

Table S4. *In-vivo* examples of dose-response correlations of individual genes per time point

Dose response correlations per time point of individual genes that belong to gene sets: KEGG nucleotide excision repair, KEGG cell cycle, KEGG extrinsic apoptosis and IARC p53 responsive elements, in the *in-vivo* experiment (set up as Table 2).

Nucleotide Excision Repair							
gene	1h	3h	6h	9h	12h	24h	48h
Pole3	-0.97			0.97	0.97	0.92	
Pold4		0.89	0.91				
Mnat1		-0.87	-0.95				
Pole4		0.86	0.93	0.85			
Ercc1		0.86		0.82	0.81		
Ercc6			0.96				
Rfc5			0.9	-0.8	0.95		0.94
Ercc5			0.88	0.81			
Rfc1			-0.86				0.85
Ccnh			-0.84			0.95	
Ddb2				-0.96			
Xpc				-0.93			
Gtf2h3				0.88			
Rfc3				0.87	0.98		0.92
Pcna				0.81	0.86		
Cdk7				0.81			
Xpa					-1		
Pole2				0.99			
Gm9840				0.97	0.92		
Gtf2h1				0.95	0.87		
Gtf2h5				0.92	0.9		
Rfc2				0.91			0.84
Rpa2				0.91	0.82		
Lig1				0.85			0.81
Rfc4				0.84			
Pole				0.8			0.92
Rbx1				0.8			
Ddb1							0.88
Ercc2							0.86
Pold3							0.84
Rpa1							0.81

Cell Cycle							
gene	1h	3h	6h	9h	12h	24h	48h
Ccnd3	1						
E2f3	-0.91						
Rb1	-0.9		-0.91				
Mad2l2	0.88						0.93
Cdc25c	-0.86						0.8
Ccnb2	-0.85		-0.86				
Fzr1	0.83		0.86				
Tgfb1	0.81		-0.84				
Smad3	0.81						
Pkmyt1		0.98					0.94
Smc3		-0.97				-0.88	
Cdc14b		-0.96	-0.88	-0.81	-0.91		
E2f5		0.93					
Myc		0.91					
Anapc7		0.88			0.91		
Anapc4		-0.86					
Gadd45g		0.83					
Cdc14a		0.82				-0.81	
Tgfb3			-0.93				0.83
Mdm2			0.92	0.89	0.8	0.88	
Cdkn1a			0.91	0.86	0.95	0.94	0.85
Chek2			-0.89		0.9		
Stag2			-0.88				
Atr			-0.86		-0.94		
Mad2l1			-0.85		0.96		
Ccna2			-0.85				0.88
Cdkn2d			-0.85				
Bub1b			-0.84	-0.9			
Ccnh			-0.84				0.95
Ccnd2			-0.83				
Cdk6			-0.83			0.85	
Cdc20			-0.82		0.8		0.92
Cdc25b			-0.81				
Cdc7			0.8		0.93		0.94
Stag1			-0.8				
Rbl2				-0.97		-0.86	
Bub1				-0.91			0.86
Rbl1				-0.91			0.92
Tfdp2				-0.89		-0.85	
Ywhab				0.85		0.85	
Cdc16				0.82			
Pcna				0.81	0.86		
Cdk7				0.81			
Mcm4					0.98		0.86
Gm9840					0.97	0.92	
Ccnd1					0.95		
Cdkn1c					-0.95		
Gm2423					0.94		
Ccnb3					-0.94		
Mcm7					0.93		0.94
Dbf4					0.92		0.84
Cdc23					0.92		
Cdc2a					0.91		0.87
Ywhaq					0.91		
Bub3					0.89		0.88
Cdc45l					0.89		0.95
Ep300					0.88	0.89	
Prkdc					-0.85	-0.86	
Orc6l					0.85		
Ywhag					0.84		
Orc3l					-0.84		
Orc2l					0.83		
Cdkn2b					0.82		
Orc1l					0.82		0.88
Orc4l					0.82	0.82	
Rbx1					0.8		
E2f4						0.81	
Ccne2							0.84
Mcm3							0.94
Mcm2							0.85
Mcm5							0.9
Tfdp1							0.8
Tgfb2						-0.92	
Trp53						0.88	
Pttg1							0.85
Sfn						0.86	
Anapc11							0.82

Extrinsic Apoptosis							
gene	1h	3h	6h	9h	12h	24h	48h
Ilrak4	0.93						
Il1rap	-0.91	-0.82	-0.9	-0.85	-0.8		
Akt3	0.89				-0.92		
Ppp3cc	-0.87		-0.82				
Akt2	0.86						
Pik3cb	-0.83	-0.83		-0.85	-0.92	-0.8	
Prkar1b	0.83						
Csf2rb	0.82						
Bcl2	-0.81		-0.87			-0.94	
Ilrak2		-0.93					
Ngf		0.9	0.9	0.81	0.99	0.84	
Dffa		-0.87	-0.91		-0.88		
Prkacb		-0.87	-0.85				-0.9
Bid		0.87					
Pik3cd		-0.87					
Prkar1a		0.85		0.9			
Fas		-0.85					
Tnfsf10		-0.83					
Pik3r5		-0.8				0.81	
Tnfrsf10b			0.9	0.89	0.86	0.89	
Myd88			0.9			0.82	
Pik3ca			-0.9				
Ppp3ca			-0.89	-0.89	-0.98		
Pik3r2			-0.87				
Nfkb1			-0.85				
Tradd			0.84		0.85		
Birc7				-0.85			
Il1a					-0.87		
Traf2					0.85		
Fasl					0.83		
Ppp3r1					0.83		
2010110P09Rik					-0.82		
Chuk							-0.8
Csf2rb2						0.85	
Dffb						0.92	
Ntrk1						0.85	
Trp53						0.88	
1500003O03Rik							0.92

p53 Responsive Elements							
gene	1h	3h	6h	9h	12h	24h	48h
Msh2	-0.97						
Btg2	0.95						
Igfbp3	-0.87	-0.86		-0.8	-0.86	-0.86	
Sesn1		-0.98		-0.87	-1		
S100a2		0.94					
Pmaip1		0.87	0.81		0.97		
Fas		-0.85					
Mdm2			0.92	0.89	0.8	0.88	
Cdkn1a			0.91	0.86	0.95	0.94	0.85
Gml			0.91				
Tnfrsf10b			0.9	0.89	0.86	0.89	
Pcna				0.81	0.86		
Sfn						0.86	

location of the sweet spot