6254: The discovery and preclinical characterization of the SAM-competitive PRMT5 inhibitor UCT-000445

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BACKGROUND

- PRMT5 is a methyltransferase that symmetrically dimethylates arginine residues using a cofactor called SAM
- PRMT5 dimethylates several histone proteins and regulates the expression of certain genes; dysregulation of PRMT5 causes epigenetic changes that are implicated in many cancers
- Inhibition of PRMT5 can reverse these epigenetic changes and has antiproliferative effects in many human cancer models
- Some cancers harboring an *MTAP*-deletion genotype have higher levels of the cell metabolite MTA, and MTA can partially occupy the SAM-binding site of PRMT5
- Inhibition of the methyltransferase activity of PRMT5 has been achieved with three modalities: (1) small molecule occupation of the argininebinding site ("substrate-competitive"); (2) small molecule occupation of the SAM-binding site ("SAM-competitive"); and (3) small molecules that cooperatively occupy the SAM-binding site with MTA ("MTA-cooperative")
- We evaluated the therapeutic potential of each modality by screening benchmark inhibitors against 500+ human cancer cell lines of various histological origins
- We found that the SAM competitive modality had the greatest therapeutic potential because of its more potent anti-proliferative effect against a broader range of cell lines
- We developed a novel, potent and more selective SAM-competitive PRMT5 inhibitor (UCT-000445) using our proprietary chemistry platform and structure-based design
- JNJ-64619178, a SAM-competitive PRMT5 small molecule inhibitor that is currently in Phase 1 clinical evaluation, was used as a benchmark

SPECIFICITY, POTENCY & OTHER PROPERTIES OF UCT-000445 **INHIBITION (%) OF METHYLTRANSFERASE**

	PRMT5 IC ₅₀	PRMT 1	PRMT 3	PRMT 4	PRMT 5	PRMT 6	PRMT 7	PRMT 8	PRMT 9	G9a
UCT-000445	25 nM	6	0	13	100	2	3	8	0	14
JNJ-64619178	32 nM	14	13	3	99	6	4	0	77	31

Table 1. IC₅₀ against PRMT5 methyltransferase activity was measured in a cell-free enzymatic activity assay. Enzymatic inhibition of other MTs was assessed with similar cell-free assays at various concentrations of each inhibitor (inhibition at 200 nM presented) and are reported as % inhibition.

	k _a (1/Ms) "k _{on} "	k _d (1/s) "k _{off} "	К _D (М)
JNJ-64619178	2.06E+06	1.66E-04	8.09E-11
UCT-000445	1.99E+06	9.85E-05	4.96E-11

	Avg. Cell IC ₅₀ (nM)	Hepatocyte T _{1/2} (mins)	P _{app} (10 ⁻⁶ cm/s) A to B
UCT- 000445	2.6	70 (Rat) 113 (Mouse)	0.360
JNJ- 64619178	3.1	31 (Rat) 38 (Mouse)	0.805

Table 2. Characterization of the association between inhibitors and a recombinant, chemically biotinylated PRMT5:MEP50 complex using SPR. UCT-000445 has enhanced retention time (" k_{off} ") as compared to the benchmark. R_{max} and Chi² were within acceptable ranges.

Table 3. Evaluation of in vitro and cellular properties. The "Avg. Cell IC_{50} " values were calculated from 50 cell lines of various histological origins evaluated in a 7-day growth assay. P_{app} data were obtained using a Caco-2 cell permeability assay (WuXi AppTec, Wuhan, China).

- PRMT5, 7 & 9 are Type II PRMTs; the role of PRMT7 & 9 in cancer cell proliferation is not well established, and therefore, targeting them may not be necessary
- Our inhibitor spares PRMT9 while potently inhibiting PRMT5 (Table 1); additionally, enzymatic inhibition of 42 recombinant human histone or methyltransferase complexes was evaluated, and none showed greater than 25% inhibition at 200 nM UCT-000445 (data not shown)
- The greater specificity of UCT-000445 may give it a wider therapeutic index then that seen with JNJ-64619178
- Notably, UCT-000445 has the same anti-proliferative effect as the benchmark (Table 3), suggesting that PRMT9 and G9a inhibition are dispensable for therapeutic action



Figure 1. 325+ human cancer cell lines of various histological origins were treated continuously during a 7day growth assay. Anti-proliferative effects are reported as IC₅₀s (μ M). "WT" are *MTAP*^{+/+} cell lines and "HD" are *MTAP^{-/-}* cell lines. (A) Results from cell lines treated with UCT-000445. Similar results were obtained with a chemically distinct SAM-competitive PRMT5 inhibitor (data not shown). (B) Results from cell lines treated with an MTA-cooperative PRMT5 inhibitor. Similar results were obtained with a chemically distinct SAM-competitive PRMT5 inhibitor (data not shown).

- and/or a major loss of cell proliferation^{1,2}
- sensitivity to PRMT5 inhibition
- **MAT2A inhibitors** (data not shown)
- lines

UCT-000445 REDUCES DIMETHYLATION OF PRMT5 SUBSTRATES B



Figure 2. Dimethylation of SmD1/D3 proteins ("SmD1/D3 Me₂") was assessed by western blot as a 0.0 0.0 0.0 pharmacodynamic marker of PRMT5 inhibition, which was previously validated^{6,7}. SmD3 and β -actin are NCI-H1048 NCI-H1155 NCI-H1435 loading controls. (A) Schematic of methyl transfer to target arginine by Type II PRMTs. Note that Type I PRMTs can mono-methylate target arginine before a Type II PRMT transfers the second methyl. (B) FaDu Figure 3. CD-1 athymic nude mice subcutaneously implanted with (A) human colon cancer cell cells were treated for the indicated time and proteins of interest were measured by western blot. "JNJ" = line models, (B) human squamous cell carcinoma models or (C) human lung cancer cell line JNJ-64619178 and "445" = UCT-000445. (C) CD-1 athymic nude mice bearing UPCISCC125 xenografts were models were treated with vehicle or with the indicated dose of UCT-000445 PO QD. All studies treated QD with vehicle control (Ctrl) or 12.5 mg/kg UCT-000445 (445) for 21 days and tumor lysates were were performed with 8 animals per arm. Tumor volumes were determined using caliper collected. Lysates from two animals of the Ctrl group and two animals of the 445 group are presented. measurements and recorded in mm³.

325+ human cancer cell lines of various histological origins

Knockdown or knockout of PRMT5 in an MTAP^{-/-} background leads to cell death

• This observation suggests that the MTAP-/- genotype could be used as a marker of

Paradoxically, this phenotype was not reproduced with a substrate-competitive inhibitor¹, nor could we reproduce it with several SAM-competitive inhibitors or

• *MTAP*^{-/-} genotype is predictive of sensitivity to MTA-cooperative inhibitors^{3,4,5}

We profiled MTA-cooperative inhibitors and found that only about 30% of 75 MTAP^{-/-} human cancer cell lines had an IC₅₀ of less than 0.5 μ M, and antiproliferative effect was seen in a surprisingly large number of MTAP-normal cell



Figure 4. (A) Reticulocytes are the immediate precursor to erythrocytes and a surrogate for early hemolytic anemia. (B – D) CD-1 athymic nude mice subcutaneously implanted with FaDu. (B) Reticulocyte levels were quantified in triplicate from animals treated with UCT-000445 either QD, 4 days on/3 days off, 7 days on/7 days off or 14 days on/7 days off. Blood samples were taken from animals at the indicated timepoints and CBCs were performed for each sample. Note the rapid rebound in reticulocyte levels upon inhibitor holiday. (C) Tumor response data. (D) Animal weight change data. The box below the line plot indicates the dosing periods of each arm.

UCT-000445 PRODUCES ANTI-TUMOR EFFECTS IN SUMMARY & CONCLUSIONS VARIOUS SOLID TUMOR MODELS



Squamous Cell Carcinoma Models 800.0



200.0 UMSCC11A UPCISCC125 **Non-Small Cell Lung Cancer Models**

600.0

400.0





- UCT-000445 is a highly selective and potent SAM-competitive PRMT5 inhibitor
- *MTAP*-deletion is not predictive of sensitivity to SAM-competitive PRMT5 inhibitors
- Even at sub-MTD doses, UCT-000445 has a significant anti-tumor effect against multiple solid tumor types
- The anti-proliferative effects of UCT-000445 in xenograft models are sustained for approximately 2 weeks after drug withdrawal
- This sustained effect allows intermittent dosing regimens to have efficacy equivalent to that of continuous dosing
- Moreover, drug holiday allows erythrocyte precursor cells to rebound, reversing the hemolytic anemia observed in animal models
- This study suggests that SAM-competitive PRMT5 inhibitors have the potential to provide benefit to patients with certain solid tumor types regardless of their tumors' MTAP gene status
- Our preclinical models support the use of intermittent dosing strategies to ameliorate potential side effects like cytopenias
- 1200 Pharma is preforming IND-enabling studies and plans to open a Phase 1 clinical trial in early 2024

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