

Supplementary Table 1. List of primers used for real-time PCR analysis

Gene Symbol	Forward (5' -> 3')	Reverse (5' -> 3')
<i>FGFRL1</i>	acacagccctccaagatgag	gcaggttcttcaggctcagt
<i>NDUFA4</i>	agcttgatccccctctttgt	ctggacgttccttcttcage
<i>GPR177</i>	aggcatctatgggatgtgga	ggaatattcgaagcgctga
<i>LRP5L</i>	ctcaaagctgtgaacgtgga	gcggctctactggtgaagac
<i>GAPDH</i>	gagtcaacggatttggctcgt	ttgattttggaggatctcg

Supplementary Table 2. The list represents 222 genes decreased by more than five-fold in miR-210-transfected cells compared with ncRNA-transfected cells				
Gene Symbol	description	ncRNA	miR-210	Log₂(ratio)
<i>IGHG3</i>	Immunoglobulin heavy chain C gene segment [Source:IMGT/GENE-DB;Acc:IGHG3]	41	1	-5.36
<i>CYB5R3</i>	NADH-cytochrome b5 reductase (EC 1.6.2.2) (B5R) (Diaphorase-1) (Cytochrome b5 reductase 3) [Contains: NADH-cytochrome b5 reductase membrane-bound form; NADH-cytochrome b5 reductase soluble form]. [Source:Uniprot/SWISSPROT;Acc:P00387]	40	1	-5.32
<i>FAM19A4</i>	[Source:Uniprot/SWISSPROT;Acc:Q96LR4]	39	1	-5.29
<i>COBL</i>	Protein cordon-bleu. [Source:Uniprot/SWISSPROT;Acc:O75128]	37	1	-5.21
<i>BST2</i>	Bone marrow stromal antigen 2 precursor (BST-2) (CD317 antigen) (HM1.24 antigen). [Source:Uniprot/SWISSPROT;Acc:Q10589]	35	1	-5.13
<i>ONCO_HUMAN</i>	Oncomodulin (OM) (Parvalbumin beta). [Source:Uniprot/SWISSPROT;Acc:P32930]	35	1	-5.13
<i>POU3F1</i>	POU domain, class 3, transcription factor 1 (Octamer-binding transcription factor 6) (Oct-6) (POU domain transcription factor SCIP). [Source:Uniprot/SWISSPROT;Acc:Q03052]	35	1	-5.13
<i>PTPRT</i>	Receptor-type tyrosine-protein phosphatase T precursor (EC 3.1.3.48) (R-PTP-T) (RPTP-rho). [Source:Uniprot/SWISSPROT;Acc:O14522]	34	1	-5.09
<i>PROM1</i>	Prominin-1 precursor (Prominin-like protein 1) (Antigen AC133) (CD133 antigen). [Source:Uniprot/SWISSPROT;Acc:O43490]	33	1	-5.04
<i>ARTS1_HUMAN</i>	(Aminopeptidase PILS) (Puromycin-insensitive leucyl- specific aminopeptidase) (PILS-AP) (Type 1 tumor necrosis factor receptor shedding aminopeptidase regulator). [Source:Uniprot/SWISSPROT;Acc:Q9NZ08]	29	1	-4.86
<i>Q96I54_HUMAN</i>	-	28	1	-4.81
<i>CALN1</i>	Calneuron-1 (Calcium-binding protein CaBP8). [Source:Uniprot/SWISSPROT;Acc:Q9BXU9]	28	1	-4.81
<i>UGT2B4</i>	UDP-glucuronosyltransferase 2B4 precursor (EC 2.4.1.17) (UDPGT) (Hyodeoxycholic acid) (HLUG25) (UDPGT-1). [Source:Uniprot/SWISSPROT;Acc:P06133]	27	1	-4.75
<i>GABRB1</i>	Gamma-aminobutyric-acid receptor subunit beta-1 precursor (GABA(A) receptor subunit beta-1). [Source:Uniprot/SWISSPROT;Acc:P18505]	26	1	-4.7
<i>DFFB</i>	DNA fragmentation factor subunit beta (EC 3.-.-.-) (DNA fragmentation factor 40 kDa subunit) (DFF-40) (Caspase-activated deoxyribonuclease) (Caspase-activated DNase) (CAD) (Caspase-activated nuclease) (CPAN). [Source:Uniprot/SWISSPROT;Acc:O76075]	26	1	-4.7
<i>GABRD</i>	Gamma-aminobutyric-acid receptor subunit delta precursor (GABA(A) receptor subunit delta). [Source:Uniprot/SWISSPROT;Acc:O14764]	26	1	-4.7
<i>Q5JV89_HUMAN</i>	-	25	1	-4.64
<i>MANEAL</i>	mannosidase, endo-alpha-like isoform 2 [Source:RefSeq_peptide;Acc:NP_689709]	25	1	-4.64
<i>MEGF6</i>	Multiple epidermal growth factor-like domains 6 precursor (EGF-like domain-containing protein 3) (Multiple EGF-like domain protein 3). [Source:Uniprot/SWISSPROT;Acc:O75095]	24	1	-4.58
<i>TIMP4</i>	Metalloproteinase inhibitor 4 precursor (TIMP-4) (Tissue inhibitor of metalloproteinases 4). [Source:Uniprot/SWISSPROT;Acc:Q99727]	23	1	-4.52
<i>TNFAIP6</i>	Tumor necrosis factor-inducible protein TSG-6 precursor (TNF- stimulated gene 6 protein) (Hyaluronate-binding protein). [Source:Uniprot/SWISSPROT;Acc:P98066]	23	1	-4.52
<i>IL24</i>	Interleukin-24 precursor (Suppression of tumorigenicity 16 protein) (Melanoma differentiation-associated gene 7 protein) (MDA-7). [Source:Uniprot/SWISSPROT;Acc:Q13007]	23	1	-4.52
<i>Q8NHA6_HUMAN</i>	Seven transmembrane helix receptor. [Source:Uniprot/SPTREMBL;Acc:Q8NHA6]	23	1	-4.52
<i>PDE1B</i>	Calcium/calmodulin-dependent 3',5'-cyclic nucleotide phosphodiesterase 1B (EC 3.1.4.17) (Cam-PDE 1B) (63 kDa Cam-PDE). [Source:Uniprot/SWISSPROT;Acc:Q01064]	23	1	-4.52
<i>EHBPI1</i>	Signal-induced proliferation-associated protein 1 (Sipa-1) (GTPase- activating protein Spa-1) (p130 SPA-1). [Source:Uniprot/SWISSPROT;Acc:Q96FS4]	22	1	-4.46
<i>C14orf174</i>	C14orf174 protein. [Source:Uniprot/SPTREMBL;Acc:Q2M3P3]	43	2	-4.43
<i>PLEKHG1</i>	[Source:Uniprot/SWISSPROT;Acc:Q9ULL1]	21	1	-4.39
<i>ASTN1</i>	Astrotactin-1 precursor. [Source:Uniprot/SWISSPROT;Acc:O14525]	20	1	-4.32
<i>PSG1</i>	Pregnancy-specific beta-1-glycoprotein 1 precursor (PSBG-1) (Pregnancy-specific beta-1 glycoprotein C/D) (PS-beta-C/D) (Fetal liver non-specific cross-reactive antigen 1/2) (FL-NCA-1/2) (PSG95) (CD66f antigen). [Source:Uniprot/SWISSPROT;Acc:P11464]	20	1	-4.32
<i>SLC6A20</i>	Sodium- and chloride-dependent transporter XTRP3 (Solute carrier family 6 member 20) (Neurotransmitter transporter rB21A homolog). [Source:Uniprot/SWISSPROT;Acc:Q9NP91]	20	1	-4.32
<i>LRRN1</i>	Leucine-rich repeat neuronal protein 1 precursor (Neuronal leucine- rich repeat protein 1) (NLRR-1). [Source:Uniprot/SWISSPROT;Acc:Q6UXK5]	39	2	-4.29
<i>IFT57</i>	estrogen-related receptor beta like 1 [Source:RefSeq_peptide;Acc:NP_060480]	19	1	-4.25
<i>Q96LL3_HUMAN</i>	CDNA FLJ25404 fis, clone TST02888 (Hypothetical protein FLJ25404). [Source:Uniprot/SPTREMBL;Acc:Q96LL3]	19	1	-4.25
<i>COCH</i>	Cochlin precursor (COCH-5B2). [Source:Uniprot/SWISSPROT;Acc:O43405]	19	1	-4.25
<i>NP_683701.2</i>	GREB1 protein isoform b [Source:RefSeq_peptide;Acc:NP_149081]	19	1	-4.25
<i>Q8NAW6_HUMAN</i>	CDNA FLJ34651 fis, clone KIDNE2018167. [Source:Uniprot/SPTREMBL;Acc:Q8NAW6]	19	1	-4.25
<i>TAF6L</i>	TAF6-like RNA polymerase II p300/CBP-associated factor-associated factor 65 kDa subunit 6L (PCAF-associated factor 65 alpha) (PAF65- alpha). [Source:Uniprot/SWISSPROT;Acc:Q9Y6J9]	18	1	-4.17
<i>TMPRSS12</i>	transmembrane protease, serine 12 [Source:RefSeq_peptide;Acc:NP_872365]	18	1	-4.17
<i>Q8NAT4_HUMAN</i>	CDNA FLJ34815 fis, clone NT2NE2007786. [Source:Uniprot/SPTREMBL;Acc:Q8NAT4]	17	1	-4.09
<i>SLC12A3</i>	Solute carrier family 12 member 3 (Thiazide-sensitive sodium-chloride cotransporter) (Na-Cl symporter). [Source:Uniprot/SWISSPROT;Acc:P55017]	17	1	-4.09
<i>CDON</i>	Cell adhesion molecule-related/down-regulated by oncogenes precursor. [Source:Uniprot/SWISSPROT;Acc:Q4KMG0]	17	1	-4.09
<i>SGCG</i>	Gamma-sarcoglycan (Gamma-SG) (35 kDa dystrophin-associated glycoprotein) (35DAG). [Source:Uniprot/SWISSPROT;Acc:Q13326]	17	1	-4.09
<i>DCDC2</i>	[Source:Uniprot/SWISSPROT;Acc:Q9UHG0]	17	1	-4.09

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Gene Symbol	description	ncRNA	miR-210	Log₂(ratio)
<i>PTF1A</i>	Pancreas transcription factor 1 subunit alpha (Pancreas-specific transcription factor 1a) (bHLH transcription factor p48) (p48 DNA-binding subunit of transcription factor PTF1) (PTF1-p48). [Source:Uniprot/SWISSPROT;Acc:Q7RTS3]	34	2	-4.09
<i>IGSF5</i>	IGSF5 protein (Fragment). [Source:Uniprot/SPTREMBL;Acc:Q9NSI5]	17	1	-4.09
<i>TECTA</i>	Alpha-tectorin precursor. [Source:Uniprot/SWISSPROT;Acc:O75443]	17	1	-4.09
<i>LRTM2</i>	Leucine-rich repeat and transmembrane domain-containing protein 2 precursor. [Source:Uniprot/SWISSPROT;Acc:Q8N967]	16	1	-4
<i>MYT1L</i>	Myelin transcription factor 1-like protein (MyT1L protein) (MyT1-L). [Source:Uniprot/SWISSPROT;Acc:Q9UL68]	16	1	-4
<i>Q9H5H6_HUMAN</i>	CDNA: FLJ23429 fis, clone HRC10578. [Source:Uniprot/SPTREMBL;Acc:Q9H5H6]	16	1	-4
<i>LIX1</i>	Protein limb expression 1 homolog. [Source:Uniprot/SWISSPROT;Acc:Q8N485]	32	2	-4
<i>FSTL5</i>	Follistatin-related protein 5 precursor (Follistatin-like 5). [Source:Uniprot/SWISSPROT;Acc:Q8N475]	16	1	-4
<i>SEMG2</i>	Semenogelin-2 precursor (Semenogelin II) (SGII). [Source:Uniprot/SWISSPROT;Acc:Q02383]	15	1	-3.91
<i>Q16653-6</i>	[Source:RefSeq_peptide;Acc:NP_996536]	15	1	-3.91
<i>Q4VXH5_HUMAN</i>	XAGE-4 protein (Fragment). [Source:Uniprot/SPTREMBL;Acc:Q8WWM0]	15	1	-3.91
<i>SERPINA2</i>	Alpha-1-antitrypsin-related protein precursor. [Source:Uniprot/SWISSPROT;Acc:P20848]	15	1	-3.91
<i>GSDMDC1</i>	Gasdermin domain-containing protein 1. [Source:Uniprot/SWISSPROT;Acc:P57764]	15	1	-3.91
<i>Q8TBU5_HUMAN</i>	-	14	1	-3.81
<i>ZNF83</i>	Zinc finger protein 83 (Zinc finger protein HPF1). [Source:Uniprot/SWISSPROT;Acc:P51522]	14	1	-3.81
<i>REG1A</i>	Lithostathine 1 alpha precursor (Pancreatic stone protein) (PSP) (Pancreatic thread protein) (PTP) (Islet of Langerhans regenerating protein) (REG) (Regenerating protein I alpha) (Islet cells regeneration factor) (ICRF). [Source:Uniprot/SWISSPROT;Acc:P05451]	14	1	-3.81
<i>GRIP1</i>	[Source:Uniprot/SWISSPROT;Acc:Q9Y3R0]	28	2	-3.81
<i>FGFRL1</i>	Fibroblast growth factor receptor-like 1 precursor (FGF receptor-like protein 1) (Fibroblast growth factor receptor 5) (FGFR-like protein) (FGF homologous factor receptor). [Source:Uniprot/SWISSPROT;Acc:Q8N441]	277	21	-3.72
<i>SHBG</i>	Sex hormone-binding globulin precursor (SHBG) (Sex steroid-binding protein) (SBP) (Testis-specific androgen-binding protein) (ABP) (Testosterone-estrogen-binding globulin) (Testosterone-estradiol-binding globulin) (TeBG). [Source:Uniprot/SWISSPROT;Acc:P04278]	13	1	-3.7
<i>PPIL2</i>	Peptidyl-prolyl cis-trans isomerase-like 2 (EC 5.2.1.8) (PP1ase) (Rotamase) (Cyclophilin-60) (Cyclophilin-like protein Cyp-60). [Source:Uniprot/SWISSPROT;Acc:Q13356]	13	1	-3.7
<i>IGSF21</i>	Immunoglobulin superfamily member 21 precursor. [Source:Uniprot/SWISSPROT;Acc:Q96ID5]	26	2	-3.7
<i>IL21</i>	Interleukin-21 precursor (IL-21) (Za11). [Source:Uniprot/SWISSPROT;Acc:Q9HBE4]	13	1	-3.7
<i>CLIC6</i>	Chloride intracellular channel 6. [Source:Uniprot/SWISSPROT;Acc:Q96NY7]	13	1	-3.7
<i>SLITRK2</i>	SLIT and NTRK-like protein 2 precursor. [Source:Uniprot/SWISSPROT;Acc:Q9H156]	12	1	-3.58
<i>RIMS1</i>	Regulating synaptic membrane exocytosis protein 1 (Rab3-interacting molecule 1) (RIM 1). [Source:Uniprot/SWISSPROT;Acc:Q86UR5]	12	1	-3.58
<i>LAYN</i>	Layilin precursor. [Source:Uniprot/SWISSPROT;Acc:Q6UX15]	12	1	-3.58
<i>ASB17</i>	Ankyrin repeat and SOCS box protein 17 (ASB-17). [Source:Uniprot/SWISSPROT;Acc:Q8WXJ9]	12	1	-3.58
<i>RAD21L1</i>	Syntaphilin. [Source:Uniprot/SWISSPROT;Acc:O15079]	12	1	-3.58
<i>FGF23</i>	Fibroblast growth factor 23 precursor (FGF-23) (Tumor-derived hypophosphatemia-inducing factor). [Source:Uniprot/SWISSPROT;Acc:Q9GZV9]	12	1	-3.58
<i>CAMTA1</i>	Calmodulin-binding transcription activator 1. [Source:Uniprot/SWISSPROT;Acc:Q9Y6Y1]	23	2	-3.52
<i>APOA2</i>	Apolipoprotein A-II precursor (ApoA-II) (ApoA-II) [Contains: Apolipoprotein A-II(1-76)]. [Source:Uniprot/SWISSPROT;Acc:P02652]	11	1	-3.46
<i>NP_861450.1</i>	LOC283537 protein (OTTHUMP0000018184). [Source:Uniprot/SPTREMBL;Acc:Q6P9B3]	11	1	-3.46
<i>NDUFA5</i>	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 (EC 1.6.5.3) (EC 1.6.99.3) (NADH-ubiquinone oxidoreductase 13 kDa-B subunit) (Complex I-13kD-B) (CI-13kD-B) (Complex I subunit B13). [Source:Uniprot/SWISSPROT;Acc:Q16718]	11	1	-3.46
<i>DDO</i>	D-aspartate oxidase (EC 1.4.3.1) (DASOX) (DDO). [Source:Uniprot/SWISSPROT;Acc:Q99489]	22	2	-3.46
<i>BCORL2</i>	BCoR-like protein 2 (BCL-6 corepressor-like protein 2). [Source:Uniprot/SWISSPROT;Acc:Q8N888]	11	1	-3.46
<i>NM_207504</i>	CDNA FLJ46365 fis, clone TESTI4051054. [Source:Uniprot/SPTREMBL;Acc:Q6ZRG8]	11	1	-3.46
<i>HLA-DPA1</i>	major histocompatibility complex, class II, DP alpha 1	22	2	-3.46
<i>CDKN1C</i>	Cyclin-dependent kinase inhibitor 1C (Cyclin-dependent kinase inhibitor p57) (p57KIP2). [Source:Uniprot/SWISSPROT;Acc:P49918]	11	1	-3.46
<i>DNALI1</i>	Axonemal dynein light intermediate polypeptide 1 (Inner dynein arm light chain, axonemal) (hp28). [Source:Uniprot/SWISSPROT;Acc:O14645]	11	1	-3.46
<i>MCHR2</i>	(MCH-2R) (MCH2) (G-protein coupled receptor 145) (GPRv17). [Source:Uniprot/SWISSPROT;Acc:Q969V1]	21	2	-3.39
<i>DNAH2</i>	dynein heavy chain domain 3 [Source:RefSeq_peptide;Acc:NP_065928]	21	2	-3.39
<i>RTN2</i>	Reticulon-2 (Neuroendocrine-specific protein-like 1) (NSP-like protein 1) (NSPLI). [Source:Uniprot/SWISSPROT;Acc:O75298]	21	2	-3.39
<i>ASIP</i>	[Source:Uniprot/SWISSPROT;Acc:P42127]	10	1	-3.32
<i>AMPH</i>	Amphiphysin. [Source:Uniprot/SWISSPROT;Acc:P49418]	10	1	-3.32
<i>FBXL21</i>	F-box and leucine-rich repeat protein 21 [Source:RefSeq_peptide;Acc:NP_036291]	10	1	-3.32
<i>ANGPTL2</i>	[Source:Uniprot/SWISSPROT;Acc:Q9UKU9]	30	3	-3.32
<i>Q96RZ4_HUMAN</i>	-	30	3	-3.32
<i>C12orf50</i>	C12orf50 protein (Fragment). [Source:Uniprot/SPTREMBL;Acc:Q6P674]	20	2	-3.32

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<i>MSMB</i>	Beta-microseminoprotein precursor (Prostate secreted seminal plasma protein) (Prostate secretory protein PSP94) (PSP-94) (Seminal plasma beta-inhibin) (Immunoglobulin-binding factor) (IGBF) (PN44). [Source:Uniprot/SWISSPROT;Acc:P08118]	20	2	-3.32
<i>HKR2_HUMAN</i>	Krüppel-related zinc finger protein 2 (Protein HKR2) (Zinc finger protein 50) (Zinc finger and SCAN domain-containing protein 22) (Fragment). [Source:Uniprot/SWISSPROT;Acc:P10073]	10	1	-3.32
<i>HIVEP3</i>	[Source:RefSeq_peptide;Acc:NP_078779]	19	2	-3.25
<i>GPR35</i>	Probable G-protein coupled receptor 35. [Source:Uniprot/SWISSPROT;Acc:Q9HC97]	19	2	-3.25
<i>C1orf145</i>	C1orf145 protein (Fragment). [Source:Uniprot/SPTREMBL;Acc:Q8N372]	19	2	-3.25
<i>CBX6</i>	Chromobox protein homolog 6. [Source:Uniprot/SWISSPROT;Acc:O95503]	28	3	-3.22
<i>Q9H353_HUMAN</i>	-	37	4	-3.21
<i>C11orf69</i>	Uncharacterized protein C11orf69. [Source:Uniprot/SWISSPROT;Acc:Q8TAY4]	37	4	-3.21
<i>MYF5</i>	Myogenic factor 5 (Myf-5). [Source:Uniprot/SWISSPROT;Acc:P13349]	9	1	-3.17
<i>PRKAR2B</i>	[Source:Uniprot/SWISSPROT;Acc:P31323]	9	1	-3.17
<i>NOX1</i>	NADPH oxidase homolog 1 (NOX-1) (NOH-1) (NADH/NADPH mitogenic oxidase subunit P65-MOX) (Mitogenic oxidase 1) (MOX1). [Source:Uniprot/SWISSPROT;Acc:Q9Y5S8]	9	1	-3.17
<i>ELOVL3</i>	Elongation of very long chain fatty acids protein 3 (Cold-inducible glycoprotein of 30 kDa). [Source:Uniprot/SWISSPROT;Acc:Q9HB03]	45	5	-3.17
<i>BCL2L10</i>	Apoptosis regulator Bcl-B (Bcl-2-like 10 protein) (Bcl2-L-10) (Anti- apoptotic protein NrH). [Source:Uniprot/SWISSPROT;Acc:Q9HD36]	18	2	-3.17
<i>SEPT8</i>	Septin-8. [Source:Uniprot/SWISSPROT;Acc:Q92599]	18	2	-3.17
<i>C17orf57</i>	C17orf57 protein. [Source:Uniprot/SPTREMBL;Acc:Q49AG9]	18	2	-3.17
<i>SPANXN2</i>	SPANX-N2 protein [Source:RefSeq_peptide;Acc:NP_001009615]	9	1	-3.17
<i>CHST5</i>	(GlcNAc6ST-3) (Intestinal GlcNAc-6- sulfotransferase) (Intestinal N-acetylglucosamine-6-O-sulfotransferase) (I-GlcNAc6ST) (hIGn6ST) (Galactose/N- acetylglucosamine [Source:Uniprot/SWISSPROT;Acc:Q9GZS9]	26	3	-3.12
<i>DUOXA2</i>	[Source:Uniprot/SWISSPROT;Acc:Q1HG44]	26	3	-3.12
<i>FAM26A</i>	Protein FAM26A. [Source:Uniprot/SWISSPROT;Acc:Q86XJ0]	69	8	-3.11
<i>NEURL2</i>	Neutralized-like protein 2. [Source:Uniprot/SWISSPROT;Acc:Q9BR09]	43	5	-3.1
<i>TERT</i>	Telomerase reverse transcriptase (EC 2.7.7.49) (Telomerase catalytic subunit) (HEST2) (Telomerase-associated protein 2) (TP2). [Source:Uniprot/SWISSPROT;Acc:O14746]	34	4	-3.09
<i>NDUFA4</i>	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 (EC 1.6.5.3) (EC 1.6.99.3) (NADH-ubiquinone oxidoreductase MLRQ subunit) (Complex I-MLRQ) (CI-MLRQ). [Source:Uniprot/SWISSPROT;Acc:O00483]	4846	593	-3.03
<i>ART1</i>	GPI-linked NAD(P)(+)-arginine ADP-ribosyltransferase 1 precursor (EC 2.4.2.31) (Mono(ADP-ribosyl)transferase) (CD296 antigen). [Source:Uniprot/SWISSPROT;Acc:P52961]	8	1	-3
<i>NP_001001684.1</i>	CDNA FLJ45831 fis, clone NT2RP8007416. [Source:Uniprot/SPTREMBL;Acc:Q6ZS49]	8	1	-3
<i>AKAP3</i>	A-kinase anchor protein 3 (Protein kinase A-anchoring protein 3) (PRKA3) (A-kinase anchor protein 110 kDa) (AKAP 110) (Sperm oocyte- binding protein) (Fibrousheathin-1) (Fibrousheathin I) (Fibrous sheath protein of 95 kDa) (FSP95). [Source:Uniprot/SWISSPROT;Acc:O75969]	8	1	-3
<i>GTF2IRD2</i>	GTF2I repeat domain containing 2 [Source:RefSeq_peptide;Acc:NP_775808]	8	1	-3
<i>SPATA16</i>	spermatogenesis associated 16 [Source:RefSeq_peptide;Acc:NP_114161]	23	3	-2.94
<i>SPINLW1</i>	Eppin precursor (Epididymal protease inhibitor) (Serine protease inhibitor-like with Kunitz and WAP domains 1) (WAP four-disulfide core domain protein 7) (Protease inhibitor WAP7). [Source:Uniprot/SWISSPROT;Acc:O95925]	23	3	-2.94
<i>OR2L2</i>	Olfactory receptor 2L2 (HTPCRH07). [Source:Uniprot/SWISSPROT;Acc:Q8NH16]	23	3	-2.94
<i>HTR5A</i>	5-hydroxytryptamine 5A receptor (5-HT-5A) (Serotonin receptor 5A) (5- HT-5). [Source:Uniprot/SWISSPROT;Acc:P47898]	30	4	-2.91
<i>CF015_HUMAN</i>	[Source:Uniprot/SWISSPROT;Acc:Q6UXA7]	30	4	-2.91
<i>Q9H7Y2_HUMAN</i>	CDNA FLJ14100 fis, clone MAMMA1000855. [Source:Uniprot/SPTREMBL;Acc:Q9H7Y2]	30	4	-2.91
<i>ZFP42</i>	zinc finger protein 42 [Source:RefSeq_peptide;Acc:NP_777560]	15	2	-2.91
<i>C1orf112</i>	C1orf112 protein. [Source:Uniprot/SPTREMBL;Acc:Q3KNQ1]	15	2	-2.91
<i>EDAR</i>	Tumor necrosis factor receptor superfamily member EDAR precursor (Anhidrotic ectodysplasin receptor 1) (Ectodysplasin-A receptor) (EDA- A1 receptor) (Ectodermal dysplasia receptor) (Downless homolog). [Source:Uniprot/SWISSPROT;Acc:Q9UNE0]	15	2	-2.91
<i>RNF11</i>	RING finger protein 11 (Sid 1669). [Source:Uniprot/SWISSPROT;Acc:Q9Y3C5]	949	129	-2.88
<i>CLDN14</i>	Claudin-14. [Source:Uniprot/SWISSPROT;Acc:O95500]	22	3	-2.87
<i>SGCZ</i>	Zeta-sarcoglycan (Zeta-SG) (ZSG1). [Source:Uniprot/SWISSPROT;Acc:Q96LD1]	29	4	-2.86
<i>PLAC8</i>	Placenta-specific gene 8 protein (Protein C15). [Source:Uniprot/SWISSPROT;Acc:Q9NZF1]	121	17	-2.83
<i>LRRC17</i>	[Source:Uniprot/SWISSPROT;Acc:Q8N6Y2]	7	1	-2.81
<i>PCDH7</i>	[Source:Uniprot/SWISSPROT;Acc:O60245]	35	5	-2.81
<i>TRAV22</i>	T-cell receptor alpha V gene segment [Source:IMG/GENE-DB;Acc:TRAV22]	7	1	-2.81
<i>XAB2</i>	XPA-binding protein 2 (HCNP protein). [Source:Uniprot/SWISSPROT;Acc:Q9HCS7]	35	5	-2.81
<i>FCGR2A</i>	(IgG Fc receptor II-a) (Fc-gamma-RIIa) (CD32 antigen) (CDw32). [Source:Uniprot/SWISSPROT;Acc:P12318]	21	3	-2.81
<i>SEC14L5</i>	-	21	3	-2.81
<i>C10orf53</i>	Uncharacterized protein C10orf53. [Source:Uniprot/SWISSPROT;Acc:Q8N6V4]	21	3	-2.81
<i>LOC729516</i>	-	21	3	-2.81
<i>SLFN1</i>	schlafen-like 1 [Source:RefSeq_peptide;Acc:NP_659427]	14	2	-2.81

Supplementary Table 2. The list represents 222 genes decreased by more than five-fold in miR-210-transfected cells compared with ncRNA-transfected cells				
Gene Symbol	description	ncRNA	miR-210	Log₂(ratio)
<i>LRAT</i>	Lecithin retinol acyltransferase (EC 2.3.1.135) (Phosphatidylcholine--retinol O-acyltransferase). [Source:Uniprot/SWISSPROT;Acc:O95237]	7	1	-2.81
<i>GPR177</i>	Integral membrane protein GPR177 precursor (Protein wntless homolog) (Putative NFkB-activating protein 373). [Source:Uniprot/SWISSPROT;Acc:Q5T9L3]	817	121	-2.76
<i>ADAMTS13</i>	ADAMTS-13 precursor (EC 3.4.24.-) (A disintegrin and metalloproteinase with thrombospondin motifs 13) (ADAM-TS 13) (ADAM-TS13) (von Willebrand factor-cleaving protease) (vWF-cleaving protease) (vWF-CP). [Source:Uniprot/SWISSPROT;Acc:Q76LX8]	47	7	-2.75
<i>PTCHD1</i>	patched domain containing 1 [Source:RefSeq_peptide;Acc:NP_775766]	20	3	-2.74
<i>ACPL2</i>	acid phosphatase-like 2 [Source:RefSeq_peptide;Acc:NP_001032249]	20	3	-2.74
<i>TFEC</i>	transcription factor EC isoform b [Source:RefSeq_peptide;Acc:NP_001018068]	33	5	-2.72
<i>STMN1</i>	phosphoprotein p18) (pp17) (Prosolin) (Metablastin) (Protein Pr22). [Source:Uniprot/SWISSPROT;Acc:P16949]	1003	153	-2.71
<i>Q8WYR5_HUMAN</i>	Tumor rejection antigen. [Source:Uniprot/SPTREMBL;Acc:Q8WYR5]	557	85	-2.71
<i>Q8NHT0_HUMAN</i>	MGC32805 protein. [Source:Uniprot/SPTREMBL;Acc:Q8NHT0]	26	4	-2.7
<i>C20orf39</i>	UPF0338 protein C20orf39. [Source:Uniprot/SWISSPROT;Acc:Q9H7V2]	26	4	-2.7
<i>PSG10</i>	Pregnancy-specific beta-1-glycoprotein 10 precursor (PSBG-10) (PSBG-12). [Source:Uniprot/SWISSPROT;Acc:Q15235]	13	2	-2.7
<i>GJA5</i>	Gap junction alpha-5 protein (Connexin-40) (Cx40). [Source:Uniprot/SWISSPROT;Acc:P36382]	13	2	-2.7
<i>POT15_HUMAN</i>	[Source:Uniprot/SWISSPROT;Acc:Q6S5H4]	13	2	-2.7
<i>NP_001008784.1</i>	CD200 cell surface glycoprotein receptor isoform 2 [Source:RefSeq_peptide;Acc:NP_001008784]	13	2	-2.7
<i>PER3</i>	Period circadian protein homolog 3 (Circadian clock protein PERIOD 3) (hPER3). [Source:Uniprot/SWISSPROT;Acc:P56645]	45	7	-2.68
<i>TEDDM1</i>	putative membrane protein HE9 [Source:RefSeq_peptide;Acc:NP_741997]	44	7	-2.65
<i>DOCK4</i>	Dedicator of cytokinesis protein 4. [Source:Uniprot/SWISSPROT;Acc:Q8N1I0]	25	4	-2.64
<i>RIMS3</i>	Regulating synaptic membrane exocytosis protein 3 (Nim3) (Rab-3-interacting molecule 3) (RIM 3) (RIM3 gamma). [Source:Uniprot/SWISSPROT;Acc:Q9UJD0]	43	7	-2.62
<i>FCRL3</i>	Fc receptor-like 3 precursor [Source:RefSeq_peptide;Acc:NP_443171]	49	8	-2.61
<i>GCKR</i>	Glucokinase regulatory protein (Glucokinase regulator). [Source:Uniprot/SWISSPROT;Acc:Q14397]	6	1	-2.58
<i>NP_631912.2</i>	Na ⁺ /H ⁺ exchanger like domain containing [Source:RefSeq_peptide;Acc:NP_631912]	24	4	-2.58
<i>C2orf51</i>	CDNA FLJ25369 fis, clone TST01830 (Hypothetical protein FLJ25369) (Chromosome 2 open reading frame 51). [Source:Uniprot/SPTREMBL;Acc:Q96LM6]	48	8	-2.58
<i>EN2</i>	Homeobox protein engrailed-2 (Hu-En-2). [Source:Uniprot/SWISSPROT;Acc:P19622]	30	5	-2.58
<i>RFPL1</i>	Ret finger protein-like 1 (RING finger protein 78). [Source:Uniprot/SWISSPROT;Acc:O75677]	30	5	-2.58
<i>GDF3</i>	Growth/differentiation factor 3 precursor (GDF-3). [Source:Uniprot/SWISSPROT;Acc:Q9NR23]	36	6	-2.58
<i>NP_775916.1</i>	CDNA FLJ37357 fis, clone BRAMY2023060 (FLJ37357 protein) (Hypothetical protein FLJ37357). [Source:Uniprot/SPTREMBL;Acc:Q8N1W6]	30	5	-2.58
<i>ISX</i>	intestine-specific homeobox [Source:RefSeq_peptide;Acc:NP_001008494]	24	4	-2.58
<i>CX3CR1</i>	CX3C chemokine receptor 1 (C-X3-C CKR-1) (CX3CR1) (Fractalkine receptor) (G-protein coupled receptor 13) (V28) (Beta chemokine receptor-like 1) (CMK-BRL-1) (CMKBRL1). [Source:Uniprot/SWISSPROT;Acc:P49238]	24	4	-2.58
<i>MIA2</i>	Melanoma inhibitory activity protein 2 precursor. [Source:Uniprot/SWISSPROT;Acc:Q96PC5]	18	3	-2.58
<i>O10D4_HUMAN</i>	Olfactory receptor 10D4. [Source:Uniprot/SWISSPROT;Acc:Q8NGN7]	18	3	-2.58
<i>OR7G2</i>	Olfactory receptor 7G2 (Olfactory receptor 19-13) (OR19-13) (OST260). [Source:Uniprot/SWISSPROT;Acc:Q8NG99]	18	3	-2.58
<i>LRP2</i>	Low-density lipoprotein receptor-related protein 2 precursor (Megalin) (Glycoprotein 330) (gp330). [Source:Uniprot/SWISSPROT;Acc:P98164]	12	2	-2.58
<i>C2orf46</i>	Putative uncharacterized protein C2orf46. [Source:Uniprot/SWISSPROT;Acc:Q6ZSB3]	12	2	-2.58
<i>Q6ZMT9_HUMAN</i>	[Source:Uniprot/SPTREMBL;Acc:Q6ZMT9]	6	1	-2.58
<i>CIQTNF2</i>	Complement C1q tumor necrosis factor-related protein 2 precursor. [Source:Uniprot/SWISSPROT;Acc:Q9BXJ5]	6	1	-2.58
<i>SERHL2</i>	Serine hydrolase-like protein 2 (EC 3.1.-.-). [Source:Uniprot/SWISSPROT;Acc:Q9H4I8]	64	11	-2.54
<i>USHBP1</i>	USH1C-binding protein 1 (Usher syndrome type-1C protein-binding protein 1) (MCC-2) (AIE-75-binding protein). [Source:Uniprot/SWISSPROT;Acc:Q8N6Y0]	29	5	-2.54
<i>VSTM2</i>	[Source:Uniprot/SWISSPROT;Acc:Q8TAG5]	23	4	-2.52
<i>AGC1</i>	sulfate proteoglycan core protein 1 [Contains: Aggrecan core protein 2]. [Source:Uniprot/SWISSPROT;Acc:P16112]	23	4	-2.52
<i>BPY2C</i>	[Source:Uniprot/SWISSPROT;Acc:O14599]	23	4	-2.52
<i>DDOST</i>	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit precursor (EC 2.4.1.119) (Oligosaccharyl transferase 48 kDa subunit) (DDOST 48 kDa subunit). [Source:Uniprot/SWISSPROT;Acc:P39656]	2238	391	-2.52
<i>IRX1</i>	Iroquois-class homeodomain protein IRX-1 (Iroquois homeobox protein 1) (Homeodomain protein IRXA1). [Source:Uniprot/SWISSPROT;Acc:P78414]	34	6	-2.5
<i>CD4</i>	T-cell surface glycoprotein CD4 precursor (T-cell surface antigen T4/Leu-3). [Source:Uniprot/SWISSPROT;Acc:P01730]	17	3	-2.5
<i>Q8NAH5_HUMAN</i>	CDNA FLJ35343 fis, clone PROST2015932. [Source:Uniprot/SPTREMBL;Acc:Q8NAH5]	17	3	-2.5
<i>PSG2</i>	Pregnancy-specific beta-1-glycoprotein 2 precursor (PSBG-2) (Pregnancy-specific beta-1 glycoprotein E) (PS-beta-E). [Source:Uniprot/SWISSPROT;Acc:P11465]	28	5	-2.49
<i>THBS1</i>	Thrombospondin-1 precursor. [Source:Uniprot/SWISSPROT;Acc:P07996]	206	37	-2.48
<i>ANKRD20A3</i>	Ankyrin repeat domain-containing protein 20A3. [Source:Uniprot/SWISSPROT;Acc:Q5VUR7]	22	4	-2.46

Supplementary Table 2. The list represents 222 genes decreased by more than five-fold in miR-210-tansfected cells compared with ncRNA-tansfected cells				
Gene Symbol	description	ncRNA	miR-210	Log₂(ratio)
<i>FNBP1L</i>	Formin-binding protein 1-like (Transducer of Cdc42-dependent actin assembly protein 1) (Toca-1). [Source:Uniprot/SWISSPROT;Acc:Q5TON5]	33	6	-2.46
<i>HBB</i>	Hemoglobin subunit beta (Hemoglobin beta chain) (Beta-globin). [Source:Uniprot/SWISSPROT;Acc:P68871]	22	4	-2.46
<i>PIWIL3</i>	Piwi-like protein 3. [Source:Uniprot/SWISSPROT;Acc:Q7Z3Z3]	22	4	-2.46
<i>MAGEA11</i>	Melanoma-associated antigen 11 (MAGE-11 antigen). [Source:Uniprot/SWISSPROT;Acc:P43364]	22	4	-2.46
<i>KCNGB1</i>	Potassium voltage-gated channel subfamily G member 1 (Voltage-gated potassium channel subunit Kv6.1) (KH2). [Source:Uniprot/SWISSPROT;Acc:Q9UIX4]	11	2	-2.46
<i>ADRA1A</i>	Alpha-1A adrenergic receptor (Alpha 1A-adrenoceptor) (Alpha 1A- adrenoceptor) (Alpha-1C adrenergic receptor) (Alpha adrenergic receptor 1c). [Source:Uniprot/SWISSPROT;Acc:P35348]	11	2	-2.46
<i>NP_001001343.1</i>	MGC27121 gene (MGC27121), mRNA [Source:RefSeq_dna;Acc:NM_001001343]	38	7	-2.44
<i>GPNMB</i>	Transmembrane glycoprotein NMB precursor (Transmembrane glycoprotein HGFIN). [Source:Uniprot/SWISSPROT;Acc:Q14956]	54	10	-2.43
<i>C1orf173</i>	C1orf173 protein. [Source:Uniprot/SPTREMBL;Acc:Q6GMR8]	27	5	-2.43
<i>NTS</i>	Neurotensin/neuromedin N precursor [Contains: Large neuromedin N (NmN- 125); Neuromedin N (NmN) (NN); Neurotensin (NT); Tail peptide]. [Source:Uniprot/SWISSPROT;Acc:P30990]	27	5	-2.43
<i>XKR4</i>	XK-related protein 4. [Source:Uniprot/SWISSPROT;Acc:Q5GH76]	16	3	-2.42
<i>C12orf59</i>	C12orf59 protein. [Source:Uniprot/SPTREMBL;Acc:Q4KMG9]	159	30	-2.41
<i>USP29</i>	specific-processing protease 29) (Deubiquitinating enzyme 29). [Source:Uniprot/SWISSPROT;Acc:Q9HBJ7]	21	4	-2.39
<i>PLA2G7</i>	acylhydrolase) (LDL-associated phospholipase A2) (LDL-PLA(2)) (2-acetyl-1-alkylglycerophosphocholine esterase) (1-alkyl-2-acetyl-glycerophosphocholine esterase) [Source:Uniprot/SWISSPROT;Acc:Q13093]	21	4	-2.39
<i>NP_001006656.1</i>	MSTP119. [Source:Uniprot/SPTREMBL;Acc:Q7Z2S5]	21	4	-2.39
<i>CEECAM1</i>	cerebral endothelial cell adhesion molecule 1 [Source:RefSeq_peptide;Acc:NP_057258]	21	4	-2.39
<i>ZDHHC22</i>	Putative palmitoyltransferase ZDHHC22 (EC 2.3.1.-) (Zinc finger DHHC domain-containing protein 22) (DHHC-22). [Source:Uniprot/SWISSPROT;Acc:Q8N966]	21	4	-2.39
<i>ZNF101</i>	Zinc finger protein 101 (Zinc finger protein HZF12). [Source:Uniprot/SWISSPROT;Acc:Q8IZC7]	47	9	-2.38
<i>ADRA1B</i>	Alpha-1B adrenergic receptor (Alpha 1B-adrenoceptor) (Alpha 1B- adrenoceptor). [Source:Uniprot/SWISSPROT;Acc:P35368]	26	5	-2.38
<i>ACOX3</i>	Acyl-coenzyme A oxidase 3, peroxisomal (EC 1.3.3.6) (Pristanoyl-CoA oxidase) (Branched-chain acyl-CoA oxidase) (BRCAcox). [Source:Uniprot/SWISSPROT;Acc:O15254]	31	6	-2.37
<i>Q8N7S0_HUMAN</i>	CDNA FLJ40424 fis, clone TESTI2039026. [Source:Uniprot/SPTREMBL;Acc:Q8N7S0]	31	6	-2.37
<i>PHKG1</i>	Phosphorylase b kinase gamma catalytic chain, skeletal muscle isoform (EC 2.7.11.19) (Phosphorylase kinase subunit gamma 1). [Source:Uniprot/SWISSPROT;Acc:Q16816]	31	6	-2.37
<i>LRP5L</i>	low density lipoprotein receptor-related protein 5-like [Source:RefSeq_peptide;Acc:NP_872298]	67	13	-2.37
<i>AAK1</i>	AP2-associated protein kinase 1 (EC 2.7.11.1) (Adaptor-associated kinase 1). [Source:Uniprot/SWISSPROT;Acc:Q2M218]	221	43	-2.36
<i>KIAA0195</i>	KIAA0195 (KIAA0195), mRNA [Source:RefSeq_dna;Acc:NM_014738]	40	8	-2.32
<i>Q8NA59_HUMAN</i>	CDNA FLJ35816 fis, clone TESTI2006109. [Source:Uniprot/SPTREMBL;Acc:Q8NA59]	30	6	-2.32
<i>EMID1</i>	EMI domain-containing protein 1 precursor (Protein Emu1) (Emilin and multimerin domain-containing protein 1). [Source:Uniprot/SWISSPROT;Acc:Q96A84]	5	1	-2.32
<i>C6orf122</i>	Uncharacterized protein C6orf122. [Source:Uniprot/SWISSPROT;Acc:Q5T6M2]	20	4	-2.32
<i>ERC1</i>	ELKS/RAB6-interacting/CAST family member 1 (RAB6-interacting protein 2) (ERC protein 1). [Source:Uniprot/SWISSPROT;Acc:Q8IUD2]	20	4	-2.32
<i>USP6</i>	Ubiquitin carboxyl-terminal hydrolase 6 (EC 3.1.2.15) (Ubiquitin thioesterase 6) (Ubiquitin-specific-processing protease 6) (Deubiquitinating enzyme 6) (Proto-oncogene TRE-2). [Source:Uniprot/SWISSPROT;Acc:P35125]	20	4	-2.32
<i>GRB14</i>	Growth factor receptor-bound protein 14 (GRB14 adapter protein). [Source:Uniprot/SWISSPROT;Acc:Q14449]	15	3	-2.32
<i>ALG10</i>	(Asparagine-linked glycosylation protein 10 homolog A). [Source:Uniprot/SWISSPROT;Acc:Q5BKT4]	15	3	-2.32
<i>SPTB</i>	Spectrin beta chain, erythrocyte (Beta-I spectrin). [Source:Uniprot/SWISSPROT;Acc:P11277]	15	3	-2.32
<i>C3AR1</i>	[Source:Uniprot/SWISSPROT;Acc:Q16581]	10	2	-2.32
<i>HTR2C</i>	5-hydroxytryptamine 2C receptor (5-HT-2C) (Serotonin receptor 2C) (5- HT2C) (5-HTR2C) (5HT-1C). [Source:Uniprot/SWISSPROT;Acc:P28335]	10	2	-2.32
<i>O95724_HUMAN</i>	Reverse transcriptase (Fragment). [Source:Uniprot/SPTREMBL;Acc:O95724]	5	1	-2.32

The fifteen genes predicted as a miR-210 target gene by microCosm, TargetScan or PicTar were highlighted in gray.

Supplementary Table 3. The list represents 811 genes predicted as a miR-210 target gene by microCosm, TargetScan and PicTar

<i>ABCC1</i>	<i>C10orf122</i>	<i>CD8A</i>	<i>EFNA3</i>	<i>GIT2</i>	<i>IMP3</i>
<i>ABCC6P2</i>	<i>C11orf2</i>	<i>CD99</i>	<i>EGFL9</i>	<i>GJA7</i>	<i>IQCK</i>
<i>ABCD1</i>	<i>C11orf58</i>	<i>CDC2L1</i>	<i>EGR3</i>	<i>GLI1</i>	<i>IRGC</i>
<i>ABCD4</i>	<i>C11orf59</i>	<i>CDC2L2</i>	<i>EGR4</i>	<i>GLIS1</i>	<i>IRX6</i>
<i>ABHD2</i>	<i>C11orf63</i>	<i>CDH26</i>	<i>EHD2</i>	<i>GLS2</i>	<i>ISCA2</i>
<i>ABL2</i>	<i>C12orf34</i>	<i>CDKN1C</i>	<i>EIF3S9</i>	<i>GNA15</i>	<i>ISCU</i>
<i>ABTB1</i>	<i>C12orf48</i>	<i>CDX2</i>	<i>ELA2</i>	<i>GNAT1</i>	<i>ITGA8</i>
<i>AC006273.1</i>	<i>C14orf148</i>	<i>CEND1</i>	<i>ELAC2</i>	<i>GNG3</i>	<i>ITGB4</i>
<i>AC008772.1</i>	<i>C15orf43</i>	<i>CETN3</i>	<i>ELFN2</i>	<i>GNG8</i>	<i>IWS1</i>
<i>AC008898.1</i>	<i>C15orf52</i>	<i>CFHR2</i>	<i>ELOVL6</i>	<i>GOLGA1</i>	<i>JOSD2</i>
<i>AC009967.5</i>	<i>C15orf62</i>	<i>CHAD</i>	<i>EMID1</i>	<i>GOLGA2LY1</i>	<i>KAAG1</i>
<i>AC020916.6</i>	<i>C16orf35</i>	<i>CHD1L</i>	<i>EML2</i>	<i>GOLPH3</i>	<i>KCMF1</i>
<i>AC040977.2</i>	<i>C16orf70</i>	<i>CHEK2</i>	<i>ENPP2</i>	<i>GPD1L</i>	<i>KCNH5</i>
<i>AC109322.1</i>	<i>C17orf57</i>	<i>CHES1</i>	<i>ENSA</i>	<i>GPR153</i>	<i>KCNMB1</i>
<i>AC114498.5</i>	<i>C17orf64</i>	<i>CHN1</i>	<i>EPB41L1</i>	<i>GPR17</i>	<i>KCNN2</i>
<i>AC133485.1</i>	<i>C17orf83</i>	<i>CHRD1</i>	<i>EPGN</i>	<i>GPR177</i>	<i>KCNQ2</i>
<i>AC141586.2</i>	<i>C18orf10</i>	<i>CHRM3</i>	<i>EPHA2</i>	<i>GPR19</i>	<i>KIAA0664</i>
<i>AC226119.1</i>	<i>C18orf34</i>	<i>CHRN1</i>	<i>EPHB2</i>	<i>GPR39</i>	<i>KIAA0748</i>
<i>ACP1</i>	<i>C19orf16</i>	<i>CHRN2</i>	<i>EPS15</i>	<i>GPR87</i>	<i>KIAA1622</i>
<i>ACTA1</i>	<i>C19orf34</i>	<i>CHST1</i>	<i>ERP27</i>	<i>GRIA2</i>	<i>KIAA1751</i>
<i>ACTC1</i>	<i>C19orf59</i>	<i>CHST12</i>	<i>EVL</i>	<i>GRIK2</i>	<i>KIAA1755</i>
<i>ACTG1</i>	<i>C1orf102</i>	<i>CHUK</i>	<i>EVPL</i>	<i>GRIN3B</i>	<i>KIAA2013</i>
<i>ACTL7A</i>	<i>C1orf110</i>	<i>CIRBP</i>	<i>EXOSC10</i>	<i>GRINL1A</i>	<i>KIF13B</i>
<i>ADA</i>	<i>C1orf111</i>	<i>CLDN15</i>	<i>F11R</i>	<i>GRM5</i>	<i>KIF20A</i>
<i>ADAMTS7</i>	<i>C1orf112</i>	<i>CLEC11A</i>	<i>F7</i>	<i>GRM6</i>	<i>KIR2DL1</i>
<i>ADCY7</i>	<i>C1orf116</i>	<i>CLEC19A</i>	<i>FAHD1</i>	<i>GSC</i>	<i>KLHDC4</i>
<i>AGPAT2</i>	<i>C1orf128</i>	<i>CNGB1</i>	<i>FAM105B</i>	<i>GSTA1</i>	<i>KLHL35</i>
<i>AGTRAP</i>	<i>C1orf133</i>	<i>COL16A1</i>	<i>FAM108A1</i>	<i>GSTA2</i>	<i>KLRA1</i>
<i>AGTRL1</i>	<i>C1orf167</i>	<i>COL4A2</i>	<i>FAM108A2</i>	<i>GSTTP2</i>	<i>KLRC4</i>
<i>AGXT2L2</i>	<i>C1orf184</i>	<i>COL4A3</i>	<i>FAM116A</i>	<i>GSTZ1</i>	<i>KRTAP5-4</i>
<i>AHRH</i>	<i>C1orf86</i>	<i>COL6A2</i>	<i>FAM120A</i>	<i>H2AFY</i>	<i>LAIR2</i>
<i>AIFM3</i>	<i>C1orf88</i>	<i>COL9A3</i>	<i>FAM12A</i>	<i>HAAO</i>	<i>LAMA3</i>
<i>AKAP7</i>	<i>C1QTNF4</i>	<i>COMMD4</i>	<i>FAM149A</i>	<i>HAS1</i>	<i>LAMA5</i>
<i>AKAP9</i>	<i>C1S</i>	<i>COX10</i>	<i>FAM25A</i>	<i>HCN4</i>	<i>LAMC3</i>
<i>AKR1CL2</i>	<i>C20orf103</i>	<i>CPNE2</i>	<i>FAM25B</i>	<i>HDLBP</i>	<i>LCE3E</i>
<i>AL132661.1</i>	<i>C20orf195</i>	<i>CRI1</i>	<i>FAM25C</i>	<i>HEATR7B2</i>	<i>LCK</i>
<i>AL161662.2</i>	<i>C2orf32</i>	<i>CREB3L3</i>	<i>FAM5B</i>	<i>HELZ</i>	<i>LCN8</i>
<i>ALDH3A1</i>	<i>C2orf46</i>	<i>CRHBP</i>	<i>FAM73B</i>	<i>HHIP</i>	<i>LDHB</i>
<i>ALDH4A1</i>	<i>C3orf21</i>	<i>CRLF3</i>	<i>FAM84A</i>	<i>HIF3A</i>	<i>LEAP2</i>
<i>ALKBH3</i>	<i>C3orf30</i>	<i>CRYBA4</i>	<i>FAM90A1</i>	<i>HIRIP3</i>	<i>LG3</i>
<i>ALLC</i>	<i>C4orf23</i>	<i>CS</i>	<i>FAM90A10</i>	<i>HIST1H1B</i>	<i>LHPP</i>
<i>AMBN</i>	<i>C6orf125</i>	<i>CSPP1</i>	<i>FAM90A18</i>	<i>HIST1H2AK</i>	<i>LIPE</i>
<i>ANAPC7</i>	<i>C6orf145</i>	<i>CST3</i>	<i>FAM90A3</i>	<i>HK3</i>	<i>LIP1</i>
<i>ANK1</i>	<i>C6orf184</i>	<i>CTBP2</i>	<i>FAM90A7</i>	<i>HLXB9</i>	<i>LMAN1L</i>
<i>ANKRD24</i>	<i>C6orf191</i>	<i>CUEDC2</i>	<i>FAM90A8</i>	<i>HMG20B</i>	<i>LMTK2</i>
<i>ANKRD47</i>	<i>C6orf57</i>	<i>CUL9</i>	<i>FAM90A9</i>	<i>HOXA3</i>	<i>LMX1B</i>
<i>ANKS3</i>	<i>C9</i>	<i>CYB5R2</i>	<i>FAM9B</i>	<i>HPCA</i>	<i>LOC399947</i>
<i>AP000355.2</i>	<i>C9orf129</i>	<i>CYGB</i>	<i>FANCB</i>	<i>HPCAL1</i>	<i>LOC643224</i>
<i>AP000889.1</i>	<i>C9orf134</i>	<i>CYP11B1</i>	<i>FANCE</i>	<i>HRH2</i>	<i>LRFN1</i>
<i>APOBEC3G</i>	<i>C9orf78</i>	<i>CYP2F1</i>	<i>FANK1</i>	<i>HSD17B1</i>	<i>LRP5L</i>
<i>ARFRP1</i>	<i>C9orf85</i>	<i>DAB1</i>	<i>FBXL12</i>	<i>HSD17B7</i>	<i>LRPAP1</i>
<i>ARHGAP17</i>	<i>C9orf86</i>	<i>DARC</i>	<i>FBXL16</i>	<i>HSPA8</i>	<i>LRRC62</i>
<i>ARHGFEF17</i>	<i>C9orf93</i>	<i>DBN1</i>	<i>FBXL17</i>	<i>HSPBAP1</i>	<i>LRRC68</i>
<i>ARMC1</i>	<i>CACNA2D2</i>	<i>DCHS1</i>	<i>FBXW5</i>	<i>HTATSF1</i>	<i>LRRC8A</i>
<i>ARMC4</i>	<i>CALCOCO1</i>	<i>DCTN1</i>	<i>FER1L4</i>	<i>HTRA1</i>	<i>LRRC8D</i>
<i>ARSD</i>	<i>CAMK2G</i>	<i>DDX24</i>	<i>FEZF1</i>	<i>HYAL1</i>	<i>LRRC8D</i>
<i>ASCC1</i>	<i>CAPN10</i>	<i>DDX51</i>	<i>FGD1</i>	<i>IDI2</i>	<i>LRSAM1</i>
<i>ASCL1</i>	<i>CAPN9</i>	<i>DEAF1</i>	<i>FGF10</i>	<i>IGF1R</i>	<i>LY6H</i>
<i>ASL</i>	<i>CARD9</i>	<i>DEC1</i>	<i>FGF18</i>	<i>IGHA2</i>	<i>LYN</i>
<i>ATG4D</i>	<i>CASQ1</i>	<i>DEFB118</i>	<i>FGF22</i>	<i>IGHV3-13</i>	<i>LYPLA2</i>
<i>ATN1</i>	<i>CCBP2</i>	<i>DGKA</i>	<i>FGFRL1</i>	<i>IGHV3-16</i>	<i>MAG12</i>
<i>ATP12A</i>	<i>CCDC146</i>	<i>DGKG</i>	<i>FHIT</i>	<i>IGHV3-23</i>	<i>MAN1B1</i>
<i>ATP1B1</i>	<i>CCDC17</i>	<i>DHRS3</i>	<i>FHOD1</i>	<i>IGHV3-35</i>	<i>MAP6</i>
<i>ATP2B3</i>	<i>CCDC24</i>	<i>DHX58</i>	<i>FKBP9L</i>	<i>IGHV3-38</i>	<i>MAP7D1</i>
<i>ATP6V0C</i>	<i>CCDC28A</i>	<i>DIMT1L</i>	<i>FMO1</i>	<i>IGHV3-47</i>	<i>MARCH4</i>
<i>AVP</i>	<i>CCDC28B</i>	<i>DNAH11</i>	<i>FNTA</i>	<i>IGHV3-66</i>	<i>MARK1</i>
<i>AVPR1B</i>	<i>CCDC53</i>	<i>DNAJC16</i>	<i>FOXD2</i>	<i>IGKV4-1</i>	<i>MAST1</i>
<i>B3GALT5</i>	<i>CCDC95</i>	<i>DNAJC4</i>	<i>FOXP3</i>	<i>IGLC7</i>	<i>MBD5</i>
<i>B4GALT5</i>	<i>CCDC97</i>	<i>DNAJC8</i>	<i>FRAP1</i>	<i>IGLL3</i>	<i>MCCD1</i>
<i>B4GALT7</i>	<i>CCKBR</i>	<i>DPEP2</i>	<i>FRY</i>	<i>IGSF21</i>	<i>MCM4</i>
<i>BAIAP3</i>	<i>CCNB1IP1</i>	<i>DPYSL5</i>	<i>FTSJ2</i>	<i>IHPK2</i>	<i>MCM8</i>
<i>BCAS2</i>	<i>CCNL2</i>	<i>DRD5</i>	<i>FUT8</i>	<i>IHPK3</i>	<i>MDGA1</i>
<i>BCL2L12</i>	<i>CCT3</i>	<i>DTX1</i>	<i>GOS2</i>	<i>IKBKKG</i>	<i>MEF2D</i>
<i>BDNF</i>	<i>CD180</i>	<i>DUOXA1</i>	<i>GAK</i>	<i>IL11RA</i>	<i>MEGF6</i>
<i>BICD1</i>	<i>CD22</i>	<i>DUSP12</i>	<i>GARNL1</i>	<i>IL17RC</i>	<i>MEIS1</i>
<i>BRP44L</i>	<i>CD276</i>	<i>DUSP28</i>	<i>GAS6</i>	<i>IL18</i>	<i>MFAP3</i>
<i>BTBD10</i>	<i>CD300LD</i>	<i>E2F3</i>	<i>GBGT1</i>	<i>IL3RA</i>	<i>MFSB4</i>
<i>BX324178.1</i>	<i>CD55</i>	<i>EBF3</i>	<i>GDAP1L1</i>	<i>ILVBL</i>	<i>MID1IP1</i>

Supplementary Table 3. The list represents 811 genes predicted as a miR-210 target gene by microCosm, TargetScan and PicTar

<i>MINA</i>	<i>PCP4</i>	<i>RP4-565E6.1</i>	<i>SURF5</i>	<i>WNT9B</i>
<i>MLL2</i>	<i>PCSK7</i>	<i>RPS27</i>	<i>SUSD1</i>	<i>XCR1</i>
<i>MPV17L</i>	<i>PDCL</i>	<i>RSC1A1</i>	<i>SUZ12</i>	<i>XKR5</i>
<i>MRPL36</i>	<i>PDE3A</i>	<i>RUFY1</i>	<i>SYNE2</i>	<i>XRRA1</i>
<i>MRPS30</i>	<i>PDLIM3</i>	<i>RUVBL2</i>	<i>SYNGAP1</i>	<i>YIPF3</i>
<i>MS4A8B</i>	<i>PDX1</i>	<i>SACM1L</i>	<i>SYT15</i>	<i>YY1</i>
<i>MT4</i>	<i>PDZD4</i>	<i>SAMD13</i>	<i>TAC4</i>	<i>Z97180.2</i>
<i>MUC4</i>	<i>PEX10</i>	<i>SAP30L</i>	<i>TAF6</i>	<i>ZBTB12</i>
<i>MX1</i>	<i>PEX13</i>	<i>SARNP</i>	<i>TBC1D16</i>	<i>ZCCHC11</i>
<i>MXD4</i>	<i>PEX5L</i>	<i>SCGB1C1</i>	<i>TBC1D28</i>	<i>ZDHHC12</i>
<i>MYH11</i>	<i>PHF15</i>	<i>SCN1B</i>	<i>TBC1D8</i>	<i>ZDHHC4</i>
<i>MYO15B</i>	<i>PHF23</i>	<i>SCN9A</i>	<i>TCF7L2</i>	<i>ZFPM1</i>
<i>MYOHD1</i>	<i>PHKG2</i>	<i>SCOC</i>	<i>TCHP</i>	<i>ZHX2</i>
<i>MYT1L</i>	<i>PHTF1</i>	<i>SCRT1</i>	<i>TFAP2A</i>	<i>ZMAT4</i>
<i>NAT14</i>	<i>P11KAP2</i>	<i>SCYL1</i>	<i>TFCP2</i>	<i>ZMIZ2</i>
<i>NDUFA4</i>	<i>PIK3R5</i>	<i>SDCCAG8</i>	<i>TGFBRAP1</i>	<i>ZMYM2</i>
<i>NEFM</i>	<i>PIK4CA</i>	<i>SDF2</i>	<i>THOC5</i>	<i>ZMYM6</i>
<i>NEK3</i>	<i>PKNOX2</i>	<i>SDHALP1</i>	<i>TIAM1</i>	<i>ZNF227</i>
<i>NEUROD2</i>	<i>PLA1A</i>	<i>SDHALP2</i>	<i>TIGD2</i>	<i>ZNF274</i>
<i>NEUROG3</i>	<i>PLCB3</i>	<i>SEC24C</i>	<i>TIGD5</i>	<i>ZNF397</i>
<i>NFIB</i>	<i>PLCD3</i>	<i>SEH1L</i>	<i>TIMM17B</i>	<i>ZNF403</i>
<i>NFIL3</i>	<i>PLK1</i>	<i>SENP8</i>	<i>TIMP1</i>	<i>ZNF418</i>
<i>NKX2-5</i>	<i>PLXNC1</i>	<i>SEPT12</i>	<i>TLX1</i>	<i>ZNF45</i>
<i>NKX2-8</i>	<i>PODN</i>	<i>SERPINA12</i>	<i>TMEM142A</i>	<i>ZNF462</i>
<i>NOMO1</i>	<i>POF1B</i>	<i>SERPINA3</i>	<i>TMEM16B</i>	<i>ZNF467</i>
<i>NOMO3</i>	<i>POLR2I</i>	<i>SETD2</i>	<i>TMEM16D</i>	<i>ZNF583</i>
<i>NOX4</i>	<i>POU2AF1</i>	<i>SF3B1</i>	<i>TMEM194B</i>	<i>ZNF585B</i>
<i>NP_660343.1</i>	<i>PPAP2A</i>	<i>SF3B4</i>	<i>TMEM195</i>	<i>ZNF720</i>
<i>NP_683701.2</i>	<i>PIIE</i>	<i>SF3B5</i>	<i>TMEM204</i>	<i>ZNF827</i>
<i>NPHP4</i>	<i>PPP1CC</i>	<i>SH2D7</i>	<i>TMEM208</i>	<i>ZRANB3</i>
<i>NPM3</i>	<i>PPP1R12C</i>	<i>SH3BGR</i>	<i>TMEM40</i>	<i>ZSCAN20</i>
<i>NPR1</i>	<i>PPP1R16A</i>	<i>SH3BGRL</i>	<i>TNFRSF13C</i>	
<i>NPSR1</i>	<i>PPP3R1</i>	<i>SH3KBP1</i>	<i>TNIP1</i>	
<i>NPTX1</i>	<i>PRHOXNB</i>	<i>SHC3</i>	<i>TNIP3</i>	
<i>NQO2</i>	<i>PRICKLE4</i>	<i>SHCBP1</i>	<i>TNPO3</i>	
<i>NR6A1</i>	<i>PRIMA1</i>	<i>SIDT2</i>	<i>TOM1L2</i>	
<i>NRG4</i>	<i>PRMT2</i>	<i>SIL1</i>	<i>TOR1A</i>	
<i>NRGN</i>	<i>PRPF38B</i>	<i>SIN3B</i>	<i>TP53TG3</i>	
<i>NSDHL</i>	<i>PRR11</i>	<i>SIPAIL3</i>	<i>TPST1</i>	
<i>NSUN5</i>	<i>PRRG2</i>	<i>SIX1</i>	<i>TRAFD1</i>	
<i>NT5C1B</i>	<i>PRSS16</i>	<i>SLC12A8</i>	<i>TRAV4</i>	
<i>NT5DC1</i>	<i>PSG2</i>	<i>SLC15A1</i>	<i>TRIM14</i>	
<i>NTN4</i>	<i>PSME3</i>	<i>SLC16A14</i>	<i>TRIM17</i>	
<i>NUDT6</i>	<i>PTAFR</i>	<i>SLC20A1</i>	<i>TSEN2</i>	
<i>NUP133</i>	<i>PTCHD3</i>	<i>SLC25A28</i>	<i>TSNAXIP1</i>	
<i>NUPL2</i>	<i>PTGES2</i>	<i>SLC26A3</i>	<i>TSPAN10</i>	
<i>NUPR1</i>	<i>PTS</i>	<i>SLC2A1</i>	<i>TSPAN14</i>	
<i>NYX</i>	<i>PYDC1</i>	<i>SLC37A1</i>	<i>TTC12</i>	
<i>OAF</i>	<i>PYY</i>	<i>SLC38A5</i>	<i>TTC13</i>	
<i>OBSCN</i>	<i>Q6ZRP6_HUMAN</i>	<i>SLC43A1</i>	<i>TTC24</i>	
<i>ODF3</i>	<i>Q9BZU2_HUMAN</i>	<i>SLC4A11</i>	<i>TTC4</i>	
<i>ODZ2</i>	<i>Q9HBR8_HUMAN</i>	<i>SLC6A19</i>	<i>TTF1</i>	
<i>ODZ3</i>	<i>RAB26</i>	<i>SLITRK5</i>	<i>TLL1</i>	
<i>OGG1</i>	<i>RAB27B</i>	<i>SMARCA4</i>	<i>TXNL6</i>	
<i>OLFML2A</i>	<i>RABGAP1L</i>	<i>SMCHD1</i>	<i>UBASH3A</i>	
<i>OLIG3</i>	<i>RAD52</i>	<i>SORBS2</i>	<i>UBE2O</i>	
<i>OR10G8</i>	<i>RALGDS</i>	<i>SOX21</i>	<i>UBN2</i>	
<i>OR2T8</i>	<i>RANBP5</i>	<i>SOX30</i>	<i>UBQLN1</i>	
<i>OR2V2</i>	<i>RASAL2</i>	<i>SPACA3</i>	<i>UMODL1</i>	
<i>OR4K14</i>	<i>RASSF1</i>	<i>SPACA4</i>	<i>UNC45A</i>	
<i>OR4P1P</i>	<i>RASSF6</i>	<i>SPO11</i>	<i>UNC5A</i>	
<i>OR5M13P</i>	<i>RBM3</i>	<i>SPRR2F</i>	<i>URM1</i>	
<i>OSBPL2</i>	<i>RBMY1C</i>	<i>SPTB</i>	<i>UROS</i>	
<i>OTOS</i>	<i>RBPJL</i>	<i>SQLE</i>	<i>USP21</i>	
<i>OTP</i>	<i>RC3H1</i>	<i>SREBF1</i>	<i>USP6NL</i>	
<i>OTUB1</i>	<i>RECK</i>	<i>SRMS</i>	<i>VIT</i>	
<i>P2RX1</i>	<i>RETN</i>	<i>SRPX</i>	<i>VRK1</i>	
<i>P2RY10</i>	<i>RG9MTD1</i>	<i>ST3GAL3</i>	<i>VSIG6</i>	
<i>P2RY11</i>	<i>RG54</i>	<i>ST6GALNAC6</i>	<i>VWA5B1</i>	
<i>P4HA3</i>	<i>RIBC2</i>	<i>STAB1</i>	<i>WDFY2</i>	
<i>PABPC1L</i>	<i>RLN3</i>	<i>STARD3NL</i>	<i>WDR20</i>	
<i>PADI1</i>	<i>RNF207</i>	<i>STAT6</i>	<i>WDR22</i>	
<i>PANX3</i>	<i>RNF208</i>	<i>STIP1</i>	<i>WDR38</i>	
<i>PAOX</i>	<i>RNF212</i>	<i>STT3B</i>	<i>WDR5B</i>	
<i>PARK2</i>	<i>RP1-163G9.1</i>	<i>STX11</i>	<i>WDR6</i>	
<i>PBX1</i>	<i>RP11-181G12.1</i>	<i>STX1B2</i>	<i>WDR64</i>	
<i>PCBP4</i>	<i>RP11-191L9.1</i>	<i>STXBP5</i>	<i>WDR66</i>	
<i>PCDH21</i>	<i>RP11-397P14.1</i>	<i>SUMO3</i>	<i>WISP2</i>	