Supporting Information

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Fig. S1. Paired inhibitor combinations perturb CVB3-induced phosphorylation signatures nonadditively. (*A*) Measured inhibitor pairs were reprinted from Fig. 1 and compared with (*B*) an additive model, in which single inhibitor time courses from Fig. 1 were used to predict paired inhibitor signatures by adding the net perturbation of each inhibitor compared with the DMSO control. Differences between the measured and modeled signatures were evaluated by R^2 goodness of fit (*Right*). The median R^2 value was 0.6.



Fig. 52. Single and paired inhibitors specifically inhibit CVB3-induced cardiotoxicity. Cell death as measured by MTS assay is shown for (*A*) sham-infected cardiomyocytes or (*B*) CVB3-infected cardiomyocytes at 16 and 24 h p.i. HL1 cells were pretreated and infected as described in Fig. 1. Data are presented as the mean of three independent replicates. Cell death was normalized to the DMSO-treated control.



Euclidean Distances

Fig. S3. Time-dependent hierarchical clustering of the nine-protein signature based on inhibitor data. (*A–F*) Data were analyzed as described in Fig. 3. HL1 cells were pretreated and infected as described in Fig. 1. Using a Euclidean distance metric and average linkage, phosphoproteins were clustered at the indicated time points. The dendrograms were built from the complete original dataset of single and double inhibitors (original) or subsampled as single, double, or single + double inhibitor subsets as described in *Methods*.

Variables	Assays		
Akt (p\$473)	Phospho-ELISA		
ATF2 (pTpT69/71)	Phospho-ELISA		
CREB (pS133)	Phospho-ELISA		
ERK1/2 (pTpY185/187)	Phospho-ELISA		
GSK3β (pS9)	Phospho-ELISA		
Hsp27 (pS82)	Phospho-ELISA		
ΙκΒα (pS32)	Phospho-ELISA		
JNK1/2 (pTpY183/185)	Phospho-ELISA		
p38 MAPK (pTpY180/182)	Phospho-ELISA		
Src (pY418)	Phospho-ELISA		
Viral protein (CVB3 VP1)	Western blot		
Virion progeny release	Plaque assay		

Table S1.	Measured	signaling	molecules	and	virus	replication
indicators						

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Table S2. Scientific support for the CVB3 partial correlation network shown in Fig. 4B

Edge	Partial correlation	n Reported interaction		
Hsp-27–CREB	0.5	Direct phosphorylation of Hsp-27 and CREB by RSK2	(1)	
JNK–ΙκΒα	-0.5	Inhibition of JNK activation by NF-κB-mediated induction of GADD45B and XIAP	(2, 3)	
р38–ІкВα	0.5	p38-mediated phosphorylation of MSK1 promotes NF-KB function, which induces IKBA	(4, 5)	
ERK–CREB	0.5	Direct phosphorylation of CREB through ERK-mediated phosphorylation of RSK	(6)	
Akt–GSK3β	0.4	Direct phosphorylation of GSK3 ^β by Akt	(7)	
JNK-ATF-2	-0.3	Direct phosphorylation of ATF-2 by JNK (negative correlation unexplained)	(8)	
lκBα–ATF-2	-0.3	Negative correlation unexplained	_	
GSK3β–ΙκΒα	0.3	GSK3 β required for normal NF- κ B function, which induces <i>IKBA</i>	(9–11)	
Akt–lκBα	0.3	Direct phosphorylation of $I\kappa B\alpha$ through Akt-mediated phosphorylation of IKK	(12, 13)	
p38–Hsp-27	0.3	Direct phosphorylation of Hsp-27 through p38-mediated phosphorylation of MK2	(14, 15)	

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