

Metformin Inhibits Small Intestinal Neuroendocrine Tumor (SI-NET) Proliferation In Vivo

F. Axling, P. Hellman, O. Norlén, P. Stålberg, E. Barazeghi
Endocrine Surgery, Department of Surgical Sciences, Uppsala University, Sweden

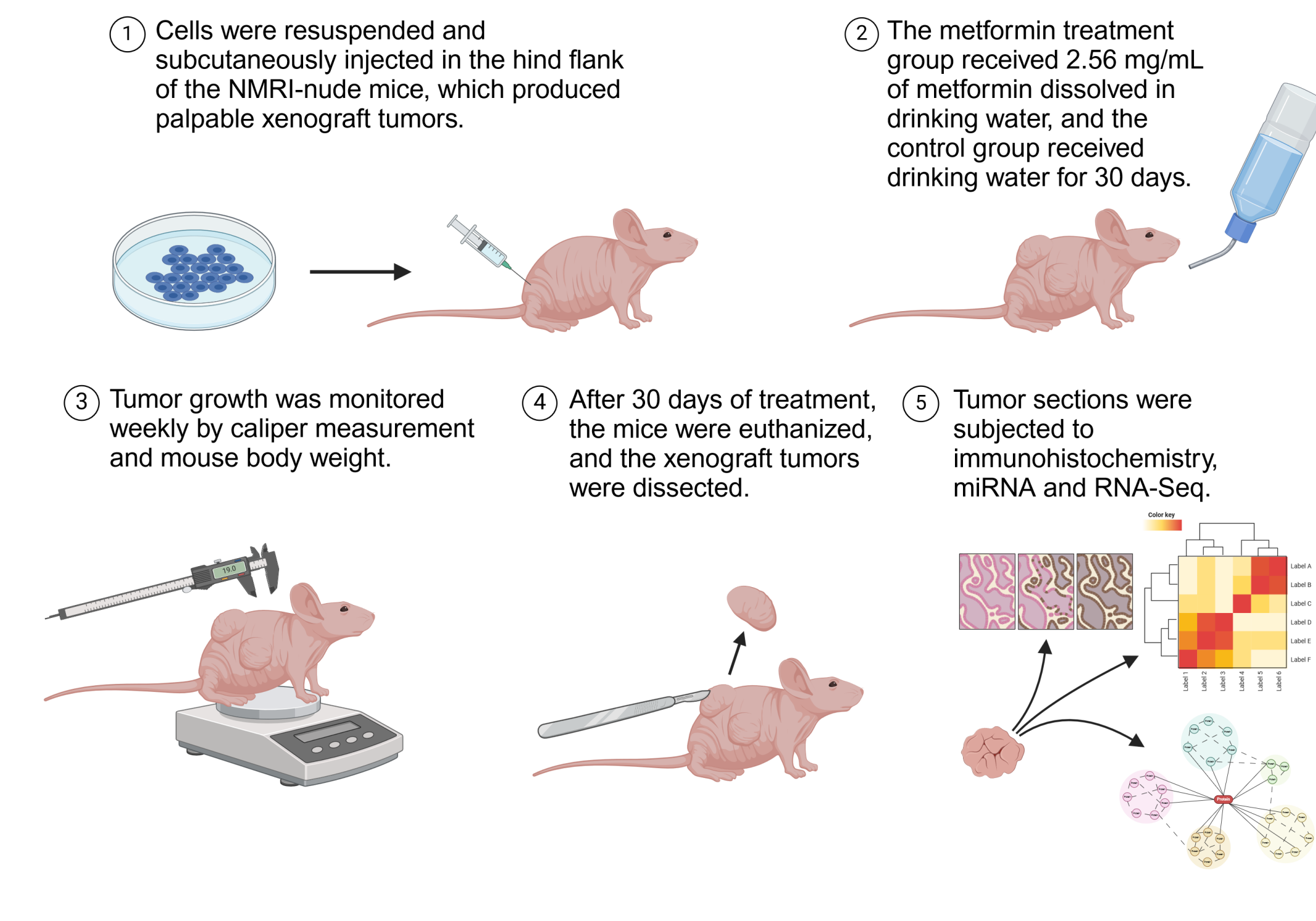
Introduction

- Small intestinal neuroendocrine tumors (SI-NETs) are rare and slow-growing, with most patients being diagnosed at a late stage with distant metastases.
- Metformin has been hypothesized as a potential anti-tumor agent by several studies in the past few years.
- Confirmed by experimental studies in which metformin has inhibited cancer cell growth in different cancers.
- We recently demonstrated that metformin treatment repressed the cell viability of SI-NET cells and inhibited the proliferation of cell spheroids.

Aim

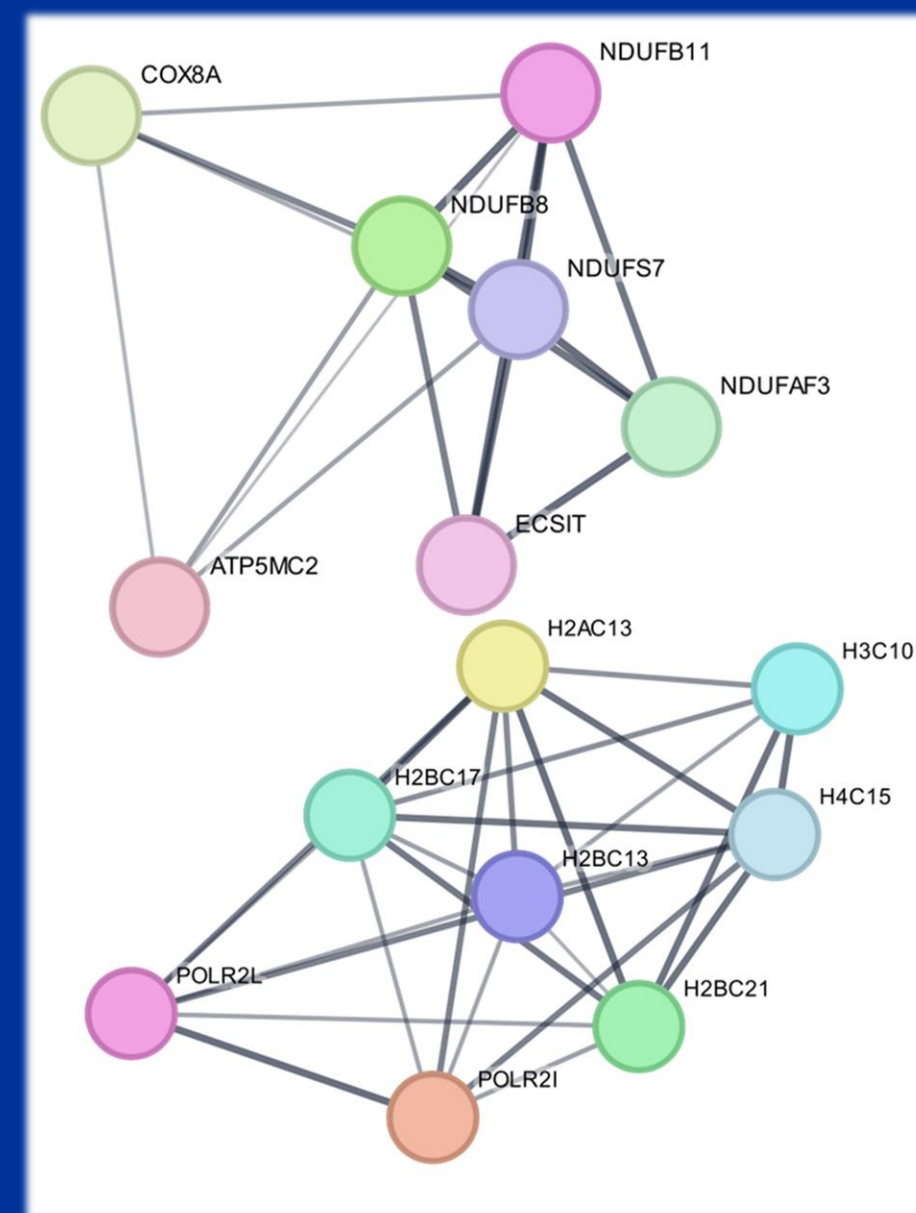
- To investigate the effect of metformin on the proliferation of SI-NETs in vivo, to identify novel molecular targets, and to unravel mechanisms responsible for the anti-tumor activity of metformin.

Materials and methods



Conclusion

- Metformin inhibited proliferation and Ki-67 expression in SI-NET xenografted tumors.
- CST3 (Cystatin C) expression was found to be induced by metformin.
- CST3 is inhibited in SI-NET compared to EC cells.
- CST3 is a potential clinically relevant target for further functional studies.
- Overlapping gene clusters:
 - Oxidoreductase
 - Nucleosome-RNA polymerase II



Results

- A smaller tumor size could be observed in the metformin treatment groups (Fig. A).
- All the tumors ($n = 22$) stained positively for the neuroendocrine cell marker **Synaptophysin**, and the expression of **Ki-67** was reduced in the tumor tissues while no effect on **activation of caspase-3** was detected (Fig. B).
- Significantly ($p \leq 0.05$) inhibited **CD44** expression were identified in both **CNDT2.5** and **GOT1** xenografted tumors, indicating successful metformin treatment.
- 227 significantly ($p \leq 0.05$) differentially expressed genes (DEGs) overlapping between **CNDT2.5** and **GOT1** xenografted tumors could be detected, and one significantly ($p \leq 0.05$) differentially expressed miRNA: **hsa-miR-99a-5p**.
- We could observe two overlapping gene networks: an **oxidoreductase** and a **nucleosome-RNA polymerase II** cluster.
- We compared our 227 genes against overlapping significantly differentially expressed genes in our in-house (unpublished) dataset consisting of comparisons between patient-derived enterochromaffin (EC) cells and SI-NET cells by single-cell RNA-seq (scRNA-seq).
- This revealed **CST3** (Cystatin C) as a potential clinically relevant target for further functional genomic studies. This is due to its expression being significantly inhibited in SI-NET, and our results revealed that **CST3** expression levels were significantly induced in both our **CNDT2.5** and **GOT1** xenografted tumors following metformin treatment (Fig. C).

