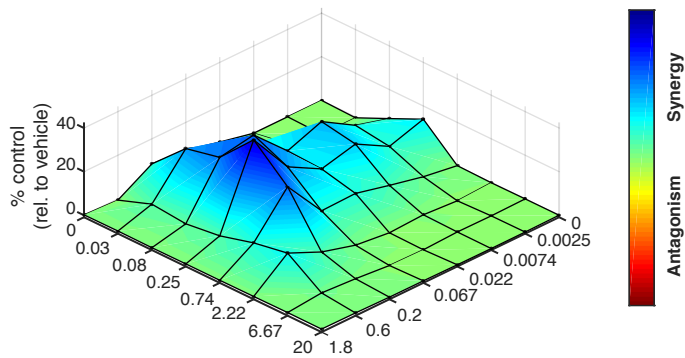
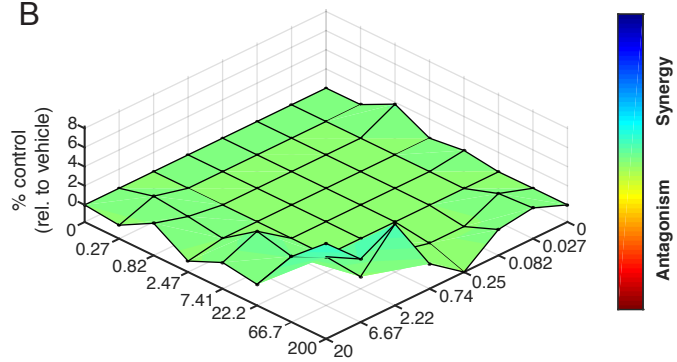


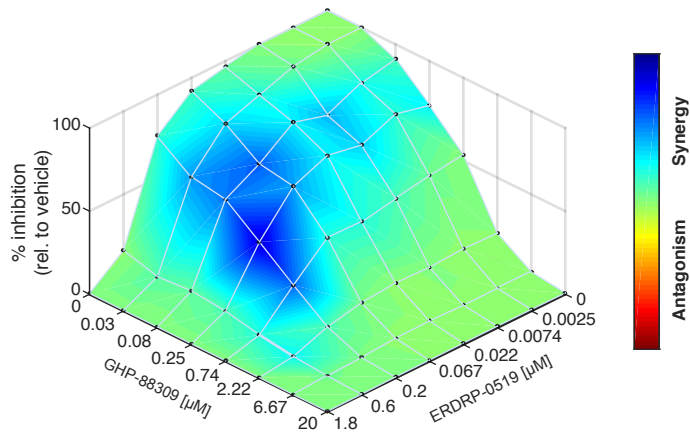
A



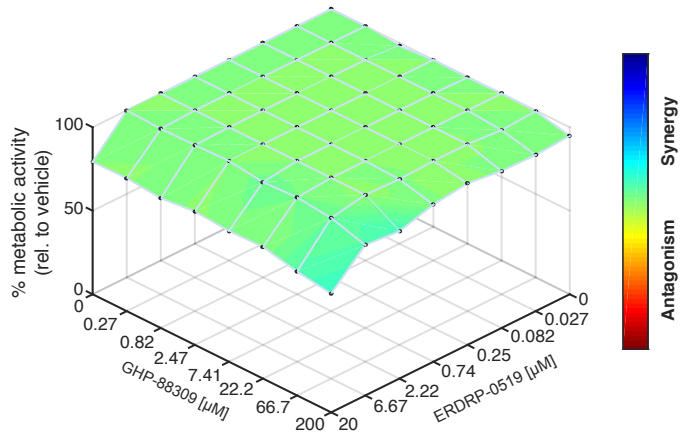
B



C



D



E

		GHP-88309 [ $\mu$ M]						
		0.03	0.08	0.25	0.74	2.22	6.67	20
ERDRP-0519 [ $\mu$ M]	0.0025	0 +/-0 ***	11 +/-17	9 +/-25	3 +/-20	-1 +/-5	0 +/-1	0 +/-1
	0.0074	3 +/-6	20 +/-17	18 +/-21	6 +/-22	0 +/-5	0 +/-2	0 +/-1
	0.022	10 +/-8	14 +/-17	14 +/-17	10 +/-12	-1 +/-6	-1 +/-2	0 +/-1
	0.067	17 +/-18	29 +/-22	22 +/-18	10 +/-20	1 +/-6	0 +/-2	0 +/-1
	0.2	23 +/-19	27 +/-14 *	42 +/-11 *	28 +/-9 *	6 +/-1 *	1 +/-0 *	0 +/-1
	0.6	5 +/-4	4 +/-5	6 +/-4	10 +/-2 *	13 +/-1 **	2 +/-1 *	0 +/-1
	1.8	0 +/-0	0 +/-0	0 +/-1	0 +/-1	0 +/-0	1 +/-0	0 +/-1
			HSA synergy and antagonism					

F

		GHP-88309 [ $\mu$ M]						
		0.27	0.82	2.47	7.41	22.2	66.7	200
ERDRP-0519 [ $\mu$ M]	0.027	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	1 +/-3	1 +/-5
	0.082	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	-1 +/-0	1 +/-6
	0.25	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	-1 +/-0	-2 +/-2
	0.74	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-2	1 +/-5
	2.22	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	0 +/-0 ***	-1 +/-0	7 +/-17
	6.67	1 +/-2	1 +/-1	0 +/-0 ***	2 +/-4	2 +/-3	3 +/-6	5 +/-6
	20	0 +/-8	2 +/-5	-1 +/-11	1 +/-6	0 +/-8	5 +/-8	8 +/-5 *
			HSA synergy and antagonism					