

Supplementary Table. 6. Primes and probes.

| | strand | sequences |
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| Cloning Primers* | hCD81-F | GCGGGATCCGCCACCATGGGAGTGGAGGGCTGC |
| | hCD81-R | GCGCTCGAGTCAGTACACGGAGCTGTTC |
| | hOCLN-F | GCGGTTCGACGCCACCATGTCATCCAGGCCTTTG |
| | hOCLN-R | GCGCTCGAGCTATGTTTTCTGTCTATCATAG |
| Genotyping primers | hCD81-F | GCCATGACCCGCAGACCACC |
| | hCD81-R | CCGGATGCCACAGCACAGCA |
| | hOCLN-F | TGTGTGGCCTCCACGCTTGC |
| | hOCLN-R | GGACTCGCCGCCAGTTGTGT |
| miR-122 | RT | GTCGTATCCAGTGCAGGGTCCGAGGTATTCGCACTGGATACGACCAAACA |
| HCV negative strand primers | RT | TGTCATGGTGGCGAATAAGCCATGGCGTTAGTAT |
| | 1-F | TGTCATGGTGGCGAATAA |
| | 1-R | GGGACCGCTCCGAAGTTTT |
| | 2-F | GCCGAGTAGCGTTGGGTT |
| | 2-R | ACAAGTATACTCCGCCAACG |
| Molecular beacon probes | POSITIVE | TRAMRA-CCGGTGAagagccatagtggtctgcgaaTCACCGG-DABCYL |
| | NEGATIVE | TRMRA-CCGGTGA [†] tcgagaccactatggctctTCACCGG-DABCYL |
| Real-time PCR primers [†] | hCD81_F | AACATCATCAGCAACCT <u>CTTCA</u> |
| | hCD81_R | CAATGCCGATGAGGTACAG |
| | hOCLN_F | ATATGGTTCACAAATATAT <u>GCCC</u> |
| | hOCLN_R | CCCCAGTACAATGGCAATG |
| | HCV(5'UTR)_F | ATCACTCCCCTGTGAGGAAT |
| | HCV(5'UTR)_R | GCGGGTTGATCCAAGAAAGG |
| | HCV(1b)_F | CGGGAGAGCCATAGTGGTCTGCG |
| | HCV(1b)_R | CTCGCAAGCACCTATCAGGCAGTA |
| | HCV(2a)_F | GTCTAGCCATGGCGTTAGTATGAG |
| | HCV(2a)_R | ACCCTATCAGGCAGTACCACAAG |
| | ACTIN_F | AAGGAGCCCACGAGAAAAAT |
| | ACTIN_R | ACCGAACTTGCATTGATTCCAG |
| | ApoE_F | CTGACAGGATGCCTAGCCG |
| | ApoE_R | CGCAGGTAATCCCAGAAGC |
| | miR-122_F | GCGGCGGTGGAGTGTGACA |
| | miR-122_R | ATCCAGTGCAGGGTCCGAGG |
| | U6 snRNA_F | CGCTTCGGCAGCACATATAC |
| | U6 snRNA_R | TTCACGAATTTGCGTGTGTCAT |
| | mGAPDH_F | ACGGCCGCATCTTCTTGTGCA |
| | mGAPDH_R | ACGGCCAATCCGTTACACC |
| | mViperin_F | TGCTGGCTGAGAATAGCATTAGG |
| | mViperin_R | GCTGAGTGTGTTCCCATCT |
| | mIif27_F | GCTTGTGGGAACCCTGTTTG |
| | mIif27_R | GGATGGCATTGTTGATGTGGAG |
| | mIif44_F | AACTGACTGCTCGCAATAATGT |
| | mIif44_R | GTAACACAGCAATGCCTCTTGT |
| | mMx1_F | GACCATAGGGTCTTGACCAA |
| | mMx1_R | AGACTTGCTCTTTCTGAAAAGCC |
| | mEif2ak2_F | ATGCACGGAGTAGCCATTACG |
| | mEif2ak2_R | TGACAAATCCACCTGTTTTCGT |
| | mOas1a_F | ATGGAGCACGGACTCAGGA |
| | mOas1a_R | TCACACACGACATTGACGGC |
| | mIifnb1_F | CAGCTCCAAGAAAGGACGAAC |
| mIifnb1_R | GGCAGTGTAACTCTTCTGCAT | |
| mIip10_F | CCAAGTGCTGCCGTCATTTTC | |
| mIip10_R | GGCTCGCAGGGATGATTTCAA | |
| mCypA_F | GACTGAATGGCTGGATGGC | |
| mCypA_R | GGTGATCTTCTGCTGGTCTTG | |
| mIifna4_F | AGGACAGGAAGGATTTTGA | |
| mIifna4_R | GCTGCTGATGGAGGTCATT | |
| mIifn3_F | CTCTTCATACACTGAGCCACATTC | |
| mIifn3_R | AACAGGTTGGAGGTGACAGA | |

* the underline indicates restriction sites

[†] the underline indicates specific sites in human