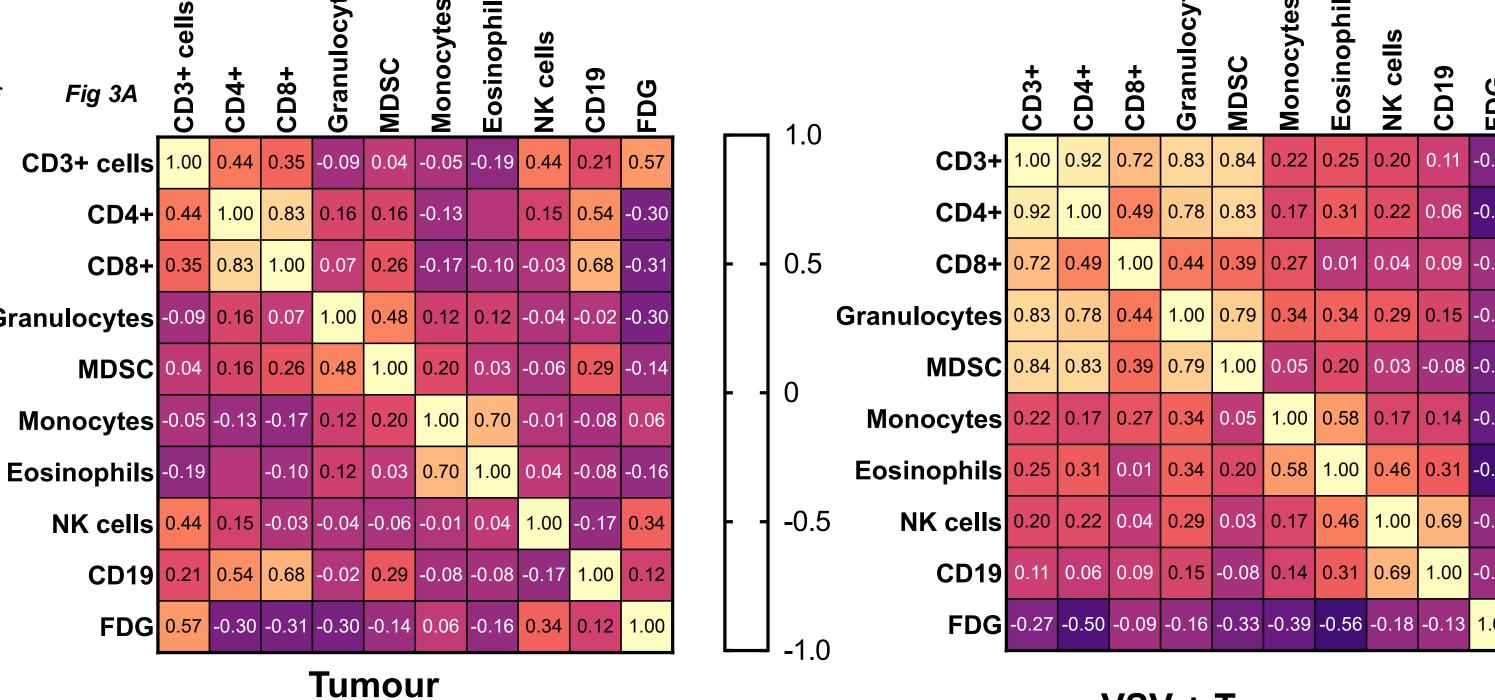
concentration, with a P-value of 0.0467 between groups (ANOVA).

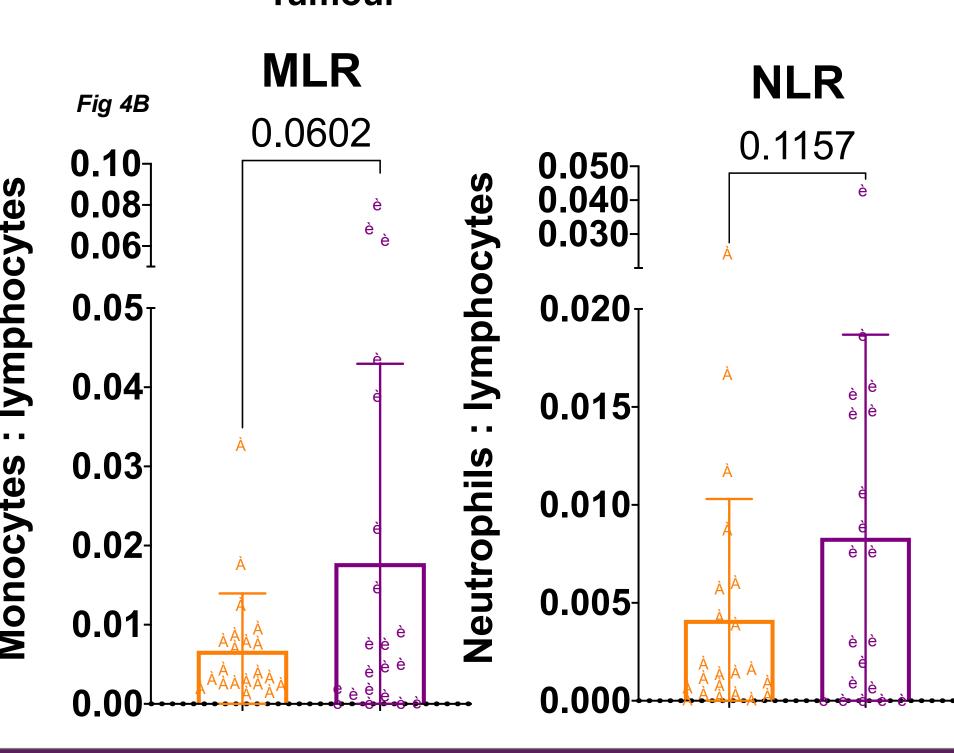
Figure 2: Tumour volume changes over time.

Volumes were calculated using caliper measurements and MRI data. Tumour volumes were initially monitored by caliper; however, MRI data was needed to determine internal tumour volume.

The combination group showed slower initial tumour growth than those with EO771.lmb tumours only.







VSV + Tumour

Figure 4 Immune cell changes
between experimental groups.

A) Pearson correlation matrix for
both experimental groups showing
the correlation between cell types.
The combination group showed an
increased correlation between
granulocytes and T cells
B) Neutrophil: lymphocyte and
monocyte: lymphocyte ratios were
increased for the combination
group. These ratios are indicators of
systemic inflammation and are often

reported in cancer and infections,

in combination groups.

suggesting increased inflammation

Conclusions & Future Directions

CONCLUSIONS

- Tumour growth was delayed in combination group
- Increased FDG concentration in combination group possible overall inflammatory state?
- Combination group has increased monocytes but decreased lymphocytes
- Increased levels of NK cells in combination

FUTURE WORK

- Compare to VSVΔG "empty" as a mock control and to mice infected only with VSVΔG omicron
- Test different VSV infection times relative to tumour implantation
- Perform tumour resection to extend survival time to observe long term changes
- Test different variants of SARS-CoV-2
- Test with full SARS-CoV-2 in a CL3 setting

References and Acknowledgements

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