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| Supplemental Table 1. Primers used in this study |  |  |  |
| Name | Primer sequence (5′–3′) | Fragment size (bp) | Position | Purpose |
| APMV F-1F | TATACACCTCRTCBCARACRGG | 403 | 4698–4720a | Detection |
| APMV F-1R | ACRAAYTGCTGCATCTTCCC | 5080–5100a |
| NDV M-F | TAGCAAATGCCTCTCCCC | 1247 | 4206–4223b | Sequencing |
| NDV M-R | GGTGGCACGCATATTAT | 5436–5452b |
| NDV F-F | GGGAAGATGCAGCAGTTTG | 1093 | 5087–5105b |
| NDV F-R | GGGTATTATTCCCAAGCC | 6162–6179b |
| NDV HN-1F | GACCCTCCTGGTATCATATC | 1066 | 5756–5775b |
| NDV HN-1R | CCCCCGATATAATCTGGG | 6804–6821b |
| NDV HN-2F | GCAAAGAACACATGGCG | 551 | 6466–6482b |
| NDV HN-2R | GAGTGATCTCTGCAACC | 7000–7016b |
| NDV HN-3F | GGTTGCACTCGGATACCC | 980 | 6928–6945b |
| NDV HN-3R | GCTTGTTCATCATCAAGC | 7890–7907b |
| NDV L-F | CCCTTGCCAGGCATCAGC | 825 | 7776–7793b |
| NDV L-R | GTTTTATCATTCTCTCTGTG | 8581–8600b |

a Numbers correspond to positions within the LaSota strain genome (JF950510).

b Numbers correspond to positions within the Chicken/China/Liaoning/02/2005 genome (KC542893).