Development and Biological Evaluation of Novel α -Helix Mimetic Prodrugs as Leads for Prostate Cancer Treatment

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CANCER RESEARCH

NTRODUCTION

The androgen receptor (AR) transcription pathway plays a critical role in the proliferation of prostate cancer (PCa) cells.¹ Consequently, traditional PCa treatments focus on androgen suppression and deprivation by the use of AR antagonists and castration. These reduce AR activity however the onset of resistance is inevitable.²



SPIVEY α -HELIX MIMETIC

Our research has focused on the design and development of a novel small drug-like molecule, a coactivator binding inhibitor (CBI), targeting the activating function 2 domain (AF2) within the AR ligand-binding domain.³

To assess binding of **CBI-1** to the AR, a Fluorescence Polarisation (FP) assay

FP BINDING ASSAYS



Fig. 2a: Protein α-helix structure. **b**: **CBI** plug and play scaffold. **c**: **CBI-1**, an Asp-Phe-Glu-X-Leu-Phe (FXXLF) Spivey α-helix mimetic structure. **d**: Computationally optimized energy minimized 3D model of **CBI-1**.

CBI-1 contains an α -helix mimetic scaffold imitating the FXXLF epitope of AF2binding coactivators. With a relatively rigid 3D structure, it is anticipated to be resistant to proteolysis, and to bind to the target with high affinity and at a reduced entropic cost, inhibiting the AR and hence the development of PCa.



was developed. Exploiting the inverse relationship between molecular size and rotational speed, the extent of light depolarisation by a fluorophorelabelled ligand was monitored and correlated to AR binding.



Fig. 4: Schematic of the FP assay where decrease in polarisation indicates binding to the AR. Alongside **CBI-1**, several positive control ligands known to bind to the AR were screened, all of which resulted in a decrease in polarisation indicating successful binding to the AR.



Fig. 3: Optimized synthetic route of the α -helix mimetic **CBI-1**.

IN-CELL LR ACTIVITY ASSAYS

To assess the activity of CBI-1 inside cells, a Luciferase Reporter (LR) assay was employed where AR activity can be directly correlated to luciferase activity (luminescence) using genetically engineered cells.



Fig. 5: AR FP binding assay results showing light polarisation against ligand concentration. 5a: Testosterone, an endogenous AR agonist. **5b**: Dihydrotestosterone, an endogenous AR agonist. **5c**: Enzalutamide, a clinical AR antagonist drug. 5d: Katzenellenbogen-CBI, a published AR CBI. 5e: 3-(Indolin-2-yl)-1H-indole, a published AR antagonist targeting the BF3 domain. **5f: CBI-1** α -helix mimetic.





Positive control ligands showed a decrease in luciferase activity with increasing ligand concentration, however **CBI-1** did not, indicating a lack of inhibition.

REFERENCES

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ACKNOWLEDGEMENTS

CBI-2 Concentration (µM)

20 10 5.0 2.5 1.0 0.1 0.01

Thank you to Sue Powell for help with assay development and Imperial College London (DTP) and CRUK for the funding of this project.

7.5 17.5 2.5 We hypothesise that inhibition is not **CBI-2** Concentration (µM) observed in cells due to a lack of cell Fig. 8: CBI-1 prodrug derivative approach. permeability. This can be overcome by Initial cell viability assays of the masking the polar carboxylic acid ethyl ester prodrug showed activity groups as esters to give prodrugs. in prostate cancer cells (LNCaP) with Fig. 9: CBI-2 viability assay a decrease in cell growth. well plate with crystal violet.

> Future work involves investigating receptor selectivity, tests on prostate tumour explant models, and X-ray crystallography studies to confirm mode of binding.