

GeneSymbol	pmID	LCFVector	shRNAClone	tarSeq
0.GFP control	n/a	shGFP (#444)	n/a	n/a
0.GFP scrambled control	n/a	shSCR-GFP	n/a	n/a
0.IPTG scrambled control	n/a	shSCR-IPTG	n/a	n/a
0.Neo/G418 scrambled control	n/a	shSCR-neo	n/a	n/a
0.Puro scrambled control	n/a	shSCR (#665)	n/a	n/a
Ager	NM_007425	sh311.1	1	GCCACTTGTGCTAAGCTGTAA
Ager	NM_007425	sh311.2	2	CGAGTCCGAGTCTACCAGATT
Ager	NM_007425	sh311.3	3	GAGCTGAATCAGTCAGAGGAA
Ager	NM_007425	sh311.4	4	TGGCAAAGAAACACTCGTGAA
Ager	NM_007425	sh311.5	5	GCAGCTAGAATGGAAACTGAA
Anapc7	NM_019805	sh418.1	1	CCTGCCTGTTTCCAAGTAGTA
Anapc7	NM_019805	sh418.2	2	CCATCTGAAATTGAAGTGAAA
Anapc7	NM_019805	sh418.3	3	GCCTTAAGTAAGACTTCCAAA
Anapc7	NM_019805	sh418.4	4	GCTGCTCACCATGAGTAACAA
Anapc7	NM_019805	sh418.5	5	CGAAGGTCTCATAGAGTGTTA
Atg5	NM_053069	sh123.1	1	GCCAAGTATCTGTCTATGATA
Atg5	NM_053069	sh123.2	2	CCTTGGAACATCACAGTACAT
Atg5	NM_053069	sh123.3	3	GCAGAACCATACTATTTGCTT
Atg5	NM_053069	sh123.4	4	GCATCTGAGCTACCCAGATAA
Atg5	NM_053069	sh123.5	5	CCCTGAAATGGCATTATCCAA
Atg7	NM_028835	sh122.1	1	CCAGCTCTGAACTCAATAATA
Atg7	NM_028835	sh122.2	2	GCCTGGCATTGATAAATGTA
Atg7	NM_028835	sh122.3	3	CGTGACACATAGCATCATCTT
Atg7	NM_028835	sh122.4	4	CCTAAAGAAGTACCACTTCTA
Atg7	NM_028835	sh122.5	5	CCAAGGTCAAAGGACAAAGAT
Atp13a2	NM_029097	sh124.1	1	GAACTCTCTCAAGAGACAGTA
Atp13a2	NM_029097	sh124.2	2	GCCGAAACACTCGTTATAGAA
Atp13a2	NM_029097	sh124.3	3	GCTCATTCTCTACACAATCAA
Atp13a2	NM_029097	sh124.4	4	CACGCCGAAACACTCGTTATA
Atp13a2	NM_029097	sh124.5	5	GCCGACCACTACTACTGGTAT
Atr	XM_147046	sh284.1	1	GATTATTGAATGGGTGAACAA
Atr	XM_147046	sh284.2	2	CATTCCCTGATCCTACATCAT
B2m	NM_009735	sh160.1	1	CGCCTCACATTGAAATCCAAA
B2m	NM_009735	sh160.2	2	GCCGAACATACTGAACTGCTA

B2m	NM_009735	sh160.3	3	GCAGAGTTAAGCATGCCAGTA
B2m	NM_009735	sh160.4	4	CCAAATGCTGAAGAACGGGAA
B2m	NM_009735	sh160.5	5	CGGCCTGTATGCTATCCAGAA
Becn1	NM_019584	sh202.1	1	CCACATGTTTACAATACCAAA
Becn1	NM_019584	sh202.2	2	CCCTATGGAAATCATTCTAT
Becn1	NM_019584	sh202.3	3	GCGGGAGTATAGTGAGTTTAA
Becn1	NM_019584	sh202.4	4	AGCGGGAGTATAGTGAGTTTA
Becn1	NM_019584	sh202.5	5	CGGACAGTTTGGCACAATCAA
Cav1	NM_007616	sh148.1	1	CCAGTTAGATTTAGGGATTTA
Cav1	NM_007616	sh148.2	2	CCGCTTGTTGTCTACGATCTT
Cav1	NM_007616	sh148.3	3	CGACGTGGTCAAGATTGACTT
Cav1	NM_007616	sh148.4	4	TGAAGCTATTGGCAAGATATT
Cav1	NM_007616	sh148.5	5	GCTTCCTGATTGAGATTCAGT
Cnp	NM_009923	sh326.1	1	CCCAGATGCATCCAAGAACTT
Cnp	NM_009923	sh326.2	2	CGACACTTTATTTCTGGAGATC
Cnp	NM_009923	sh326.3	3	CGCTGATGCTTACAAGATCAT
Cnp	NM_009923	sh326.4	4	CGAGTGCAAGACGCTATTCAT
Cnp	NM_009923	sh326.5	5	GACTAAGAAGATGGAGGTCAA
Cxcl10	NM_021274	sh199.1	1	GCATTGTATATGGAAGAACTT
Cxcl10	NM_021274	sh199.2	2	CCGCTGCAACTGCATCCATAT
Cxcl10	NM_021274	sh199.3	3	GATGTCTGAATCCGGAATCTA
Cxcl10	NM_021274	sh199.4	4	GCCAGTGAGAATGAGGGCCAT
Cxcl10	NM_021274	sh199.5	5	TCCGGAATCTAAGACCATCAA
Cxcl11	NM_019494	sh200.1	1	CGCCTCATAATGCAGGCAATA
Cxcl11	NM_019494	sh200.2	2	CTTCTGTAATTTACCCGAGTA
Cxcl11	NM_019494	sh200.3	3	GTTACTATGAAGGCTCATAAA
Cxcl11	NM_019494	sh200.4	4	CAAGGCTTCCTTATGTTCAA
Cxcl11	NM_019494	sh200.5	5	TGCGACAAAGTTGAAGTGATT
Cxcl9	NM_008599	sh162.1	1	CCATCTTCAGAGCTTATTCTA
Cxcl9	NM_008599	sh162.2	2	GCTACACTGAAGAACGGAGAT
Cxcl9	NM_008599	sh162.3	3	GTCGTCGTTCAAGGAAGACTA
Cxcl9	NM_008599	sh162.4	4	AGCTGATGAAAGAATGGGAAA
Cxcl9	NM_008599	sh162.5	5	GATCCACTACAAATCCCTCAA
Cxcr3	NM_009910	sh157.1	1	GCCGATGTTCTGCTGGTGTTA
Cxcr3	NM_009910	sh157.2	2	CCCGCAACTGTGGTCGAGAAA

Cxcr3	NM_009910	sh157.3	3	GCCTGAACTTTGACAGAACCT
Cxcr3	NM_009910	sh157.4	4	GCCTCGGACTTTGCCTTTCTT
Cxcr3	NM_009910	sh157.5	5	CCGTCATCTTCACGGAGAGAA
Cybb	NM_007807	sh277.1	1	GCTGCCAAATGTCCCAGTAAT
Cybb	NM_007807	sh277.2	2	CCTGCCTGAATTTCAACTGTA
Cybb	NM_007807	sh277.3	3	CCAAGTTAGAATGGCACACAT
Cybb	NM_007807	sh277.4	4	CGGAGAGTTTGAAGAGCATA
Cybb	NM_007807	sh277.5	5	GCCTGGAACTACCTAAGATA
Edg1	NM_007901	sh286.1	1	CCATGTAAACTGGGTCAAGTT
Edg1	NM_007901	sh286.2	2	CGGATCGTATCTTGTTGCAAA
Edg1	NM_007901	sh286.3	3	CGTCTGGAAACGTCAATTCTT
Edg1	NM_007901	sh286.4	4	CATGGAATTTAGCCGCAGCAA
Edg1	NM_007901	sh286.5	5	GTCTCTGACTATGGGAACTAT
Efnb2	NM_010111	sh161.1	1	CGGGTGTTACAGTAGCCTTAT
Efnb2	NM_010111	sh161.2	2	GCACAATTAAGAAGGAGAATA
Efnb2	NM_010111	sh161.3	3	CGCATCAGGATGCATCATCTT
Efnb2	NM_010111	sh161.4	4	GTGACGTTATCATACCACTAA
Efnb2	NM_010111	sh161.5	5	CCAAGATGTGAAATTCACCAT
EII2	XM_484289	sh036.1	1	GCCACAAGAATTTAATTCCTT
EII2	XM_484289	sh036.2	2	CCAATCCTCCTCAGACTGTAA
EII2	XM_484289	sh036.3	3	CCACCGACATTAATGGTTAT
EII2	XM_484289	sh036.4	4	GCCTGCATTCTCCTGTGATAA
Ephb4	NM_010144	sh156.1	1	CGGATCTGAAATGGGTGACTT
Ephb4	NM_010144	sh156.2	2	CCTGTGTCGAATTGGGTATTA
Ephb4	NM_010144	sh156.3	3	CGGTTATGATCCTCACGGAAT
Ephb4	NM_010144	sh156.4	4	CGAGGTCAAGTATCATGAGAA
Ephb4	NM_010144	sh156.5	5	GTCTTCTATTACGAGAGCGAA
Foxo3a	NM_019740	sh201.1	1	GAGTCCATCATCCGTAGTGAA
Foxo3a	NM_019740	sh201.2	2	CCCGGACAAACGGCTCACTTT
Foxo3a	NM_019740	sh201.3	3	GCTTCGCAACGATCCAATGAT
Foxo3a	NM_019740	sh201.4	4	CGGCACCATGAATCTGAATGA
Foxo3a	NM_019740	sh201.5	5	GTTTGGACCTTCGTCTCTGAA
Fyn	NM_008054	sh138.1	1	GCTCGGTTGATTGAAGACAAT
Fyn	NM_008054	sh138.2	2	CCTGTATGGAAGGTTACAAT
Fyn	NM_008054	sh138.3	3	CCTTTGGAAACCCAAGAGGTA

Fyn	NM_008054	sh138.4	4	GCTCTGAAGTTGCCAAACCTT
Fyn	NM_008054	sh138.5	5	CATCCCGAACTACAACAACCTT
Gad1	NM_008077	sh147.1	1	CCCTCTTTACAGAACCAGAAT
Gad1	NM_008077	sh147.2	2	GTAGACATACTCCTCAACTAT
Gad1	NM_008077	sh147.3	3	GCCCTACAACGTATGATACTT
Gad1	NM_008077	sh147.4	4	CATTTCGATCCAATCCAGGAAA
Gad1	NM_008077	sh147.5	5	GCATTTCGATCCAATCCAGGAA
Gfi1	NM_010278	sh425.1	1	CGGCTCCTACAAATGCATCAA
Gfi1	NM_010278	sh425.2	2	CCTCATCACTCATAGCAGAAA
Gfi1	NM_010278	sh425.3	3	ACCTCATCACTCATAGCAGAA
Gfi1	NM_010278	sh425.4	4	CCAGCCCTACACGCTGCCTTT
Gfi1	NM_010278	sh425.5	5	CGCAGTTATCAGAGTAAGGAA
gzmb	NM_013542	sh174.1	1	CGAGAATGTTATCTAATGCTA
gzmb	NM_013542	sh174.2	2	CGAGTTTCTTATCCTGGATAA
gzmb	NM_013542	sh174.3	3	GCCTTACTTTTCGATCAAGGAT
gzmb	NM_013542	sh174.4	4	CCTATGGATATAAGGATGGTT
gzmb	NM_013542	sh174.5	5	CAGACTATAATCCTAAGACAT
Hck	NM_010407	sh153.1	1	GCACGAATCATCGAGGACAAT
Hck	NM_010407	sh153.2	2	CTGTACGACTATGAGGCTATT
Hck	NM_010407	sh153.3	3	GCATCACTGGTGTGTAAGATT
Hck	NM_010407	sh153.4	4	CATGACAAACTGGTGAAGCTA
Hck	NM_010407	sh153.5	5	ACCTACAACAAGCACACCAAAA
Icam1	NM_010493	sh159.1	1	CCAACCTCTTCTTGATGTATTT
Icam1	NM_010493	sh159.2	2	CGCCAGAGAAAAGATCAGGATA
Icam1	NM_010493	sh159.3	3	GCAGAGGACCTTAACAGTCTA
Icam1	NM_010493	sh159.4	4	GAACCTGTTCTTCTCATGCAA
Ifng	NM_008337	sh145.1	1	CCTTTACTTCACTGACCAATA
Ifng	NM_008337	sh145.2	2	CGTCATTGAATCACACCTGAT
Ifng	NM_008337	sh145.3	3	GAGCCAGATTATCTCTTTCTA
Ifng	NM_008337	sh145.4	4	GCAGAGCCAGATTATCTCTTT
Ifng	NM_008337	sh145.5	5	CAGTCATTGAAAGCCTAGAAA
Ikbke	NM_019777	sh192.1	1	GCAGGAGCTTTGTAATGATAT
Ikbke	NM_019777	sh192.2	2	CCAGAAGATTCAGTGTTGTTT
Ikbke	NM_019777	sh192.3	3	CGACTCATCGAACGGTTACAT
Ikbke	NM_019777	sh192.4	4	CCAAAGTTCGTCCCTAAGGTT

lkbke	NM_019777	sh192.5	5	CTGGACGATGATGAGAAGTTT
lkbke	NM_019777	sh342.1	1	GCAGGAGCTTTGTAATGATAT
lkbke	NM_019777	sh342.2	2	CCAGAAGATTCAGTGTTGTTT
lkbke	NM_019777	sh342.3	3	CGACTCATCGAACGGTTACAT
lkbke	NM_019777	sh342.4	4	CCAAAGTTCGTCCCTAAGGTT
lkbke	NM_019777	sh342.5	5	CTGGACGATGATGAGAAGTTT
lrf1	NM_008390	sh146.1	1	GCTCTGTACTAGGAGACAATT
lrf1	NM_008390	sh146.2	2	GCCGATACAAAGCAGGAGAAA
lrf1	NM_008390	sh146.3	3	CGAAGACCTTATGAAGCTCTT
lrf1	NM_008390	sh146.4	4	AGATGGACATTATACCAGATA
lrf1	NM_008390	sh146.5	5	CTCTTCTGTCTATGGAGACTT
lrf2	NM_008391	sh164.1	1	CCAGGAATAGATAAACCAGAT
lrf2	NM_008391	sh164.2	2	GCAGATAAATTCCAATACGAT
lrf2	NM_008391	sh164.3	3	CCTACGCAGAAAGCGAAACTA
lrf2	NM_008391	sh164.4	4	CCAGTGTCATCAAGAAGACAT
lrf2	NM_008391	sh164.5	5	CGGTCCTGACTTCAGCTATAA
Map1lc3b	NM_026160	sh127.1	1	GCATCCTAAGTTGCCAATAAA
Map1lc3b	NM_026160	sh127.2	2	CCCAGTGATTATAGAGCGATA
Map1lc3b	NM_026160	sh127.3	3	GCTCAATGCTAACCAAGCCTT
Met	NM_008591	sh152.1	1	CGGGATTCTTCCAAACACTT
Met	NM_008591	sh152.2	2	GCACGACAAATACGTTGAAAT
Met	NM_008591	sh152.3	3	CCCGACGTGAACACATTTGAT
Met	NM_008591	sh152.4	4	CCTTGAATAGAATCACAGATA
Mlxip	NM_177582	sh281.1	1	CCTTTCTCTAAGGCTTTACAT
Mlxip	NM_177582	sh281.2	2	CCCAGTTAAAGAAACAGTTAA
Mlxip	NM_177582	sh281.3	3	CCTGTCTTTACGATGACCTTA
Mlxip	NM_177582	sh281.4	4	CCCTGTCTTTACGATGACCTT
Mlxip	NM_177582	sh281.5	5	CCAACTGCATTGCCCTTAGTT
Mlxipl	NM_021455	sh279.1	1	CCGCAGGATACAGTTTCTGAA
Mlxipl	NM_021455	sh279.2	2	CCTCTTTGTTGCCAGAAGAAT
Mlxipl	NM_021455	sh279.3	3	GCATCCTCATCCGACCTTTAT
Mlxipl	NM_021455	sh279.4	4	ACCCACATCTTCAAACACTACAT
Mlxipl	NM_021455	sh279.5	5	CCTCTGGAATACTGGAATTAA
Mmp13	NM_008607	sh254.1	1	GCCTTACATTTGGCTTAATTT
Mmp13	NM_008607	sh254.2	2	GCTCCGAGAAATGCAATCTTT

Mmp13	NM_008607	sh254.3	3	CCGTGACCTTATGTTTATCTT
Mmp13	NM_008607	sh254.4	4	GCCCTGATGTTTCCCATCTAT
Mmp13	NM_008607	sh254.5	5	GCACTACTTGAAATCATACTA
Ncf1	NM_010876	sh287.1	1	CGGCTATTTCCCATCCATGTA
Ncf1	NM_010876	sh287.2	2	CCCATCATCCTTCAGACCTAT
Ncf1	NM_010876	sh287.3	3	CCCAGGTGGTTTGATGGACAA
Ncf1	NM_010876	sh287.4	4	GAACCGTATGTAACCATCAAA
Ncf1	NM_010876	sh287.5	5	TGTGTACATGTTCTGGTTAA
Nox1	NM_172203	sh276.1	1	CGTGATTACCAAGGTTGTCAT
Nox1	NM_172203	sh276.2	2	GAAAGAAGATTCTTGGCTAAA
Nox1	NM_172203	sh276.3	3	GCATGGAAGTAGGACAGTATA
Nox4	NM_015760	sh278.1	1	GCATCAAATAACCACCTGTAT
Nox4	NM_015760	sh278.2	2	GCCAGTATATTATTCTCCATT
Nox4	NM_015760	sh278.3	3	CCTACGCAATAAGAGTTTCTA
Nox4	NM_015760	sh278.4	4	GCATTAGTCTTAACCAGACAT
Nox4	NM_015760	sh278.5	5	GCCAGAATACTACTACATTCA
Nr3c1	NM_008173	sh348.1	1	GCTGTTTATCTCCACTGAATT
Nr3c1	NM_008173	sh348.2	2	CCCAGAGATGTTAGCTGAAAT
Nr3c1	NM_008173	sh348.3	3	CCGCAGCCAGATTTATCCAAA
Nr3c1	NM_008173	sh348.4	4	GAGGTGTTATATGCAGGATAT
Nr3c1	NM_008173	sh348.5	5	CTGGAGATACAATCTTATCAA
Prf1	NM_011073	sh165.1	1	CTATGCATAGAGAGGCCACTA
Prf1	NM_011073	sh165.2	2	GCCCATTTGGTGGTAAGCAAT
Prf1	NM_011073	sh165.3	3	CGGTGTCGTGTGGAACAATAA
Prf1	NM_011073	sh165.4	4	CCACTCCAAGGTAGCCAATTT
Prf1	NM_011073	sh165.5	5	AGGGTGAAATTCTCCTACCAT
Prickle1	NM_001033217	sh454.1	1	TTTGTCTGCTTCACGTGTAAT
Prickle1	NM_001033217	sh454.2	2	GACAACATGGTTACGAATAAG
Prickle1	NM_001033217	sh454.3	3	AGCAAGCCAAGACCGTCATTA
Prickle1	NM_001033217	sh454.4	4	TTACTGCCATGCTAGATTT
Prickle1	NM_001033217	sh454.5	5	CGGAAGCTACGACATCGAAAT
Ptgr2	NM_029880	sh436.1	1	GAGAGAGATTTACGGTATTAA
Ptgr2	NM_029880	sh436.2	2	GCCATATATCTGCTGGATCTA
Ptgr2	NM_029880	sh436.3	3	CCTGGCTTGACTTCCTTGAT
Ptgr2	NM_029880	sh436.4	4	GATGTCTACTTTGACAATGTT

Ptgr2	NM_029880	sh436.5	5	GAATGAGAACAGCCACATCAT
Ripk1	NM_009068	sh151.1	1	CCATCTTTGATAACACCACTA
Ripk1	NM_009068	sh151.2	2	CCTGAATGACATCAATGCAAA
Ripk1	NM_009068	sh151.3	3	CGTGACTTTACATTAAGATA
Ripk1	NM_009068	sh151.4	4	GCCAAATCTAAGCCAAATGTA
Ripk1	NM_009068	sh151.5	5	GCATTGTCCTTTGGGCAATAT
Sall1	NM_021390	sh416.1	1	CCCAAGATAGCCTGTCATCTT
Sall1	NM_021390	sh416.2	2	GCTGCGCTGAATTCTTTGAAT
Sall1	NM_021390	sh416.3	3	GCACTATCTGTGGAAGAGCAT
Sall1	NM_021390	sh416.4	4	GCGCTGAATTCTTTGAATTAT
Sall1	NM_021390	sh416.5	5	CCACCAAATGTCACTGCCTTT
Serpinb9	NM_009256	sh167.1	1	CCAGTAATTTATAGCCAGTTA
Serpinb9	NM_009256	sh167.2	2	GCCTCTGCCATCATAGAATTT
Serpinb9	NM_009256	sh167.3	3	GACACATATAACCTCGCCTAT
Serpinb9	NM_009256	sh167.4	4	AGAGCACTGATGTTGAGGTTT
Serpinb9	NM_009256	sh167.5	5	GTACTIONCTTAGAGTGGCCAA
Slc9a3r1	NM_012030	sh320.1	1	GACCGAATTGTGGAGGTCAAT
Slc9a3r1	NM_012030	sh320.2	2	CCCTTCAGCAATGGAGAGATA
Slc9a3r1	NM_012030	sh320.3	3	GAACGGTGAGAATGTGGAGAA
Slc9a3r1	NM_012030	sh320.4	4	GAGTTCTTCAAGAAGTGCAAA
Slc9a3r1	NM_012030	sh320.5	5	CGATACCAGTGAGGAGCTAAA
Src	NM_009271	sh154.1	1	GCAAGATCACTAGACGGGAAT
Src	NM_009271	sh154.2	2	CCTAAATGTGAAACACTACAA
Src	NM_009271	sh154.3	3	GCGGCTGCAGATTGTCAATAA
Src	NM_009271	sh154.4	4	CGAGCCGCAATATCCTAGTA
Src	NM_009271	sh154.5	5	GTGGCTTACTACTCAAACAT
Stat1	NM_009283	sh158.1	1	GCCGAGAACATACCAGAGAAT
Stat1	NM_009283	sh158.2	2	CCGAAGAACTTCACTCTCTTA
Stat1	NM_009283	sh158.3	3	GCTGCCTATGATGTCTCGTTT
Stat1	NM_009283	sh158.4	4	GCTGTTACTTTCCAGATATT
Stat1	NM_009283	sh158.5	5	GCTCACTCAGAACACTCTGAT
Stat3	NM_011486	sh163.1	1	CCTAACTTTGTGGTTCCAGAT
Stat3	NM_011486	sh163.2	2	CGACTTTGATTTCAACTACAA
Stat3	NM_011486	sh163.3	3	CACCATTCAATTGATGCAGTTT
Stat3	NM_011486	sh163.4	4	CCTGAGTTGAATTATCAGCTT

Stat3	NM_011486	sh163.5	5	GCAGGTATCTTGAGAAGCCAA
Tbk1	NM_019786	sh193.1	1	CCTCGGAGGAACAAAGAAGTA
Tbk1	NM_019786	sh193.2	2	GCTGGCCGAGAACAATCATAT
Tbk1	NM_019786	sh193.3	3	GCGTTTAAAGATAAGTCGGAA
Tbk1	NM_019786	sh193.4	4	CCAGAATCAGAATTTCTCATT
Tbk1	NM_019786	sh193.5	5	GCTGGGTGAGATTTGAGACAT
Tbk1	NM_019786	sh343.1	1	CCTCGGAGGAACAAAGAAGTA
Tbk1	NM_019786	sh343.2	2	GCTGGCCGAGAACAATCATAT
Tbk1	NM_019786	sh343.3	3	GCGTTTAAAGATAAGTCGGAA
Tbk1	NM_019786	sh343.4	4	CCAGAATCAGAATTTCTCATT
Tbk1	NM_019786	sh343.5	5	GCTGGGTGAGATTTGAGACAT
Timeless	NM_011589	sh168.1	1	CCCTGATTTGTGTTTCATATT
Timeless	NM_011589	sh168.2	2	GCTGCTTCTTTGAGTCAGCAA
Timeless	NM_011589	sh168.3	3	CCTCACCAACTACTACGAGAT
Timeless	NM_011589	sh168.4	4	GCTATACAAGAAACGGAGAAA
Timeless	NM_011589	sh168.5	5	GCAATGGAGAACGAACAGTTT
Tnf	NM_013693	sh184.1	1	GCTATCTCATACCAGGAGAAA
Tnf	NM_013693	sh184.2	2	CGATGGGTTGTACCTTGTCTA
Tnf	NM_013693	sh184.3	3	CCTCCCTCTCATCAGTTCTAT
Tnf	NM_013693	sh184.4	4	CACTCAGATCATCTTCTCAA
Traf2	NM_009422	sh166.1	1	GCCTGGTTTGCACAATAGGAA
Traf2	NM_009422	sh166.2	2	GCGATCTTCATCAAAGCTATT
Traf2	NM_009422	sh166.3	3	GAGTTGGAAGTATCCACCTAT
Traf2	NM_009422	sh166.4	4	CCTTGAAAGAATACGAGAGCT
Traf2	NM_009422	sh166.5	5	GCCCTGAGTAACAAGGTGCAA
Traf3ip2	NM_134000	sh419.1	1	CCAGTCTAAATCATGCTCAA
Traf3ip2	NM_134000	sh419.2	2	CCACCATCTAATGGAGTGGAA
Traf3ip2	NM_134000	sh419.3	3	GCTTCAGAACTCATGTTTA
Traf3ip2	NM_134000	sh419.4	4	CCCTGAACTTGGCAAAGATAT
Traf3ip2	NM_134000	sh419.5	5	CCTGAACTTGGCAAAGATATT
Trp53	NM_011640	sh312.1	1	CCGACCTATCCTTACCATCAT
Trp53	NM_011640	sh312.2	2	CGGGCGTAAACGCTTCGAGAT
Trp53	NM_011640	sh312.3	3	CGGGCGTAAACGCTTCGAGAT
Trp53	NM_011640	sh312.4	4	ACCGCCGTACAGAAGAAGAAA
Trp53	NM_011640	sh312.5	5	AGAGTATTTACCCTCAAGAT

Yes1	NM_009535	sh155.1	1	GCGGAAAGATTACTTCTGAAT
Yes1	NM_009535	sh155.2	2	CCAGGTACAATGATGCCAGAA
Yes1	NM_009535	sh155.3	3	GCTGCTCTGTATGGTCGATT
Yes1	NM_009535	sh155.4	4	GCCAGTCATTATGGAGTGGAA
Yes1	NM_009535	sh155.5	5	CCTTGTATGATTATGAAGCTA