

dISFV-NC_027999.1 -----GGAUUAAUUGUUGUUGAGAGAACAGCGAGAAGUACUACU---AUGGUUGAGAAAAACUAGAGUAGGAUCUCGCGGUUAAAAACCCGAGGGUGCAAUAGGU--UAGGCGUACUGG-----AA---NNNNNNNN-----UGGCGAACC--CCAUGGAAAAUUUAGC-U--CCAUUUGACACCAGGGAAAAAGACGGAGAAAACUUCUUGCUUACUCGCAGC--AACCACCGGAAAG-GCACAGUAUGCCUGA-AA-CAUUCGGUGGUUGAUGUAGUAAACAACACAGUU 255
MBFV-NC_008719.1 -----AUUUUCUACUGAAAGGUAGAGA--A--AAGGAGUUUUGCUUUCUACUUGGCAACAAGAGAGCAAGCAAGUCUCGGGUACAGACCCCGGGGCGCAAUAGGU--GACGGCCAAGGC-----CA---NNNNNNNN-----GAGGAGACCCUCCCGAGCAC-UAUAGC-GGAACCAUUAUGACGCCUGGG--AAAGACGGAGACACUCCUUGAUU-CUCACCUU--UCUCACCCUAAA-GCACAGAUUGCUUG--AA-UGCAGGGUGGGGAAGUUGGGAACCAACUAGUG 255
MBFV-NC_012735.1 -----UUUCUACUG-AAAGGAGUAGA--AGAAAAGGAGAUUCAUCCCAAUGGCAACAAAGGGGAUGAAUAGUCUCGGGUCGCAUCCCGAGGGCGCAAUAGGU--AGCGGCUAGAGU-----GA---NNNNNNNN-----C-CCAUCCCUCCCGAGCACACAUAGC-GGAACCAUUAUGACGCCAGGG--AAAGACGGAGACACUCCUUGAUU-CUCACCUU--UCUCGCCUAAA-GCACAGAUUGCUUG--AA-UGCAGGGCGGGGAAGUUGAGAACCAACUAGUG 255
MBFV-NC_002031.1 -----UAAUCGUUCGUUGAGCGAUUAGCAGAGAACUGACCAGAACAUGUC-----UGGUUCGUAAAGCUCAGGGAAAAACCCUGGGCGCAAUAGGU--UACGACGAGGAGUUCGCUCCUUGU-----NNNNNNNN-----G--AGACCCUCCAGGGAAACAAUAGUGGGACCAUUAUGACGCCAGGG--AAAGACGGAGUGGUUCUCUGCUU--UUCUCC--AGAGGUUGUGA-GCACAGUUUGCUCAAGAA-UAAGCAGACCUUUGGAUGACAAACACAAAACC 255
dISFV-NC_016997.1 -----A-AACAUAACGCAUUAAGAAUAGGUUUGAAAGAGUUGAAUCA--AUGAGAUCAAUGGUCUUUCGAGCUAGGAGGCCCGUUAACGGGCGCGCAUADAUAUAAACGUAAGCUGC-----NNNNNNNN-----A-GGAAACCCCCCAAAAAUGUAUAGG-CUAUUAUAUCCGACACCUAGG-AAAGACUGGAGAUAC-CUUUGGUUUCUUCGACAC-UAACUCCUAUG-GCACAGAGCGCCAU-AA-GAAACGGAGUUAGUAUGGAAAAACAUAAGUA 255
dISFV-NC_017086.1 -----AUAAACACGUUUGAAUAGGUUUUAUUGUG-AAUCGAAAUACCG--AUG-----GCGAAUAAACCCAAGAGCCCGGAAAGACGGGCUAUCGAUUAUGAGAGACGUGCGUUGCCUCGCGUCU-----NNNNNNNN-----A-GAGGAGACCCCCCAAAAAUGAAGG-GCAUUAUAUCCGACACUUGG-AAAGACAGAGGUACUCGCUGAUUCACCGCCAC-CAG-ACUACACG-GCACAGCGCGCCCG-AA-AAGGUAGUUUGGAGGUUGUAAAAACAACAAGUA 255
MBFV-NC_009028.2 U-----UUAACACAGUCUGAAUAGUUUUUU-AGCA-AGGGAUU-UCC---CAUGUCU--A---AAAAACCAGGAAGUCCCGUGCAAAAGCGGACUGCAAUAGGU--UAAAACGCGCGGCUAGCGUGU-----NNNNNNNN-----GUGGAGACCCUUCAGCAC-CAAGCGCGAA--ACAAACAGCAUUAUGACGCCUGGGA--AAGACCGGGAGAUCCUCUGCU--UUCCAUC--ACCAGCCACUAG-GCACAGAUUGCCCG--CA-AGUAGUGGCUGGUGGUGA-AAAACACAUGGA- 255
MBFV-NC_040776.1 U-----UUAACACAGUCUGAAUAGUUUUUU-AGCA-GAGGUAC-UAC---CAUGUCU--A---AAAAACCAGGAGGACCCGCGGAAAGGCGGUUGCAAUAGGU--UGAAACGCCCAGCCAGCGUUU-----NNNNNNNN-----A-GGAGACCCCUUGGCACACAAGCGCAAA-GACAAACAGCAUUAUGACCCUGGGA--AAGACCGGGAGACCCUCUGCU--CUGCAAC--ACCAGCCACUAG-GCACAGAUUGCCCG--AA-AUCUGUGGCUGGUGCUGC-AAAACACA-GGA- 255
MBFV-NC_001563.2 A-----UUAACACAGUGCGAGCUGUUUCUU-GGCACGAAGAUC-UCC---GAUGUCU--A---AGAAACCAGGAGGGCCCGGUAAAAACCGGGCUGCAAUAGGU--UAAAACGCGGUUUGCCCGCG-----NNNNNNNN-----GAGGAGACCCUUGGCA-AAAACACCA-A-AAGAAACAGCAUUAUGACCCUGGGA--UAGACUAGGGGAUUCUUGCU--CUGCACA--ACCAGCCACAG-GCACAGUGCGCCG--AC-AUAGGUGGCUGGUGGUGCUAGAACACA-GGA- 255
MBFV-NC_009942.1 A-----UUAACACAGUGCGAGCUGUUUCUU-AGCACGAAGAUC-UCC---GAUGUCU--A---AGAAACCAGGAGGGCCCGGCAAGAGCCGGGCGCAAUADGUAUAAACGCGGAUUGCCCGCG-----NNNNNNNN-----GGAGACCCCGUGCCACAAAACACCACA-ACAAAACAGCAUUAUGACCCUGGGA--UAGACUAGGAGAUUCUUGCU--CUGCACA--ACCAGCCACAG-GCACAGUGCGCCG--AC-AAUGGUGGCUGGUGGUGCGGAGAACACA-GGA- 255
MBFV-NC_007580.2 -----UAACUUGACUGCGAAACAGUUUUUU-AGCA-GGGAAUUACCC--AAUGUCU--A---AAAAACCAGGAAAAACCCGUAAGAAACCGGGUUGCAAUAGGU--UAAAACGCGGCUUCCCGCG-----NNNNNNNN-----UAGAGGAGACCCCUUGCCGUUAA--CGCAAA---CAACAAGCAUUAUGACCCUGG-A--AAGAC-AGGAGAUCCCUUGCU--UUUCAAC--ACCAGCCACAAG-GCACAGAGCGCCG--UA-AAGUGUGGCUGGUGGUGUAAAAAUACACAGGA- 255
MBFV-NC_034151.1 -----UUAACACAGUCUGAAUAGUUUUUU-AGCA-AGGGAUU-UCC---CAUGUCU--A---AAAAACCAGGAGGACCCGGAAGAAACCGGGUUGCAAUAGGU--UGAAGCGCACAUCCCGCGCUA-----NNNNNNNN-----AGACCCUUGGCAGUGAA-CGCAAA-UAAAAACAGCAUUAUGACCCUGGGA--UAGACUGGGAGACCCUCUGCU--AUUUCAAC--ACCAGCCACUG-GCACAGAACGCCG--UA-AGU-GUGGCUGGUGGUGUAAAAUAAACACAGAG 255
MBFV-NC_012534.1 CC----GUAAACACAGUUUGAAC-GUUUUUUUGAGA-GAGAU---UAU---UAUGUUU--AACAAAAAACAGGAAAAACCCGGGGCAACCCGGGUUGCAAUAGGU--UAAAGCGCGGUGCAUCCC-----NNNNNNNN-----GGAGACCCCUUGCGAGUUAAACACCAAA-ACAAAACAGCAUUAUGACCCUGGGA--GAGACUAGGAGACCCUCUGUCC--UAACAAC--ACCAGCAUCCG-GCACAGAGCGCCG--UA-AGU-GUGGCUGGUGGUGGUAGAACACAGUG-- 255
MBFV-NC_015843.2 AC----GUAAACACAGUUUGAAC-GUUUUUUUGAUA-GAGAC---AAC---UAUGUCU--A---AAAAAACCCAGGAAGACCCGGCUCAGGCCGGGUUGCAAUAGGU--UAAAGCGCGGAACGUCCC-----NNNNNNNN-----AGGAGACCCCUUGCGAGUGAAACACCACA-A-GAAACAGCAUUAUGACCCUGGGA--UAGACUAGGAGACCCUCUGUCC--UAACAAC--ACCAGCCACUUG-GCACAGAUUGCCG--AA-AGU-GUGGCUGGUGGUGGUAGAACACAGGA-- 255
MBFV-NC_000943.1 -----UUAACGCGGUUUUGAACAGUUUUUUUGAGCUUUUGAUU-UCA---AAUGUCU--A---AAAAACCAGGAGGACCCGGGAAGCCCGGGUUGCAAUADGUAUAAACGCGGAUUGCCCGCG-----NNNNNNNN-----GACCCACUCUCAAAAAGCAUC-AA---ACAACAAGCAUUAUGACCCUGGGA--AAGACUAGGAGAUUCUUGCUUAUUCCAAC-AUCAGUACAAG-GCACCAGCGCCG--AA-CACUGUGACUGAUGGGGGAGAGACCACAGGA 255
MBFV-NC_006551.1 -----UUAACACAGUGCCGGCAGUUUCUUUGAGC-GUUGAUU-UUC---AAUGUCU--A---AGAAACCAGGAGGGCCCGGAAGAAACCGGGCCAUAUAGGU--UGAAACGCGGCAUACCCCGCG-----NNNNNNNN-----ACCCCGCA-UCAUAGCAUCAA---AAAAACAGCAUUAUGACCCUGGGA--UAGACUAGGAGAUUCUUGCUUAUUCCAAC-AUCAACCACAAG-GCACAGAGCGCCG--AA-AAUUGUGGCUGGUGGGAACUAGACCACAGGA 255
MBFV-NC_001437.1 -----UUAACACAGUCUGAAUAGUUUUUUUGAAC-GAGGUAAAAAC---CAUGUCU--A---AAAAACCAGGAGGGCCCGGUAAAAACCGGGUUGCAAUAGGU--UGAAACGCGGCUAACC CGCG-----NNNNNNNN-----UAGAGGAGACCCCGCAUUUGC-AU---CAAACAAGCAUUAUGACCCUGGGA--UAGACUGGGAGAUUCUUGCUUAUUCCAAC-AUCAGUACUAG-GCACAGAGCGCCG--AA-GUAUGUAGCUGGUGGAGGAAGAACACAGGA 255
MBFV-NC_009026.2 -----AGAAAAGUUUUCGAGGAAAGACGAGAA-GCGAAUUCUCUG---AAUGGCU--A---GAAACCAGGAAGACCCGGAGGAAACCGGGUUGCAAUAGGU--UAAAGCGUACGGCGGCAACG-----NNNNNNNN-----CCCUUGGCAAAACAGUUAACGCACAA-AAGAAACAGCAUUAUGACCCUGGGA--UAGACCGGAGAUUUGCUGCC--UCGCAAC--ACCUCCACCCG-GCACAGAACGCCG--AC-A-UGGUGGAGGGGUGCUAAGACACCA--GAU 255
MBFV-NC_009029.2 UGAUUACAUAACACAGUGUGAACAGUUUUUUGAACAG---C-----AAUGACC--A---AAAAACCAGGAAGGCCCGGUAGAAAUCCGGGCGCAAUADGUAUAAAGCGGUGUCCC-----NNNNNNNN-----AGACCCUCCCAAAAGAAAGACCAACACAAAACAGCAUUAUGACCCUGGGA--AAGACUAGGGAUUUGCUGCU--UAGACU--GGAGCCGGAGGCC--UU-GA--GGAGCCGGAGGCC--AAAAACCCAGA 255
MBFV-NC_032088.1 UC-----UAUAACUCAGUGGAAACAGUUUUUUGAUCAGUGGAU-----AAUGACU--A---AAAAACCAGGAAGGCCCGGCAAUUCGUCGAGCUGCAAUADGUAUAAAGCGGAGCAUCCCGCG-----NNNNNNNN-----UAGAGGUUAGAGGAGACCCCAUACCGCAAA-ACAAAACAGCAUUAUGACCCUGGGA--UAGACUAGGAGAUCCGUGCU--CUGGAAC--CCCAGUU-CUAG-GCACAGAACGCCG--U-AA--UGAACUGGGGUGCC--AAAAACCCGAA 255
MBFV-NC_012532.1 AA-----CAGGUUUA-----AUUUGGAUUUGGAAACGAGAGUUUCUGG-UCAUGAAA--A-ACCCCAAGAAAGAAAUCCGGAGGA-UCCGGAUUGCAAUAGGU--UAAAACGC--GGAG-UAGCCC--GUGNNNNNNNN-----GGUAGAGGAGACCCCGGAAACGCACAAACAGCAUUAUGACG--UGGGA--AAGAC-CAGAGACUCCAUGAGUUUCCACCAC-GCUGGCCGCCAG-GCACAGAUUGCCG--AA-CUUCGGCGGCCGGUGUGGGGAAAUCCAUGGUU 255
MBFV-NC_035889.1 AC-----AGGUUUUA-----UUUUGGAUUUGGAAACGAGAGUUUCUGG-UCAUGAAA--A-CCCCAAAAAGAAAUCCGGAGGAUUCGGAUUGCAAUAGGU--UAAAACGC--GGAG-UAGCCC--GUGNNNNNNNN-----GUUAGAGGAGACCCCGGAAACGCACAAACAGCAUUAUGACG-CUGGGA--AAGAC-CAGAGACUCCAUGAGUUUCCACCAC-GCUGGCCGCCAG-GCACAGAUUGCCG--AA-UAGCGCGCGCGGUGUGGGGAAAUCCAUGGUU 255
MBFV-NC_012533.1 AA-----CAGUUUUC-----GUGUUGAGAAAGAAACAGAUUCUUCGG-AAUGAAA--A-----CCCAAAAAGG-----GCCGCAUCGCGGCCUAUAUAGGU--UGAAGCGCUUUGGAUCUAGCGCUGGCCNNNNNNNN-----GAGACCCCGCGCCCAUAACCAACAUAAAACAGCAUUAUGACCCUGGGA--AAGAC-CGGAGACUCUGUUGGCUUUCCCAC--ACCAGCCCCUG-GCACAGAGCGCC--A-GAUCGGGGGUUGGUGGAGCAACCCAACAGG 255
MBFV-NC_002640.1 -----UCUAACAGUUUGUUGAAUAGAGAGCAG-AUCUCUGGAAAAAUC-----A-A-CCAACGAAAAAGGUG-GUUGACCACCUUCAAUADGUAUAAACGCGAGAGAAACCGCGUAUCAANNNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCAA-CACAA-AAACAGCAUUAUGACG-CUGGGA--AAGAC-CAGAGAUCUUGCU-GUCUCUGCAAC-AUCA--AUCCAG-GCACAGAGCGCCG--CA-AGAUG--GAUUGGUUGUUGAUCCAACAGGU 255
MBFV-NC_001477.1 AC-----GUAGUUCUAACAGUUU-UUU-AUUGAGAGAGCAG-AUCUCUG-----AUGAAC--A-A-CCAACGAAAAAGACG--GGUCGACCGUCUUCAAUADGUAUAAACGCGGAGAGAAACCGCGUGU--NNNNNNNN--UAGAGGUUAGAGGAGACCCCGCACAAACAAACAGCAUUAUGACG-CUGGGA--GAGAC-CAGAGAUCUUGCU-GUCUCUACAGC-AUCA--UUCAG-GCACAGAACGCCA--AA-AAAUG--GAAUGGUGCUUGUAAUCAAACAGGU 255
MBFV-NC_001475.2 AC-----GUAGUGCUGACAGUUU-UUU-AUUGAGAGAGCAG-AUCUCUG-----AUGAAC--A-A-CCAACGGAAGAAGACG--GGAACACCGUCUAUAUAGGU--UGAAACGCGUGAGAGAAACCGUGUGU--NNNNNNNN--ACUAGAGGUUAGAGGAGACCCCGCA-AAUAA-AAACAAGCAUUAUGACG-CUGGGA--GAGAC-CAGAGAUCUUGCU-GUCUCCUACG-AUCA--UUCAG-GCACAGAACGCCA--GA-AAAUG--GAAUGGUGCUUGUAAUCAAACAGGU 255
MBFV-NC_001474.2 -C-----GUAGUUCUAACAGUUU-UUUAAUUAGAGAGCAG-AUCUCUG-----AUGAAU--A-A-CCAACGAAAAAGGCG--AAAAACACGCCUUCAAUADGUAUAAACGCGAGAGAAACCGCGUGU--NNNNNNNN--CUAGAGGUUAGAGGAGACCCCGCA-AACAAAAACAGCAUUAUGACG-CUGGGA--AAGAC-CAGAGAUCUUGCU-GUCUCCUACG-AUCA--UUCAG-GCACAGAACGCCA--GA-AAAUG--GAAUGGUGCUUGUAAUCAAACAGGU 255
dISFV-NC_024017.1 -----UCGCGCAUUUUGAU-AGAGUUGACUAGAUAAAGAAUUAUC---GAUGGUG---ACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUAAUADGUAUAAACACGCAUACCAAGCGUG-----NNNNNNNN-----CCCCACUGCUGAGAG-CAAAAUAGCAUUAUGACACUUGGGAAGAGACAGAGAUUGAUGUUGUUUGUCAAUGCU-GACAACAGCCAUGGCACAGAGCGCCUGAAGAGAGCUGUUGUUAAUUUACAACAAAUAAUA 255
dISFV-NC_040610.1 -----GAACAAUUUUAAAGAGUCAAACUAGAAAUUUAAU-UC---AAUGGUU---AGACAACUUGGAAAGACCGUUGCAAGACGGUCUGCAAUADGUAUAAAGCGGAUUAACACCCCGCU-----NNNNNNNN-----ACCCCUCAUUUUUAUUAACAAGCAUUAUGACACCAGG-AUAGACGGAGAAU-ACAAAUUUCUGGGCACUAAACACUACAAGCACAGAGCGCUUCGAGAAGUAGUUGUUCUUGACGAGAGAAUAGUAUAUCU 255
MBFV-NC_018705.3 CC----GUAAACACAGUUUGAACGUUUUUUGGAGAGAGACUUC-----AUGUCA--AACAAAAAACAGGAAGACCCGGGGCAACCCGGGUUGCAAUADGUAUAAAGCGCGGCAUCCC-----NNNNNNNN-----CCCGCGGAAUCAAAGCAAGGCCCAACUGCAAUAGGU--UAAAACAGGGGAAGGACUAGAGGUUAGAGGAGACCCUUGCG--AGUGAACACCAC-ACAAAACAGCAU-----AUUGACACCCUGGGAAAGACUAGGAGACCCUC
.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280.....290.....3

