

Table S1. The list of primers used in this study.

Primer	Gene Accession Number	Sequence		Amplicon Size (bp)
		Forward (5→3)	Reverse (5→3)	
<i>Gapdh</i>	NM_008084.2	atcaagaaggtggtgaagca	gacaacctggtcctcagtgt	75
<i>Il1b</i>	NM_008361.3	agttgacggaccccaaaag	agctggatgctctcatcagg	75
<i>Il6</i>	NM_031168.1	aacgatgatgcacttgacaga	ccagaggaaatttcaataggc	113
<i>Il17a</i>	NM_010552.3	cagggagagcttcatctgtgt	gctgagctttgagggatgat	94
<i>Il17c</i>	AF458061.1	cctctagctggaacacagtgc	gcggttctcatctgtgtcg	114
<i>Tnf</i>	NM_013693.2	catcttctcaaaattcgagtgaca	tgggagtagacaaggtacaacc	175
<i>Il23a</i>	NM_031252.2	caccagcgggacatatgaa	ccttggtgggtcacaccat	100
<i>Il22</i>	NM_016971.2	tgacgaccagaacatccaga	aatcgcttgatctctccac	85
<i>Il19</i>	NM_001009940.1	tggagaacctcaggagcatt	gaatgtcagcaggttgttg	65
<i>Cxcl1</i>	NM_008176.3	cttgacctgaagctccctt	gttgtcagaagccagcgttc	127
<i>Cxcl3</i>	NM_203320.3	ccccaggctcagataatca	tctgattagaatgcaggctcctt	110
<i>Cxcl5</i>	NM_009141.3	tagagcccaatctccacac	ggagctggaggctcattgt	88
<i>Ccl20</i>	NM_016960.2	aactgggtgaaaagggtgt	gtccaattccatccccaaaa	86
<i>Defb4</i>	NM_019728.4	cagtcatgaggatccattacctt	aatttgggtaagggtgcaa	77
<i>S100a8</i>	NM_013650.2	gtgtcctcagttgtgcagaat	tgatgaccacaccacttt	126
<i>S100a9</i>	NM_001281852.1	gacacctgacacctgag	tgaggcttcatttctctctc	96
<i>Krt16</i>	NM_001313958.1	atgagctgacctgtccaga	ctcaaggcaagcatctctc	106
<i>GAPDH</i>	NM_002046.5	agccacatcgtcagacac	aatacgaccaatccgtgact	62
<i>CCL20</i>	NM_004591.2	gtgctgctactccacctctg	gcattgatgtcacagccttca	143
<i>CXCL1</i>	NM_001511.3	catcgaagatgctgaacagt	ataaggcagggtcctcct	94
<i>CXCL8</i>	NM_000584.3	cagagacagcagagcacaca	gcactccttgcaaaaactgc	167
<i>IL1B</i>	NM_000576.2	aaagcttggatgtctggtc	ggacatggagaacaccacttg	89
<i>TNF</i>	NM_000594.3	agccatgttagcaaac	tctcagctccagccatt	94
<i>IL19</i>	NM_153758.2	cagagtcacatgacaactatgat	cgagctctcccagggtatt	71
<i>IL17RA</i>	NM_014339.6	catcctgctcatctgtgc	gccatcgggtgatttgggtgt	85
<i>IL17RC</i>	NM_153460.3	ttgaaccacacagacctggtt	tctaacggagtcagggtcc	70
<i>TNFRSF1A</i>	NM_001065.3	cacaagccacagagcctaga	gaattcctccagcgaac	87
<i>LEPR</i>	NM_002303.5	aaaaatgtcacttacttgggaagc	ttccacatctctgacca	119