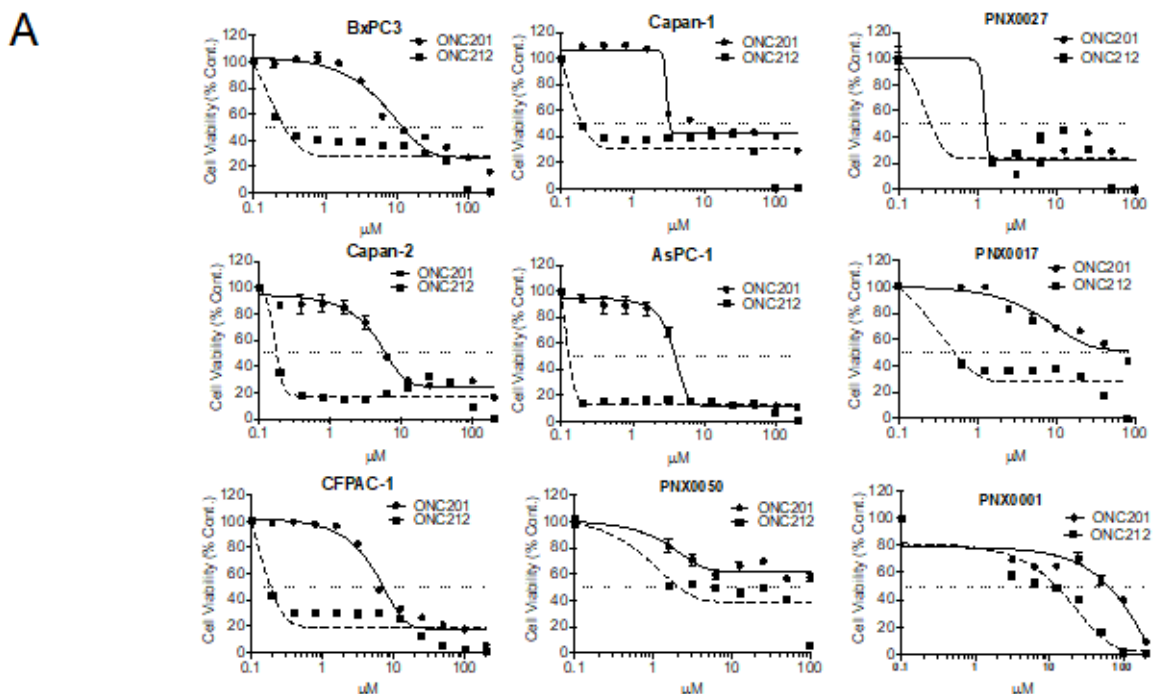


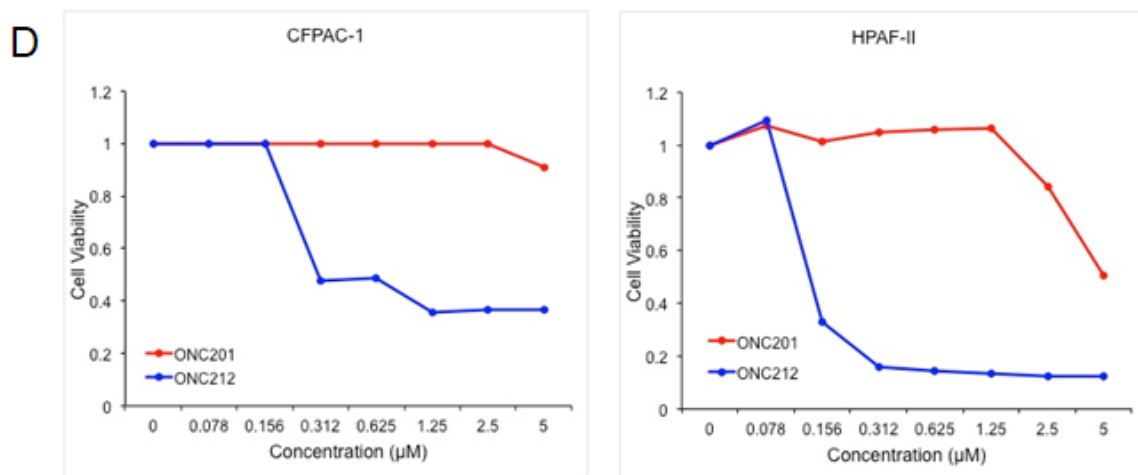
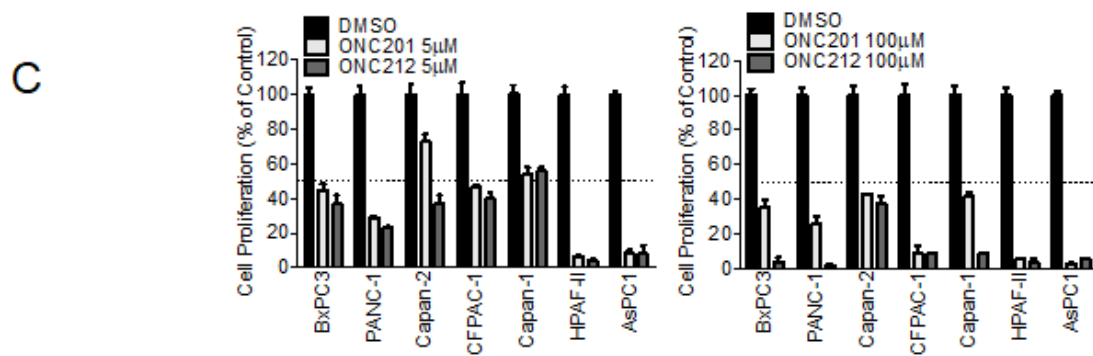
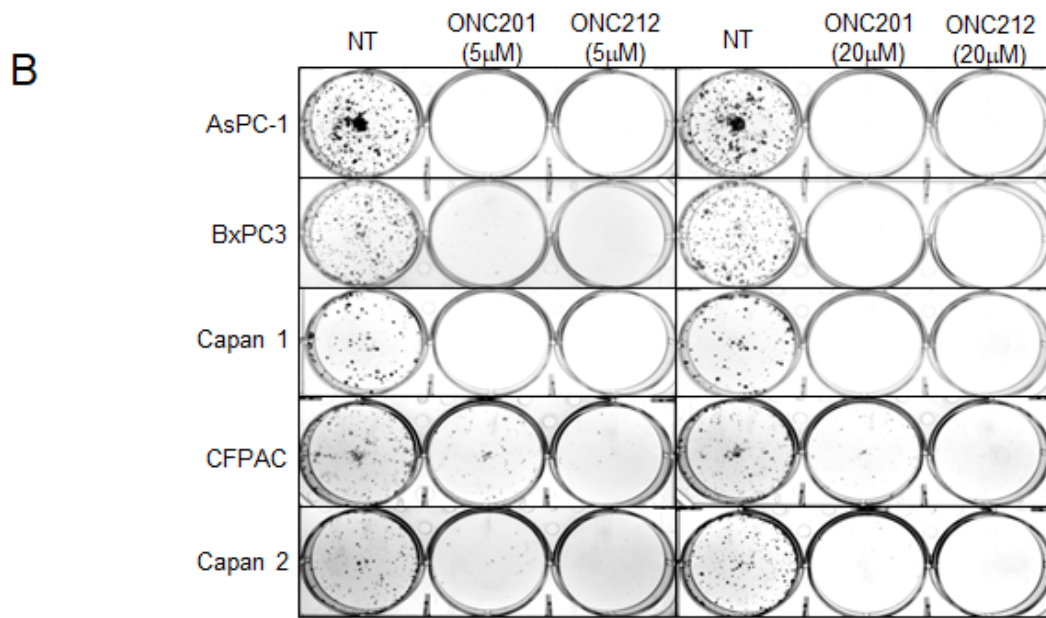
Anti-pancreatic cancer activity of ONC212 involves the unfolded protein response (UPR) and is reduced by IGF1-R and GRP78/BIP

SUPPLEMENTARY MATERIALS

Supplementary Table 1: GI₅₀ doses of ONC201 and ONC212 in a panel of seven human pancreatic cancer cell lines and nine low-passage PDX pancreatic cancer cell lines

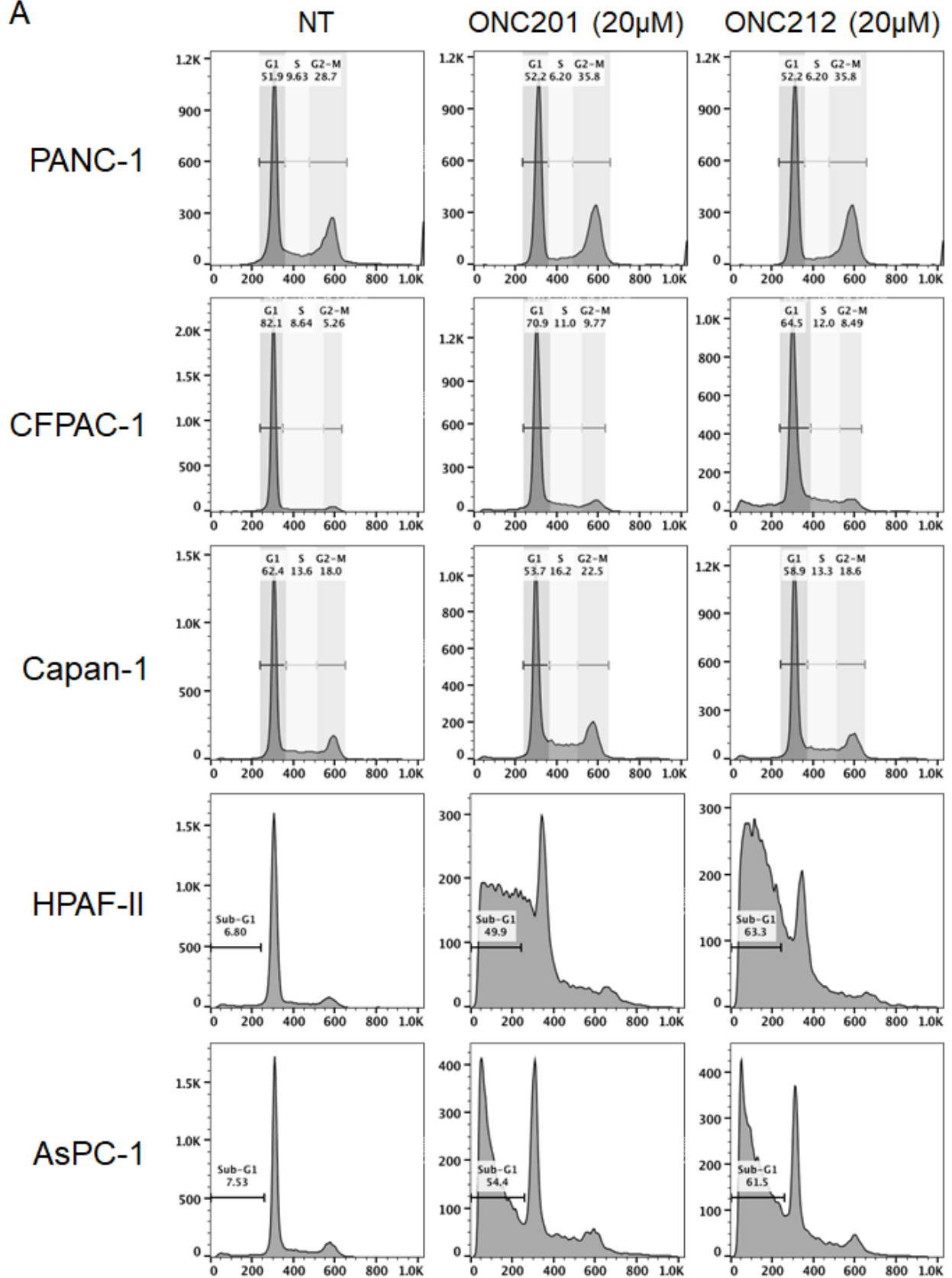
Cell Line	GI ₅₀ (μM)		PDX Cell line	GI ₅₀ (μM)	
	ONC201	ONC212		ONC201	ONC212
BxPC3	12.5	0.25	PNX0001	70	14
PANC-1	3.125	0.17	PNX0050	10	2
Capan-2	6.25	0.18	PNX0027	1.5	0.28
CFPAC-1	7	0.19	PNX0015	3.75	0.4
Capan-1	3	0.2	PNX0017	50	0.55
HPAF-II	4.5	0.15	PNX0007	100	16
AsPC-1	4	0.14	PNX0016	3	20
			PNX0029	100	25
			PDSE-JH101	18	9



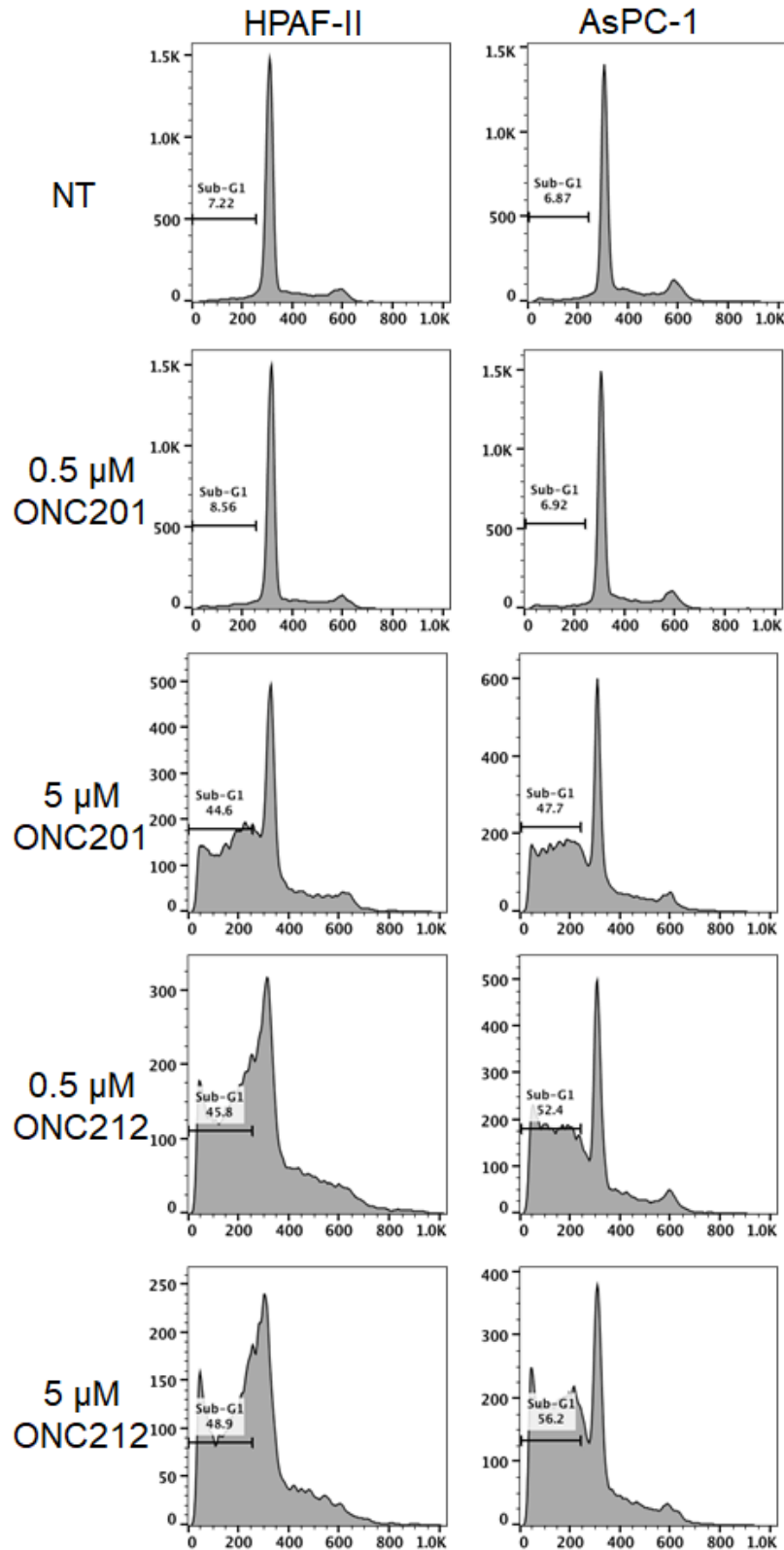


Supplementary Figure 1: (A) Dose-response curves for five pancreatic cancer lines (BxPC3, Capan-2, CFPAC-1, Capan-1 and AsPC-1) and PDX cell lines treated with 0-100 µM of ONC201 or ONC212. (B) Representative image of colony formation in ONC201/ONC212 treated pancreatic cancer cell lines. (C) Short-term cell proliferation measured by MTT assay 72 hours post ONC201 or ONC212. Graphs represent percent proliferation normalized to untreated control at 5 µM (Left) and 20 µM (Right) doses of ONC201 or ONC212. (D) Cell viability assays comparing ONC201 and ONC212 in selected pancreatic cancer cell lines. The data in Panel D was generated through the GDSC screen.

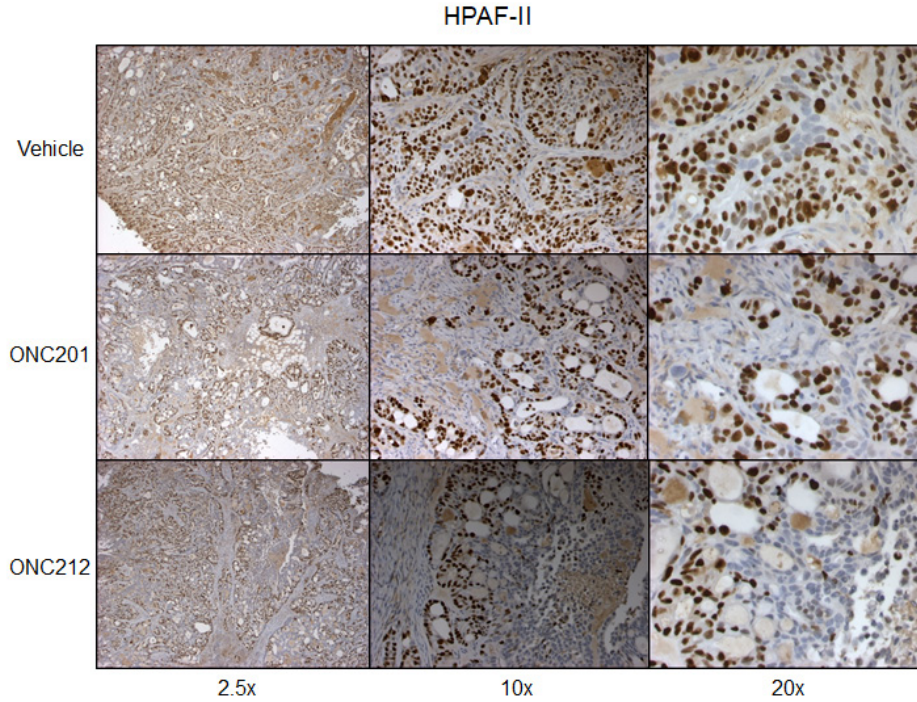
A



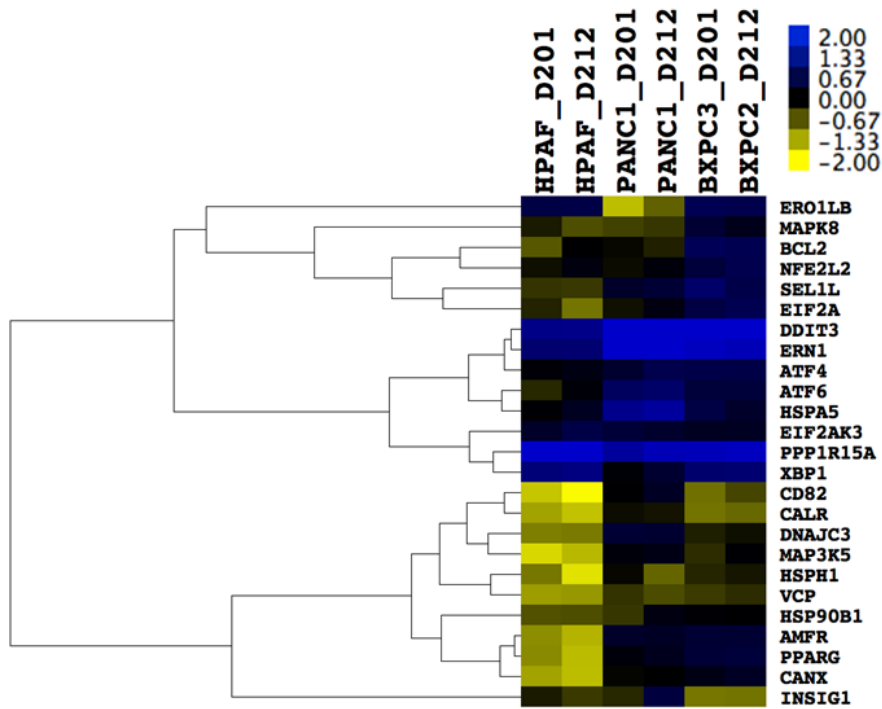
B



Supplementary Figure 2: (A) Histograms showing changes in different phases of the cell cycle in five human pancreatic cancer cell lines, 72 hours post-treatment. (B) Histograms indicating sub-G1 percentage in HPAF-II and AsPC-1 cell lines 72 hours post-treatment at the indicated doses of ONC201 or ONC212.



Supplementary Figure 3: Ki67 staining of HPAF-II tumor sections treated with either ONC201 or ONC212.



Supplementary Figure 4: Gene expression profiling by microarray analysis in HPAF-II, PANC-1 and BxPC3 cells 48 hours post treatment with 20 μ M ONC201 or ONC212 in comparison to untreated cells. Representative heat map shows expression of UPR genes relative to control for each cell line.