Allosteric mitochondrial ClpP agonist ONC206 alters stress response, metabolic and epigenetic profiles to elicit anti-cancer efficacy in high-grade gliomas

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Introduction

- Dordaviprone (ONC201), a first-in-class imipridone, is an oral, blood-brain barrier penetrating, selective Figure 4. ClpP knockout impairs dordaviprone and ONC206 efficacy in glioma cells in vitro small molecule antagonist of dopamine receptor D2 (DRD2) and agonist of the mitochondrial protease caseinolytic mitochondrial matrix peptidase proteolytic subunit (ClpP).¹⁻⁵
- Dordaviprone has demonstrated tolerability and durable tumor regressions in patients with H3 K27Mmutant glioma.⁶
- ONC206, a chemical derivative of dordaviprone, is the second imipridone to be developed and is currently in Phase 1 clinical development.⁷
- Relative to dordaviprone, ONC206 has demonstrated differentiated DRD2 receptor pharmacology⁸, improved potency, enhanced absorption/tissue distribution, and preclinical anti-cancer activity in vitro and in vivo.⁹⁻¹³

Results

structure relative to the dordaviprone-bound or apo complex



Figure 9. ONC206 induces expression of ATF4, CHOP, and H3 K27me3 while reducing ClpX in parental, but not resistant cells



Results

Western blot results for SF8628 parental and resistant lines. 201-A16 were selected with 16 µM dordaviprone, 201-C2 were selected with 2 µM dordaviprone, 206-B1.6 were selected with 1.6 μ M ONC206. Cells were exposed to vehicle or 0.55 μ M ONC206 for 72 h prior to protein extraction and Western blotting. ClpX and GAPDH antibodies were from AbCam. ATF-4, CHOP, and K27me3 antibodies were from Cell Signaling Technology.

Table 1. Positions of ClpP mutations identified in dordaviprone/ONC206 resistant glioma cells

Parental Line	Name	Selection (µM)	ClpP Mutation(s)	Mutation Location	Mutant Proportion
T98G, GBM	201-R	Dordaviprone (16 µM) _	A131D	ClpP monomer interfaces	17-20%
			R226G	ClpX interface	12-21%
T98G, GBM	206-R	ONC206 (1.6 μM)	R226G	ClpX interface	43-54%
SF8628, DIPG K27M	201-A16	Dordaviprone (16 μM)	I193V	ClpX interface	50-67%
SF8628, DIPG K27M	201-C2	Dordaviprone (2 μM)	N172Y	ClpP monomer interfaces	22-33%
SF8628, DIPG K27M	206-B1.6	ONC206 (1.6 μM)	A188D	ClpX interface	~50%

Figure 1. Co-crystallization with ClpP revealed distinctions in the ONC206-ClpP resolved X-ray crystal



Cell viability for 5000 SF8628 or T98G cells, with or without ClpP KO, treated with dordaviprone or ONC206 at indicated concentrations Viability was measured on Day 5 using CellTiter-Glo. CRISPR-mediated ClpP knockout was made using Edit-R All-in-one Lentiviral sgRNA vector (Dharmacon, Horizon) and confirmed using Western blot. ClpP antibody was from AbCam. Actin antibody was from Santa Cruz Biotechnology. KO, knock-out; NTC, non-targeted control; WB, western blot; WT, wild-type.

Figure 5.DRD2 knockout does not impair dordaviprone and ONC206 efficacy in glioma cells in vitro



Figure 2. ONC206 exhibits nanomolar potency in cell-free proteolysis assays and against glioma cell lines

ClpP Activity (AMC-peptide) ClpP Activity (AMC-peptide) EC₅₀ = 2.3 μM EC₅₀ = 0.31 μM -2.5 -2.0 -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5 2.0 -2.5 -2.0 -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5 2.0 ONC206 log[µM] Dordaviprone log[µM] ClpP Activity (FITC-casein) ClpP Activity (FITC-casein)





ClpP crystal structure is from PDB ID: 6DL7 and displayed using iCn3D (NCBI). Whole genome/whole exome sequencing was performed at Novogene.

Figure 10. ClpP with resistance mutations are catalytically impaired and reduce dordaviprone-mediated agonism



A. Cell-free degradation activities of recombinant human wild-type and mutated ClpP on FITC-casein and AMCpeptide substrates were determined. **B.** Agonism of wild-type and mutated ClpP was determined in the same assay with dordaviprone added at increasing concentrations.

Figure 11. Expression of ClpP mutants is sufficient for imipridone resistance while over-expression of wild-type ClpP partially reverses resistance





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Protein localization-

Metabolomics results of SF8628 cells treated with dordaviprone or ONC206 at the indicated concentrations for 24 h are shown. 2X IC50 concentrations were 4.5, 0.55, and 0.12 μM, respectively. Only metabolites with consistent directional change, fold change >2, and significant difference by t-test (p<0.05) from control for at least 2 out of 3 imipridones are shown.

Figure 7. Proteomics analysis in response to ONC206 treatment in SF8628 cells¹⁴



ONC206). Reactome pathway analysis for significantly (p<0.05) downregulated proteins with dordaviprone and ONC206 treatment for 24h compared to vehicle treatment shown (FDR: false discovery rate)







2 100-

50-

- 24 Hours

48 Hours

→ 72 Hours

→ 96 Hours

→ 120 Hours

2 100-

50-

Cell viability for 5000 SF8628 cells treated with dordaviprone or ONC206 at indicated concentrations and duration before removal of drugcontaining media and replenishment with standard media. Viability was measured on Day 7 using CellTiter-Glo. Bar graphs indicate % cell viability after treatment with 10µM dordaviprone or ONC206 for the same duration across multiple cell lines.

5 or 2 μM dordaviprone, ely. The SF8628-A16 ND derived from A16 and without dordaviprone at for 25 days. T98G- d T98G-206R lines were up to 16 μM dordaviprone M ONC206, respectively. lity for 5000 cells treated ordaviprone or ONC206 ated concentrations was d on Day 5 using CellTiter-	0 				
	Conclusions				
	 ONC206 exhibits nanomolar potency in glioma cell lines ClpP expression and agonism is key for the anti-cancer efficacy of ONC206 in vitro Acquired resistance to ONC206 in vitro is associated with alterations in ClpP ONC206 treatment is associated with alterations in metabolism, ISR, and histone methylation 				



Conflicts of Interest

SF, AL, CM, JEA, and VVP are employees of and have stock ownership in Chimerix, Inc. JEA and VVP are shareholders of Oncoceutics, Inc.

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