Investigating *SSTR1* as an Actionable Drug Target in Prostate Cancer Resistant to Androgen Receptor Signaling Inhibitors

Tatyanah Farsh^{1,2}, <u>Xiaolin Zhu^{1,3}</u>*, Tianyi Liu^{1,2}, Jun Zhu^{1,2}, Emma Dolan^{1,2}, Jin Shin^{1,2}, Rahul R. Aggarwal^{1,3}, Eric J. Small^{1,3}, David A. Quigley^{1,4}, Haolong Li^{1,2,5}, Felix Y. Feng^{1,2}

¹Helen Diller Family Comprehensive Cancer Center, University of California San Francisco, San Francisco, CA

²Department of Radiation Oncology, University of California San Francisco, San Francisco, CA ³Department of Medicine, Division of Hematology and Oncology, University of California San Francisco, San Francisco, CA

⁴Department of Urology, University of California San Francisco, San Francisco, CA ⁵Human Biology Division, Fred Hutchinson Cancer Center, Seattle, WA *Presenting author

Background: Resistance to androgen receptor signaling inhibitors (ARSIs), such as abiraterone or enzalutamide, is a major obstacle to improving outcomes for patients with prostate cancer. We previously analyzed paired RNA-seq data generated from metastatic biopsies before and after ARSI therapy in patients with metastatic castration-resistant prostate cancer (mCRPC) and identified *SSTR1* as the most significantly downregulated gene in ARSI-resistant tumors. *SSTR1* encodes somatostatin receptor type 1, one of the five somatostatin receptors, which, as a class, mediate anti-proliferative, anti-invasive, and anti-angiogenic effects when binding to the endogenous hormone, somatostatin, or exogenous somatostatin analogues. We were motivated to study the biology of *SSTR1* in prostate cancer, particularly its relationship with ARSI-resistance and therapeutic potential. We hypothesized that *SSTR1* has anti-tumor effects in prostate cancer, and its downregulation after ARSI treatment may promote tumor growth and ARSI resistance. We further assessed *SSTR1* as a potential drug target for overcoming ARSI resistance.

Method: To assess *SSTR1* expression in mCRPC, we selected the enzalutamide-resistant 22Rv1 cells as a model for ARSI-resistant prostate cancer. To assess the effects of genetic manipulation of *SSTR1*, we generated a stable knockdown cell line using CRISPR interference and a stable lentiviral overexpression cell line and then conducted proliferation assays. We also performed RNA-seq experiments on these cell lines to understand the transcriptomic impact of modulating *SSTR1*. To assess the impact of AR signaling on *SSTR1* expression, we treated 22Rv1 cells with dihydrotestosterone (DHT) for 48 hours and measured *SSTR1* mRNA. To evaluate the therapeutic potential of targeting SSTR1, we tested pasireotide, the only FDA-approved somatostatin analogue with SSTR1 agonist activity.

Results: Incucyte experiments confirmed the anti-proliferative effects of *SSTR1* overexpression and the growth-promoting effects of *SSTR1* knockdown. RNA-seq analysis showed that EGFR signaling was the most significantly upregulated pathway after *SSTR1* knockdown, with an upregulation of *SSTR5*, potentially as a compensatory mechanism. *SSTR1* overexpression induced no significant change in *AR* mRNA but a remarkable decrease in canonical AR target genes such as *KLK3*, *STEAP1*, and *TMPRSS2*, as well as significantly downregulated FGFR signaling. DHT treatment for 48 hours did not significantly change *SSTR1* mRNA expression. Pasireotide treatment significantly suppressed the proliferation of 22Rv1 cells.

Conclusions: *SSTR1* is anti-proliferative in enzalutamide-resistant 22Rv1 cells. SSTR1 signaling impacts the expression of AR target genes and EGFR and FGFR signaling pathways. Pasireotide treatment suppresses 22Rv1 cell proliferation. Further work using additional cell lines and *in vivo* models is warranted to further investigate SSTR1 in prostate cancer, particularly as a potential drug target for overcoming ARSI resistance.

Funding Acknowledgments:

X. Zhu was funded by a Prostate Cancer Foundation Young Investigator Award under the sponsorship of Dr. Elliot and Nan Abramowitz and a Department of Defense Prostate Cancer Research Program Physician

Research Award. F.Y. Feng was funded by Prostate Cancer Foundation Challenge Awards. Additional funding was provided by a UCSF Benioff Initiative for Prostate Cancer Research award.

Conflicts of Interest:

R. Aggarwal reports grants from Janssen, Amgen, Zenith Epigenetics, and Xynomic Pharmaceuticals; grants and personal fees from AstraZeneca, Merck, and Novartis; personal fees from Dendreon, Elsevier, Exelixis, Jubilant Therapeutics, Bayer, Pfizer, and Alessa Therapeutics outside the submitted work. E.J. Small reports other support from Fortis, Harpoon, Teon, Janssen, Johnson & Johnson, and Ultragenyx during the conduct of the study; other support from Fortis, Harpoon, Teon, Janssen, Johnson & Johnson & Johnson, and Ultragenyx outside the submitted work. F.Y. Feng reports personal fees from Janssen Oncology, Bayer, PFS Genomics, Myovant Sciences, Roivant Sciences, Astellas Pharma, Foundation Medicine, Varian, Bristol Myers Squibb (BMS), Exact Sciences, Clearnote Health, Novartis, and Tempus; other support from Serimmune and Artera outside the submitted work.