

dISFV-NC_016997.1 AAACA----UAACGCAUUAAGAAUAGGUUU----GAAAGAGUUAGAAUCAUAGAGAUCAAUG 52
dISFV-NC_017086.1 ---A----UAACACGUUUUGAAUAGGUUUUAUUGUGAAUACCAUG----- 43
MBFV-NC_012532.1 AACAGGUUU----AAUUUGGAAUUUGGAAACGAGAGUUUCUGG--UCAUG---AAAAAC 49
MBFV-NC_035889.1 ACAGGUUUU----AUUUUGGAAUUUGGAAACGAGAGUUUCUGG--UCAUG---AAAAA- 48
MBFV-NC_012533.1 AACAGUUUU-----CGUUGUGAGAAAGAAAACAGAUUCUUCUGG--AAAUG---AAAA- 47
MBFV-NC_002640.1 -----UCUAACAGUUUUGUUUGAAUAGAGAGCAG-AUCUCUGGAAAAAUG-----AA- 45
MBFV-NC_001477.1 ACGUAGUUUCUAACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUG---AACAA- 48
MBFV-NC_001475.2 ACGUAGUGCUGACAGUUU-UUU-AUUAGAGAGCAG-AUCUCUG-----AUG---AACAA- 48
MBFV-NC_001474.2 -CGUAGUUUCUAACAGUUU-UUUAAUUAGAGAGCAG-AUCUCUG-----AUG---AAUAA- 48
MBFV-NC_009028.2 U--U----UAACACAGUCUGAAUAGUUUUUUU-AGCAAGGGA-UUUUCCAUG---UCUA-- 47
MBFV-NC_040776.1 U--U----UAACACAGUCUGAAUAGUUUUUUU-AGCAGAGGU-ACUACCAUG---UCUA-- 47
MBFV-NC_034151.1 -----UAACACAGUGUGAAUAGUUUUUUUGAACAGAGGUGAAAAACAUG---UCUA-- 47
MBFV-NC_007580.2 ---U----AACUUGACUGCGAACAGUUUUUUU-AGCAGGGAAUUAACCAUG---UCUA-- 47
MBFV-NC_001437.1 ---U----UAGUGCAGUUUAAACAGUUUUUUU-AGAACGGAAGAUAAACCAUG---ACUA-- 47
MBFV-NC_006551.1 ---U----UAACACAGUGCGCGAGCUGUUUCUU-UGAGCGUUGAUUUUCAAUG---UCUA-- 47
MBFV-NC_001563.2 A--U----UAACACAGUGCGAGCUGUUUCUU-GGCACG-AAGAUCUCGAUG---UCUA-- 47
MBFV-NC_009942.1 A--U----UAACACAGUGCGAGCUGUUUCUU-AGCACG-AAGAUCUCGAUG---UCUA-- 47
MBFV-NC_012534.1 CCGU----AAACACAGUUUGAACGUUUUUUGGAGA---GAGAUUUAUUG---UUUAAAC 49
MBFV-NC_015843.2 ACGU----AAACACAGUUUGAACGUUUUUUGGAUA---GAGACAACUAUG---UCUAAC 49
MBFV-NC_000943.1 -----UAACGCGGUUUGAACAGUUUUUUUGGAGCUUUUGAUUUCAAUG---UCUA-- 47
MBFV-NC_009026.2 A-----GAAAGUUUUUGAACAGAGAGAGAGAG-AGCGAAUUCUCUGAAUG---GCUA-- 47
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAGUUUUUUUGAAC----AG----CAAUG---ACCA-- 47
MBFV-NC_032088.1 UC-----UAUAACUCAGUGUGAACAAAGUUUUUUUGAUC----AGUGGAUAAUG---ACUA-- 47
GGAU-----UAAUUGUUGUUAGAGAGACAGCGAGAGAGUAUACU---AUGGUUGAGAAA 52
AUUU-----CUACUGAAAGG---GUAGAGAAAAGGAGUUUUGCUUCUCAUGGCAACAAGA 52
MBFV-NC_012735.1 -UUU-----CUACUGAAAGG---AGUAGAAGAAAGGAGAUUCAUUCUCCAAUGGCAACAAAG 52
MBFV-NC_002031.1 -----UAAUCGUUCGUGAGCGAUUAGCAGAGAACUGACAGAAUGUC----- 45
dISFV-NC_040610.1 --GAACAAUUUUUAAAGAAGUCAAAACU-----AGAAAUUUAAUU-UCAAUGGUUAGAC-- 50
dISFV-NC_024017.1 UCGCGCAUUUUUGAUAGAGUU-----GACU-----AGAUAAAGAAUUAUCGAUGGUGACUA- 50
MBFV-NC_018705.3 CCGUA-----AACACAGUUUGAACGUUUUUUGGAGA---GAGACUACUAUG---UCAAAC 49
.....10.....20.....30.....40.....50.....6



.....((((.....((.....((((.....((((.....((((..... 110
dISFV-NC_016997.1 GUCUUUCGAGCUGAGGAGGCCGCUUAAACGGGGCCGUCGAUAUAUAAAACGUAAGCUGC-- 110
dISFV-NC_017086.1 GCGAAUAAACCCCAAGAACCCGGAGACCGGCUAUCGAUAUAGAGAGCGUGCGUUGCCU 103
MBFV-NC_012532.1 CCCAAAGAAGAAAU--CCGGAGGA-UCCGGAUUGUCAUAUGCUAAAACGCGGAGUAGCC 106
MBFV-NC_035889.1 CCCAAAAAAGAAAU--CCGGAGGAUUCGCGGAUUGUCAUAUGCUAAAACGCGGAGUAGCC 106
MBFV-NC_012533.1 CCCAAAAAAGG-----GCCGCAUCGCGGCCUAUCAAUAUGCUAAGAGCGCUUUGGAUCU 100
MBFV-NC_002640.1 -CCAACGAAAAAAG--GUG--GUUAGACCACCUUUCAAUAUGCUGAAACGCGAGAGAAAC 100
MBFV-NC_001477.1 -CCAACGAAAAAAG--ACG--GGUCGACCUCUUUCAAUAUGCUGAAACGCGCGAGAAAC 103
MBFV-NC_001475.2 -CCAACGAAAAAAG--ACG--GGAAACCGCUAUCAAUAUGCUGAAACGCGGAGAGAAAC 103
MBFV-NC_001474.2 -CCAACGAAAAAAG--GCG--AAAAACACGCCUUUCAAUAUGCUGAAACGCGGAGAAAC 103
MBFV-NC_009028.2 -AAAAACCAGGAAAGUCCGCGUGCAAAGCGGACUGUCAUAUGCUAAAACGCCUGGCUAGC 106
MBFV-NC_040776.1 -AAAAACCAGGAGGACCCCGGGAAGGCGGGUUGUCAUAUGCUGAAACGCCCGCAGCAGC 106
MBFV-NC_034151.1 -AAAAACCAGGAGGACCCGGAAGAAACCCGGGUCUCAUAUGCUGAAACGCCGCAUACCC 106
MBFV-NC_007580.2 -AAAAACCAGGAAACCCCGUAGAAACCCGGGUUGUCAUAUGCUAAAACGCCGCGUAUCC 106
MBFV-NC_001437.1 -AAAAACCAGGAGGCCCGGUAAAAAACCCGGGCUAUCAAUAUGCUGAAACGCCGCCUACC 106
MBFV-NC_006551.1 -AGAAACCAGGAGGGCCGGAAGAAACCCGGGCCUCAUAUGCUGAAACGCCGCAUACCC 106
MBFV-NC_001563.2 -AGAAACCAGGAGGGCCCGUAAAAACCCGGGCUGUCAUAUGCUAAAACGCCGUAUGCCC 106
MBFV-NC_009942.1 -AGAAACCAGGAGGGCCCGCAAGAGCCGGGCUGUCAUAUGCUAAAACGCCGGAUAGCCC 106
MBFV-NC_012534.1 AAAAAACCAGGAAACCCGGGGCAACCCGGGUUGUCAUAUGCUAAAACGCCGGUGCAUCC 109
MBFV-NC_015843.2 AAAAAACCAGGAAGACCCGGCUCAGGCCGGGUUGUCAUAUGCUAAGCGCGGAACGUCC 109
MBFV-NC_000943.1 -AAAAACCAGGAGACCCGGGAAGCCCGGGUCUGUCAUAUGCUAAAACGCCGCAUACCC 106
MBFV-NC_009026.2 -AAAAACCAGGAAGACCCGGAGAAACCCGGGUCUCAUAUGCUAAAACGCCGCGGCGCC 106
MBFV-NC_009029.2 -AAAAACCAGGAAGGCCCGGUAGAAUCCGGGCCUGUCAUAUGCUAAGCGCGGUGCGUCC 106
MBFV-NC_032088.1 -AAAAACCAGGAAGUCCGGCAUUCGUCGAGCUGUCAUAUGCUAAGCGCGGAGCAUCC 106
dISFV-NC_027999.1 ACUAGAGUAGGAUCUCGGGUUAAACCCGAGGUGUCAUAUGCUAAGCGGUAUCUG----- 107
MBFV-NC_008719.1 GGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAUAUGGUGACGGCCAAGG----- 107
MBFV-NC_012735.1 GGGAUGAUUAAGUCUCGGGUCGAUCCCGAGGCGUCAUAUGGUGACGGGCUAGAG----- 107
MBFV-NC_002031.1 --UGGUGUAAAGCUCAGGGAAAAACCCUGGGCGUCAUAUGGUAACGACGAGGAGUUCGC 103
dISFV-NC_040610.1 ----AACUUGGAAAGACCGUUGCAAGACGGUCUGUCAUAUGGGAAGCGGAUUACCACC 106
dISFV-NC_024017.1 ----AAUCAAGAAAGCCCGCCGAAAGCCGGGUCUCAUAUGGUAAACACGCAUACCCAAG 106
MBFV-NC_018705.3 AAAAAACCAGGAAGACCCGGGCACCCCGGGUUGUCAUAUGCUAAGCGCGGCGCAUCC 109
0.....70.....80.....90.....100.....110.....1

