

| Viroid | Starting Nucleotide | Primer Name | Sequence (5'-3') |
|------------|---------------------|--|---|
| PLMVd (+) | 91 | T7_PLMVd_F | TAATACGACTCACTATAGGTGCCCTATTCGGAGCACTGCAG |
| | | PLMVd_R | CCAAGGTGGAGGGGCTGAGA |
| | 157 | T7_PLMVd_F2 | TAATACGACTCACTATAGGACTTTTCCCTAGGGTTTGAGCGG |
| | | PLMVd_R2 | CACCTTACGTCATTGCGAGGTGC |
| 256 | T7_PLMVd_F3 | TAATACGACTCACTATAGGTCCCCCTCAGAGGTGACTC | |
| | PLMVd_R3 | GGGTTTGAATCCCGGGTAG | |
| PLMVd (-) | 97 | T3_PLMVd(-)_R | AATTAACCCTCACTAAAGGGTAGGCGTCGTAATCCAGTTTC |
| | | PLMVd(-)_F | GGGATTCAAACCCGGTCCCC |
| | 269 | T3_PLMVd(-)_R2 | AATTAACCCTCACTAAAGGTAACCACTCTCTCATAAGTCTGG |
| | | PLMVd(-)_F2 | TCTCAGCCCCCTCCACCTTGG |
| GHVd (+) | 112 | T7_GHVd_F | TAATACGACTCACTATAGGACTTTTGTCTCGTCAGAG |
| | | GHVd_R | GAATCCGTGGTACCCCTAACGC |
| | 259 | T7_GHVd_F2 | TAATACGACTCACTATAGGGGGAAGGTTAGCTGCCATTCTTG |
| | | GHVd_R2 | GGAGGAGGCTTCTCCGAAGTCCCC |
| GHVd (-) | 172 | GHVd (-)F2 | ACGAAGTTTAACTTCGTAAGTCG |
| | | T3_GHVd_R2 | AATTAACCCTCACTAAAGGGAAAAGGTTACCGGCTAAGGC |
| | 372 | GHVd (-)F | GTGGAGAATAGTTCCACCTCTCGCC |
| | | T3_GHVd_R | AATTAACCCTCACTAAAGGTCAGTCACTGTGTAGTGGCTCTG |
| CChMVd (+) | 71 | T7_CChMVd_F | TAATACGACTCACTATAGGGAGTAAAGCTTTCGCTCTCTCC |
| | | CChMVd_R | GCACAAGCCGAATCTGGAAGA |
| | 205 | T7_CChMVd_F2 | TAATACGACTCACTATAGGGACCCGGAACAGATCCAGTTC |
| | | CChMVd_R2 | AACCCCTTCTCCCGGGGTTGGAC |
| CChMVd (-) | 78 | CChMVd(-)_F | TCTTCCAGATTCGGCTTGTGC |
| | | T3_CChMVd_R | AATTAACCCTCACTAAAGGATTTCGATCCTGTGATGGATC |
| | 347 | CChMVd(-)_F2 | GTCCAACCCCGGAGGAAAGGGTT |
| | | T3_CChMVd_R2 | AATTAACCCTCACTAAAGGAGGTATTTACCTCTACTCCG |
| | | CChMVd(-)3_6FAM/NED | GGAACAGATCCAGTTCCGG |
| ELVd (+) | 140 | T7_ELVd_F | TAATACGACTCACTATAGGATCGGCGGATGTACCGATTTCGC |
| | | ELVd_R | GTACTTACGCAAGTAAGGGATTGG |
| | 268 | T7_ELVd_F3 | TAATACGACTCACTATAGGGTGAGTGTACCTCTTTTCG |
| | | ELVd_R3 | TCCTAGGGAACACATCCTTG |
| ELVd (-) | 84 | ELVd_F | CCAATCCCTTACTTTCGTAAGTAC |
| | | T3_ELVd_R | AATTAACCCTCACTAAAGGGAAGAGCTTTGGAGGAACG |
| | 229 | T3_ELVd_R3 | AATTAACCCTCACTAAAGGAAGACCGGAGAAATCCGAAAAG |
| | | ELVd(-)F3 | GAGAGAGACAGAGGACGGCCTCT |
| ASBVd (+) | 26 | T7_ASBVd_F3_G | TAATACGACTCACTATAGATTAAACTTTGTTTGACGAATCCAGG |
| | | ASBVd_R3 | CATATCCTCACTTCTTGTCTAATAAAC |
| | 114 | T7_ASBVd_F4_G | TAATACGACTCACTATAGTTTTATTAATAAAATAGTTCACTCG |
| | | ASBVd_R4 | GTTACCACGACTCCTCCTTCTCTC |
| ASBVd (-) | 2 | ASBVd(-)_F3 | GTTTATTAGAACAAGAAGTGAGGATATG |
| | | T3_ASBVd(-)_R3_G | AATTAACCCTCACTAAAGAAGATTTTGTAATAAAACAATGAAG |
| | 158 | ASBVd(-)_F4 | GAGAGAAGGAGGAGTCGTGGTGAAC |
| | | T3_ASBVd(-)_R4_G | AATTAACCCTCACTAAAGACAAGTCGAATCTCAGAGTCGGAAAGTCG |

Figure S8
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