

Table S2. Bergamaschi et al

Name	Sequence	Start bp	Stop bp	Tm
Fw_COL4A1_Hs	CGTAACTAACACACCCTGCTTCAT	5230	5253	58
Rev_COL4A1_Hs	CACTATTGAAAGCTTATCGCTGTCTT	5304	5279	58
Probe_COL4A1_Hs	ACCTCTACTTGCTGAAGGA	5256	5274	69
Fw_COL4A1_Mm	GGCGGTACACAGTCAGACCATT	4805	4826	60
Rev_COL4A1_Mm	GGAATAGCCGATCCACAGTGA	4877	4857	59
Probe_COL4A1_Mm	AGTGCCCTAACGGTTGG	4837	4853	70
Fw_CTSB_Hs	AGACTTGTAGCTGCTGTCTCTCTTTG	2193	2218	59
Rev_CTSB_Hs	AAGAGTCGCAAGAACATGCAGTT	2265	2243	59
Probe_CTSB_Hs	CCTGCAGAGAATCACGT	2221	2237	70
Fw_CTSB_Mm	GTCAAAAGCAGGCTAGGGATGT	3344	3365	58
Rev_CTSB_Mm	AAGGGTCTTGCTTGGACAATATG	3412	3390	58
Probe_CTSB_Mm	ATGGTAGAGTGTTTGTCC	3370	3387	68
Fw_MMP11_Hs	GATGGCAGAGGCCCTAAAGG	409	428	60
Rev_MMP11_Hs	GCCCTCGTGACCTCAGTA	475	457	58
Probe_MMP11_Hs	TGGAGCGATGTGACGC	431	446	68
Fw_MMP11_Mm	ACCGGAGAGTCACCGTCATC	127	146	59
Rev_MMP11_Mm	ATTAGGCAGAGCTGCATGCA	193	174	58
Probe_MMP11_Mm	AAGAAAGGGCCTCGGT	155	171	70
Fw_PECAM1_Hs	GCAGCATCGTGGTCAACATAAC	1157	1178	59
Rev_PECAM1_Hs	CCAGATGTGTGAAGGAAGATTCC	1225	1203	59
Probe_PECAM1_Hs	AACTATTTTCCAAGCCCG	1181	1198	69
Fw_PECAM1_Mm	CCCCCAGAACATGGATGTAGAAT	2242	2264	60
Rev_PECAM1_Mm	CAGAGCTTGGTGAGGCTCAAG	2308	2288	59
Probe_PECAM1_Mm	AGAAGTGGAAAGTGTCCCTC	2269	2286	69
Fw_SPARC_Hs	TTTCACATTAGGCTGTTGGTTCA	1674	1696	58
Rev_SPARC_Hs	GCTGACCACTTCCCAGAGAACT	1741	1720	58
Probe_SPARC_Hs	TTGGGAGCACGGACTG	1702	1717	70
Fw_SPARC_Mm	TGGCTGTTGGTTTTAGTTTTGGT	1639	1661	59
Rev_SPARC_Mm	CCCGGCAAGAACCTGAAA	1721	1704	58
Probe_SPARC_Mm	ACATCACTCAACTGCAATT	1681	1699	70
Fw_GAPDH_Hs	GACCTGACCTGCCGTCTAGAAA	832	853	59
Rev_GAPDH_Hs	CCTGCTTCAACCACCTTCTTGA	895	875	59
Probe_GAPDH_Hs	ACCTGCCAAATATGATGAC	855	873	69
Fw_GAPDH_Mm	CATGGCCTTCCGTGTTCCCTA	734	753	60
Rev_GAPDH_Mm	GCGGCACGTCAGATCCA	788	772	59
Probe_GAPDH_Mm	CCCCAATGTGTCCGTC	755	770	69