

Supplementary Table S1. Primers used for PCR.

| N | Gene | Sequence (5' → 3') | Amplicon Size (bp) | PrimerBank or PubMed ID | N | Gene | Sequence (5' → 3') | Amplicon Size (bp) | PrimerBank or PubMed ID |
|----|--------------|--|--------------------|-------------------------|----|----------------|--|--------------------|-------------------------|
| 1 | <i>GAPDH</i> | F: TGCACCACCAACTGCTTAGC R: GGCATGGACTGTGGTCATGAG | 87 | PMID: 12184808 | 19 | <i>PDE7A</i> | F: CGTATGCTAGGAGATGTACGTGT R: TGAACCCGACGTACCACGAAA | 201 | 341823662c1 |
| 2 | <i>PDE1A</i> | F: ATGGGGTCTAGTGCCACAGAG R: GCACAGATGCCGCATATTCAAT | 187 | 51102298c1 | 20 | <i>PDE7B</i> | F: GGTTGAGAGGTGTGGCGAAAT R: CCTTAGCTCGTATATCTCCAGCA | 85 | 57242789c1 |
| 3 | <i>PDE1B</i> | F: CTGCGCTACATGGTGAAGCA R: CAAGATTTGCCGTGTCTCATCTA | 123 | 260436978c1 | 21 | <i>PDE8A</i> | F: TCCAGCCAGAGACGACACT R: ACAGGCATGGGACTACTTTCC | 116 | 341823697c2 |
| 4 | <i>PDE1C</i> | F: ATAAGACAGGAGTGGCGAACT R: GTATGCTCGTAGTCATGGATGG | 85 | 300796870c3 | 22 | <i>PDE8B</i> | F: ACGCAGGCTTCAACAGGAG R: CGTGGTCATCGCTTGTATTCT | 177 | 300244576c3 |
| 5 | <i>PDE2A</i> | F: GACCGCAAGATCCTCCAACCTG R: CCGAGCACTTTGTCTCCGA | 176 | 344925850c2 | 23 | <i>PDE9A</i> | F: GACTCCTCGACGCGATGTTT R: TTTCTGTAGTTGTCGTGGACG | 222 | 48762733c1 |
| 6 | <i>PDE3A</i> | F: CCACGGCCTCATTACCGAC R: TTGCTCACGGCTCTCAAGG | 78 | 347658968c1 | 24 | <i>PDE10A</i> | F: GAGACAACCAGCTACTCTCT R: ACAGGCTATTATTGCACTCTCCA | 105 | 359465520c2 |
| 7 | <i>PDE3B</i> | F: GTGCCGCCGAAGAAAAAGTG R: CAACTGCCATAGTAAGTGGCTG | 94 | 219879808c1 | 25 | <i>PDE11A</i> | F: TGATGACTTTTCTCTCGACGTTG R: AAGCCACTACACAGTGTCTC | 115 | 116536086c1 |
| 8 | <i>PDE4A</i> | F: ACACACCTGTCAGAAATGAGC R: GGGTGATGGGATCTCCACTTC | 99 | 341572546c1 | 26 | <i>PDE12</i> | F: GTAGTGCGCTGCGTACCTT R: GCGCTGCATGTTCTTGTGG | 81 | 189027128c1 |
| 9 | <i>PDE4B</i> | F: AACGCTGGAGGAATTAGACTGG R: GCTCCCGGTTCCAGATTCT | 110 | 82799485c1 | 27 | <i>GUCY1A2</i> | F: GAGGACTGCTTCGAGATTGTATC R: TGGTTCTAATCACAAACGGGGTA | 91 | 374093197c3 |
| 10 | <i>PDE4C</i> | F: GAGACGCTAGACGAGCTGGA R: GTGGGTCAACTCCCGGTTT | 120 | 341604763c2 | 28 | <i>GUCY1A3</i> | F: AGTGAGGAGATGGGACAACCTC R: GAGCCCTCGTCTGTAAAATC | 148 | 374253805c3 |
| 11 | <i>PDE4D</i> | F: GACCAATGTCTCAGATCAGTGG R: GTCAAGGGCCGGTTACAG | 187 | 308387383c1 | 29 | <i>GUCY1B3</i> | F: CTGCTTACGTCTCAAGGGTCA R: ATGTCACTTAGATACAGCCCTCT | 123 | 221139770c3 |
| 12 | <i>PDE5A</i> | F: CGGCCTCTTAGACCCATTGTT R: AGGGAATAGCGGTCAGCAGAT | 239 | 61744429c2 | 30 | <i>GUCY2F</i> | F: GAGGTTGCTGCGCGATTAG R: CCAGGGTTGGTAGGTCCAATAA | 173 | 134152693c2 |
| 13 | <i>PDE6A</i> | F: CATCGAACGACAGTTCCACAA R: GGTCATGTCTAAGAGACCCACAG | 88 | 170650673c3 | 31 | <i>GUCY2D</i> | F: TCGTGCTGGTTCCGATCCAAAC R: GTAGCCGGTGCCCTCACATAAT | 133 | 169791019c3 |
| 14 | <i>PDE6B</i> | F: ACACCCTCAGCCGATCACT R: ACATTTCTGTCAGCGGAAGCAT | 103 | 223718033c2 | 32 | <i>ADCY2</i> | F: GGGACCAGGTATCGTTCTTCC R: GTGAGGACGCTGGCAATGAT | 94 | 115387101c2 |
| 15 | <i>PDE6C</i> | F: ATGGACAAGCAAAGTGGTATG R: TGGTGATGAAGCCTTAGGAT | 187 | 57364938c2 | 33 | <i>ADCY6</i> | F: CTCCTGGTCCCTAAAGTGGAT R: GGAGGCAGCTCATATAGCGG | 124 | 168480141c1 |
| 16 | <i>PDE6D</i> | F: TTCAAAGGGCAATGCCTAGAAG R: TTCCCAGTTAAGACGCTTGCT | 134 | 56676309c3 | 34 | <i>ADCY8</i> | F: CCTGCGGCACCAAAGTCTT R: CGAGTTGCTAGGGGCACAG | 110 | 168480145c1 |
| 17 | <i>PDE6G</i> | F: AAGGCTAGAGATGACCTCGG R: GGGGTCCCTAAGTGAAAGC | 246 | PMID: 20655036 | 35 | <i>ADCY10</i> | F: GTGACGATTGTGTTTGTGAACC R: GATGTGCATATAGGCATCCTGG | 90 | 268370068c3 |
| 18 | <i>PDE6H</i> | F: GCAGACTCGCCAATTCAAGAG R: TCTGTTCTAGCCCCTCCATT | 90 | 291045288c1 | | | | | |