

# The Role of *Ndufs4* and *Slc2a1* on D4M-UV2 Melanoma Tumor Metabolism and Growth

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## Introduction

- Melanoma is one of the most aggressive and metastasis-prone forms of cancer.
- Melanoma brain metastases (MBMs) are a common and devastating complication of advanced melanoma.
- Oxidative Phosphorylation (OXPHOS) has been observed to be particularly elevated in MBMs.
- Previous studies have found that tumor metabolism may be involved in suppressing immune responses to cancer.
- The role of tumor metabolism in immune responses remains unclear.

## Hypothesis

We hypothesize that elevated tumor metabolism suppresses the immune response in MBMs.

## Aim

- To develop models of High- and Low-OXPHOS or glycolysis using CRISPR technology.
- To better understand the role of tumor metabolism in MBM pathogenesis.

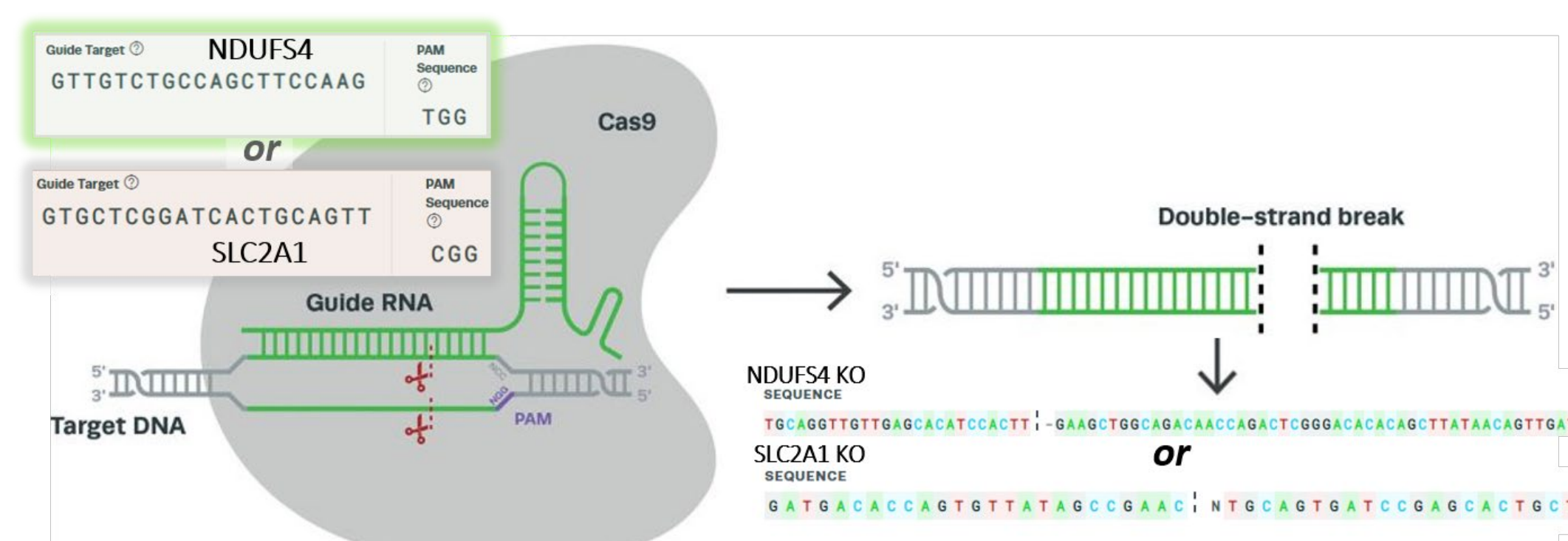
## Methods and Materials

**Cell Culture:** D4M-UV2 cells were cultured in DMEM media containing 10% FBS and 1% NEAA. Cells were incubated at 37°C and 5% CO<sub>2</sub> and were grown *in vitro*.

**Cellular Metabolism Analysis:** The MitoStress Test was run on Seahorse 96-well XF Analyzer to measure oxygen consumption rate (OCR) and extracellular acidification rate (ECAR) of the D4M-UV2 cell line.

**Cell Proliferation Assay:** D4M-UV2 cells were incubated with Cell Titer Blue reagent until color change was observed. Then, fluorescence of cell media was analyzed on a Tecan plate reader at 560 nm excitation / 590 nm emission.

## CRISPR Knockouts



## Results

### Metabolic Effects on NDUFS4 KO

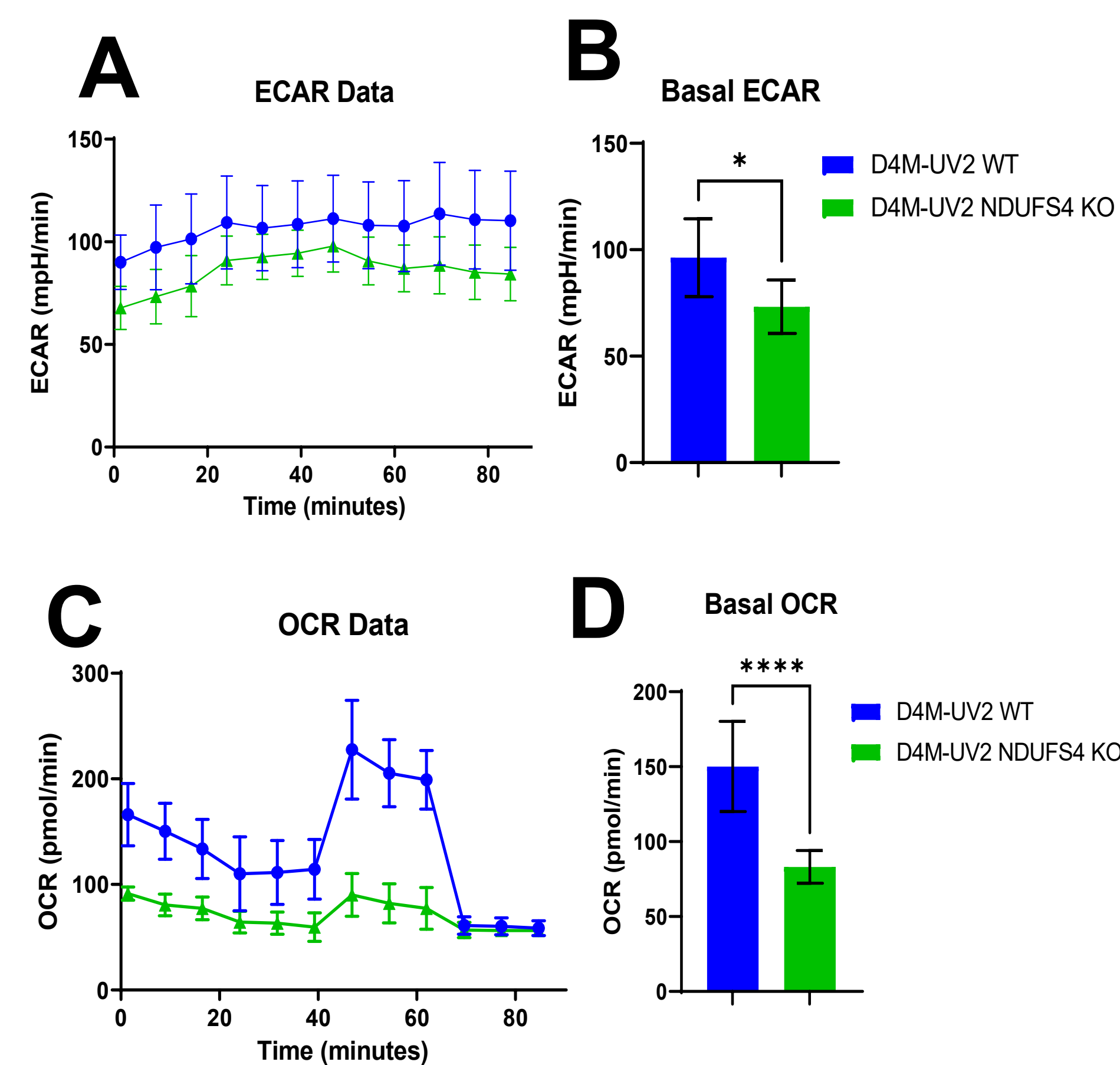


Figure 1: Observation of Metabolic Effects on NDUFS4 KO conducted by MitoStress Test. (A) Tumor glycolysis is represented by the Extracellular Acidification Rate (ECAR) of treated cells (n = 5). (B) KO of NDUFS4 resulted in a small 20.6% decrease in basal ECAR (p = 0.0439). (C) Tumor mitochondrial OXPHOS is represented by the Oxygen Consumption Rate (OCR) of treated cells (n = 5). (D) KO of NDUFS4 decreased basal OCR by 44.6% (p < 0.0001). Dots in MitoStress Test curves represent individual replicates (n = 3) and lines represent SD. Bars in ECAR and OCR histograms represent mean values and lines represent SD.

### Metabolic Effects on SLC2A1 KO

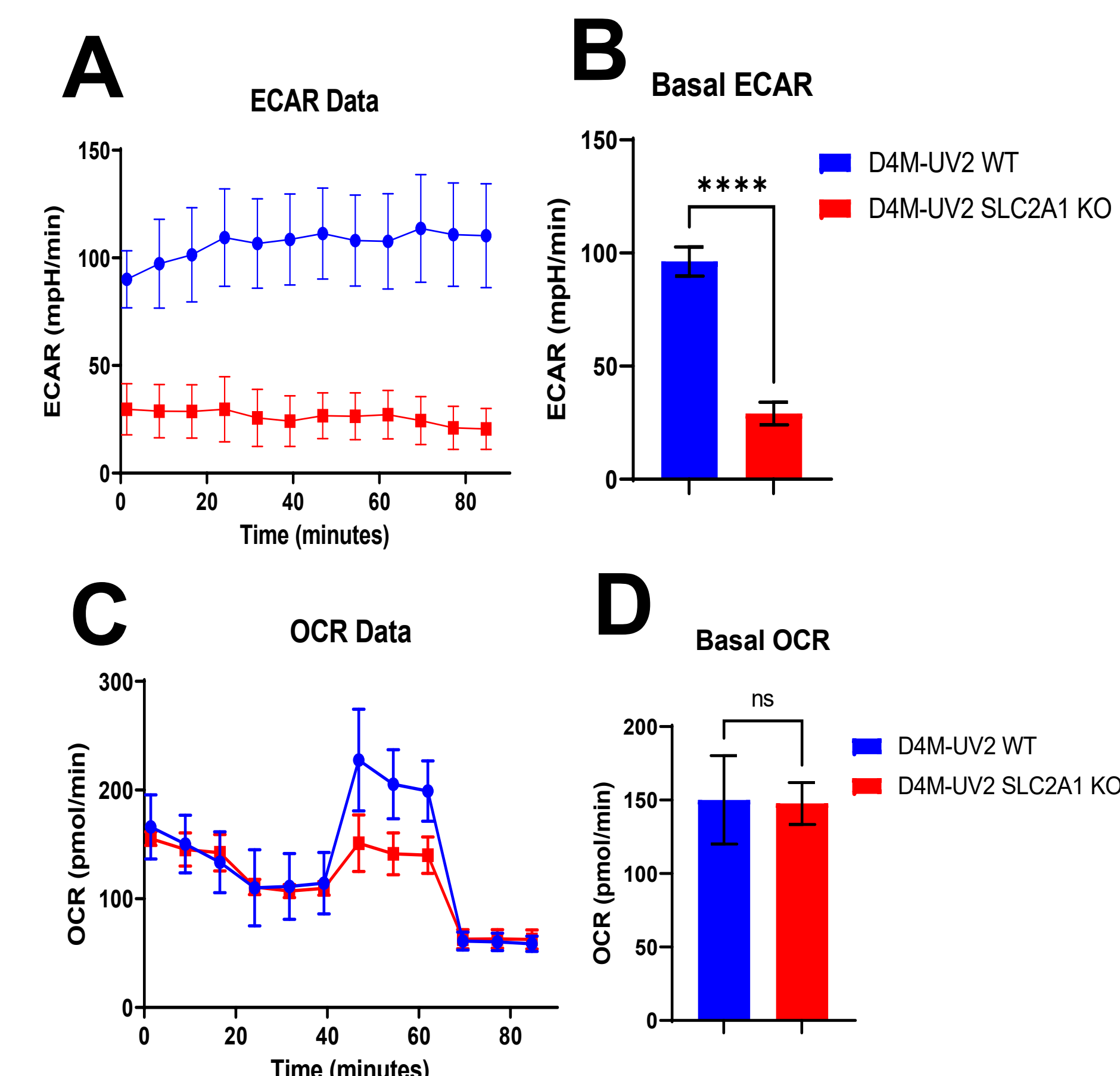


Figure 2: Observation of Metabolic Effects on SLC2A1 KO conducted by MitoStress Test. (A) Tumor glycolysis is represented by the Extracellular Acidification Rate (ECAR) of treated cells (n = 5). (B) KO of SLC2A1 resulted in a 69.8% decrease in basal ECAR (p < 0.0001). (C) Tumor mitochondrial OXPHOS is represented by the Oxygen Consumption Rate (OCR) of treated cells (n = 5). (D) KO of SLC2A1 did not significantly change Basal OCR (p = 0.8891). Dots in MitoStress Test curves represent individual replicates (n = 3) and lines represent SD. Bars in ECAR and OCR histograms represent mean values and lines represent SD.

### Effects on Cell Growth

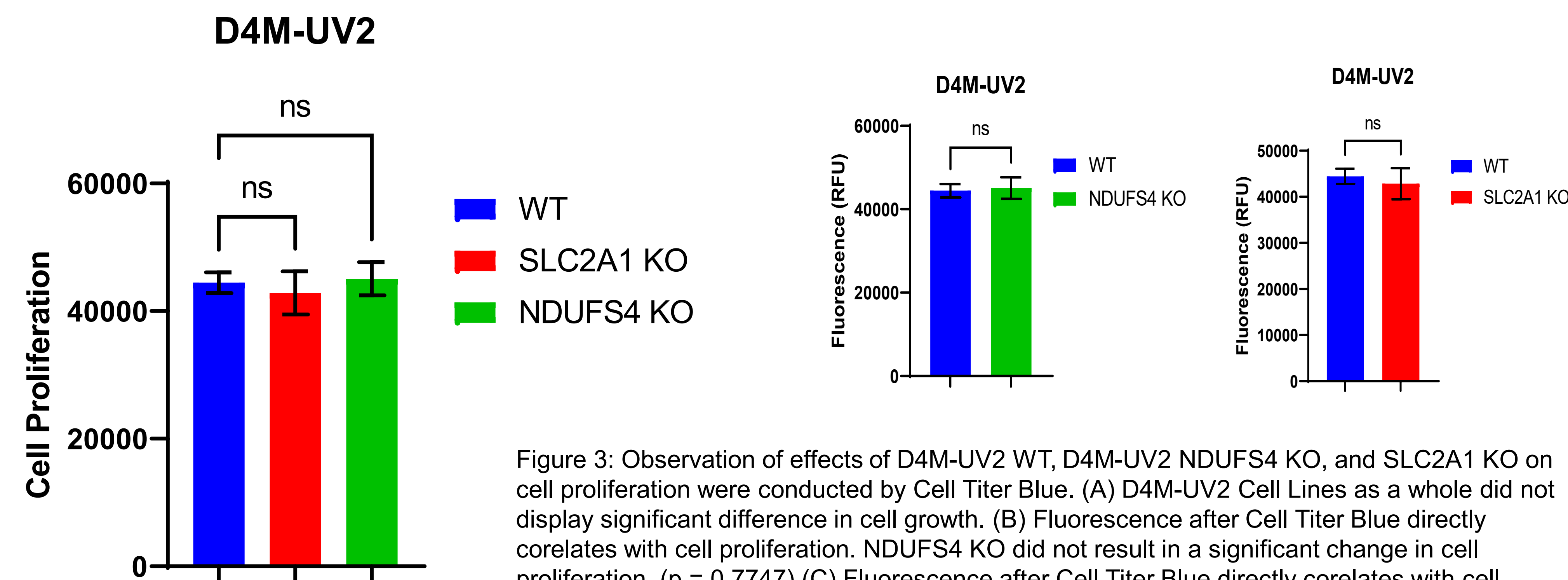
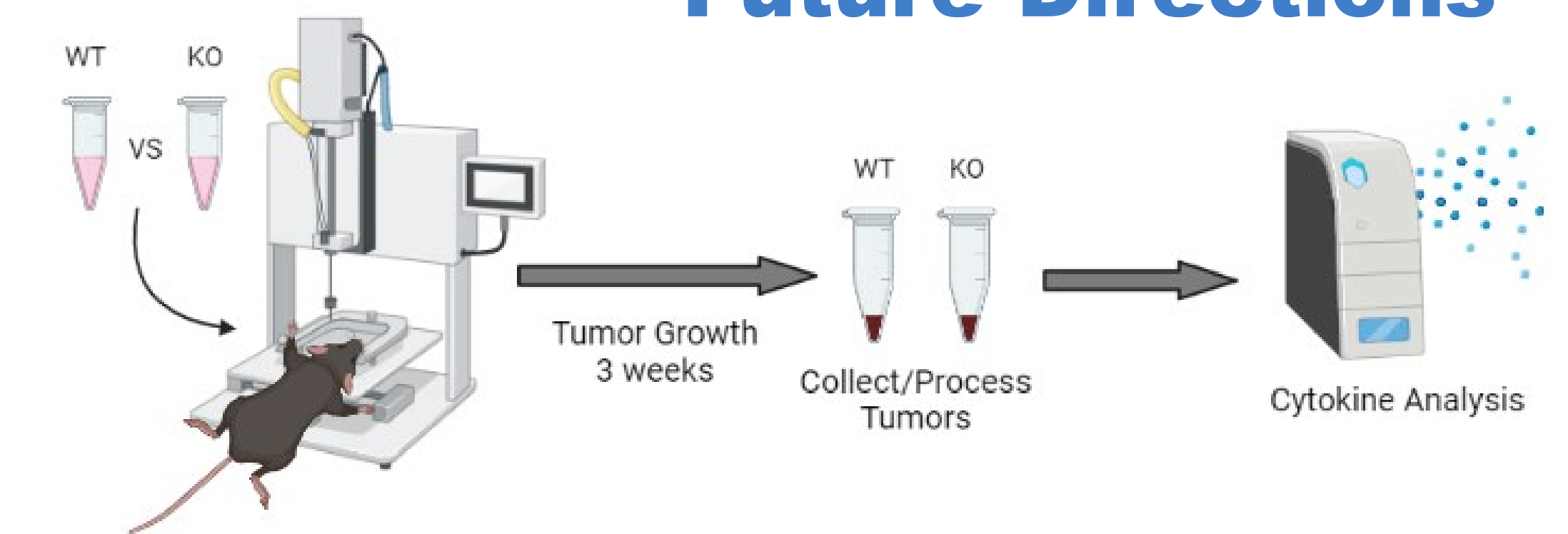


Figure 3: Observation of effects of D4M-UV2 WT, D4M-UV2 NDUFS4 KO, and SLC2A1 KO on cell proliferation were conducted by Cell Titer Blue. (A) D4M-UV2 Cell Lines as a whole did not display significant difference in cell growth. (B) Fluorescence after Cell Titer Blue directly correlates with cell proliferation. NDUFS4 KO did not result in a significant change in cell proliferation. (p = 0.7747) (C) Fluorescence after Cell Titer Blue directly correlates with cell proliferation. SLC2A1 KO did not result in a significant change in cell proliferation. (p = 0.1716)

## Conclusions

- Gene KO of NDUFS4 strongly inhibited OXPHOS and slightly inhibited glycolysis *in vitro*.
- Gene KO of SLC2A1 resulted in specific inhibition of glycolysis *in vitro*.
- KO of NDUFS4 OR SLC2A1 does not impact the proliferation of tumor cells *in vitro*.
- We plan to study these models of high and low OXPHOS or glycolysis and how it can apply to tumor metabolism and anti-tumor immune response.
- D4M-UV2 Cell Lines will be evaluated for *in vivo* growth in mice.
- Future studies will evaluate the rationale for development of combinatorial treatment strategies targeting tumor metabolism and immunotherapy.

## Future Directions



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Figure 4: Evaluation of D4M-UV2 WT and KO Cell Lines can be conducted in mice models via intracranial injections.

## Acknowledgements

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## References

- Hasanov M et al. Predictors of overall survival (OS) in patients (pts) with melanoma brain metastasis (MBM) in the modern era. *Journal of Clinical Oncology*. 2021; 39(15):9540.
- DePeaux K & GM Delgoffe. Metabolic barriers to cancer immunotherapy. *Nature Reviews Immunology*. 2021 Apr 29;21:785-797.
- Fischer et al. Clinical, molecular, metabolic, and immune features associated with oxidative phosphorylation in melanoma brain metastases. *Neurooncol Adv*. 2021 Jan 6;3(1):vdad177.
- Najjar YG et al. Tumor cell oxidative metabolism as a barrier to PD-1 blockade immunotherapy in melanoma. *JCI Insight*. 2019 Feb;4(5):e124989.