

miRNA	gene	snps	total	$MAF_{avg}$	$p_{MIN}$	chr	predicted	pathways
hsa-mir-221	FGF14	13	309	4.88E-01	1.59E-03	13	TRUE	4
hsa-mir-141	EDA2R	10	34	4.93E-01	4.76E-02	X	TRUE	1
hsa-mir-141	FOXO1	10	38	4.74E-01	2.91E-02	13	TRUE	3
hsa-mir-642a	MTOR	10	36	4.60E-01	2.68E-02	1	FALSE	9
hsa-mir-125b-2	LRRC4C	9	437	4.85E-01	9.62E-03	11	FALSE	1
hsa-mir-222	FGF14	8	309	4.88E-01	7.10E-04	13	TRUE	4
hsa-mir-429	C7	7	53	5.15E-01	2.43E-02	5	FALSE	3
hsa-mir-642a	NUP210	7	164	5.16E-01	6.81E-02	3	FALSE	1
hsa-mir-24-2	TUSC3	7	218	4.96E-01	8.21E-03	8	TRUE	3
hsa-mir-134	CHSY3	6	67	4.72E-01	5.22E-02	5	FALSE	2
hsa-mir-218-2	COL11A1	6	115	4.89E-01	5.33E-02	1	TRUE	4
hsa-mir-3200	GSPT2	6	14	5.09E-01	5.92E-02	X	FALSE	1
hsa-mir-758	HTR1B	6	233	5.03E-01	3.24E-02	6	FALSE	1
hsa-let-7c	LRRC4C	6	437	4.85E-01	2.00E-03	11	FALSE	1
hsa-mir-190b	NUP210	6	164	5.16E-01	2.43E-02	3	FALSE	1