

Table S1. Oligonucleotides used in PCR.

Oligo name	Tm	Length	Sequence	PCR Product (bp)	Accession no.
M-GAPDH-3-F	79	22	GTGCAGTGCCAGCCTCGTCCCG	111	NM_008084.2
M-GAPDH-3-R	68.7	22	CTCCACTTTGCCACTGCAAATG		
hGAPDH-F	65.1	20	CTCCTGTTCGACAGTCAGCC	101bp	NM_002046.4
hGAPDH-R	67.2	21	ACCAAATCCGTTGACTCCGAC		
hmGAPDH-F	59.2	18	CCCAGAAGACTGTGGATG	76bp	NM_002046.4
hmGAPDH-R	61.8	18	GCAGGGATGATGTTCTGG		
mTGFB2 F1	66.5	20	AATGGCTCTCCTTCGACGTG	180	NM_009367.3
mTGFB2 R1	66.4	21	AGGTGCCATCAATACCTGCAA		
hTGFB F	69.3	20	TCTTCCCCTCCGAAAATGCC		NM_001135599
hTGFB R	66.9	20	GCTCAATCCGTTGTTCAAGC		
mGDF15 F1	65.9	20	TCCAGAGGTGAGATTGGGGT	155	NM_011819.2
mGDF15 R1	65.9	20	GCTTCAGGGCCTAGTGATG		
hGDF15 F	65.7	19	GTCCGGAT ACTCACGCCAG		NM_004864
hGDF15 R	68.7	20	TCACGTCCCACGACCTTGAC		
hTGFB1 F1	65.7	20	TGGGCTCTGCTTTGTCTCTG	199	NM_001130916.1
hTGFB1 R1	60.4	23	TGGTAAACCAGTAGTTGGAAGTT		
hTGFB2 F1	67.4	21	CGTTCAGAAGTCGGATGTGGA	87	NM_001024847.2
hTGFB2 R1	65.9	20	TGTCTCAGTGGATGGGCAGT		
hTGFB3 F1	67.4	20	CTGGTGAAGTGGGCTTTGGA	116	NM_001195683.1
hTGFB3 R1	66.8	24	TCCTCATCTCCCATCTCCTCATTA		
hFGF2-F	66.5	20	GCTGTACTGCAAAAACGGGG	94	NM_002006.4
hFGF2-R	65.3	20	TAGCTTGATGTGAGGGTCGC		
hFGF3-F	64.4	20	ATGCTTCGGAGCACTACAGC	160	NM_005247.2
hFGF3-R	64.8	20	CCGTTACAGACACGTACCA		
hFGF4-F	66.4	20	CTATGGCTCGCCCTTCTTCA	120	NM_002007.2
hFGF4-R	68.5	20	CCATTCTTGCTCAGGGCGAT		
hFGF6-F	66.2	20	GCTAGCTGGAGAGATTGCCG	169	NM_020996.1
hFGF6-R	66.2	21	AATTTCCAGCAGGCTGTAGGG		
hFGF7-F	66.5	24	TGCAAAGAAAGAATGCAATGAAGA	104	NM_002009.3
hFGF7-R	67.2	20	TTTCCCCTCCGTTGTGTGTC		
hFGF9-F	68.8	20	AGCCGATTTGGCATTCTGGA	85	NM_002010.2
hFGF9-R	64.3	20	TCCCGAGGTAGAGTCCACTG		
hFGF19-F	64.7	20	AGATCAAGGCAGTCGCTCTG	106	NM_005117.2
hFGF19-R	63.9	20	GAGTACTGAAGCAGCCCCTG		
hFGFR1-F	61.7	20	GGCAGTGACACCACCTACTT	144	NM_023110.2
hFGFR1-R	66.8	21	CCAATATGGAGCTACGGGCAT		
hFGFR2-F	68.1	20	CCTCAAGTTTCTCAAGGCCG	104	NM_023029.2
hFGFR2-R	65.9	20	AGCACGTATATTCCCCAGCG		
hFGFR3-F	69.7	19	TACCGTGCTCAAGACGGCG	75	NM_000142.4
hFGFR3-R	65.7	20	AAGGTGACGTTGTGCAAGGA		
hFGFR4-F	66.9	19	TGAGGAGGAGCCAGGAAGG	161	NM_002011.3
hFGFR4-R	64.7	20	GCTCAAGCTCCACTTCTCTCA		
mFGFR1-F	63	20	GGCAGCGATACCACCTACTT	137	NM_010206.2
mFGFR1-R	64.3	20	GGAGCTACAGGCCTACGGTT		

mFGFR2-F	67.4	20	CCAGAAGAGCCACCAACCAA	118	NM_000141.4
mFGFR2-R	66.2	20	TAGTCCAAGTATGATCACGGCG		
mFGFR3-F	66.3	20	GCGACAGGTGTCCTTGGAAT	96	NM_008010.5
mFGFR3-R	65.1	20	GCCAGAACAGGACCTTCTCC		
mFGFR4-F	66.5	20	GCCCGACAGTTCTCTTTGGA	141	NM_008011.2
mFGFR4-R	68.6	20	TTCCCAAAGCGGATCGAGAG		
H-KL F1	69.4	20	ATGGAATCGATGACGGGCTG	101	NM_004795.3
H-KL R1	64.3	21	ACCATCCAGTATGTGGGCTTT		
M-KL F1	69.1	20	AATCGATGATGACCCCCACG	96	NM_013823.2
M-KL R1	64.5	20	TCGTCCAACACGTAGGCTTT		
H-KLB F1	62.7	21	TGTTCTTAAGCCCGAGTCTGT	140	NM_175737.3
H-KLB R1	66.8	20	AATCTGTGCATTGAGCGGGT		
M-KLB F1	64.6	21	GACACAACCTGATCAAGGCAC	72	NM_031180.2
M-KLB R1	68.2	20	CCTTCTGATGAGGGCGGAAG		
M-CXCL1-2-F	64.3	20	GCCTCTAACCAGTTCCAGCA	144	NM_008176.3
M-CXCL1-2-R	72.9	20	GCTCATTGGCGATAGGCGCC		
H-CXCL1-2-F	59.6	21	CTCTACCTGCACACTGTCCTA	151	NM_001511.3
H-CXCL1-2-R	68.4	20	GATGTTGCAGGCTCCTCAG		
H-CXCR2-2-F	67.9	21	CTAAGTGGCACCTGTCCTGGG	238	NM_001557.3
H-CXCR2-2-R	65.3	21	CAGAAGAGCAGCTGTGACCTG		
M-CXCR2-1-F	57.4	19	GCTGATACTCACAACAGCA	112	NM_009909.3
M-CXCR2-1-R	60.8	19	GCACTTCTCATTGTGGAC		