

Table S1 Primers used in RT-qPCR

Gene	GenBank accession	Primer sequence (5'→3')	Fragment length (bp)
<i>β-actin</i>	KJ018754.1	F- GGGAGTCATGGTTGGTATGG R- TCCATATCGTCCCAGTTGGT	126
<i>CYP380C12</i>	MT268652.1	F- GATGATTGGTGGTAGTGAAA R- AATAATGGTGCTGTAGGGTA	216
<i>CYP6DAI</i>	KX945359.1	F- CGTAGAAGTCATCCATTTCGCTG R- GCTTTCCACTGATCTCCTCG	225
<i>CYP6DA2</i>	MF471386.1	F- AAACACAGAATACCAACGAG R- ACCTAAATAAGTCCACAAGC	113
<i>CYP4G15</i>	MF471388.1	F- CACCTATTTAAATGTGGATCGGGCC R- CGATCAATTTTCTGTGGGCTCTC	189
<i>CYP4CK1</i>	MF471405.1	F- CTATTTTGCCGGGGACATCG R- TCGGTCCTGCACTAAATGGT	175
<i>CYP6CY7</i>	MF471381.1	F- CTGATGTTACGCGATCCAGAGCT R- GATCTGTTTCATGCGTGTCTT	207
<i>CYP6CY18</i>	MF471377.1	F- CAGGTGAAGGAGTGCAGCGA R- GAGTTTGGACCTCAGCGACGG	215
<i>CYP6CY19</i>	KX945360.1	F- GATTCTTCTGTTTGTGCGCCGGTG R- GGGTAAGTCTTAGTTGCTTCTC	197
<i>CYP4CJ5</i>	MT268653.1	F- CTCATCAACGCATCAGACAATGG R- CTCTGCCAATATTACTAAGGTTG	157
<i>CYP6CY22</i>	KX950717.1	F- GATGTACTACTTATATCCGAACGCC R- GTGTCCTTTAGCTTTCCAGATGTG	255
<i>CYP380C6</i>	MF471383.1	F- CAAGGCCCTGGAGAAAGACC R- GGTGTCCGCTATATAGTCCCAAAG	211
<i>CYP6CY14</i>	MF687493.1	F- GATGGTTGGCTTAGAACGGC R- TCGTCAATGTCTATACCGTGG	186
<i>CYP6A14</i>	KR028424.1	F- CGGTGCTCAATTCTTCTCCG R- ATAAAGTCATTCCGGCGCAC	181
<i>CYP6CY13</i>	MF471380.1	F- CCGGAACCTCATAAACAACG R- ATCGTGACGTCGAACTGATC	244
<i>CYP305E1</i>	MF471390.1	F- GACTCAGGACAATGGGTAAACG R- CTTAGTGCCCGCTGTCAGTTCC	251
<i>CYP3323AI</i>	KY827388.1	F- CCCGAGCCTGAAGAGTTCCA R- TGACCACCTTCAAGCGTACT	238
<i>CYP18AI</i>	MF471394.1	F- TGGCGCTCAGATCATACCTT R- TTCTCCGACCAACTCCGAAA	152
<i>CYP6DBI</i>	MF471384.1	F- TGCCGCACCGCTATTATTAC R- GAAATTCAATCGGCGGTCCA	230
<i>CYP6G1</i>	JN989967.1	F- GGTGCGCGATCCTGAAATAA R- CGCCACACTCTTCGATTGTA	235
<i>CYP6CY5</i>	MF471387.1	F- GACACCTGGTTTCACATCGG R- TGGTGTCCAACCTTGAGTCCA	191
<i>CYP4CH1</i>	MF443134.1	F- ACCATCATGATTGGAGGCCA R- GGGCAAAGGAGGGAACAATC	222
<i>CYP4CJ2</i>	MF471385.1	F- TCTGGGACGAGATGCTGTT R- GTTCAGCCACGGTTTCAACA	227
<i>CYP6CY21</i>	MT268677.1	F- TTGACAAAACCTGTTGGGAATC R- GAAGGAACATCAGAATCGCA	223
<i>CYP315A1</i>	MF471375.1	F- TGCAGTGGTGATACTACTG R- GGTGCCAACGGATACATTCT	171
<i>CYP6CY4</i>	MF687492.1	F- TGGGAAAAGCTCAATGTGCC R- AGGCGTCTCATCTGAAACA	159
<i>CYP306A1</i>	MF471382.1	F- GTGCACAAATGGCGGGATAA R- ATGCGGTATCCCTCAAGTGT	153