



Determinants of Gemcitabine-Pemetrexed Synergism in Pancreatic Cancer Cell Lines

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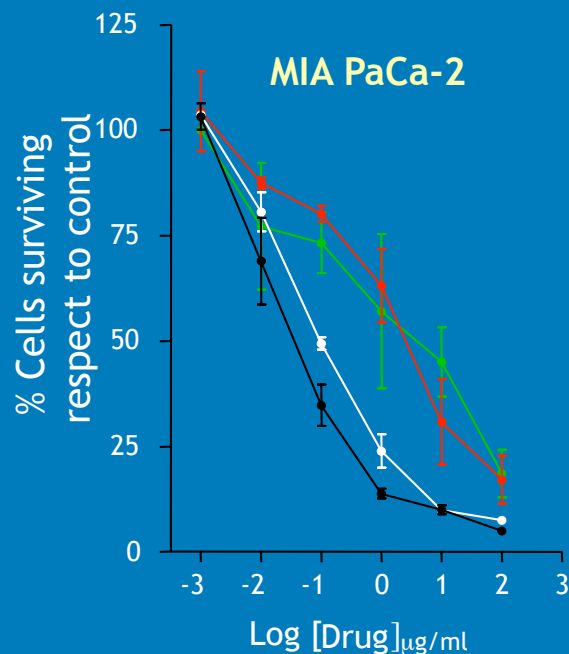
Pancreas Cancer 2004 International Symposium
Tirrenia, Pisa, 24-26 April 2004

DNA

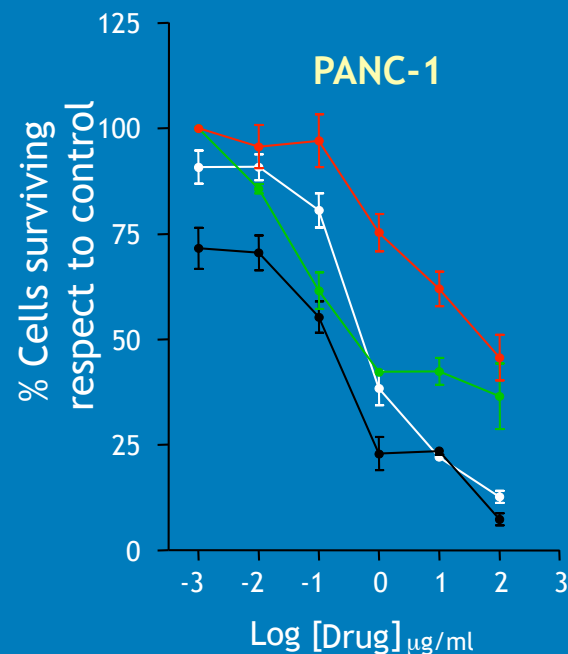


Tonkinson et al., Cancer Res 1999; 59:3671
Tesei et al., Clin Cancer Res 2002; 8:233

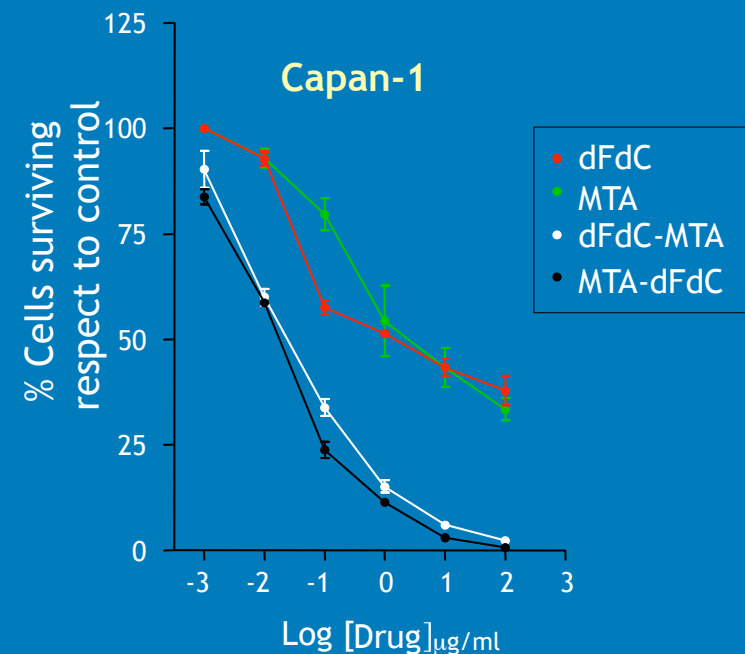
Cytotoxicity and pharmacologic interaction between gemcitabine and pemetrexed



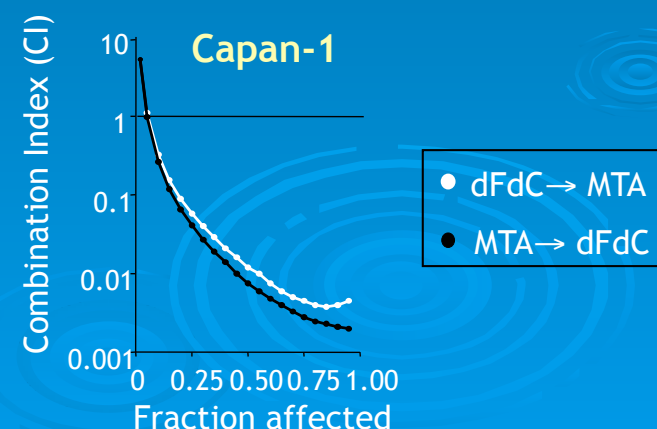
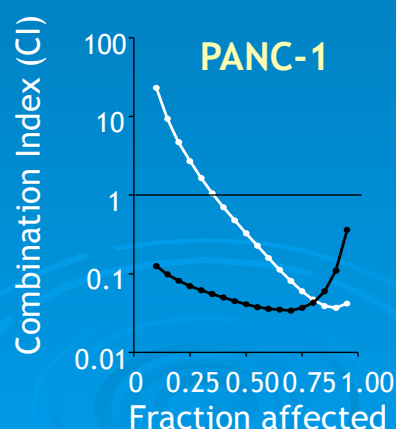
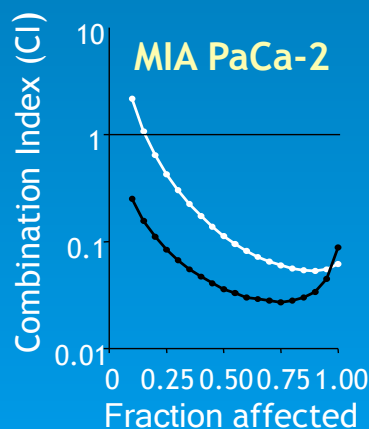
dFdC (IC_{50})= 2.90 μ g/ml
 MTA (IC_{50})= 1.58 μ g/ml
 dFdC-MTA (IC_{50})= 0.12 μ g/ml
 MTA-dFdC (IC_{50})= 0.04 μ g/ml



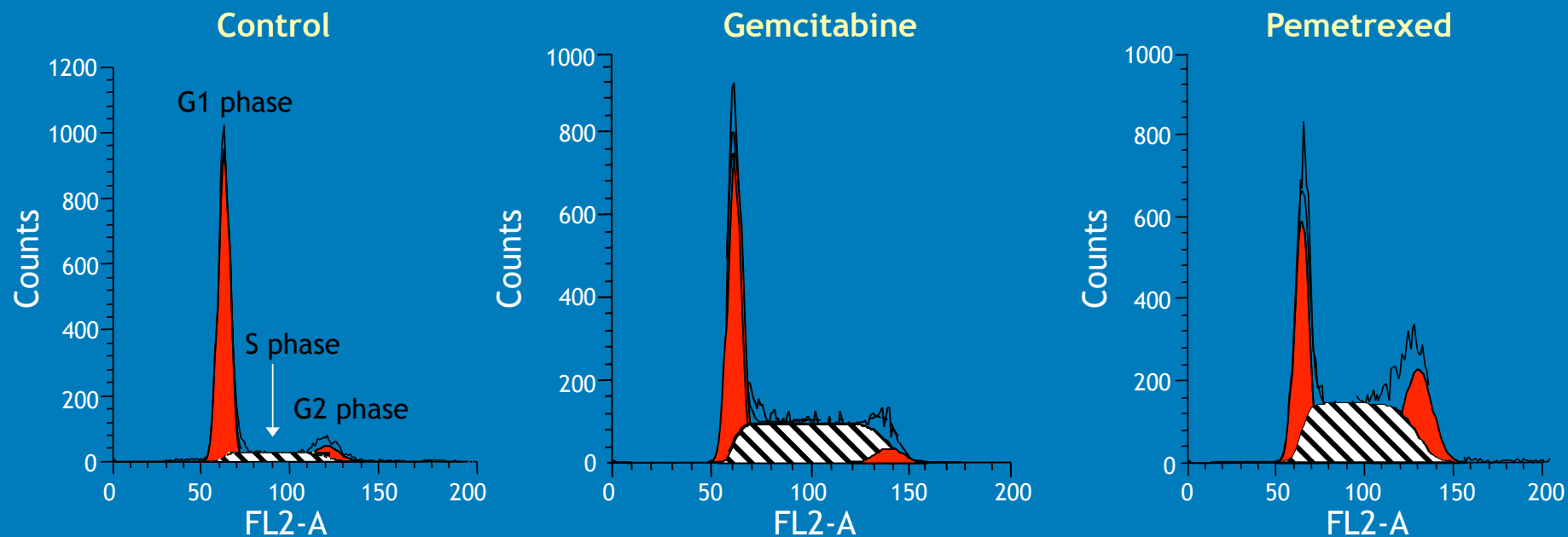
dFdC (IC_{50})= 42.21 μ g/ml
 MTA (IC_{50})= 2.42 μ g/ml
 dFdC-MTA (IC_{50})= 0.75 μ g/ml
 MTA-dFdC (IC_{50})= 0.09 μ g/ml



dFdC (IC_{50})= 4.75 μ g/ml
 MTA (IC_{50})= 7.33 μ g/ml
 dFdC-MTA (IC_{50})= 0.03 μ g/ml
 MTA-dFdC (IC_{50})= 0.02 μ g/ml



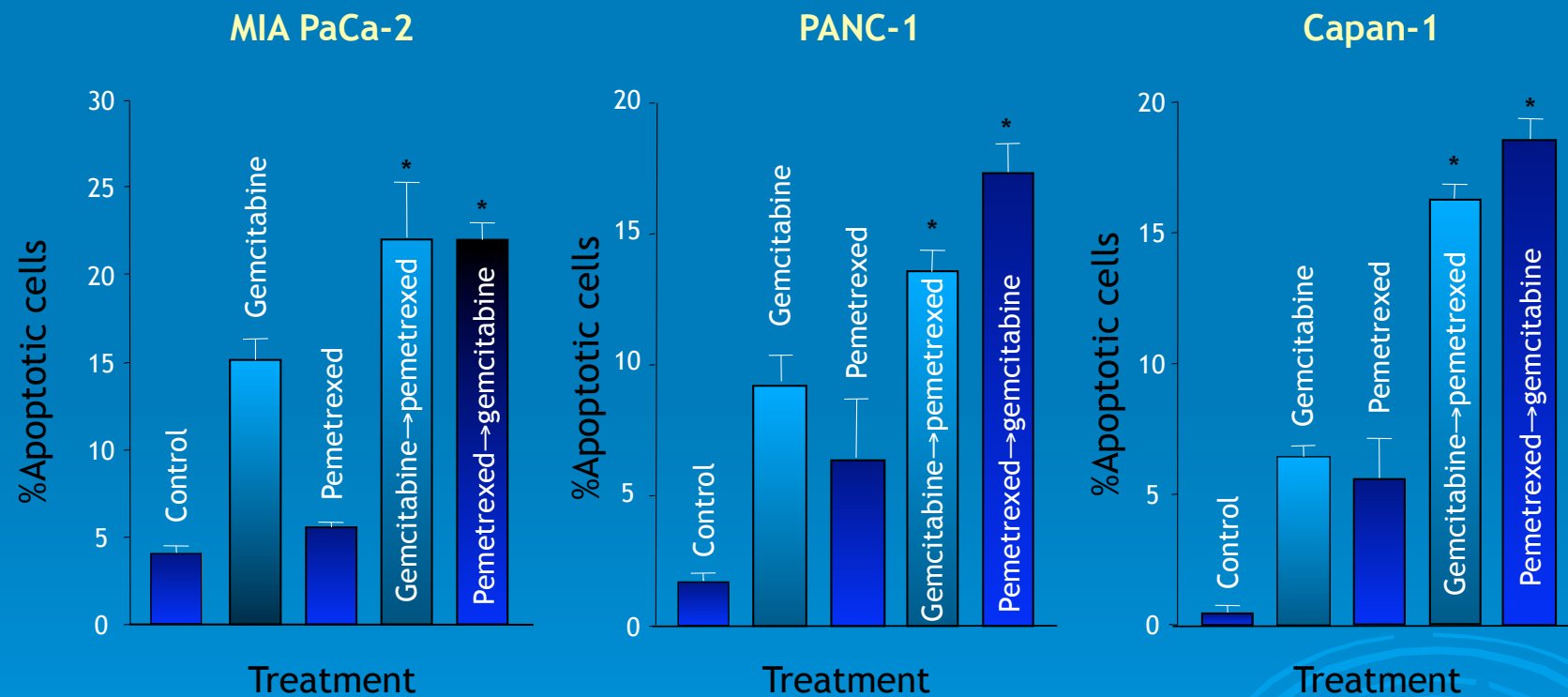
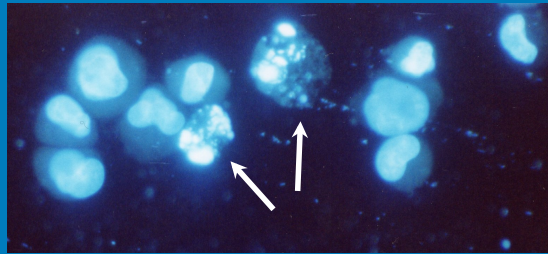
Cell cycle modulation by gemcitabine and pemetrexed



	Treatment	G1 (%) ^a	S (%)	G2 (%)
MIA PaCa-2	Control	77.01	15.30	7.69
	Gemcitabine	46.36	49.29	4.35
	Pemetrexed	30.12	46.63	23.25
PANC-1	Control	88.25	10.55	1.20
	Gemcitabine	66.98	29.29	3.73
	Pemetrexed	17.21	80.13	2.66
Capan-1	Control	50.73	31.13	18.14
	Gemcitabine	54.19	36.40	9.41
	Pemetrexed	31.10	63.21	5.69

^a Mean percent values of total number of cells examined in three independent experiments

Induction of apoptosis by gemcitabine, pemetrexed and their combination



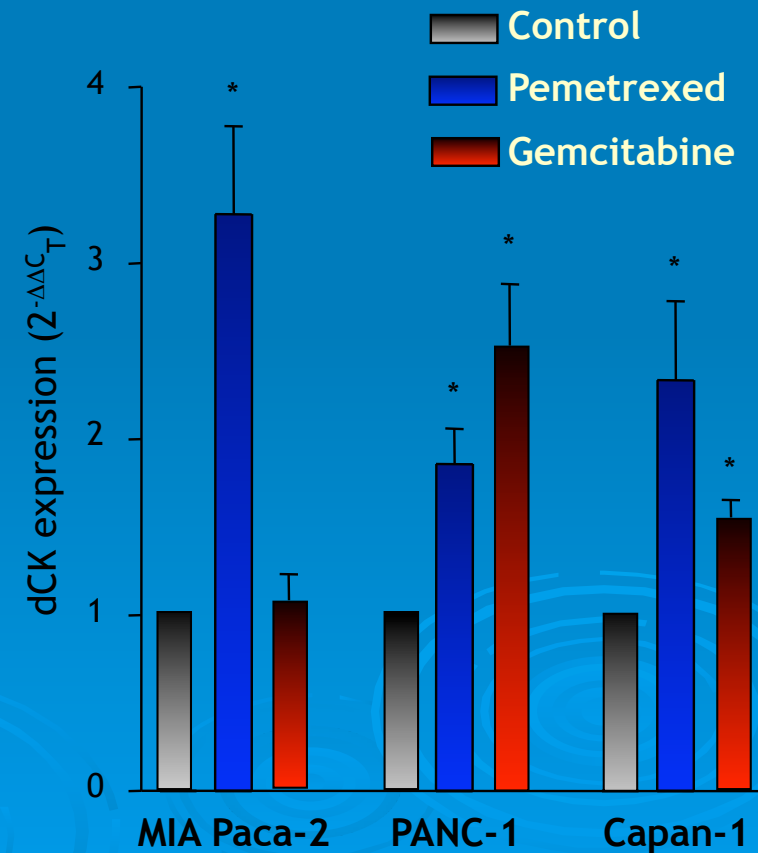
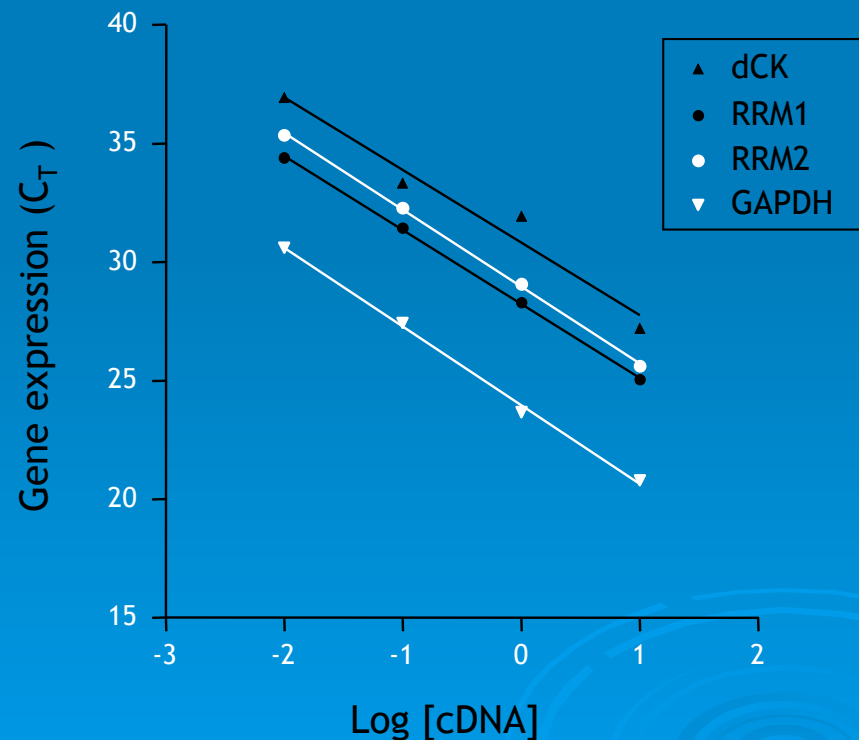
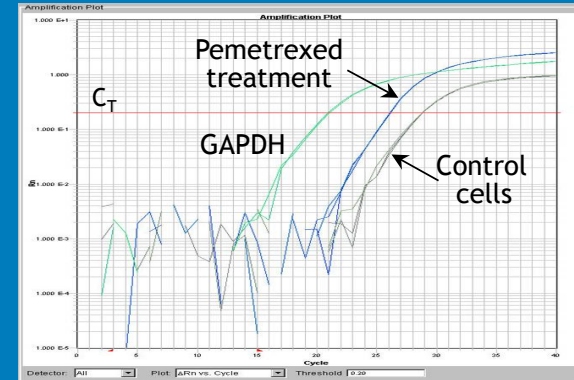
Columns, mean values obtained from three independent experiments; bars, SE

*Statistically significant different from controls ($P < 0.05$)

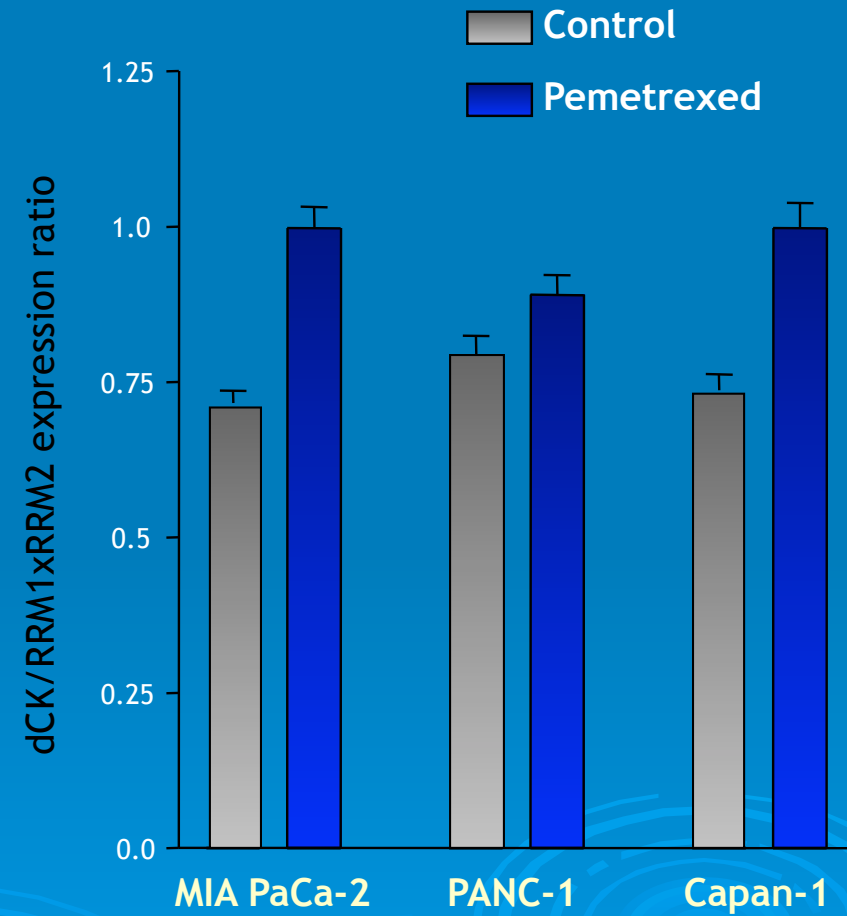
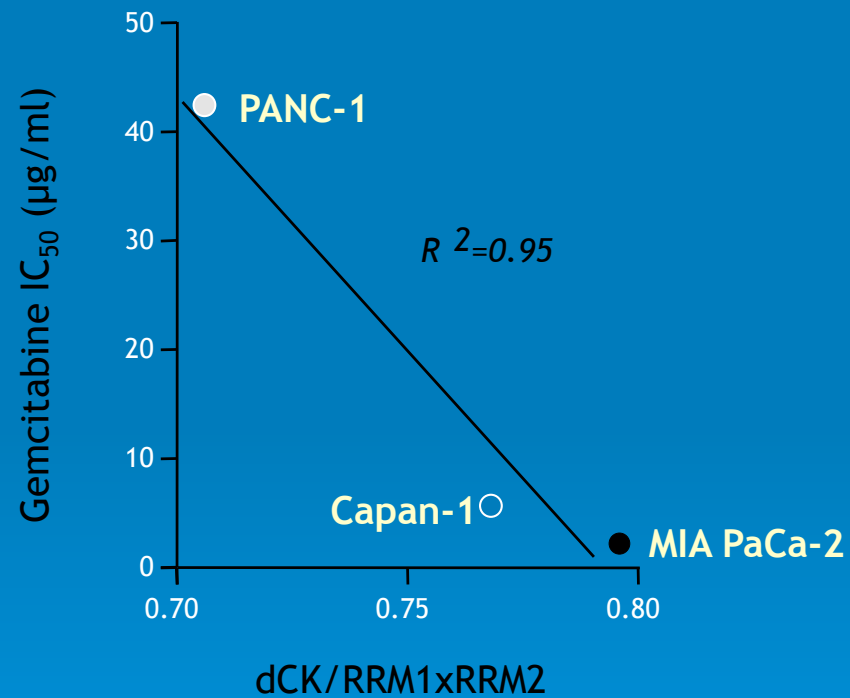
Modulation of dCK expression by pemetrexed

	IC ₅₀ (ng/ml) ^a			
	Gemcitabine	+dCyd	+DEPC	+THU
MIA PaCa-2	12.29	86.11	10.15	7.54
PANC-1	53.43	503.97	28.03	10.52
Capan-1	29.90	272.53	13.71	9.40

^a Mean values ±SE of at least three independent experiments



Enhancement of dCK/RRM1×RRM2 expression ratio after pemetrexed treatment



Conclusions

Gemcitabine and pemetrexed were cytotoxic against MIA PaCa-2, PANC-1 and Capan-1 cells and the combination index demonstrated that the drug sequence showing the maximum degree of synergism was pemetrexed→gemcitabine in all cell lines

Flow cytometric studies demonstrated that pemetrexed and gemcitabine enhanced cellular population in S phase in all cell lines

Gemcitabine-pemetrexed combinations increased the occurrence of apoptosis

Quantitative RT-PCR analysis showed that pemetrexed significantly enhanced dCK expression in all cell lines, while there was only a minor increase of RR expression

These data provide evidence that the combination of gemcitabine and pemetrexed displays schedule-dependent synergistic cytotoxic activity against various pancreatic cancer cells, associated with favorable modulation of cell cycle, induction of apoptosis and inducible dCK gene expression.