

**Table S3. Arrays analysis of miRNAs up- or down-regulated in HOB cells following treatment with 1,25D (n=3 separate culture preparations)**

miRNA	P value (group)	P value (treated vs vehicle)	Ratio (treated vs vehicle)	Fold-change (treated vs vehicle)	Direction of fold-change (treated vs vehicle)	F (group)	SS (group)	SS (error)
hsa-miRPlus-E1148	0.00203882	0.00203882	0.580467	-1.72275	down	50.9307	0.923667	0.072543
mmu-miR-450b-5p	0.0185626	0.0185626	0.656984	-1.52211	down	14.6956	0.550982	0.149972
ebv-miR-BART4	0.0260463	0.0260463	0.654783	-1.52722	down	11.9049	0.559818	0.188097
hsv1-miR-H2*	0.0517376	0.0517376	1.58618	1.58618	up	7.52487	0.664453	0.353204
hsa-miR-637	0.0590673	0.0590673	2.47297	2.47297	up	6.84172	2.55942	1.49636
hsa-miRPlus-E1186	0.0654071	0.0654071	1.5718	1.5718	up	6.34676	0.638482	0.402399
mcmv-miR-m88-1*	0.072164	0.072164	1.57476	1.57476	up	5.89348	0.643801	0.436959
kshv-miR-K12-9*	0.0722943	0.0722943	1.65881	1.65881	up	5.88538	0.799677	0.543501
hsa-miRPlus-E1110	0.07985	0.07985	1.50257	1.50257	up	5.45029	0.517613	0.379879
rno-miR-20b-5p	0.0868239	0.0868239	0.599272	-1.66869	down	5.10085	0.818557	0.641899
mmu-miR-466e-5p	0.0872131	0.0872131	0.469537	-2.12976	down	5.08261	1.78441	1.40433
mmu-miR-690	0.0960418	0.0960418	1.66931	1.66931	up	4.69954	0.819746	0.697725
hsa-miR-1228	0.0988068	0.0988068	1.95651	1.95651	up	4.59043	1.40635	1.22546