

SUPPLEMENTAL TABLE 1. Short Tandem Repeat (STR) fingerprinting report of cell lines used in the study.

Sample Name	AMEL	CSF1PO	D16S539	D18S51	D21S11	D3S1358	D5S818	D7S820
XP12ROSV	X,Y	12	9,11	13,22	30.2	15,16	11,12	12
XPAXP12ROSV	X,Y	12	9,11	13,22	30.2	15,16	11,12	12
XP20SSV	X	10,11	12	13,16	29,30	16	11	8,12
XPAXP20SSV	X	10,11	12	13,16	29,30	16	11	8,12
KO142	X	9,10	9,10	16	27,28	15,18	11,12	8,12
KO38	X	9,10	9,10	16	27,28	15,18	11,12	8,12
HeLa S3	X	9,10	9,10	16	27,28	15,18	11,12	8,12

The repeats of 8 STR fingerprinting markers are given. A total of 14 markers were used for the STR fingerprinting. All had matching scores for the pairs of cell lines, showing that the pairs are isogenic.

Functional analysis of genes upregulated by RA -log(p-values) for each cell line pair

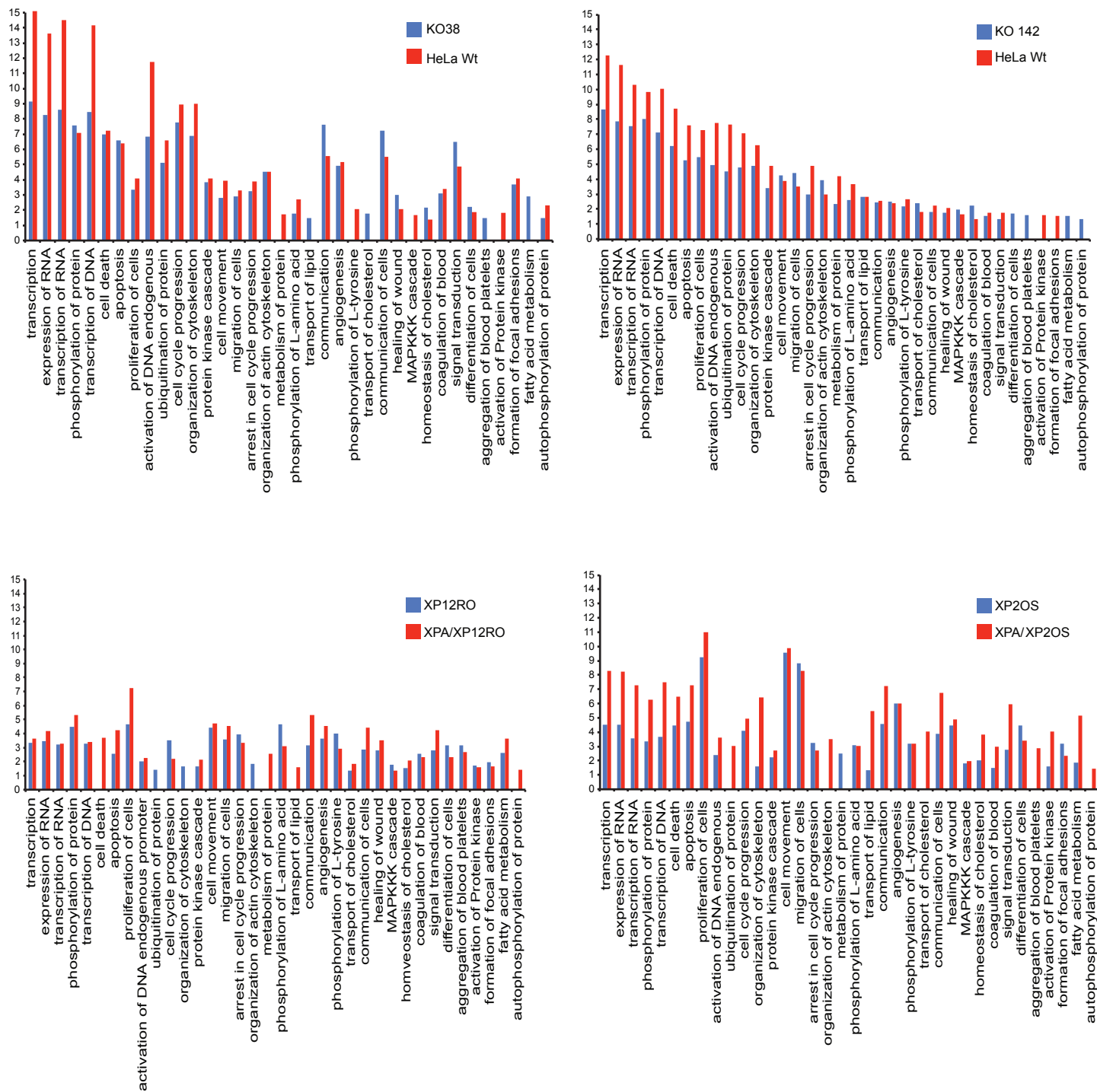


Figure S1