

Table SI. List of miRNAs that are upregulated in HepG2 co-culture cells in comparison to the monoculture HepG2 cells (with reads in TPM).

Name	Monoculture	Co-Culture	Fold change
hsa-miR-122-5p	34	1218	35.82
hsa-miR-10b-5p	55	1489	27.07
hsa-miR-483-3p	14	213	15.21
hsa-miR-483-5p	9	74	8.22
hsa-miR-126-3p	8	56	7.00
hsa-miR-210-3p	30	106	3.53
hsa-let-7d-5p	18	60	3.33
hsa-miR-885-5p	3	10	3.33
hsa-miR-10a-5p	1	3	3.00
hsa-miR-20a-5p	102	257	2.52
hsa-miR-365a-3p	13	32	2.46
hsa-miR-32-5p	22	54	2.45
hsa-let-7g-5p	1064	2266	2.13
hsa-miR-30b-5p	34	70	2.06
hsa-miR-98-5p	21	43	2.05
hsa-miR-365b-3p	14	28	2.00
hsa-miR-19b-3p	10	20	2.00
hsa-miR-193b-5p	3	6	2.00
hsa-miR-2277-5p	2	4	2.00
hsa-miR-323a-3p	1	2	2.00
hsa-miR-3173-5p	1	2	2.00
hsa-miR-3613-5p	1	2	2.00

miR/miRNA, microRNA.

Table SII. Selection of miRNA of interest (with reads in TPM).

Name	Monoculture		Co-Culture	
	LX2_cell	LX2_EV	LX2_cell	LX2_EV
hsa-miR-122-5p	6	5001	1	640
hsa-miR-10b-5p	29	852	26	179
hsa-miR-483-3p	0	0	0	6
hsa-miR-483-5p	0	0	0	17
hsa-miR-126-3p	58	1066	84	301

miR/miRNA, microRNA.