

**Supplementary Table 1.** List of primers with the corresponding target genome region used to detect infectious agents associated with bovine respiratory disease.

| Organism                           | Target genome region                 | Primer sequence  | Fragment length (bp) | Reference |
|------------------------------------|--------------------------------------|--|----------------------|-----------|
| Bovine viral diarrhea virus        | 5' untranslated region (5'-UTR)      | 324-F: 5'-ATGCCCWTAGTAGGACTAGGCA-3'<br>326-R: 5'-TCAACTCCATGTGCCATGTAC-3'  | 288                  | 8         |
| Bovine respiratory syncytial virus | G attachment glycoprotein gene       | B5A-F: 5'-CCACCCTAGCAATGATAACCTTGAC-3'<br>B6A-R: 5'-AAGAGAGGATGCCTTGCTGTGG-3'<br>B7A-F: 5'-CATCAATCCAAAGCACCACACTGTC-3'<br>B8-R: 5'-GCTAGTTCTGTGGTGGATTGTTGTC-3' | 371                  | 1         |
| Bovine coronavirus                 | Nucleocapsid protein gene            | BCoV1-F: 5'-CGATGAGGCTATTCCGAC-3'<br>BCoV2-R: 5'-TGTGGGTGCGAGTTCTGC-3'<br>BCoV3-F: 5'-TTGCTAGTCTTGTCTGGC-3'  | 251                  | 6         |
| Bovine parainfluenza 3 virus       | Hemagglutinin-neuraminidase gene     | HNfwd-F: 5'-GAATGACTCATGATAGAGGTAT-3'<br>HNseq1-R: 5'-AGGACAACCAGTTGTATTACAT-3'  | 647                  | 9         |
| Bovine alphaherpesvirus 1          | Glycoprotein D gene                  | P3-F: 5'-GCTGTGGGAAGCGGTACG-3'<br>P4-R: 5'-GTCGACTATGGCCTTGTGTGC-3'<br>P5-F: 5'-ACGGTCATATGGTACAAGGACAGCG-3'   | 425                  | 4         |
| <i>Pasteurella multocida</i>       | Open reading frame of the clone KMT1 | KMT1SP6-F: 5'-GCTGTAAACGAACTCGCCAC-3'<br>KMT1T7-R: 5'-ATCCGCTATTTACCCAGTGG-3'  | 460                  | 7         |
| <i>Histophilus somni</i>           | 16S ribosomal (r)DNA gene            | HS453-F: 5'-GAAGGCGATTAGTTTAAGAG-3'<br>HS860-R: 5'-TTCGGGCACCAAGTRTTCA-3'  | 408                  | 2         |
| <i>Mannheimia haemolytica</i>      | lktA-artJ intergenic region          | Mh-F: 5'-GTCCCTGTGTTTTTATTATAAG-3'<br>Mh-R: 5'-CACTCGATAATTATTTCTAAATTAG-3'  | 385                  | 3         |
| <i>Mycoplasma bovis</i>            | 16S-23S rDNA intergenic region       | Molli-F: 5'-CCGTCAAACYATGGGAGC-3'<br>Molli-R: 5'-GTGYCCCCGCMTACTCAGG-3'<br>nMb-F: 5'-GTACACTTGTCTTTTATCACTATA-3'<br>nMb-R: 5'-AAGGTATCTCGCTTTATGTCCT-3'          | 488                  | 5         |

## References

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