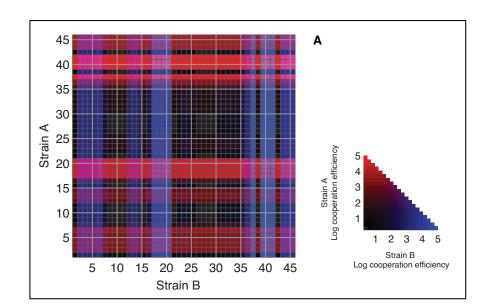
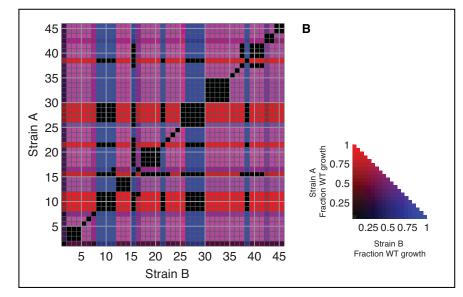


#	Mutation	#	Mutation	#	Mutation
1	ррс	17	hisB	33	pyrE
2	argB	18	hisC	34	pyrF
3	argC	19	hisD	35	lysA
4	argE	20	hisl	36	tyrA
5	argH	21	metA	37	trpA
6	proC	22	guaA	38	trpB
7	icd	23	guaB	39	trpC
8	nadC	24	purA	40	trpD
9	panB	25	purC	41	trpE
10	panD	26	purD	42	ilvC
11	serC	27	purF	43	ilvE
12	cysC	28	purL	44	leuB
13	cysE	29	purM	45	leuD
14	cysH	30	pyrB	46	leuC
15	glnA	31	pyrC		
16	serA	32	pyrD]	

Supplementary figure 1 Mutants in shared pathways have similar cooperation profiles. Red intensity corresponds to growth of the labelled row strain, blue intensity indicates growth of the numbered column strain. Fold growth is reported relative to the initial inoculation density. In most cases, correlated cooperation profiles are evident.



Supplementary figure 2 Model predictions reflect measured synergistic growth. (A) The cooperation efficiency is the ratio of the fitness benefit a strain receives from a given metabolite to the fitness cost for the conjugate partner in producing that metabolite. Cooperation efficiencies of zero are shown as black on the log scale. (B) Growth rates, as calculated from the joint flux model applying a MOMA-type objective function. Predicted growth rates are normalized to the FBA prediction for the wild type growth rate. Note the gaps along the diagonal reflecting the general failure of mutants in shared metabolic pathways to complement growth. The derivation of model predictions is detailed in Materials and Methods.



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8	nadC	24	purA	40	trpD
9	panB	25	purC	41	trpE
10	panD	26	purD	42	ilvC
11	serC	27	purF	43	ilvE
12	cysC	28	purL	44	leuB
13	cysE	29	purM	45	leuD
14	cysH	30	pyrB	46	leuC
15	glnA	31	pyrC		
16	serA	32	pyrD		