

Table S1. Primers used for qPCR

Primers without references were designed using Primer Express® software v2.0 (Applied Biosystems)

Gene	Forward primer	Reverse primer	Reference
Gapdh	AGCCAAACGGGTCATCATCTC	GGGGGCTAAGCAGTTGGTG	(1)
Cyp1a2	TGGAGCTGGCTTTGACACAG	CGTTAGGCCATGTCACAAGTAGC	(2)
Cyp2c29	AAGGTTTTGGCGTTGTTTTAGC	AGCCTTTGGTTTTCTCAGTTCTT	(3)
Cyp2d9	CCTCCAAGGTCAGAAGTCCTTCA	CGATTTCTTGTGGACTCTGCG	(4)
Cyp2e1	AAGCGCTTCGGGCCAG	TAGCCATGCAGGACCACGA	(2)
Cyp3a11	GGATGAGATCGATGAGGCTCTG	CAGGTATTCCATCTCCATCACAGT	(2)
Cyp3a13	CCTCTGCCTTTCTTGGGGACGAT	CCGCCGTTTGTGAAGGTAGAGT	(3)
Cyp3a25	TTGGAGGCCTGAAGTCTAAAG	CTGTGGGCCCTCATAAAACC	(1)
Cyp4a10	TGTCCCAGGCATTGTCAGAGA	CCTTCGGGTTGTGGTGGAGA	(3)
Cyp4a14	CCCAAAGGTATCACAGCCACAA	CAGCAATTCAAAGCGGAGCAG	(3)
Cyp4f18	AGAGCCTGGTGCGAACCTT	TGGAATATGCGGATGACTGG	(5)
Cyp7a1	TGTCTGCGAGGGGCTGGAGCA	CCAGCCTGGGATGCTATGGGC	(6)
Fmo3	CAGCATTTACCAATCGGTCTTC	TGACTTCCCATTGCGCAGTAG	(7)
Sult1b1	TTGAAGAGTTCCAGAGCA	GAACCTTTGGAGGTGATAACA	(8)
Sult1d1	GAGGGAGTTAGTGGATGTT	TCACTGACCCAAGTTGTTC	(8)
Sult2b1	CAGTCTGGCTGAGAACACTA	TTCAGGATTAAGCAGACGAT	(8)
Sult3a1	CATCGGAACAGAAGTGAACA	TATGGAATGTGGGAAGTGAAGA	(9)
Sult3a2	GATATGGATGCTGTTGTGAGACAA	TCCAGTCTCCAACGATACCTTTG	(9)
Ugt1a1	CCAGCAGAAGGGGCACGAAGTTG	TGACCACGCGCAGCAGAAAAGAAT	(10)
Ugt1a2	TGCCCTCCGAGGAATCTCAG	CTCCCGCACAAACATCCCTCATG	(10)
Ugt1a6	CTGGCTGATGGTGGCTGACTG	ACTGAGGCCCAAAGCACTAGGAA	(10)
Ugt1a9	CAGAGGGCATGAGGTTGTGGA	TCCTGCAGTGTGAAAATGTTAGTT	(10)
Ugt2b5	ACCCAGCAACTTTAGGACACAAT	TGATAGATCGCCTCGTAGACACCA	(10)
Car	CAGACCCGGAGTTACCCAAAGAGA	GAGCAGCGGCGTCATAGCAGA	(1)
Fxr	ACGGGGGCAACTGCGTGATG	CGCCCTTCGCTGTCGTCCTC	(1)
Lxra	CTCAATGCCTGATGTTTCTCCT	TCCAACCCTATCCCTAAAGCAA	(11)
Ppara	GGGGGTGGGAGCATTTC	GGAGGGGCTAGAGGAGCATTAC	(1)
Pxr	CGCGGAGAAGACGGCAGCATC	CCCAGGTTCCCGTTTCCGTGTC	(1)
Rxra	GGCGCCATCTTTGACAGGGTGCTA	CCGGCTGCTCAGGGTACTTGTGTT	(1)
Nos2	CTTCACGGGTGAGAGCCACAGTCC	GAGCCAAAGCCAAACACAGCATA	(1)
Hmox1	AGGAGCTGCACCGAAGGGCT	GTGGCTGGCGTGCAAGGGAT	
Abca1	GGTTTGGAGATGGTTATACAATAGT TGT	TTCCCGAAACGCAAGTC	
Srbp1c	GATGTGCGAACTGGACACAG	CATAGGGGGCGTCAAACAG	(12)
Abcg1	AGTCTGACGATGACCAGTGTGAG	CTTCTAGTTAGAAAAGGAAACGAA CAGT	
Apoe	GCCGTGCTGTTGGTCACA	TGCCACTCGAGCTGATCTGT	
IL1b	CAACCAACAAGTGATATTCTCCATG	GATCCACACTCTCCAGCTGCA	(13)
IL6	CAACCACGGCCTTCCCTACT	TCATTTCCACGATTTCCAGAG	(3)
Ifng	AACGCTACACACTGCATCTTGG	GCCGTGGCAGTAACAGCC	(14)
Tnfa	GACAAGGCTGCCCGACTACG	CTTGGGGCAGGGGCTCTTGAC	(3)
Agp	CTTCGGGAGTCTCAAACAATAGGT	GACAGTCCCCGTTTCTTCTCATCC	(3)
Fga	AGCAGCCCAGCCACAAGAAAAGAG	CGTCGAGACTGCCGCTAAAAGTGT	(3)
Hapt	TGAACACAGTCGCTGGAGAG	GCTGCCTTTGGCATCCATAG	(15)
Saa1/2	GTAATTGGGGTCTTTGCC	TTCTGCTCCCTGCTCCTG	(16)
Sap	AGGGCAGAGACAATGAGCTACT	GGACTCCCAAGTGGTACATAGG	(17)
PccAS MSP1	ACAGTAACACAAGAAGGAAC	GATACTTGTGTTGATGCTGG	(18)

## References

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