Number	Gene	Forward Primer Sequence	Reverse Primer Sequence
P1	*Cdkn2a (p16^{Ink4a})*	GAACTCTTTCGGTCGTACCC	AGTTCGAATCTGCACCGTAGT
P2	*Cdkn1a (p21)*	GAACATCTCAGGGCCGAAAA	TGCGCTTGGAGTGATAGAAATC
P3	*Cdkn1b (p27)*	TCCGCCTGCAGAAATCTCTT	CCAAGTCCCGGGTTAGTTCTT
P4	*Cdkn2b (p15)*	AGATCCCAACGCCCTGAAC	CCCATCATCATGACCTGGATT
P5	*Cdkn2d (p19)*	GCGATAAGAGAGGGCCATAGC	CCTGTGGTGGAGATCAGATTCA
P6	*Ccl1*	GACCAGGTTGGGCAAAGAGA	GGCATCCTGGACCCACTTCT
P7	*Ccl2*	GTCTGTGCTGACCCCAAGAAG	TGGTTCCGATCCAGGTTTTTA
P8	*Ccl3*	TCCCAGCCAGGTGTCATTTT	TTGGAGTCAGCGCAGATCTG
P9	*Ccl4*	AGGGTTCTCAGCACCAATGG	CCGGGAGGTGTAAGAGAAACAG
P10	*Ccl5*	GCCCACGTCAAGGAGTATTTCT	ACAAACACGACTGCAAGATTGG
P11	*Ccl6*	GATCGTCGCTATAACCCTCCAA	TGGGATCTGTGTGGCATAAGAG
P12	*Ccl7*	CCCTGGGAAGCTGTTATCTTCA	CTGATGGGCTTCAGCACAGA
P13	*Ccl8*	CCACACAGAAGTGGGTCAGTGA	TTCAAGGCTGCAGAATTTGAGA
P14	*Ccl9*	GGCCGGGCATCATCTTTATC	TGCATCTCTGAACTCTCCGATCA
P15	*Ccl11*	GACCAGGTTGGGCAAAGAGA	GGCATCCTGGACCCACTTCT
P16	*Csf1*	ATTGCCAAGGAGGTGTCAGAA	GGACCTTCAGGTGTCCATTCC
P17	*Cxcl1*	CCGAAGTCATAGCCACACTCAA	CAAGGGAGCTTCAGGGTCAAG
P18	*Cxcl2*	TCAAGGGCGGTCAAAAAGTT	CAGTTAGCCTTGCCTTTGTTCA
P19	*Cxcl5*	GAGCTGCGTTGTGTTTGCTTAA	CAGGGATCACCTCCAAATTAGC
P20	*Cxcl9*	AACTCAGCTCTGCCATGAAGT	ACACTGCTCCAGGAAGATGAT
P21	*Cxcl10*	TGAATCCGGAATCTAAGACCA	TTTTTGGCTAAACGCTTTCAT
P22	*Cxcl12*	GCCAACGTCAAGCATCTGAAA	CAGCCGTGCAACAATCTGAA
P23	*Cxcl13*	AGCACAGCAACGCTGCTTCT	AATACCGTGGCCTGGAGAGA
P24	*Cxcl16*	GCACCCCTGCACATAGTCAGA	AGGACAGTGCTCCTGATGGAA
P25	*Cxcr4*	CTGGACCGGTACCTCGCTATT	GCAGTTTCCTTGGCCTCTGA
P26	*Cxcr5*	CCTTCGCTGGCGTAAAGTTC	ACAGCCCAGCTTGGTCAGAA
P27	*Cxcr6*	TGGGAGTCTCAAGTCAATGGAA	TGTGGGAGGCAGAACAAGTCT
P28	*Cxcr7*	TGTGCTCTTTACAGCGTTGCAT	TTGACACAGCAGTGCACCAA
P29	*Icam1*	GTGGCGGGAAAGTTCCTGTT	GTCCAGCCGAGGACCATACA
P30	*Igfbp4*	GCAACTTCCACCCCAAACAGT	CCTGTCTTCCGATCCACACA
P31	*Il1a*	AAGAGACCATCCAACCCAGATC	CCTGACGAGCTTCATCAGTTTG
P32	*Il1b*	TCAGGCAGGCAGTATCACTCA	CACGGGAAAGACACAGGTAGCT
P33	Il-1rn	CCACCACCAGCTTTGAGTCA	ACGGTCAGCCTCTAGTGTTGTG
P34	*Il6*	ACCACGGCCTTCCCTACTTC	TTGGGAGTGGTATCCTCTGTGA
P35	*Il10*	TGGCTCAGCACTGCTATGCT	TGTACTGGCCCCTGCTGATC
P36	*Il11*	GCTCCCCTCGAGTCTCTTCAG	GGGATCGGGTTAGGAGAACAG
P37	*Il12a*	ATCCTGCTTCACGCCTTCAG	GATAGCCCATCACCCTGTTGA
P38	*Il12b*	GCCAGTACACCTGCCACAAAG	TGTGGAGCAGCAGATGTGAGT
P39	*Il15*	GGCATTCATGTCTTCATTTTGG	TCCAGTTGGCCTCTGTTTTAGG
P40	*Il18*	GGCTGTGACCCTCTCTGTGAA	AGGTGGATCCATTTCCTCAAAG

P41	*Lyn*	AGCACAAAGGTGGCTGTGAAG	GGAATGCCTGCACAGACATG
P42	*Mif*	GCCACCATGCCTATGTTCATC	GGGTGAGCTCCGACAGAAAC
P43	*Mmp2*	TGTGGGTGGAAATTCAGAAGGT	ACTTGTTGCCCAGGAAAGTGA
P44	*Mmp12*	GTGCCCGATGTACAGCATCTT	GGTACCGCTTCATCCATCTTG
P45	*Mmp13*	TGAGGAAGACCTTGTGTTTGCA	GCAAGAGTCGCAGGATGGTAGT
P46	*Nfkb1*	GGCTTTGCAAACCTGGGAAT	TCCGTGCTTCCAGTGTTTCA
P47	*Pf4*	TGGGATCCATCTTAAGCACATCA	CCATTCTTCAGGGTGGCTATGAG
P48	*Serpinb2*	TTCCGCATACTGGAAACATCAG	GGATGCGTCCTCAATCTCATC
P49	*Serpine1*	GGACACCCTCAGCATGTTCA	CGGAGAGGTGCACATCTTTCT
P50	*Tnfsf10*	CTCTCGGAAAGGGCATTCATT	TCGATGACCAGCTCTCCATTC
P51	*Vcam1*	GGCTCCAGACATTTACCCAGTT	CATGAGCTGGTCACCCTTGAA
P52	*Actb*	AATCGTGCGTGACATCAAAGAG	GCCATCTCCTGCTCGAAGTC
P53	*Tuba1*	GGTTCCCAAAGATGTCAATGCT	CAAACTGGATGGTACGCTTGGT