

	name		primer		name		primer
1	Ubc	F	CCGCCTTCTCAGTATGCAGA		42 Itga2b (CD41)	F	GAGTTTTCCGGGAGACAAGC
		R	TGCTTCGGATATAGCCCTGG			R	TTCTCAATTGGAAACGCGAGC
2	Ef2	F	CTTTGAACGGAGGGCAGATG		43 Pf4	F	TCCCGAAGAAAGCGATGGAG
		R	CGCCTCTGTTCCGCTTTG			R	CCATTCTCAGGGTGCGTATGA
3	Actb	F	GGCTGTATTCCCTCCATCG		44 Gata1	F	TATGGCAAGACGGCACTCTAC
		R	CCAGTTGGTAACAATGCCATGT			R	GGTGTCCAAGAACGTGTGTT
4	Mcm6	F	GGACGAGTCGGCACTAAAGA		45 Klf1	F	CTGCGCCAAGCCTATTACA
		R	AACGTGATCGTAGTGTGAGA			R	GGCGAACTCTTGTCACAGC
5	Cdkn1c (p57)	F	CAGCCAGCCTTCGACCAT		46 Sdpr	F	CCAGGTGAAGAGACTGGAGAAC
		R	TACGCGTATCACTGGGAAG			R	TCCTTCAAAACACTGGCA
6	Cdkn1a (p21)	F	CGG TGT CAG AGT CTA GGG GA		47 Vwf	F	TGTATGGGCAAGTGAACCCCTT
		R	ATC ACC AGG ATT GGA CAT GG			R	GAAGTCGTCCCAAGAGGCG
7	Ccne1 (Cyclin	F	GGCAAAATGGCGCTGTTTT		48 Selp	F	CCCCATTCTGTGAAGCGTTG
		R	GGATGAAAGAGCAGGGGTCC			R	ACCAAGAGCGAGCCTTCATC
8	Ccne2 (Cyclin	F	GGCCCCGCTATATATTGAGTT		49 Gfi1b	F	GTTGCTGAACACAGAGCCTTC
		R	TAAACGGCTCGCTTTGACA			R	CAGGAGAAGCTGGGCTTGA
9	Cdc6	F	TTTTGGAAAGTTGATGGGAAC		50 Epb4.1	F	GCAGGTTGAGGTTGCTCTT
		R	GGGTCAAAGCAGCAAAGAG			R	AATGTCTCGCGAAGCTGAA
10	Cdk2	F	AAGAAGATCCGGCTCGACAC		51 Epb4.2	F	ACTGCATTAGGAAGAGCCC
		R	ACATCCAGCAGCTTGACGATA			R	CAACAAGGCGCCCAAC
11	E2f4	F	CCCTTCTACCTCTTTGAGCC		52 Pbx1	F	ACAGTCTCCAGGTGGATACC
		R	CATGCACTCTCGTGGGG			R	CCTGCCAACCTCCATTAGCA
12	Cdkn2d (p19)	F	GCCTTGCAGGTCAATGATGTTT		53 Fli1	F	TGTCAGGAGAGGAGCATGGGA
		R	GAACCTTCAGGGTGTCCAGG			R	GATAAGGATCTGGCTGGGGC
13	Cdk6	F	CCTTACCTCGTGGTCTC		54 Hba-a1	F	CAACTTCAAGCTCCTGAGCCA
		R	GAACCTCCACGAAAAGAGGCT			R	AACGGTACTGGAGGTGAGC
14	Cdkn2c (p18)	F	GGGGCACTAGAGCACTTACT		55 Hbb-b1	F	TGCTTGCATAGTTGTGTTGAC
		R	AAATTGGGATTAGCACCTCTGAG			R	AACACGAGCGCTGCC
15	Ccng2 (Cyclin	F	TTCTGAGCTGCCACTATC		56 Hbb-b2	F	AACCTGGACAACCTCAAGGG
		R	AAGGTGCACTCTGATCGC			R	GGTGAATCTTGCCAGGT
16	Ccnf (Cyclin F	F	CGCGCTTGAGAGCGGT		57 Bpgm	F	GCTGTCTCGGCTAGGTAC
		R	GTAGGATAACAGAAACACTTGGCAC			R	AGCCTAAAGAAACGGCAGCT
17	Mki67 (Ki 67)	F	ATCATTGACCGCTCCTTAGGT		58 Plid6	F	CGCAAGGCAGGGATACAGG
		R	GCTCGCCTTATGTTTCT			R	GTGTGGTCCAGTTGAGGGAG
18	Cena2 (Cyclin	F	AAGAAACAGAGTGTGAAGATGCC		59 Chac1	F	GGAAAGCGGACTTTGCTAT
		R	AGACATATCCATAGCATGTGGTGA			R	CGGTCTCAAGGAGGTCAC
19	Ccnb1 (Cyclin	F	GTGAGTGACGTAGACGACAGA		60 Atp1b2	F	CACACCAGGCTTGATGATTCG
		R	CACTTACAGACCTGATGGT			R	CTGTGCTTGGATGGGTCGT
20	Ccnb2 (Cyclin	F	CGACGGTGTCCAGTGATTTG		61 Epor	F	GGGCTCCGAAGAACT
		R	TCTGAGGTTTCTTCGCCACC			R	ATGACTTTCGTGACT
21	Cdc20	F	GTGCTCACTCTGTTCTC		62 SCC40A1	F	GTCTATGGACTGGTGGTGGC
		R	ATTGTAAGACCCAGGACCCG			R	GTGACGTCTGGCCACTTTA
22	Anxa3	F	CAGGTCAAGCAGGCAGATGA		63 Gfi1	F	TCCGAGTTGAGGACTTTTG
		R	CGAGAGCTTTCCGGAAGTCT			R	CATGCATAGGGCTTGAAGG
23	Ly6A (Sca1)	F	CCTTCTCTGAGATGGACACTT		64 Mpo	F	ACTGCAACAACAGACGAAGC
		R	AGTCCCTGAGCTTTCTGC			R	GAAGCCATTGCAGTTACTCC
24	Cd48	F	TGCCAAGTCAGCAATCCTGT		65 Myo1F	F	AACAATTCAGCGCTTTGG
		R	ACTAGCCAAGTTGCAGTCCA			R	GCATGACCCACGAGACTTTTC
25	Cd34	F	ACATCACCCACCGAGCCATA		66 Ctsg	F	GGGTTTCTGGTGCAGAGAAGA
		R	AAACTCCTCACAACTAGATGCTTCA			R	AGTTGTGGGTCCTTTCTCG
26	Cd74	F	CACCGGTGTCTGTCTG		67 Alox5ap	F	CGTTCTTTGCCCAAGAGTG
		R	GGAGGTGATGGTCACTTGT			R	GGGTACGCATCTACGCAGTT
27	Cd11a (ItgaL)	F	ACTTCCACTTCCCGATCTGC		68 Tlr2	F	GGGGCTTCACTTCTCTGCTT
		R	CTCAGATAGGCTGCATGGC			R	AGAGCCAAAGAGCTCGTAGC
28	Cxcr 4	F	ATGGAACCGATCAGTGTAGT		69 Crebbp	F	AAGATGCGCAACTGGAGAC
		R	TGAAGTAGATGGTGGGCAGG			R	GCTTGAACAAGTCCCGAGG
29	Dnmt3a	F	AAACTTCGGGGCTTCTCTG		70 Trim10	F	TGAAACCAGAAAGTGCCGGA
		R	GCTTCTCCACACACACCACT			R	GCTGGCTCATAGTCCCACTCA
30	Malat 1	F	CCTTTGCAGAGCCTCATTTT		71 Elane	F	CGTCTGTGAACGTATGCAC
		R	CCTCCACATCCAGGAAAAA			R	ATCCAGATCCACAGCCTCCT
31	Mpl	F	CTGTATGCCTACCGAGGAGAG		72 Fcgr3 (CD16/3	F	GCTGCTGTTGCTTTTGAG
		R	TCTGGTTGAGGGACACTTCT			R	TCCCTTCGCACATCAGTGT
32	Meis 1	F	TAGACCAGTCCAACCGAGCA		73 Mphosph (MP	F	AGACCAGATTGAGGAGCAGC
		R	CATTCCACTCATAGTCTCGGC			R	GAGCCAGTTACGGTGTGAT
33	Meg3	F	ACTTTCGCACATTAGCCC		74 Csf1r	F	CAGTACTTCAGGCGGCTCTC
		R	CACAGGAAGACGCGACAG			R	TCGGCTTCTGCTTGTACTGT
34	Gata3	F	CTTATCAAGCCAAAGCGAAG		75 Alas2	F	GCTGGAGGAACTCCAAGAC
		R	CCATTAGCGTTCTCTCCT			R	ATGAATGGCTCGGGTGGTT
35	Gata2	F	CCTGGTTCCAAAGCACAGT		76 Spi1 (PU.1)	F	CTTGATCCCAACCGAAGCAG
		R	CTGTGCTGGCTCATGTAGTT			R	CAAGTCACTCCGATGAGGGG
36	Tal1 (Scl)	F	GTTGTTTACCAGCAGGAC		77 Cma1	F	CCTGCAAACTTCAACCACTT
		R	AGAGACCTACTCGGCTGTT			R	GATGCAATGCCTTGGGTATC
37	HoxA9	F	CCGGACGGCAGTTGATAGAG				
		R	CCTTTCGAGTGGAGCGAG				
38	Flt3	F	AGCACTGCGCCAGTTTCA				
		R	GACCACAGATTCCCTCGGAC				
39	Casp3	F	TGGTGATGAAGGGGTCATTTATG				
		R	TTCGGCTTCCAGTCAGACTC				
40	Zfpn1	F	ACCAGAGCCTTATCCCTGA				
		R	AGGGCGTCATCCTTCTGTA				
41	Cd9	F	GGGATTGTTCTCGGTTCTCT				
		R	GGTTCATCCTGTCCGTA				