

Table SI. mRNA targets of hsa-miR-200c-3p which are the 20 mRNAs having the highest target score using the miR target prediction and functional annotation database (miRDB).

Target Rank	Target score	Gene symbol	Gene
1	100	VASH2	Vasohibin 2
2	100	HIPK3	Homeodomain interacting protein kinase 3
3	100	MAP2	Microtubule associated protein 2
4	100	ERRFI1	ERBB receptor feedback inhibitor 1
5	100	ZEB1	Zinc finger E-box binding homeobox 1
6	100	NR5A2	Nuclear receptor subfamily 5 group A member 2
7	100	ZEB2	Zinc finger E-box binding homeobox 2
8	99	RECK	Reversion including cysteine rich protein with kazal motifs
9	99	SLIT2	Slit guidance ligand 2
10	99	WIPF1	WAS/WASL interacting protein family member 1
11	99	C11orf95	Chromosome 11 open reading frame 95
12	99	FAM8A1	Family with sequence similarity 8 member A1
13	99	LHFPL6	LHFPL tetraspan subfamily member 6
14	99	AP1S2	Adaptor related protein complex 1 subunit sigma 2
15	99	SEC23A	Sec23 homolog A, coat complex II component
16	99	TBX18	T-box 18
17	99	PTPN21	Protein tyrosine phosphatase, non-receptor type 21
18	99	DNAJC3	DnaJ heat shock protein family (Hsp40) member C3
19	99	QKI	QKI, KH domain containing RNA binding
20	99	RIC1	RIC1 homolog, RAB6A GEF complex partner 1

miR, microRNA.