

Gene	Function	References	ENSEMBL ID	Forward primer	Reverse primer
Reference genes					
<i>b2m</i>	Beta-2-microglobulin	(Hibbeler et al., 2008)	ENSGACT000000025544	GAAGATGTGTTGAATAGAAGCTGG	GAAGATGTGTTGAATAGAAGCTGG
<i>ef1a</i>	Elongation factor 1 α	(Hibbeler et al., 2008)	ENSGACT000000002893	CCACCGTTGCCTTTGTCC	TGGGACTGTTCCAATACCTCC
<i>rp13a</i>	L13A ribosomal binding protein	(Hibbeler et al., 2008)	ENSGACT000000012319	CACCTTGGTCAACTTGAACAGTG	TCCCTCCGCCCTTACGAC
<i>ubc</i>	Ubiquitin	(Hibbeler et al., 2008)	ENSGACT000000010662	AGACGGGCATAGCACTTGC	CAGGACAAGGAAGGCATCC
Innate					
<i>cd97</i>	Promotor of granulocyte and neutrophil migration, required for activation of the innate immune response	(Leemans et al., 2004; Rhodes et al., 2009)	ENSGACT000000024871	CTCGTGGCACTTACGACATGAAG	CAGCCCTATCTTGGTGACCAGTTG
<i>csf3r</i>	Granulocyte colony-stimulating factor 3 receptor; role in differentiation and proliferation of granulocytes	(Birrer et al., 2012; Brunner, 2016; Maxson et al., 2013; Tabbara, 1993)	ENSGACT000000018254	TGGGATTCGTCTCTTCTCAG	TGGGTCAAACCTTGGCTGCAC
<i>il-1β</i>	Interleukin 1 β ; cytokine with function in early response proinflammatory signaling	(Brunner et al., 2017; Zhu et al., 2013)	ENSGACT0000000019325	TGACGATGAAGCAGGTGGTCAAC	ACAGCGTCACGATCTCCTCTTC
<i>marco, RON</i>	Macrophage receptor with collagenous structure; mediates macrophage recognition and clearance of pathogens	(Kissick et al., 2014; Kraal et al., 2000)	ENSGACT0000000001965	CCCTTTCGACCTTCACTGCC	TGTTTACCCCAACCCCTCCA

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Innate (continued)					
<i>mif1</i>	Macrophage migration inhibitory factor; stops random macrophage migration through tissue, proinflammatory mediator of the innate immune system	(Brunner et al., 2017; Calandra and Roger, 2003)	ENSGACT000000023656	ATCAGCGGAGCTCACAACAAGC	TCAGGAGAGATGCTCAGGTGTTTG
<i>mst1ra</i>	Macrophage stimulating 1 receptor α ; plays an important role in macrophage regulation	(Huang et al., 2016; Wang et al., 2002)	ENSGACT00000013997	ATGGCCATCGAAAGCTTGCA	TGATGTCGTACGGGTCCACA
<i>nkef-β, peroxiredoxin 1</i>	Natural killer cell enhancing factor; enhances cytotoxicity of NK cells, also protects against oxidative damage	(Shau et al., 1993; Stutz et al., 2015)	ENSGACT000000021380	ACTTCTCCCACTTTGCATGG	CAATGCCTTCATCCTCCTTC
<i>p22^{phox}</i>	NADPH oxidase component p22phox; part of the reactive oxygen species production machinery	((Bedard and Krause, 2007; Mayumi et al., 2008)	ENSGACT000000021084	GCCTCGGGACTCATTCTCT	TGGCCCTCTTGCTTCTTGGA
<i>saal1</i>	Serum amyloid A; acute phase protein during inflammation response, mediates release of TNF- α and IL-1 β	(Brunner et al., 2017; Haarder et al., 2013; Kovacevic et al., 2015)	ENSGACT000000007599	TCGCAGTGAGGGCAAGATGAG	AAATCTGCCACCCTGTCCTTGG
<i>slal1</i>	Src-like-adaptor; necessary for maturation and activation of monocyte and dendritic cells, functions in T-cell signaling and B-cell development and function	(Brunner et al., 2017; Marton et al., 2015)	ENSGACT000000007895	ACAGAGTCGGCTCCTTCATGATAC	TCACAGAGAGCGGAATACAGACCTC

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<i>tnfr1</i>	Tumor necrosis factor receptor 1; functions in regulation of inflammation, mediates cellular apoptosis and differentiation	(Brunner et al., 2017; Zhu et al., 2013)	ENSGACT000000013502	AACTACTACAGAGCCCAAGGGCAAG	ACGGCACTCAGCGGTACAATTC
Adaptive					
<i>cd83</i>	Marker for mature dendritic cells, expressed on activated B and T cells, costimulatory to activate naive and memory T-cells	(Aerts-Toesgaert et al., 2007; Stutz et al., 2015)	ENSGACT000000000428	AGGACCCAGCGTATAAATGG	CCCTGGTGATTTTCCTCATC
<i>foxp3;</i> <i>forkhead box N2b</i>	Transcription factor; regulates functions important for the establishment of the T-reg lineage, key mediator of T-cell activation	(Kasheta et al., 2017; Rao and Naqvi, 2011; Robertson et al., 2015)	ENSGACT000000007261	GTTGACCCATGCAATTC CGA	CTGCTGTAGTTGTGGTCCTG
<i>igm</i>	Immunoglobulin heavy constant mu (IgM); antibody molecule, part of the humoral immune response	(S. Hibbeler, unpublished; Rønneseth et al., 2015; Zhu et al., 2013)	ENSGACT00000016907	AAGGCAGGAGAAATGAAACCTTGG	CCGAGTGAGCAGACAGGGACTGG
<i>il-16</i>	Interleukin 16; cytokine with function in T-cell migration and expansion, chemoattractant for monocytes and eosinophils	(Brunner et al., 2017; Wen et al., 2006; Zhu et al., 2013)	ENSGACT000000016499	CTGGTCTGGGCTTCAGTATTGC	CTGGGAAACACTCTGTGGACTG
<i>mhcII</i>	Major histocompatibility complex class IIb exon 2; pathogen recognizing protein of the adaptive immune response	(Lenz et al., 2009)	ENSGACT000000000425	GTCTTTAACTCCACGGAGCTGAAGG	ACTCACCGGACTTAGTCAG

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Adaptive (continued)					
<i>stat4</i>	Signal transducer and activator of transcription 4; required for TH1-cell differentiation, opposes TH2 and TH17 like responses	(Kaplan, 2005; Premachandra et al., 2013; Wang and Secombes, 2013)	ENSGACT00000003538	CTCTCAGTTTCGAGGCTTGCTT	GGCAGTTGGCTCACATTGG
<i>stat6</i>	Signal transducer and activator of transcription 6; required for TH2-cell differentiation, regulates expression of TH2 relevant cytokine IL-4	(Robertson et al., 2015; Wang and Secombes, 2013)	ENSGACT00000011232	CTCAGCCACAGTTCACACCGTTC	GTCGGATGTTCTGGACCTCGAGT
<i>tcr-β</i>	T-cell receptor β-chain; function in binding of MHC-peptide ligands to initiate adaptive immune response	(Smith-Garvin et al., 2009; Stutz et al., 2015; Yanagi et al., 1984)	ENSGACT00000016457	GAGGGCAAAAATTTCACCTG	TAGGAGAATCTGGCCGTTTG
<i>tgf-β</i>	Transforming growth factor β; cytokine with functions in cell growth, migration, differentiation and proliferation of T and B-cells	(Robertson et al., 2015; Zhu et al., 2013)	ENSGACT00000016968	TCCCGCTTCGTACCAACCA	ACGTCTGTCTGGCCACATTAC
Complement					
<i>c7</i>	Complement component 7; initializing function in the membrane attack complex of the complement system	(Brunner et al., 2017; Haase et al., 2014; Zhu et al., 2013)	ENSGACT00000009181	TGGCTCAAGCTCAGCACAACAG	AGCGACACGTGTTTGTTTGATCG
<i>c9</i>	Complement component 9; structural part of the membrane attack complex of the complement system	(Brunner et al., 2017; Haase et al., 2014; Zhu et al., 2013)	ENSGACT00000020968	CCGTGACGAACAAGACTCAGTTG	TCTGACCGATGTCAGCACCTTG

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Complement (continued)					
<i>cfb</i>	Complement factor B; activating complement component of the alternative pathway	(Brunner, 2016; Haase et al., 2014; Zhu et al., 2013)	ENSGACT00000027346	GAGCGTCGCACAAATACAGGTTG	TACCACCGGAAGCGCACAAATC

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