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MBFV-NC_009028.2     U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAAGGGA-UUUCCCAUGUCU---AAA 49
MBFV-NC_040776.1     U--U----UAACACAGUCUGAAUAGUUUUUU-AGCAGAGGU-ACUACCAUGUCU---AAA 49
MBFV-NC_034151.1     --------UAACACAGUGUGAAUAGUUUUUUGAACAGAGGUGAAAACCAUGUCU---AAA 49
MBFV-NC_001563.2     A--U----UAACACAGUGCGAGCUGUUUCUUGGCAC-GAAGAUCUC-GAUGUCU---AAG 49
MBFV-NC_009942.1     A--U----UAACACAGUGCGAGCUGUUUCUUAGCAC-GAAGAUCUC-GAUGUCU---AAG 49
MBFV-NC_000943.1     --------UAACGCGGUUUGAACAGUUUUUUGGAGCUUUUGAUUUCAAAUGUCU---AAA 49
MBFV-NC_006551.1     ---U----UAACACAGUGCCGGCAGUUUCUUUGAGC-GUUGAUUUUCAAUGUCU---AAG 49
MBFV-NC_001437.1     ---U----UAGUGCAGUUUAAACAGUUUUUUAGAACGGAAGAUAACC-AUGACU---AAA 49
MBFV-NC_007580.2     ---U----AACUUGACUGCGAACAGUUUUUUAGCAGGGAAU-UACCCAAUGUCU---AAA 49
MBFV-NC_012534.1     CCGU----AAACACAGUUUGAAC-GUUUUUUGGAGAGAGAUUAUU---AUGUUUAACAAA 52
MBFV-NC_015843.2     ACGU----AAACACAGUUUGAAC-GUUUUUUGGAUAGAGACAACU---AUGUCUAACAAA 52
MBFV-NC_009029.2     UGAUUACAUAACACAGUGUGAACAAGUUUUUUGAACAG----C----AAUGACC---AAA 49
MBFV-NC_032088.1     UC----UAUAACUCAGUGUGAACAAGUUUUUUGAUCAGUGGAU----AAUGACU---AAA 49
MBFV-NC_009026.2     AG-------AAAAGUUUUCGAGGAAAGACGAGAAGCGAAUUCUCUG-AAUGGCU---AGA 49
dISFV-NC_024017.1    UC--------GCGCAUUUUGAUAGAGUUGACUAGAUAACGAAUUAUCGAUGGUG---ACU 49
dISFV-NC_017086.1    AU--------AACACGUUUGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUGGCG---AAU 49
MBFV-NC_002640.1     -------UCUA-ACAGUUUGUUUGAAUAGAGAGCAGAUCUCUGGAAAAAUG---AA--CC 47
MBFV-NC_001474.2     -CGUAGUUCUA-ACAGUUU-UUUAAUUAGAGAGCAGAUCUCUG-----AUGAAUAA--CC 50
MBFV-NC_001477.1     ACGUAGUUCUA-ACAGUUU-UUU-AUUAGAGAGCAGAUCUCUG-----AUGAACAA--CC 50
MBFV-NC_001475.2     ACGUAGUGCUG-ACAGUUU-UUU-AUUAGAGAGCAGAUCUCUG-----AUGAACAA--CC 50
MBFV-NC_012532.1     AACAGGUUU-------AAUUUGGAUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAACCCC 52
MBFV-NC_035889.1     ACAGGUUUU-------AUUUUGGAUUUGGAAACGAGAG-UUUCUGGUCAUGAAAAA-CCC 51
MBFV-NC_012533.1     AACAGUUUU-------CGUGUUGAGAAAGAAAACAGAU-UCUUCGGAAAUGAAAA--CCC 50
dISFV-NC_040610.1    --------GAACAAUUUUAAAGAAGUCAAAACUAGAAAUUUAAUUUCAAUGGUU---AGA 49
MBFV-NC_002031.1     U-------AAUCGUUCGUUGAGCG-AUUAGCAGAGAACUGACCAGAACAUGUC------- 45
dISFV-NC_016997.1    A-------AACAUAACG-CAUUAGAAUAGGUUUGAAAGAGUUGAAUCAAUGAGAUCAAUG 52
MBFV-NC_008719.1     A-------UUUCUACUGAAAGG-GUAGAGAAAAGGAGUUUUGCUUCUCAUGGCAACAAGA 52
MBFV-NC_012735.1     --------UUUCUACUGAAAGGAGUAGAAGAAAGGAGAUUCAUUCCCAAUGGCAACAAAG 52
dISFV-NC_027999.1    G-------GAUUAAUUGUUGUUUGAGAGAACAGCGAGAAGUACUACU-AUGGUUGAGAAA 52
MBFV-NC_018705.3     CCGU----AAACACAGUUUGAACGUUUUUUGGAGAGAGACUACU----AUGUCAAACAAA 52
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MBFV-NC_009028.2     AAACCAGGAAAGUCC---GCUGCAAAGCGGACUGUCAAUAUGCUAAAACGCCUGGCUAGC 106
MBFV-NC_040776.1     AAACCAGGAGGACCC---GCCGGAAGGCGGGUUGUCAAUAUGCUGAAACGCCCAGCCAGC 106
MBFV-NC_034151.1     AAACCAGGAGGACCC---GGAAAGAACCGGGUCGUUAAUAUGCUGAAGCGCACAUCCCGC 106
MBFV-NC_001563.2     AAACCAGGAGGGCCC---GGUAAAAACCGGGCUGUCAAUAUGCUAAAACGCGGUAUGCCC 106
MBFV-NC_009942.1     AAACCAGGAGGGCCC---GGCAAGAGCCGGGCUGUCAAUAUGCUAAAACGCGGAAUGCCC 106
MBFV-NC_000943.1     AAACCAGGAGGACCC---GGGAAGCCCCGGGUCGUCAAUAUGCUAAAACGCGGCAUACCC 106
MBFV-NC_006551.1     AAACCAGGAGGGCCC---GGAAGAAACCGGGCCAUCAAUAUGCUGAAACGCGGCAUACCC 106
MBFV-NC_001437.1     AAACCAGGAGGGCCC---GGUAAAAACCGGGCUAUCAAUAUGCUGAAACGCGGCCUACCC 106
MBFV-NC_007580.2     AAACCAGGAAAACCC---GGUAGAAACCGGGUUGUCAAUAUGCUAAAACGCGGCGUAUCC 106
MBFV-NC_012534.1     AAACCAGGAAAACCC---GGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGUGCAUCC 109
MBFV-NC_015843.2     AAACCAGGAAGACCC---GGCUCAGGCCGGGUUGUCAAUAUGCUAAAGCGCGGAACGUCC 109
MBFV-NC_009029.2     AAACCAGGAAGGCCC---GGUAGAAAUCGGGCCGUCAAUAUGCUAAAGCGCGGUGCGUCC 106
MBFV-NC_032088.1     AAACCAGGAAAGUCC---GGCAAUCGUCGGACUGUCAAUAUGCUAAAGCGCGGAGCAUCC 106
MBFV-NC_009026.2     AAACCAGGAAGACCC---GGAGGAAACCGGGUCGUCAAUAUGCUAAAGCGUACGGCGGCC 106
dISFV-NC_024017.1    AAAUCAAGAAAGCCC---GCCGCAAAGCGGGCUGUUAAUAUGCUAACAC--GCAUCACCA 104
dISFV-NC_017086.1    AAACCCAAGAAGC------CCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCU 103
MBFV-NC_002640.1     AACGAAAAAAG--GU---G--GUUAGACCACCUUUCAAUAUGCUGAAACGCGAGAGAAAC 100
MBFV-NC_001474.2     AACGGAAAAAG--GC---G--AAAAACACGCCUUUCAAUAUGCUGAAACGCGAGAGAAAC 103
MBFV-NC_001477.1     AACGGAAAAAG--AC---G--GGUCGACCGUCUUUCAAUAUGCUGAAACGCGCGAGAAAC 103
MBFV-NC_001475.2     AACGGAAGAAG--AC---G--GGAAAACCGUCUAUCAAUAUGCUGAAACGCGUGAGAAAC 103
MBFV-NC_012532.1     AAAGAAGAAAU--CC---GGAGGA-UCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCC 106
MBFV-NC_035889.1     AAAAAAGAAAU--CC---GGAGGAUUCCGGAUUGUCAAUAUGCUAAAACGCGGAGUAGCC 106
MBFV-NC_012533.1     AAAAAAGG----------GCCGCAUCGCGGCCUAUCAAUAUGCGAAGCGUUUGGAUCU 100
dISFV-NC_040610.1    CAACUUGGAAAGACC---GUUGCAAGACGGUCUGUCAAUAUGCGAAAGCGGAUUACCACC 106
MBFV-NC_002031.1     --UGGUCGUAAAGCUCAGGGAAAAACCCUGGGCGUCAAUAUGGUACGACGAGGAGUUCGC 103
dISFV-NC_016997.1    GUCUUUCGAGCUAGGAGGCCCGUUAAACGGGCCGUCGAUAUAAUAAAACGUAAGC----- 107
MBFV-NC_008719.1     GGAGCAAGCAAGUCUCGGGUCACGACCCGGGGCGUCAAUAUGGUGACGGCCAAGG----- 107
MBFV-NC_012735.1     GGGAUGAAUAAGUCUCGGGCUCGAUCCCGAGGCGUCAAUAUGGUAGCGGCUAGAG----- 107
dISFV-NC_027999.1    ACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUACUG----- 107
MBFV-NC_018705.3     AAACCAGGAAGACCC---GGGGCAACCCGGGUUGUCAAUAUGCUAAAGCGCGGCGCAUCC 109
                     0........70........80........90........100.......110.......1
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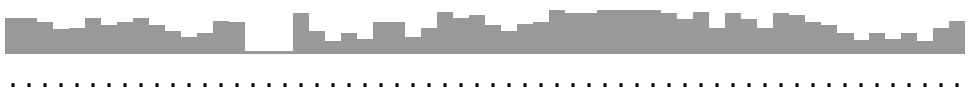

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                     ............................................................
MBFV-NC_009028.2     GUGU------NNNNNNN---GUGGAGACCCCUCAGCAC-CAAGCGCGAA--ACAAACAGC 154
MBFV-NC_040776.1     GUUU------NNNNNNN---A-GGAGACCCCUUGGCACACAAGCGCAAA-GACAAACAGC 155
MBFV-NC_034151.1     GCUA------NNNNNNN-------AGACCCCUUGGCAGUGAA-CGCAAA-UAAAAACAGC 151
MBFV-NC_001563.2     CGCG------NNNNNNN---GAGGAGACCCCGUGCCA-AAAACACCA-A-AAGAAACAGC 154
MBFV-NC_009942.1     CGCG------NNNNNNN-----GGAGACCCCGUGCCACAAAACACCACA-ACAAAACAGC 154
MBFV-NC_000943.1     CGCG------NNNNNNN--------GACCCCACUCUCAAAAGCAUC-AA-AC--AACAGC 148
MBFV-NC_006551.1     CGCG------NNNNNNN---------ACCCCGCA-UCAUAAGCAUCAAA-AA--AACAGC 147
MBFV-NC_001437.1     CGCG------NNNNNNN--------UAGAGGAGACCCCGCAUUUGCAUC-A---AACAGC 148
MBFV-NC_007580.2     CGCG------NNNNNNNUUAGAGGAGACCCCUUGCCGUUAAC-----GC-AAACAACAGC 154
MBFV-NC_012534.1     C---------NNNNNNN-----GGAGACCCCUUGCGAGUUAACACCAAA-ACAAAACAGC 154
MBFV-NC_015843.2     C---------NNNNNNN----AGGAGACCCCUUGCGAGUGAACACCACA-A-GAAACAGC 154
MBFV-NC_009029.2     CGCG------NNNNNNNAGAGG--AGACCCUCCCACAAAGAAGCGCAAACACAAAACAGC 158
MBFV-NC_032088.1     CGCG------NNNNNNNAGAGGUUAGAGGAGACCCCCACAUACCGCAAA-ACAAAACAGC 159
MBFV-NC_009026.2     AACG------NNNNNNNCCCCUUG-----GCAAAACAGUUAACGCACCAAAAGAAACAGC 155
dISFV-NC_024017.1    AGCGU-G---NNNNNNN-------CCCCACUGCU------------GAGAGCAAAAUAGC 141
dISFV-NC_017086.1    CGCGUCU---NNNNNNNAGAGGAGACCCCCCCAA------------AAAUGAAGGGCACU 148
MBFV-NC_002640.1     CGCGUAUCAANNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCCAACA--CAAAAACAGC 156
MBFV-NC_001474.2     CGCGUGU---NNNNNNN---CUAGAGGUUAGAGGAGACCCCCC-CGAAACAAAAAACAGC 156
MBFV-NC_001477.1     CGCGUGU---NNNNNNN----UAGAGGUUAGAGGAGACCCCCCGCACAACAACAAACAGC 156
MBFV-NC_001475.2     CGUGUGU---NNNNNNN--ACUAGAGGUUAGAGGAGACCCCCCG--CAAAUAAAAACAGC 156
MBFV-NC_012532.1     CGUG------NNNNNNN--------GGUUAGAGGAGACCCCCCGGAAAACGCAAAACAGC 152
MBFV-NC_035889.1     CGUG------NNNNNNN---------GUUAGAGGAGACCCCCCGGAAAACGCAAAACAGC 151
MBFV-NC_012533.1     AGCGCUGGCCNNNNNNN--------GAGACCCCCCGCGCCCAUAACCAACAUAAAACAGC 152
dISFV-NC_040610.1    CGCU------NNNNNNNA------------------CCCCCUCAUUAUUUAUAUAAACAGC 142
MBFV-NC_002031.1     UCCUUGU---NNNNNNN----------GAG---ACCCUCCAGGGAACAAAUAGUGGGACC 147
dISFV-NC_016997.1    ----UGC---NNNNNNN-----------AGGAAACCCCCCCAAAAAUGUAUAG-GCUAUU 148
MBFV-NC_008719.1     ----CCA---NNNNNNN----------GAGGAGACCCUCCCGAGCAC-UAUAG-CGGACC 148
MBFV-NC_012735.1     ----UGA---NNNNNNN----------C-CCAUUCCCUCCCGAGCACACAUAG-CGGACC 148
dISFV-NC_027999.1    ----GAA---NNNNNNN----------UGGCGAACCCCAUUGGAAAUUUA-----GCUUC 145
MBFV-NC_018705.3     C---------NNNNNNN----------CCCCGCGGAAAUCAAAGCAAGGCCCAACCUGGC 150
                     20.......130.......140.......150.......160.......170.......1
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                     ...........((((........))))...(((((...((.(((...(((...(((((((
MBFV-NC_009028.2     AUA-UUGACGCCUGGGAA--AGACCGG-GAGAUCC-UCUGCUU---UCCA--UCACCAGC 204
MBFV-NC_040776.1     AUA-UUGACACCUGGGAA--AGACCGG-GAGACCC-UCUGCUC---UGCA--ACACCAGC 205
MBFV-NC_034151.1     AUA-UUGACACCUGGGAU--AGACUGG-GAGACCC-UCUGCUUA-UUUCA--ACACCAGC 203
MBFV-NC_001563.2     AUA-UUGACACCUGGGA--UAGACUAG-GGGAUCU-UCUGCUCU---GCA--CAACCAGC 204
MBFV-NC_009942.1     AUA-UUGACACCUGGGA--UAGACUAG-GAGAUCU-UCUGCUCU---GCA--CAACCAGC 204
MBFV-NC_000943.1     AUA-UUGACACCUGGGAA-AAGACUAG-GAGAUCU-UCUGCUCUAUUCCA--ACAUCAGU 202
MBFV-NC_006551.1     AUA-UUGACACCUGGGAAUUAGACUAG-GAGAUCU-UCUGCUCUAUUCCA--ACAUCAAC 202
MBFV-NC_001437.1     AUA-UUGACACCUGGGAAU-AGACUGG-GAGAUCU-UCUGCUCUAUCUCA--ACAUCAGC 202
MBFV-NC_007580.2     AUA-UUGACACCUGGAAAGA----CAG-GAGAUCC-CCUGCUUU--UUCA--ACACCAGC 203
MBFV-NC_012534.1     AUA-UUGACACCUGGGA--GAGACUAG-GAGACCC-UCUGUCCUAA--CA--ACACCAGC 205
MBFV-NC_015843.2     AUA-UUGACACCUGGGA--UAGACUAG-GAGACCC-UCUGUCCUAA--CA--ACACCAGC 205
MBFV-NC_009029.2     AUA-UUGACACCUGGGAA--AGACUAG-GGGAUUU-GCUGCUC----UGG--ACUUCCGG 207
MBFV-NC_032088.1     AUA-UUGACACCUGGGAU--AGACUAG-GAGAUCC-GCUGCUC----UGG--AACCCCAG 208
MBFV-NC_009026.2     AUA-UUGACACCUGGGA--UAGACCGG--AGAAUU---UGCUGCCUCGCA--ACACCUCC 205
dISFV-NC_024017.1    AUA-UUGACACUUGGGAA-GAGACCAG-AGAUAUGAUGUUGUUUGUCAAUGCUGACAACA 198
dISFV-NC_017086.1    AUA-UCGACACUUGGGAA--AGACCAG-AGGUACUCGCUGAUUCACCGCC---ACCAGAC 201
MBFV-NC_002640.1     AUA-UUGACG-CUGGGAA--AGACCAGAGA-UCCUGCUGUCUC---UGCA---ACAUCAA 205
MBFV-NC_001474.2     AUA-UUGACG-CUGGGAA--AGACCAGAGA-UCCUGCUGUCUC---CUCA---GCAUCAU 205
MBFV-NC_001477.1     AUA-UUGACG-CUGGGAG--AGACCAGAGA-UCCUGCUGUCUC---UACA---GCAUCAU 205
MBFV-NC_001475.2     AUA-UUGACG-CUGGGAG--AGACCAGAGA-UCCUGCUGUCUC---CUCA---GCAUCAU 205
MBFV-NC_012532.1     AUA-UUGACG--UGGGAA--AGACCAGAGACU-CCAUGAGUUUCCACCAC---GCUGGCC 203
MBFV-NC_035889.1     AUA-UUGACG-CUGGGAA--AGACCAGAGACU-CCAUGAGUUUCCACCAC---GCUGGCC 203
MBFV-NC_012533.1     AUA-UUGACACCUGGGAA--AGACCGGAGACUCUGUUGGCUUUCC-CCAC---ACCAGCC 205
dISFV-NC_040610.1    AUA-UUGACACCAGGGA--UAGACCGG--AGAAAUACACAAUUUCUGGGCACUAAACAAC 197
MBFV-NC_002031.1     AUA-UUGACGCCAGGG-A-AAGACCGGAGUGGUUCUCUGCUUUUCCUC-C---AGAGGUC 200
dISFV-NC_016997.1    AUA-UCGACACCUAGG-A-AAGACUGGAGAUACCUUUGGUUCUUCGAC-A---CUAACUC 201
MBFV-NC_008719.1     AUA-UUGACGCCUGGG-A-AAGACCGGAGACACUCCUUGAUUCUCACC-U---UUCUCAC 201
MBFV-NC_012735.1     AUA-UUGACGCCAGGG-A-AAGACCGGAGACACUCCUUGAUUCUCACC-U---UUCUCGC 201
dISFV-NC_027999.1    AUA-UUGACACCAGGGAA-AAGACCGGAGAAACUUCUUGCUUACUCGCAG---CAACCAC 200
MBFV-NC_018705.3     AAAGCUGUAAACCAGGGGAAGGACUAGAGGUUAGAGGAGACCCCUUGCGAGUGAACAC-- 208
                     80.......190.......200.......210.......220.......230.......2
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MBFV-NC_009028.2     CACUAGGCACAGAUCGCCGC--AAGUAGUGGCUGGUG--GUGAA---AAACACAUGGAUC 257
MBFV-NC_040776.1     CACUAGGCACAGAUCGCCGA--AAUCUGUGGCUGGUG--CUGCA---AAACACA-GGAUC 257
MBFV-NC_034151.1     CACUCGGCACAGAACGCCGU--AAGU-GUGGCUGGUG--GUGAAAAUAAACACA-GAGUC 257
MBFV-NC_001563.2     CACACGGCACAGUGCGCCGA--CAUAGGUGGCUGGUG--GUGC--UAGAACACA-GGAUC 257
MBFV-NC_009942.1     CACACGGCACAGUGCGCCGA--CAAUGGUGGCUGGUG--GUGC--GAGAACACA-GGAUC 257
MBFV-NC_000943.1     CACAAGGCACCGAGCGCCGA--ACACUGUGACUGAUG--GGGGAGAAGACCACA-GGAUC 257
MBFV-NC_006551.1     CACAAGGCACAGAGCGCCGA--AAAUUGUGGCUGGUG--GGGAACUAGACCACA-GGAUC 257
MBFV-NC_001437.1     UACUAGGCACAGAGCGCCGA--AGUAUGUAGCUGGUG--GUGAGGAAGAACACA-GGAUC 257
MBFV-NC_007580.2     CACAAGGCACAGAGCGCCGU--AAAGUGUGGCUGGUG--GUGAA-AAAAUCACA-GGAUC 257
MBFV-NC_012534.1     CAUCCGGCACAGAGCGCCGU--AAG-UGUGGCUGGUG--GUG--GUAGAACACA-GUGUC 257
MBFV-NC_015843.2     CACUUGGCACAGAUCGCCGA--AAG-UGUGGCUGGUG--GUG--GUAGAACACA-GGAUC 257
MBFV-NC_009029.2     CUCUCGGCACAGAACGCCGU--UGA-GGAGCCGGAGG--CCCAA----AACAC-CAGAUC 257
MBFV-NC_032088.1     UUCUAGGCACAGAACGCCG---UAA-UGAACUGGGGU--GCCAA----AACAC-CGAAUC 257
MBFV-NC_009026.2     CACCCGGCACAGAACGCCGA--CA-UGGUGGGAGGGG--UCGUAAGACACCA---GAUUC 257
dISFV-NC_024017.1    GCCAUGGCACAGAGCGCCUG--AAGAGAGCUGUUGUUAUAAUUUACAAAAAAUAACAUAU 256
dISFV-NC_017086.1    UACACGGCACAGCGCGCCGG--AAAAGGUAGUUUGG---AGGUUGUAAAACAACAAGUAU 256
MBFV-NC_002640.1     UCCA-GGCACAGAGCGCCGC--AAGAUGGAUUGGUGU---UGU---UGAUCCAACAGGUU 256
MBFV-NC_001474.2     UCCA-GGCACAGAACGCCAG--AAAAUGGAAUGGUGC---UGU---UGAAUCAACAGGUU 256
MBFV-NC_001477.1     UCCA-GGCACAGAACGCCAA--AAAAUGGAAUGGUGC---UGU---UGAAUCAACAGGUU 256
MBFV-NC_001475.2     UCCA-GGCACAGAACGCCAG--AAAAUGGAAUGGUGC---UGU---UGAAUCAACAGGUU 256
MBFV-NC_012532.1     GCCA-GGCACAGAUCGCCGA--ACUUCGGCGGCCGGU---GUGG-GGAAAUCCAUGGUUU 256
MBFV-NC_035889.1     GCCA-GGCACAGAUCGCCGA--AUAGCGGCGGCCGGU---GUGG-GGAAAUCCAUGGGUC 256
MBFV-NC_012533.1     CCCU-GGCACAGAGCGCC----AGAUCGGGGGUUGGU---GUGG-AGCAACCCAACAGGA 256
dISFV-NC_040610.1    UAC-AAGCACAGAGCGCUUC--GAGAAGUAGUUGUUUCUUGACGAGAGAAUAUGUAUAUC 254
MBFV-NC_002031.1     UGU-GAGCACAGUUUGCUCAAGAAUAAGCAGACCUUUGGAUG----ACAAACACAAAACC 255
dISFV-NC_016997.1    CUAAUGGCACAGAGCGCCAU--AAGAAACGGAGUUAGUAUGG----AAAAAACAUAAGUA 255
MBFV-NC_008719.1     CCUUAAGCACAGAUUGCUUG--AAUGCAGGGUGGGGAAGUUG----GGAACCAACUAGUG 255
MBFV-NC_012735.1     CCUAAAGCACAGAUUGCUUG--AAUGCAGGGCGGGGAAGUUG----AGAACCAACUAGUG 255
dISFV-NC_027999.1    CGAAAGGCACAGUAUGCCUGA-AACAUUCGGUGGUUGAUGUA----GUAAACAACCAGUU 255
MBFV-NC_018705.3     -------CACACAAAAACAGCAUAUUGACACCUGGGAAAGACUAGGAGACCC-------U 254
                     40.......250.......260.......270.......280.......290.......3
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                     ....
MBFV-NC_009028.2     -U-- 258
MBFV-NC_040776.1     -U-- 258
MBFV-NC_034151.1     -U-- 258
MBFV-NC_001563.2     -U-- 258
MBFV-NC_009942.1     -U-- 258
MBFV-NC_000943.1     -U-- 258
MBFV-NC_006551.1     -U-- 258
MBFV-NC_001437.1     -U-- 258
MBFV-NC_007580.2     -U-- 258
MBFV-NC_012534.1     -U-- 258
MBFV-NC_015843.2     -U-- 258
MBFV-NC_009029.2     -U-- 258
MBFV-NC_032088.1     -U-- 258
MBFV-NC_009026.2     -U-- 258
dISFV-NC_024017.1    CU-- 258
dISFV-NC_017086.1    CU-- 258
MBFV-NC_002640.1     CU-- 258
MBFV-NC_001474.2     CU-- 258
MBFV-NC_001477.1     CU-- 258
MBFV-NC_001475.2     CU-- 258
MBFV-NC_012532.1     CU-- 258
MBFV-NC_035889.1     UU-- 258
MBFV-NC_012533.1     CU-- 258
dISFV-NC_040610.1    UUGG 258
MBFV-NC_002031.1     -ACU 258
dISFV-NC_016997.1    -UCU 258
MBFV-NC_008719.1     -UCU 258
MBFV-NC_012735.1     -UCU 258
dISFV-NC_027999.1    -UCU 258
MBFV-NC_018705.3     CUGU 258
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