

Supplementary table 1

Summary of differentially expressed genes (DEGs) and miRNAs (DEMs) between NPC and normal samples.

Upregulated DEGs

ADAM23
ANLN
BMP2
BUB1B
CCL8
CENPF
COL4A1
COL5A1
COL5A2
COL8A1
DST
FN1
GALNT11
GJA1
GPNMB
HOXA10
IDH1
IFIH1
IFIT3
KIF23
LAMB1
MMP1
MMP3
NFE2L3
NRXN1
PLAU
POSTN
PRC1
PTGS2
RCN1
SOX4
SSTR2
TNFAIP6
TOP2A
VRK2
XG
ZIC2

Downregulated DEGs

LTF

TEKT1

PIGR

MSLN

ROPN1L

CR2

SPAG6

AK7

MSMB

SCGB1A1

CLU

KRT4

TMC5

DNAI1

SLC27A6

ZMYND10

MS4A1

UPK1B

C22orf23

STATH

TEKT2

CD22

PTGDS

VPREB3

PROM1

CHST9

CAPS

CASC1

RIBC2

FAM3D

BANK1

RGS13

ATP12A

CD19

RRAD

AGR2

RIBC1

EFHC2

CH25H

RAB37

ZMYND12

CYP4B1

PZP

MEF2C

CD1D

CRYM

SERPINB7

WFDC2
KLRB1
HRASLS2
IQCD
CDH26
SLPI
BCAS1
KLF2
RALGPS2
LRMP
CXCL14
C9orf116
CEACAM6
GCNT3
OSBPL10
C3
POU2AF1
PLCG2
ALDH3A1
S100P
ALOX15
LRRC6
MUC13
SPA17
KIF9
LTB
EFHC1
ADAM28
SYNE1
MUC20
VSIG2
TNFSF11
KRT23
SLC22A4
FUT3
DUOX1
LCN2
CXXC5

DEMs

hsa-miR-1
hsa-miR-105
hsa-miR-135b
hsa-miR-144
hsa-miR-149
hsa-miR-154
hsa-miR-184
hsa-miR-18a
hsa-miR-190
hsa-miR-208
hsa-miR-211
hsa-miR-219
hsa-miR-302c
hsa-miR-30a-3p
hsa-miR-31
hsa-miR-323
hsa-miR-345
hsa-miR-346
hsa-miR-34b
hsa-miR-34c
hsa-miR-363
hsa-miR-365
hsa-miR-371
hsa-miR-372
hsa-miR-376a
hsa-miR-377
hsa-miR-380-3p
hsa-miR-383
hsa-miR-432
hsa-miR-448
hsa-miR-449
hsa-miR-452
hsa-miR-453
hsa-miR-455
hsa-miR-511
hsa-miR-514
hsa-miR-518b
hsa-miR-520a
hsa-miR-520g
hsa-miR-521
hsa-miR-523
hsa-miR-526c
hsa-miR-544
hsa-miR-9

Supplementary table 2
Four significant modules from the PPI network.

SUID	AverageShortestPathLength	ClusteringCoefficient	ClusterQuality	Degree	Eccentricity	IsSingleNode	MCODE	ClusMCODE	ModMCODE	Seed	Scanname	Neighborho	NumberofC	NumberofU	NumberofP	NumberofM	Radiality	selected	Selfloops	shared name	Stress	TopologicalCoefficient
1359	3.31395349	0.02030416	0.30175439	0.48888899	10	FALSE	Cluster 1	Clustered	5	BMP2	9	1	10	0	0	0.80717054	FALSE	0	BMP2	604	0.28470588	
1380	3.47674419	0.01466421	0.26762542	0.46666667	6	FALSE	Cluster 1	Clustered	4	TNFAIP6	10.1666667	1	6	0	0	0.79360465	FALSE	0	TNFAIP6	442	0.31770833	
1383	3.40697674	0.0250384	0.29351538	0.42857143	7	FALSE	Cluster 1	Clustered	4	MSLN	9.71428571	2	7	0	0	0.7994186	FALSE	0	MSLN	502	0.25868726	
1364	3.59069767	0.11722927	0.30386693	0.4	10	FALSE	Cluster 1	Clustered	5	MMIP3	8.5	1	10	0	0	0.80910653	FALSE	0	MMIP3	1652	0.24411765	
1366	3.56046512	0.00272452	0.29757785	0.76190476	7	FALSE	Cluster 1	Clustered	5	MMIP1	11	1	7	0	0	0.80329457	FALSE	0	MMIP1	110	0.32352941	
1334	3.29069767	0.0746891	0.30386693	0.28787879	12	FALSE	Cluster 1	Clustered	5	PTGS2	6.75	1	12	0	0	0.80910853	FALSE	0	PTGS2	1288	0.203125	
1336	4.53023256	0	0.20283019	1	5	FALSE	Cluster 1	Clustered	5	ANLN	5.6	1	5	0	0	0.67248062	FALSE	0	ANLN	0	0.7	
1337	4.40697674	0.03102371	0.22681293	0.47619048	7	FALSE	Cluster 1	Clustered	5	KIF23	4.57142857	1	7	0	0	0.71608527	FALSE	0	KIF23	778	0.42857143	
1370	2.6744186	0.49758878	0.37991304	0.19480519	22	FALSE	Cluster 1	Seeded	5	FN1	7.40909091	1	22	0	0	0.86946512	FALSE	0	FN1	10066	0.13149351	
1339	4.53023256	0	0.20283019	1	5	FALSE	Cluster 1	Clustered	5	BUB1B	5.6	1	5	0	0	0.67248062	FALSE	0	BUB1B	0	0.7	
1341	4.53023256	0	0.20283019	1	5	FALSE	Cluster 1	Clustered	5	TOP2A	5.6	1	5	0	0	0.67248062	FALSE	0	TOP2A	0	0.7	
1343	4.10465116	0.06834365	0.24362606	0.66666667	6	FALSE	Cluster 1	Clustered	5	CENPF	5.66666667	1	6	0	0	0.74127907	FALSE	0	CENPF	1622	0.42307692	
1345	4.53023256	0	0.20283019	1	5	FALSE	Cluster 1	Clustered	5	PRC1	5.6	1	5	0	0	0.67248062	FALSE	0	PRC1	0	0.7	
1348	3.41860465	0.03974044	0.29251701	0.5	8	FALSE	Cluster 1	Clustered	5	TNFSF11	8.75	2	8	0	0	0.79844961	FALSE	0	TNFSF11	622	0.27822581	
1382	3.27906977	0.04918838	0.30496454	0.27777778	9	FALSE	Cluster 1	Clustered	4	C3	8.22222222	2	9	0	0	0.81007752	FALSE	0	C3	986	0.19512195	
1383	3.56976744	0	0.28013029	1	4	FALSE	Cluster 1	Clustered	4	RGN1	11.25	1	4	0	0	0.79585271	FALSE	0	RGN1	96	0.41666667	
1320	3.27906977	0.11058734	0.30496454	0.28571429	7	FALSE	Cluster 1	Clustered	4	PLAU	9	1	7	0	0	0.81007752	FALSE	0	PLAU	2824	0.24489796	

module 2

SUID	AverageShortBetweennessClosenessCg	ClusteringCg	Degree	Eccentricity	isSingleNode	MCODE_Clus	MCODE_Nod	MCODE_Seqname	Neighborhood no	NumberOfDir	NumberOfTur	Partno	QIMU	Radiality	selected	SelfLoops	shared name	Stress	TopologicalCoefficient
1460	4.41860465	0	0.22631579	4	FALSE	Cluster 2	Seed	4 COL5A1	7.25	1	4	0	0	0.71511628	FALSE	0	COL5A1	0	0.725
1464	3.52325581	0.00743913	0.26392838	7	FALSE	Cluster 2	Clustered	3.42857143 COL5A1	9.28571429	1	7	0	0	0.78972868	FALSE	0	COL5A1	324	0.37142857
1466	3.59302326	0.00441138	0.27831715	6	FALSE	Cluster 2	Clustered	3.42857143 COL5A2	9.33333333	1	6	0	0	0.78391473	FALSE	0	COL5A2	244	0.4057971
1373	3.54651163	0.00715907	0.26196721	8	FALSE	Cluster 2	Clustered	3.42857143 COL4A1	8.025	1	8	0	0	0.7877907	FALSE	0	COL4A1	346	0.359375

Module 3

SUID	AverageShortBetweennessClosenessCg	ClusteringCg	Degree	Eccentricity	isSingleNode	MCODE_Clus	MCODE_Nod	MCODE_Scan	Neighborhoon no	NumberOfDir	NumberOfTur	ParmsCINuI	Radiality	selected	SelfLoops	shared name	Stress	TopologicalCoefficient
1440	3.5	0.02450341	0.28571429	0.5	FALSE	Cluster 3	Sead	3 RAB37	7	2	4	0	0.79166667	FALSE	0	RAB37	902	0.3375
1400	3.52325581	0.08686964	0.28392838	0.14285714	FALSE	Cluster 3	Clustered	3 LRMP	342867143	2	7	0	0.78972868	FALSE	0	LRMP	1584	0.20408183
1323	2.86511623	0.38692897	0.3372549	0.14285714	FALSE	Cluster 3	Clustered	3 CEACAM6	6.875	2	8	0	0.83824031	FALSE	0	CEACAM6	11774	0.17667588
1389	3.47674419	0.03106455	0.28762542	0.28666667	FALSE	Cluster 3	Clustered	3 PI3R	6.33333333	2	6	0	0.79500465	FALSE	0	PI3R	888	0.24666667

module 4

SUID	AverageShortBetweennessClosenessCg	ClusteringCg	Degree	Eccentricity	isSingleNode	MCODE_Cluster	MCODE_Nod	MCODE_Seqname	Neighborhood no	NumberOfDir	NumberOfTur	Partno	CINMI	Radiality	selected	SelfLoops	shared name	Stress	TopologicalCoefficient
1488	6.39534884	0.08452804	0.15636364	5	FALSE	Cluster 5	Seed	2.4 SPAG6	5.4	2	5	0	0	0.5503876	FALSE	0	SPAG6	3752	0.43333333
1484	8.23255814	0	0.12146893	2	FALSE	Cluster 5	Clustered	2 LRR06	4.5	2	2	0	0	0.39728682	FALSE	0	LRR06	0	0.64285714
1481	7.25681395	0.04626995	0.13782051	7	FALSE	Cluster 5	Clustered	2.2 DNMI1	4.14285714	2	7	0	0	0.42768217	FALSE	0	DNMI1	1924	0.41428571
1482	7.26744186	0.01200339	0.1378	6	FALSE	Cluster 5	Clustered	2.2 TER12	4.5	2	6	0	0	0.47771318	FALSE	0	TER12	962	0.48148148
1484	8.23255814	0	0.12146893	2	FALSE	Cluster 5	Clustered	2 ZMYND10	4.5	2	2	0	0	0.39728682	FALSE	0	ZMYND10	0	0.64285714

SUID	AverageShortBetweennessClosenessCg	ClusteringCg	Degree	Eccentricity	isSingleNode	MCODE_Cluster	MCODE_Nod	MCODE_Seqname	Neighborhood no	NumberOfDir	NumberOfTur	Partno	CINMI	Radiality	selected	SelfLoops	shared name	Stress	TopologicalCoefficient
1488	6.39534884	0.08452804	0.15636364	5	FALSE	Cluster 5	Seed	2.4 SPAG6	5.4	2	5	0	0	0.5503876	FALSE	0	SPAG6	3752	0.43333333
1484	8.23255814	0	0.12146893	2	FALSE	Cluster 5	Clustered	2 LRR06	4.5	2	2	0	0	0.39728682	FALSE	0	LRR06	0	0.64285714
1481	7.25681395	0.04626995	0.13782051	7	FALSE	Cluster 5	Clustered	2.2 DNMI1	4.14285714	2	7	0	0	0.42768217	FALSE	0	DNMI1	1924	0.41428571
1482	7.26744186	0.01200339	0.1378	6	FALSE	Cluster 5	Clustered	2.2 TER12	4.5	2	6	0	0	0.47771318	FALSE	0	TER12	962	0.48148148
1484	8.23255814	0	0.12146893	2	FALSE	Cluster 5	Clustered	2 ZMYND10	4.5	2	2	0	0	0.39728682	FALSE	0	ZMYND10	0	0.64285714

Supplementary table 4

Summary of candidate miRNAs predicted using miRWalk online database.

mimaid	refseqid	genesymbol	bindingp
hsa-miR-629-	NM_000064	C3	0.95
hsa-miR-1207	NM_000064	C3	0.96153846
hsa-miR-6891	NM_000064	C3	0.96538462
hsa-miR-3665	NM_000064	C3	0.98076923
hsa-miR-6763	NM_000064	C3	0.98076923
hsa-miR-6782	NM_000064	C3	0.98076923
hsa-miR-6791	NM_000064	C3	0.98717949
hsa-miR-27a-	NM_000064	C3	1
hsa-miR-210-	NM_000064	C3	1
hsa-miR-30b-	NM_000064	C3	1
hsa-miR-125a	NM_000064	C3	1
hsa-miR-296-	NM_000064	C3	1
hsa-miR-550a	NM_000064	C3	1
hsa-miR-1236	NM_000064	C3	1
hsa-miR-1304	NM_000064	C3	1
hsa-miR-3124	NM_000064	C3	1
hsa-miR-3188	NM_000064	C3	1
hsa-miR-3196	NM_000064	C3	1
hsa-miR-3198	NM_000064	C3	1
hsa-miR-4316	NM_000064	C3	1
hsa-miR-4271	NM_000064	C3	1
hsa-miR-4287	NM_000064	C3	1
hsa-miR-362C	NM_000064	C3	1
hsa-miR-3917	NM_000064	C3	1
hsa-miR-4463	NM_000064	C3	1
hsa-miR-4463	NM_000064	C3	1
hsa-miR-4471	NM_000064	C3	1
hsa-miR-4496	NM_000064	C3	1
hsa-miR-2392	NM_000064	C3	1
hsa-miR-4507	NM_000064	C3	1
hsa-miR-1587	NM_000064	C3	1
hsa-miR-4538	NM_000064	C3	1
hsa-miR-4653	NM_000064	C3	1
hsa-miR-469C	NM_000064	C3	1
hsa-miR-4704	NM_000064	C3	1
hsa-miR-4738	NM_000064	C3	1
hsa-miR-4793	NM_000064	C3	1
hsa-miR-5589	NM_000064	C3	1
hsa-miR-5787	NM_000064	C3	1
hsa-miR-6085	NM_000064	C3	1
hsa-miR-6133	NM_000064	C3	1
hsa-miR-6134	NM_000064	C3	1
hsa-miR-6134	NM_000064	C3	1
hsa-miR-675C	NM_000064	C3	1
hsa-miR-6766	NM_000064	C3	1
hsa-miR-6772	NM_000064	C3	1
hsa-miR-6773	NM_000064	C3	1
hsa-miR-6788	NM_000064	C3	1
hsa-miR-6794	NM_000064	C3	1
hsa-miR-6823	NM_000064	C3	1
hsa-miR-684C	NM_000064	C3	1
hsa-miR-6854	NM_000064	C3	1
hsa-miR-6867	NM_000064	C3	1
hsa-miR-6882	NM_000064	C3	1
hsa-miR-7154	NM_000064	C3	1
hsa-miR-4433	NM_000064	C3	1
hsa-miR-784E	NM_000064	C3	1
hsa-miR-8082	NM_000064	C3	1
hsa-miR-1211	NM_000064	C3	1
hsa-miR-443C	NM_0012780	COL5A1	0.95384615
hsa-miR-8071	NM_0012780	COL5A1	0.95384615
hsa-miR-443C	NM_000093	COL5A1	0.95384615
hsa-miR-8071	NM_000093	COL5A1	0.95384615
hsa-miR-3138	NM_0012780	COL5A1	0.96153846
hsa-miR-4464	NM_0012780	COL5A1	0.96153846
hsa-miR-3138	NM_000093	COL5A1	0.96153846
hsa-miR-4464	NM_000093	COL5A1	0.96153846
hsa-miR-1178	NM_0012780	COL5A1	0.97435897
hsa-miR-4302	NM_0012780	COL5A1	0.97435897
hsa-miR-361C	NM_0012780	COL5A1	0.97435897
hsa-miR-676-	NM_0012780	COL5A1	0.97435897
hsa-miR-1178	NM_000093	COL5A1	0.97435897
hsa-miR-4302	NM_000093	COL5A1	0.97435897
hsa-miR-361C	NM_000093	COL5A1	0.97435897
hsa-miR-676-	NM_000093	COL5A1	0.97435897
hsa-miR-1238	NM_0012780	COL5A1	0.98076923
hsa-miR-1204	NM_0012780	COL5A1	0.98076923
hsa-miR-430C	NM_0012780	COL5A1	0.98076923
hsa-miR-7111	NM_0012780	COL5A1	0.98076923
hsa-miR-1238	NM_000093	COL5A1	0.98076923
hsa-miR-1204	NM_000093	COL5A1	0.98076923
hsa-miR-430C	NM_000093	COL5A1	0.98076923
hsa-miR-7111	NM_000093	COL5A1	0.98076923
hsa-miR-3157	NM_0012780	COL5A1	0.98717949
hsa-miR-3157	NM_000093	COL5A1	0.98717949
hsa-miR-25-5j	NM_0012780	COL5A1	1
hsa-miR-197-	NM_0012780	COL5A1	1
hsa-miR-30c;	NM_0012780	COL5A1	1
hsa-miR-181a	NM_0012780	COL5A1	1
hsa-miR-183-	NM_0012780	COL5A1	1
hsa-miR-212-	NM_0012780	COL5A1	1

hsa-miR-125t NM_0012780 COL5A1	1
hsa-miR-143-1 NM_0012780 COL5A1	1
hsa-miR-125a NM_0012780 COL5A1	1
hsa-miR-34b-1 NM_0012780 COL5A1	1
hsa-miR-130b NM_0012780 COL5A1	1
hsa-miR-302d NM_0012780 COL5A1	1
hsa-miR-371a NM_0012780 COL5A1	1
hsa-miR-373-1 NM_0012780 COL5A1	1
hsa-miR-376a NM_0012780 COL5A1	1
hsa-miR-324-1 NM_0012780 COL5A1	1
hsa-miR-423-1 NM_0012780 COL5A1	1
hsa-miR-376b NM_0012780 COL5A1	1
hsa-miR-483-1 NM_0012780 COL5A1	1
hsa-miR-486-1 NM_0012780 COL5A1	1
hsa-miR-487a NM_0012780 COL5A1	1
hsa-miR-489-1 NM_0012780 COL5A1	1
hsa-miR-517c NM_0012780 COL5A1	1
hsa-miR-500a NM_0012780 COL5A1	1
hsa-miR-501-1 NM_0012780 COL5A1	1
hsa-miR-502-1 NM_0012780 COL5A1	1
hsa-miR-506-1 NM_0012780 COL5A1	1
hsa-miR-510-1 NM_0012780 COL5A1	1
hsa-miR-455-1 NM_0012780 COL5A1	1
hsa-miR-539-1 NM_0012780 COL5A1	1
hsa-miR-487b NM_0012780 COL5A1	1
hsa-miR-567-1 NM_0012780 COL5A1	1
hsa-miR-573-1 NM_0012780 COL5A1	1
hsa-miR-589-1 NM_0012780 COL5A1	1
hsa-miR-550a NM_0012780 COL5A1	1
hsa-miR-601-1 NM_0012780 COL5A1	1
hsa-miR-605-1 NM_0012780 COL5A1	1
hsa-miR-605-2 NM_0012780 COL5A1	1
hsa-miR-619-1 NM_0012780 COL5A1	1
hsa-miR-630-1 NM_0012780 COL5A1	1
hsa-miR-640-1 NM_0012780 COL5A1	1
hsa-miR-642a NM_0012780 COL5A1	1
hsa-miR-644a NM_0012780 COL5A1	1
hsa-miR-645-1 NM_0012780 COL5A1	1
hsa-miR-648-1 NM_0012780 COL5A1	1
hsa-miR-1298 NM_0012780 COL5A1	1
hsa-miR-2113 NM_0012780 COL5A1	1
hsa-miR-708-1 NM_0012780 COL5A1	1
hsa-miR-877-1 NM_0012780 COL5A1	1
hsa-miR-887-1 NM_0012780 COL5A1	1
hsa-miR-301b NM_0012780 COL5A1	1
hsa-miR-509-1 NM_0012780 COL5A1	1
hsa-miR-936-1 NM_0012780 COL5A1	1
hsa-miR-937-1 NM_0012780 COL5A1	1
hsa-miR-1236 NM_0012780 COL5A1	1
hsa-miR-1208 NM_0012780 COL5A1	1
hsa-miR-1255 NM_0012780 COL5A1	1
hsa-miR-1276 NM_0012780 COL5A1	1
hsa-miR-191C NM_0012780 COL5A1	1
hsa-miR-191D NM_0012780 COL5A1	1
hsa-miR-1973 NM_0012780 COL5A1	1
hsa-miR-2115 NM_0012780 COL5A1	1
hsa-miR-2117 NM_0012780 COL5A1	1
hsa-miR-2117 NM_0012780 COL5A1	1
hsa-miR-3121 NM_0012780 COL5A1	1
hsa-miR-3124 NM_0012780 COL5A1	1
hsa-miR-3124 NM_0012780 COL5A1	1
hsa-miR-3126 NM_0012780 COL5A1	1
hsa-miR-3126 NM_0012780 COL5A1	1
hsa-miR-3155 NM_0012780 COL5A1	1
hsa-miR-3159 NM_0012780 COL5A1	1
hsa-miR-3175 NM_0012780 COL5A1	1
hsa-miR-3177 NM_0012780 COL5A1	1
hsa-miR-3187 NM_0012780 COL5A1	1
hsa-miR-3198 NM_0012780 COL5A1	1
hsa-miR-320C NM_0012780 COL5A1	1
hsa-miR-3202 NM_0012780 COL5A1	1
hsa-miR-4265 NM_0012780 COL5A1	1
hsa-miR-4275 NM_0012780 COL5A1	1
hsa-miR-3605 NM_0012780 COL5A1	1
hsa-miR-3622 NM_0012780 COL5A1	1
hsa-miR-366C NM_0012780 COL5A1	1
hsa-miR-368C NM_0012780 COL5A1	1
hsa-miR-3688 NM_0012780 COL5A1	1
hsa-miR-3691 NM_0012780 COL5A1	1
hsa-miR-3918 NM_0012780 COL5A1	1
hsa-miR-3922 NM_0012780 COL5A1	1
hsa-miR-4428 NM_0012780 COL5A1	1
hsa-miR-4444 NM_0012780 COL5A1	1
hsa-miR-4447 NM_0012780 COL5A1	1
hsa-miR-445C NM_0012780 COL5A1	1
hsa-miR-3155 NM_0012780 COL5A1	1
hsa-miR-449C NM_0012780 COL5A1	1
hsa-miR-4495 NM_0012780 COL5A1	1
hsa-miR-2392 NM_0012780 COL5A1	1
hsa-miR-451E NM_0012780 COL5A1	1
hsa-miR-3976 NM_0012780 COL5A1	1
hsa-miR-4632 NM_0012780 COL5A1	1
hsa-miR-4633 NM_0012780 COL5A1	1
hsa-miR-4646 NM_0012780 COL5A1	1

hsa-miR-711c NM_0012780 COL5A1	1
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hsa-miR-596	NM_000779	CYP4B1	0.96153846
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hsa-miR-6762	NM_000779	CYP4B1	0.96153846
hsa-miR-324-NM_0010997	CYP4B1	0.96153846	
hsa-miR-596	NM_0010997	CYP4B1	0.96153846
hsa-miR-1908	NM_0010997	CYP4B1	0.96153846
hsa-miR-6762	NM_0010997	CYP4B1	0.96153846
hsa-miR-544t	XM_0115408	CYP4B1	0.97435897
hsa-miR-1233	NM_0013191	CYP4B1	0.97435897
hsa-miR-1233	NM_000779	CYP4B1	0.97435897
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hsa-miR-6078	NM_0013191	CYP4B1	0.98076923
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hsa-miR-132-XM_0115408	CYP4B1	1	
hsa-miR-361-XM_0115408	CYP4B1	1	
hsa-miR-1285	XM_0115408	CYP4B1	1

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hsa-miR-1211 NM_0010997	CYP4B1	1
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hsa-miR-769- NM_000954	PTGDS	1
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hsa-miR-3178 NM_000954	PTGDS	1
hsa-miR-3202 NM_000954	PTGDS	1
hsa-miR-4257 NM_000954	PTGDS	1
hsa-miR-3622 NM_000954	PTGDS	1
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hsa-miR-143- NM_000963	PTGS2	0.95384615
hsa-miR-1304 NM_000963	PTGS2	0.95384615
hsa-miR-4681 NM_000963	PTGS2	0.95384615
hsa-miR-6793 NM_000963	PTGS2	0.95384615
hsa-miR-23b- NM_000963	PTGS2	0.96153846
hsa-miR-185- NM_000963	PTGS2	0.96153846
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hsa-miR-410- NM_000963	PTGS2	0.96153846
hsa-miR-571 NM_000963	PTGS2	0.96153846
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hsa-miR-1202 NM_000963	PTGS2	0.96153846
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hsa-miR-113S NM_000963	PTGS2	0.96153846
hsa-miR-4303 NM_000963	PTGS2	0.97435897
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hsa-miR-214- NM_000963	PTGS2	1
hsa-miR-200c NM_000963	PTGS2	1
hsa-miR-200a NM_000963	PTGS2	1
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hsa-miR-4538 XM_0170116	GPNMB	0.96153846
hsa-miR-574-NM_002510	GPNMB	0.96153846
hsa-miR-4538 NM_002510	GPNMB	0.96153846
hsa-miR-1039 NM_002510	GPNMB	0.96153846
hsa-miR-574-NM_0010053	GPNMB	0.96153846
hsa-miR-4538 NM_0010053	GPNMB	0.96153846
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hsa-miR-766-XM_0170116	GPNMB	0.97435897
hsa-miR-6818 XM_0170116	GPNMB	0.97435897
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hsa-miR-6818 NM_0010053	GPNMB	0.97435897
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hsa-miR-129-XM_0170116	GPNMB	1
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hsa-miR-215-XM_0170116	GPNMB	1
hsa-miR-200b XM_0170116	GPNMB	1
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hsa-miR-409-XM_0170116	GPNMB	1
hsa-miR-512-XM_0170116	GPNMB	1

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hsa-miR-506-XM_0170116	GPMB	1	
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hsa-miR-6862	NM_0010401	SERPINB7	1
hsa-miR-6864	NM_0010401	SERPINB7	1
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hsa-miR-104C	NM_0010401	SERPINB7	1
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hsa-miR-885-	NM_0010482	C9orf116	0.96153846
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hsa-miR-3130	NM_144654	C9orf116	0.97435897
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hsa-miR-27a-	NM_175569	XG	1
hsa-miR-28-5j	NM_175569	XG	1
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hsa-miR-1212 NM_0011458 PROM1	1
hsa-miR-4644 NM_002421 MMP1	0.96153846
hsa-miR-6835 NM_002421 MMP1	0.96153846
hsa-miR-185- NM_002421 MMP1	0.98461539
hsa-miR-22-5j NM_002421 MMP1	1
hsa-miR-25-5j NM_002421 MMP1	1
hsa-miR-105- NM_002421 MMP1	1
hsa-miR-211- NM_002421 MMP1	1
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hsa-miR-138- NM_002421 MMP1	1
hsa-miR-143- NM_002421 MMP1	1
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hsa-miR-526b NM_002421 MMP1	1
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hsa-miR-449c NM_002421	MMP1	1
hsa-miR-466c NM_002421	MMP1	1
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hsa-miR-7154 NM_032257	ZMYND12	0.95384615
hsa-miR-103a NM_032257	ZMYND12	0.96153846
hsa-miR-107 NM_032257	ZMYND12	0.96153846
hsa-miR-664a NM_032257	ZMYND12	0.96153846
hsa-miR-4723 NM_032257	ZMYND12	0.96153846
hsa-miR-4742 NM_032257	ZMYND12	0.96153846
hsa-miR-4765 NM_032257	ZMYND12	0.96153846
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hsa-miR-7111 NM_032257	ZMYND12	0.96153846
hsa-miR-6825 NM_032257	ZMYND12	0.97435897
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hsa-miR-3059 NM_032257	ZMYND12	1
hsa-miR-1211 NM_032257	ZMYND12	1
hsa-miR-518a NM_018100	EFHC1	0.95384615
hsa-miR-6165 NM_018100	EFHC1	0.95384615
hsa-miR-518a NM_0011724	EFHC1	0.95384615

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hsa-miR-6133 NM_018100 EFHC1	0.95512821
hsa-miR-6133 NM_0011724 EFHC1	0.95512821
hsa-miR-18a- NM_018100 EFHC1	0.96153846
hsa-miR-222- NM_018100 EFHC1	0.96153846
hsa-let-7g-3p NM_018100 EFHC1	0.96153846
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hsa-miR-596 NM_018100 EFHC1	0.96153846
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hsa-miR-885- NM_018100 EFHC1	0.96153846
hsa-miR-3137 NM_018100 EFHC1	0.96153846
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hsa-miR-330- NM_018100 EFHC1	0.97435897
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hsa-miR-29b- NM_018100 EFHC1	0.98076923
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hsa-miR-3176 NM_018100 EFHC1	0.98076923
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hsa-miR-6750	NM_0012862	RALGPS2	0.97435897
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hsa-miR-6891 NM_014312	VSIG2	0.96153846
hsa-miR-3055 NM_014312	VSIG2	0.96153846
hsa-miR-431- NM_0013299	VSIG2	0.96153846
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hsa-miR-3935 NM_0013299	VSIG2	0.96153846
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hsa-miR-7162 NM_0013299	VSIG2	0.96153846
hsa-miR-1237 NM_014312	VSIG2	0.97435897
hsa-miR-4667 NM_014312	VSIG2	0.97435897
hsa-miR-5193 NM_014312	VSIG2	0.97435897
hsa-miR-5581 NM_014312	VSIG2	0.97435897
hsa-miR-185- NM_0013299	VSIG2	0.97435897
hsa-miR-6726 NM_0013299	VSIG2	0.97802198
hsa-miR-129- NM_014312	VSIG2	1
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hsa-miR-324- NM_014312	VSIG2	1
hsa-miR-133b NM_014312	VSIG2	1
hsa-miR-376b NM_014312	VSIG2	1
hsa-miR-675- NM_014312	VSIG2	1
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hsa-miR-3059 NM_0013299 VSIG2	1
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hsa-miR-3198 NM_0013304 IQCD	0.98076923
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hsa-miR-125a NM_138451 IQCD	1
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hsa-miR-3187	NM_033071	SYNE1	0.96153846
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hsa-miR-512-	NM_182961	SYNE1	0.96153846
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hsa-miR-133a	NM_0013477	SYNE1	0.96153846
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hsa-miR-558	NM_0013477	SYNE1	0.97435897
hsa-miR-6131	NM_0013477	SYNE1	0.97435897
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hsa-miR-4496	NM_182961	SYNE1	1
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hsa-miR-128-: NM_021810 CDH26	0.96153846
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hsa-miR-6817 NM_177980 CDH26	0.96923077
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hsa-miR-426C NM_002272	KRT4	0.96153846
hsa-miR-4508 NM_002272	KRT4	0.96153846
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hsa-miR-675C NM_002272	KRT4	0.96153846
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hsa-miR-1911 NM_009588	LTB	0.97435897
hsa-miR-4445 NM_009588	LTB	0.98076923
hsa-miR-23a- NM_002341	LTB	1
hsa-miR-107 NM_002341	LTB	1
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hsa-miR-34c- NM_009588	LTB	1
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hsa-miR-211-	NM_002644	PIGR	0.97435897
hsa-miR-6804	NM_002644	PIGR	0.97802198
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hsa-miR-305S NM_002661	PLCG2	1
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hsa-miR-1211 NM_002661	PLCG2	1
hsa-miR-1213 NM_002661	PLCG2	1
hsa-miR-489- NM_002864	PZP	1
hsa-miR-5004 NM_002864	PZP	1
hsa-miR-7151 NM_002864	PZP	1
hsa-miR-192- NM_002901	RCN1	0.96153846
hsa-miR-622 NM_002901	RCN1	0.96153846
hsa-miR-3153 NM_002901	RCN1	0.96153846
hsa-miR-6886 NM_002901	RCN1	0.96153846
hsa-miR-6764 NM_002901	RCN1	0.97435897
hsa-let-7b-5p NM_002901	RCN1	1
hsa-let-7b-3p NM_002901	RCN1	1
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hsa-miR-593-; NM_002927	RGS13	0.96153846
hsa-miR-1231 NM_144766	RGS13	0.96923077
hsa-miR-1231 NM_002927	RGS13	0.96923077
hsa-let-7d-3p NM_144766	RGS13	1
hsa-miR-17-3j NM_144766	RGS13	1
hsa-miR-329-; NM_144766	RGS13	1
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hsa-miR-8086 NM_003059	SLC22A4	1
hsa-miR-129- NM_003064	SLPI	0.96153846
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hsa-miR-6512 NM_003064	SLPI	0.96153846
hsa-miR-7106 NM_003064	SLPI	0.96923077
hsa-miR-93-3j NM_003064	SLPI	1
hsa-miR-125a NM_003064	SLPI	1
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hsa-miR-6819 NM_003064	SLPI	1
hsa-miR-6829 NM_003064	SLPI	1
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hsa-miR-1039 NM_003064	SLPI	1
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hsa-miR-1267 NM_003107	SOX4	0.96153846
hsa-miR-1288 NM_003107	SOX4	0.96153846
hsa-miR-315C NM_003107	SOX4	0.96153846
hsa-miR-4694 NM_003107	SOX4	0.96153846
hsa-miR-4716 NM_003107	SOX4	0.96153846
hsa-miR-4524 NM_003107	SOX4	0.96153846
hsa-miR-6878 NM_003107	SOX4	0.96153846
hsa-miR-3664 NM_003107	SOX4	0.96923077
hsa-miR-198 NM_003107	SOX4	0.97435897
hsa-miR-6507 NM_003107	SOX4	0.97435897
hsa-miR-6765 NM_003107	SOX4	0.97435897
hsa-miR-316C NM_003107	SOX4	0.97435897
hsa-miR-3184 NM_003107	SOX4	0.97435897
hsa-miR-191- NM_003107	SOX4	0.98076923
hsa-miR-296- NM_003107	SOX4	0.98076923
hsa-miR-519e NM_003107	SOX4	0.98076923
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hsa-miR-6771 NM_003107	SOX4	0.98717949
hsa-miR-6847 NM_003107	SOX4	0.98717949
hsa-let-7e-3p NM_003107	SOX4	1
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hsa-miR-211- NM_003107	SOX4	1
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hsa-miR-215- NM_003107	SOX4	1
hsa-miR-30b- NM_003107	SOX4	1
hsa-miR-122- NM_003107	SOX4	1
hsa-miR-137- NM_003107	SOX4	1
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hsa-miR-575 NM_003107	SOX4	1
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hsa-miR-629-	NM_003107	SOX4	1
hsa-miR-631	NM_003107	SOX4	1
hsa-miR-661	NM_003107	SOX4	1
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hsa-miR-1296	NM_003107	SOX4	1
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hsa-miR-541-	NM_003107	SOX4	1
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hsa-miR-887-	NM_003107	SOX4	1
hsa-miR-873-	NM_003107	SOX4	1
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hsa-miR-1178	NM_003107	SOX4	1
hsa-miR-1180	NM_003107	SOX4	1
hsa-miR-1228	NM_003107	SOX4	1
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hsa-miR-1238	NM_003107	SOX4	1
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hsa-miR-655-	NM_033012	TNFSF11	0.97435897
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hsa-miR-4651 NM_004058	CAPS	0.95384615
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hsa-miR-491-	NM_004289	NFE2L3	0.97435897
hsa-miR-937-	NM_004289	NFE2L3	0.97435897
hsa-miR-3659	NM_004289	NFE2L3	0.97435897
hsa-miR-525-	NM_004289	NFE2L3	0.98076923
hsa-miR-765	NM_004289	NFE2L3	0.98076923
hsa-miR-23a-	NM_004289	NFE2L3	1
hsa-miR-27a-	NM_004289	NFE2L3	1
hsa-miR-199a	NM_004289	NFE2L3	1
hsa-miR-212-	NM_004289	NFE2L3	1
hsa-miR-215-	NM_004289	NFE2L3	1
hsa-miR-143-	NM_004289	NFE2L3	1
hsa-miR-362-	NM_004289	NFE2L3	1
hsa-miR-487a	NM_004289	NFE2L3	1
hsa-miR-519e	NM_004289	NFE2L3	1
hsa-miR-520g	NM_004289	NFE2L3	1
hsa-miR-556-	NM_004289	NFE2L3	1
hsa-miR-644a	NM_004289	NFE2L3	1
hsa-miR-1185	NM_004289	NFE2L3	1
hsa-miR-937-	NM_004289	NFE2L3	1
hsa-miR-1179	NM_004289	NFE2L3	1
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hsa-miR-4714	NM_004289	NFE2L3	1
hsa-miR-4717	NM_004289	NFE2L3	1
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hsa-miR-394C NM_005980	S100P	0.97435897
hsa-miR-23a-1 NM_005980	S100P	0.98076923
hsa-miR-1227 NM_005980	S100P	0.98076923
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hsa-miR-198 NM_005980	S100P	1
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hsa-miR-31-5j NM_006235	POU2AF1	0.96703297
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hsa-miR-3158 NM_014466	TEKT2	0.96153846
hsa-miR-6867 NM_014466	TEKT2	0.96923077
hsa-miR-328- NM_014466	TEKT2	0.98076923
hsa-let-7b-3p NM_014466	TEKT2	1
hsa-let-7d-3p NM_014466	TEKT2	1
hsa-miR-18a- NM_014466	TEKT2	1

hsa-miR-24-2	NM_014466	TEKT2	1
hsa-miR-181a	NM_014466	TEKT2	1
hsa-miR-125a	NM_014466	TEKT2	1
hsa-miR-296-	NM_014466	TEKT2	1
hsa-miR-326	NM_014466	TEKT2	1
hsa-miR-329-	NM_014466	TEKT2	1
hsa-miR-92b-	NM_014466	TEKT2	1
hsa-miR-596	NM_014466	TEKT2	1
hsa-miR-618	NM_014466	TEKT2	1
hsa-miR-634	NM_014466	TEKT2	1
hsa-miR-636	NM_014466	TEKT2	1
hsa-miR-1224	NM_014466	TEKT2	1
hsa-miR-744-	NM_014466	TEKT2	1
hsa-miR-941	NM_014466	TEKT2	1
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hsa-miR-147C	NM_014466	TEKT2	1
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hsa-miR-4683	NM_014466	TEKT2	1
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hsa-miR-7848	NM_014466	TEKT2	1
hsa-miR-1211	NM_014466	TEKT2	1
hsa-miR-6891	NM_015653	RIBC2	0.96153846
hsa-miR-149-	NM_015653	RIBC2	1
hsa-miR-513a	NM_015653	RIBC2	1
hsa-miR-1236	NM_015653	RIBC2	1
hsa-miR-4428	NM_015653	RIBC2	1
hsa-miR-470C	NM_015653	RIBC2	1
hsa-miR-4717	NM_015653	RIBC2	1
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hsa-miR-6833	NM_015653	RIBC2	1
hsa-miR-6845	NM_015653	RIBC2	1
hsa-miR-6851	NM_015653	RIBC2	1
hsa-miR-18b-	NM_016270	KLF2	0.95384615
hsa-miR-372-	NM_016270	KLF2	0.96153846
hsa-miR-424-	NM_016270	KLF2	0.96153846
hsa-miR-483-	NM_016270	KLF2	0.96153846
hsa-miR-3170	NM_016270	KLF2	0.96153846
hsa-miR-3685	NM_016270	KLF2	0.96153846
hsa-miR-4732	NM_016270	KLF2	0.96153846
hsa-miR-5587	NM_016270	KLF2	0.96153846
hsa-miR-6133	NM_016270	KLF2	0.96153846
hsa-miR-1224	NM_016270	KLF2	0.97435897
hsa-miR-3191	NM_016270	KLF2	0.97435897
hsa-miR-6773	NM_016270	KLF2	0.97435897
hsa-miR-4695	NM_016270	KLF2	0.98076923
hsa-miR-4776	NM_016270	KLF2	0.98076923
hsa-miR-651C	NM_016270	KLF2	0.98076923
hsa-miR-6822	NM_016270	KLF2	0.98076923
hsa-miR-569C	NM_016270	KLF2	0.99145299
hsa-let-7f-2-3j	NM_016270	KLF2	1
hsa-miR-20a-	NM_016270	KLF2	1
hsa-miR-26a-	NM_016270	KLF2	1
hsa-miR-33a-	NM_016270	KLF2	1
hsa-miR-30c-	NM_016270	KLF2	1
hsa-miR-30c-	NM_016270	KLF2	1
hsa-miR-222-	NM_016270	KLF2	1
hsa-miR-200b	NM_016270	KLF2	1
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hsa-miR-205-1 NM_017425	SPA17	0.96153846
hsa-miR-1231 NM_017425	SPA17	0.96153846
hsa-miR-1908 NM_017425	SPA17	0.96153846
hsa-miR-3129 NM_017425	SPA17	0.96153846
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hsa-miR-3192 NM_017425	SPA17	0.96153846
hsa-miR-4326 NM_017425	SPA17	0.96153846
hsa-miR-4501 NM_017425	SPA17	0.96153846
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hsa-miR-8085 NM_017425 SPA17	0.96153846
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hsa-miR-760 NM_017425 SPA17	0.96923077
hsa-miR-476S NM_017425 SPA17	0.96923077
hsa-miR-520a NM_017425 SPA17	0.97435897
hsa-miR-647 NM_017425 SPA17	0.97435897
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hsa-miR-6895 NM_017425 SPA17	0.97435897
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hsa-miR-7155	NM_025184	EFHC2	0.95384615
hsa-miR-4455	NM_025184	EFHC2	0.96923077
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hsa-miR-3178	NM_025184	EFHC2	0.98076923
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hsa-miR-6883	NM_025184	EFHC2	0.98076923
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hsa-miR-152-	NM_025184	EFHC2	1
hsa-miR-370-	NM_025184	EFHC2	1
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hsa-miR-339-	NM_033049	MUC13	0.96153846
hsa-miR-605-	NM_033049	MUC13	0.96153846
hsa-miR-4794	NM_033049	MUC13	0.96153846
hsa-let-7b-5p	NM_033049	MUC13	0.97435897
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hsa-miR-578	NM_033049	MUC13	0.98076923
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hsa-let-7b-3p	NM_033049	MUC13	1
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hsa-miR-758- NM_053285	TEKT1	0.96153846
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hsa-miR-4489 NM_053285	TEKT1	0.97435897
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hsa-miR-7114 NM_053285	TEKT1	0.97435897
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hsa-miR-486- NM_053285	TEKT1	0.98076923

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hsa-miR-7107 NM_053285	TEKT1	1
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hsa-miR-7847 NM_053285	TEKT1	1
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hsa-miR-8075 NM_053285	TEKT1	1
hsa-miR-8076 NM_053285	TEKT1	1
hsa-miR-8079 NM_053285	TEKT1	1
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hsa-miR-1039 NM_053285	TEKT1	1
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hsa-miR-1052 NM_053285	TEKT1	1
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hsa-miR-1139 NM_053285	TEKT1	1
hsa-miR-114C NM_053285	TEKT1	1
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hsa-miR-1211 NM_053285	TEKT1	1
hsa-miR-1213 NM_053285	TEKT1	1
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hsa-miR-6894 NM_138805	FAM3D	0.96153846
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hsa-miR-6792 NM_138805	FAM3D	0.97435897
hsa-miR-517- NM_138805	FAM3D	0.98076923
hsa-miR-27a- NM_138805	FAM3D	1
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hsa-miR-150- NM_138805	FAM3D	1
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hsa-miR-92b- NM_138805	FAM3D	1
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hsa-miR-766	NM_138805	FAM3D	1
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hsa-miR-892b	NM_138805	FAM3D	1
hsa-miR-301b	NM_138805	FAM3D	1
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hsa-miR-1238	NM_138805	FAM3D	1
hsa-miR-1275	NM_138805	FAM3D	1
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hsa-miR-6510	NM_138805	FAM3D	1
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hsa-miR-1273	NM_138805	FAM3D	1
hsa-miR-7978	NM_138805	FAM3D	1

hsa-miR-1039	NM_138805	FAM3D	1
hsa-miR-1039	NM_138805	FAM3D	1
hsa-miR-1039	NM_138805	FAM3D	1
hsa-miR-1211	NM_138805	FAM3D	1

Supplementary table 5

Summary of hub genes.

ID	Gene:title	Gene: symbol	Gene:ID	GenBank:Acc	Chromosome	Chromosome	logFC	AveExpr	t	P:Value	adj.P.Val	sig
204748_at	prostaglandin PTGS2		5743	NM_000963	1q25.2-q25.3	Chromosome 3	1.6667657	6.93533666	3.04867631	0.00484908	0.35860098	UP
211719_x_at	fibronectin 1 FN1		2335	BC005858	2q34	Chromosome 2	1.1753329	9.18110901	2.49650052	0.01843405	0.53415589	UP
204475_at	matrix metallo MMP1		4312	NM_002421	11q22.3	Chromosome 4	4.0689709	7.17515732	3.14816202	0.0037703	0.33206949	UP
211668_s_at	plasmihogen PLAU		5328	AH007073	10q22.2	Chromosome 3	1.1102172	7.69598447	4.35734683	0.00014883	0.08503438	UP
205628_at	matrix metallo MMP3		4314	NM_002422	11q22.3	Chromosome 1	1.74777973	6.45166842	1.49350805	0.14603782	0.78579535	UP
206398_s_at	CD19 molecu CD19		930	NM_001770	16p11.2	Chromosome -1	2.940458	6.24482953	-1.5817579	0.12448311	0.76331215	DOWN
205289_at	bone morpho BMP2		650	AA583044	20p12	Chromosome 0	5.7263978	4.61484683	0.72624605	0.47347268	0.92191469	UP
204213_at	polymeric immr PLGR		5284	NM_002644	1q31-q41	Chromosome -2	1.194156	6.81087127	-2.3458694	0.02598389	0.57847579	DOWN

Supplementary table 6**Summary of molecule agents predicted using CMap.**

rank	cmap name	mean	n	enrichment	p	specificity	percent non-r
	3 phenoxybenz	-0.743	4	-0.917	0.0001	0.0091	100
	5 luteolin	-0.752	4	-0.902	0.00016	0	100
	7 thioguanosine	-0.707	4	-0.869	0.00058	0.007	100
	28 reserpine	-0.71	3	-0.863	0.00519	0.0201	100
	90 blebbistatin	-0.643	2	-0.857	0.04072	0.0608	100
	31 camptothecin	-0.707	3	-0.847	0.00709	0.1818	100
	12 apigenin	-0.771	4	-0.837	0.00123	0.0163	100
	45 caffeic acid	-0.6	3	-0.799	0.01679	0.0139	100
	51 verteporfin	-0.692	3	-0.785	0.02029	0.0756	100
	27 6-azathymine	-0.628	4	-0.78	0.00487	0.0058	100
	26 omeprazole	-0.629	4	-0.78	0.00483	0.0246	100
	9 morantel	-0.605	5	-0.776	0.00096	0.0048	100
	62 canavanine	-0.606	3	-0.763	0.02722	0.0412	100
	32 thiostrepton	-0.668	4	-0.752	0.00762	0.0463	100
	37 sitosterol	-0.59	4	-0.737	0.00949	0.0306	100
	40 daunorubicin	-0.622	4	-0.726	0.01158	0.114	100
	39 amoxicillin	-0.581	4	-0.726	0.01154	0.0444	100
	22 meticrane	-0.656	5	-0.717	0.00395	0.0195	100
	42 spectinomycin	-0.576	4	-0.714	0.01343	0	100
	11 0175029-000	-0.666	6	-0.714	0.00115	0.0775	83
	47 xylometazolin	-0.584	4	-0.696	0.01804	0.0166	100
	48 rimexolone	-0.617	4	-0.693	0.019	0.0725	100
	49 nipecotic acid	-0.54	4	-0.691	0.01971	0.0266	100
	61 monobenzonitrile	-0.572	4	-0.669	0.02715	0.0405	100
	64 epitiostanol	-0.522	4	-0.664	0.02863	0.0556	100
	66 apramycin	-0.483	4	-0.66	0.03046	0.128	75
	70 ketanserin	-0.556	4	-0.658	0.03127	0.0269	100
	71 levycycloserine	-0.629	4	-0.657	0.03149	0.0773	100
	73 fluspirilene	-0.593	4	-0.656	0.03241	0.0885	100
	25 flunisolid	-0.605	6	-0.654	0.00471	0	100
	78 parthenolide	-0.522	4	-0.652	0.03443	0.2207	100
	86 tolfenamic acid	-0.52	4	-0.642	0.03945	0.0353	100
	89 capsaicin	-0.586	4	-0.64	0.04024	0.0437	100
	93 doxazosin	-0.511	4	-0.633	0.04402	0.1846	75
	95 sulconazole	-0.496	4	-0.63	0.04611	0.1118	75
	94 chlorambucil	-0.46	4	-0.63	0.04589	0.0779	75
	54 procaine	-0.488	5	-0.615	0.02313	0.1371	80
	38 flumetasone	-0.532	6	-0.613	0.01079	0.0248	83
	59 Prestwick-665	-0.525	5	-0.607	0.02643	0.0688	80
	10 resveratrol	-0.535	9	-0.607	0.00104	0.0833	77
	44 solasodine	-0.312	6	-0.591	0.01643	0.0117	66
	76 gallamine triel	-0.485	5	-0.59	0.03394	0.037	80
	83 trimethoprim	-0.45	5	-0.582	0.03773	0.0449	80
	96 nalidixic acid	-0.404	5	-0.564	0.04796	0.0487	60
	67 medrysone	-0.611	6	-0.552	0.03049	0.3676	83
	98 0173570-000	-0.501	6	-0.521	0.0494	0.219	83
	79 methotrexate	-0.361	8	-0.477	0.03456	0.137	75
	87 vorinostat	-0.37	12	-0.388	0.03945	0.6018	66

1	trichostatin A	-0.276	182	-0.188	0	0.8317	53
97	diethylstilbest	0.277	6	0.52	0.04877	0.3053	50
74	myosmine	0.33	6	0.547	0.03297	0.0916	66
80	midodrine	0.386	5	0.59	0.03579	0.1788	80
75	lycorine	0.318	5	0.595	0.03362	0.284	80
69	trimethobenz	0.379	5	0.6	0.03108	0.2147	80
65	thioperamide	0.459	5	0.604	0.02988	0.1348	80
55	digoxigenin	0.356	5	0.62	0.02331	0.1886	80
53	biperiden	0.521	5	0.622	0.02251	0.2546	80
92	canadine	0.492	4	0.638	0.04213	0.2454	75
91	viomycin	0.521	4	0.639	0.04132	0.3526	75
88	methyl dopate	0.295	4	0.643	0.03949	0.04	75
85	chloropyrazin	0.321	4	0.643	0.03937	0.1076	50
82	furazolidone	0.497	4	0.646	0.03768	0.1161	75
29	ethisterone	0.371	6	0.646	0.00564	0	66
81	metoprolol	0.32	4	0.648	0.0368	0.0497	75
77	(+)-chelidoni	0.492	4	0.652	0.03416	0.033	75
72	0316684-000	0.46	4	0.657	0.03177	0.0064	75
68	isocorydine	0.32	4	0.659	0.03105	0.0182	50
63	pivmecillinam	0.284	4	0.667	0.02741	0.0305	50
60	oxybenzone	0.44	4	0.669	0.02678	0.0941	75
57	loxapine	0.219	4	0.676	0.02401	0.0106	50
56	streptozocin	0.53	4	0.678	0.02335	0.0417	75
35	ciprofloxacin	0.271	5	0.678	0.00893	0	60
20	aciclovir	0.394	6	0.68	0.00284	0.0062	66
52	alverine	0.262	4	0.682	0.02194	0	50
33	merbromin	0.335	5	0.684	0.00801	0.0688	80
50	cefotiam	0.455	4	0.688	0.02005	0.0633	75
18	vincamine	0.406	6	0.692	0.00224	0	66
46	furosemide	0.411	4	0.699	0.01695	0.0231	75
43	Prestwick-857	0.389	4	0.707	0.01522	0.0581	75
41	Trolox C	0.332	4	0.723	0.0117	0.0081	75
84	SC-560	0.436	3	0.732	0.03784	0.0748	66
21	dicycloverine	0.457	5	0.735	0.003	0.0055	80
36	bambuterol	0.45	4	0.737	0.00939	0	75
19	adiphenine	0.572	5	0.74	0.00278	0.1915	80
34	perhexiline	0.337	4	0.741	0.00859	0.057	50
30	cinchonine	0.43	4	0.754	0.00702	0.08	75
58	tetracaine	0.337	3	0.764	0.0263	0.0294	66
13	guanabenz	0.515	5	0.775	0.00128	0	80
23	raubasine	0.453	4	0.783	0.00412	0.0064	100
16	cloxacillin	0.551	4	0.82	0.00183	0	100
14	PHA-007675(0.519	4	0.828	0.00133	0	100
8	clorsulon	0.589	4	0.852	0.00072	0	100
4	heptaminol	0.623	5	0.854	0.00014	0.0121	100
24	vigabatrin	0.622	3	0.865	0.00453	0.0177	100
6	iopromide	0.651	4	0.88	0.00028	0	100
17	Gly-His-Lys	0.59	3	0.898	0.00206	0	100
15	carbimazole	0.627	3	0.913	0.00138	0	100
2	podophyllo	0.713	4	0.926	0.00004	0.0048	100