

Primer (mouse)

Protein	Sequenz	NT	Accesssion Nr.	Anneal
Col1a1	TGACTGGAAGAGCGGAGAGT GTTTCGGGCTGATGTACCAGT	151	NM_007742	58
Col4a3	ACGGTGTGTTCTTGTCTCC CTTTGGTCCTAGCCTTGTGC	232	NM_007734	60
Itga5	GCGACTGGAATCCTCAAGA GCTGAGTCCTGTCACCTTG	184	NM_010577	62
Itgb2	GTGGTGCAGCTCATCAAGAA GCCATGACCTTTACCTGGAA	197	NM_008404	63
Il1r1	TTTGCCACGAGCAGGAGA GGTACCTCACATGCTTCCA	231	NM_001025602	
Sox9	TCAGATGCAGTGAGGAGCAC CCAGCCACAGCAGTGAGTAA	208	NM_011448	60
Col2a1	AGGTGCTAATGGCAATCC GAGGACCATCAAGACCAG	190	NM_001113515	62,5
Col9a1	GGGAGACAGAGGCATTCAAG TTCCAGGACACCTGGTAAG	185	NM_007740	58
Col13a1	GCCTGGTTTACATGGACCAC TCCCAGGTTAGAGGCATCAC	185	NM_007731	61,5
Acan	AGGACTGAAATCAGCGGAGA AGGGACATGGTTGTTTCTGC	188	NM_007424	60
Adamts5	AGAGGGCCATATACCGTTCC GGCAGGACACCTGCATATTT	157	NM_011782	61,5
Runx2	CAGACCAGCAGCACTCCATA CAGCGTCAACACCATCATTC	178	NM_009820	63
Il17ra	GTGTAGTGCCCGCTTAGAA TGTCGGGTTATCAGGGAAAG	207	NM_008359	
Itga3	TGCCGTTCTAAATCCTCCAC CACCGGTAGTCAGGCAATTT	226	NM_013565	
Itgb3	ATATCCTGGTGGTCCTGCTG CCCGGTAGGTGATATTGGTG	207	NM_016780	63
Itga4	GTTGGGAGCATGAAGACCAT TGCAGGCAAGCTTCACTATG	184	NM_010576	61,5
Mmp13	ATCCTGGCCACCTTCTTCTT TTTCTCGGAGCCTGTCAACT	202	NM_008607	61,5
Mmp9	ATTGCATGGCCTCTGTAACC CCCAGCAAAGTCTGATGTGA	203	NM_012599	63
Mmp8	TCCTACCCAACGGTCTTCAG AATGGCTTGGACACTCCTTG	161	NM_008611	55,5
Mmp2	CTTCGCTCGTTTCTTCAAC CTTGTTCTCCTCCATCCAG	178	NM_008610	58
Mmp1a	ATTCATGCCAGAACCTGAGC CTTCCTCACAAACAGCAGCA	233	NM_032006	
Itgav	GGGTGATCATCTTGGCAG GAACTTGGAGCGGACAGA	205	NM_008402	58

Protein	Sequenz	NT	Accesssion Nr.	Anneal
Itga2	CATGTGAGGTTGGTTCATCG TCGTGAGACTGACCGAATTG	198	NM_008396	61,5
Itgb5	CCTGGAACCACTGTGGAGAT ACGGACACTTCAAAGGATGC	227	NM_010580	53
Itga6	CCCAGGGACTTACAACCTGGA CTTGGAGCACCAGACACAAA	225	NM_008297	58
Itgb4	CTTCGAGCAGCCTGAATAC CTTCACCTGCAACTCCTTC	204	NM_001005608	58
Timp1	GGTGGGTGGATGAGTAATGC AACAAGAGGATGCCAGATGC	197	NM_001044384.1	61,5
Timp2	TCCTTGCTACAGGCAGGAGT AGGCCGGCTACACAGTCTTA			62,5
Ibsp	AGAAGGCTGGAGATGCAGA CTGCACCTGCTTCAGTGAC	163	NM_008318.2	61,5
Thbs1	GCGATGATGACGATGACAA TCTGTGTCTGCTTGGTCAG	154	NM_011580.3	63
Bglap1	AAGCAGGAGGGCAATAAGG GCTGCCAGAGTTTGGCTTT	212	NM_007541.2	62,5
OPN	TGCACCCAGATCCTATAGC CTCCATCGTCATCATCATC	186	AF515708.1	61,5
Sparc	AATTTGAGGACGGTGCAGAG AAGTGGCAGGAAGAGTCGAA	167	NM_009242.4	62,5