

Table S2

Gene	Orientation	Sequence (5'→3')
ACSL3	Forward	GCGTAGCGGTTTTGACAC
	Reverse	CCAGTCCTTCCAACAACGA
E-CAD (CDH1)	Forward	GAACGCATTGCCACATACAC
	Reverse	GAATTCGGGCTTGTGTCAT
K18 (KRT18)	Forward	GAGTATGAGGCCCTGCTGAA
	Reverse	CAGACACCACTTTGCCATCC
Na/K ATPase (ATP1B1)	Forward	GCCTCCAAGAATGAGTCCT
	Reverse	ATTTGGGCTGCAGGAGTTTG
N-CAD (CDH2)	Forward	CGTTTTCATTTGAGGGCACA
	Reverse	TTGGAGCCTGAGACACGATT
SLUG (SNAI2)	Forward	CGTTTTCCAGACCCTGGTT
	Reverse	CTGCAGATGAGCCCTCAGA
VIM	Forward	GAGTCCACTGAGTACCGGAG
	Reverse	ACGAGCCATTTCTCCTTCA
HMGCR	Forward	TGATTGACCTTTCCAGAGCAAG
	Reverse	CTAAAATTGCCATTCCACGAGC
ABCA1	Forward	ACCCACCCTATGAACAACATGA
	Reverse	GAGTCGGGTAACGGAAACAGG
ABCG8	Forward	AGCCTCCTTGCTAGATGTGAT
	Reverse	GTCTCTCGCACAGTCAAGTTG
APOA1	Forward	CCCTGGGATCGAGTGAAGGA
	Reverse	CTGGGACACATAGTCTCTGCC
ACSL1	Forward	ACATTATGTTCTGGGCCCA
	Reverse	AGTCAGAAGGCCATTGTGCA
COX2	Forward	ATCACAGGCTTCCATTGACC
	Reverse	CAGGATACAGCTCCACAGCA
CASP9	Forward	CAGGCCCATATGATCGAGG
	Reverse	TCGACAACCTTTGCTGCTTGC
GAPDH	Forward	TGGTATCGTGGGAAGGACTCATGAC
	Reverse	ATGCCAGTGAGCTTCCCGTTCAGC
B2M	Forward	GATGAGTATGCCTGCCGTGT
	Reverse	TGCGGCATCTTCAAACCTCC