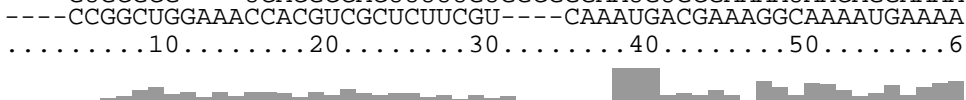


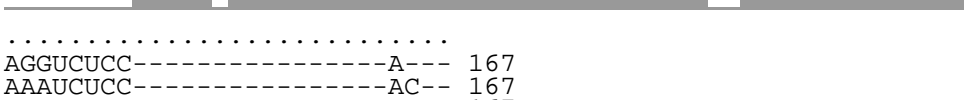
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ISFV-NC_021069.1 C--CUUAUUUACCUUCUGGUACUAAAAGUGGCACU-----AUGAAGAAGAAGGA--GUCGGG 50
ISFV-NC_012671.1 C--UUUAUUUACCUUCUGGUACUAAACGUGGCGCU-----AUGAAGAAGAAGGA--GUCUGG 50
ISFV-NC_024299.2 U--UGAAAAACUUUUUGCGACGUUGUUGAGAAG-----AUGGAGCGAAAGAAAGGAGUAA 52
ISFV-NC_027819.1 U--CACGGAAUAGCUUGGAGAAGAAAGCUUACUU-----AUGUCAAGUAAGGAAGCAGCUA 52
ISFV-NC_001564.2 G--AGAGACCGUGAGGUUUAACCCGGCAAGGA-----AUGAAGAGGAAGG--AUUUGGA 50
ISFV-NC_005064.1 A--GUUUUUCGGAAAAAGAGACUUGGUUCGU-----AUGAAUAGGAGAA--GCUUGAA 50
ISFV-NC_012932.1 A--GUUGUUUGGAAAAGAGAUUUCUGGUUCGU-----AUGAAUAGGAGG--UCUUGAA 50
ISFV-NC_034204.1 G--CGACAGAAGGAACCCAAAAAGGGAACCG-----AUGGAGCAAAACG--GAAAGUU 50
ISFV-NC_034017.1 U--AAAAAUUUUUUUUGGCGACGUACGGGAAAGU-----AUGGAGCAAUUGA--GGAAAGUU 50
ISFV-NC_027817.1 C--AAACUCUGGUGUCCACAGCUUUUGUAUUA-----AUGAGUGGAUUAGGAGCCUUC 52
TBFV-NC_003687.1 -----AGCUUAGGAGAACAAGAGCUGGGAGUGGUUAUG-----AUGACCACUUCUA 46
TBFV-NC_027709.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGGCAGGAAGUCCAUCCUGA 52
TBFV-NC_001809.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGGCAGGAAGACCAUCCUGA 52
TBFV-NC_003690.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGA-----AUGGCCGGGAAGGCCGUUCUAA 52
TBFV-NC_001672.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGUCAGGAAGGCCAUCCUGA 52
TBFV-NC_005062.1 A--AAGACAGCUUAGGAGAACAAGAGCUGGGG-----AUGGCCGGGAAGGCCAUCCUGA 52
ISFV-NC_027999.1 -----UUGUUUAGAGAGAACAGCGAGAAGUACUACU--AUGGUUGAGAAAAACUAGAGUAG 52
MBFV-NC_012735.1 -----AAGGAGUAGAAGAAAG--GAGAUUCAUUCCCAAUGGCAACAAAGGGGGAUGAAUA 52
MBFV-NC_008719.1 -----AAAGGUGUAGAGAAAAG--GAGUUUUGCUUCUCAUGGCAACAAGAGGAGCAAGCA 52
MBFV-NC_002031.1 -----UUGAGCGAU--UAGCAGAGAACUGACCAGAACUGUC-----UGGUCGUA 43
ISFV-NC_016997.1 -----CAUUAGAAUAGGUUUGAAAGAGUUGAAUC--AAUGAGAUCAUUGGUCUUUCGAG 52
MBFV-NC_002640.1 -----UGUUUGA-----AUAGAGAGCAGAUUCUCUGGAAAAAUG--AA--CCAACGAAAAA 47
MBFV-NC_001474.2 ACAGUUUUUUUA--UUAGAGAGCAGAUUCUCUG-----AUGAAUA--CCAACGAAAAA 50
MBFV-NC_001477.1 AACAGUUUUUA--UUAGAGAGCAGAUUCUCUG-----AUGAACAA--CCAACGAAAAA 50
MBFV-NC_001475.2 GACAGUUUUUA--UUAGAGAGCAGAUUCUCUG-----AUGAACAA--CCAACGAAAAA 50
MBFV-NC_012533.1 -----GUGUUGAG--AAAGAAAAACAGAU--UCUUCGGGAAAAGAAAAAC--CCAA--AAAAG 47
MBFV-NC_012532.1 -----AUUUGGA--UUUGGAAACGAGAGUUUCUGGUCUAGAAAAACCCCAAAGAAAGAA 52
MBFV-NC_035889.1 -----UUUUGGA--UUUGGAAACGAGAGUUUCUGGUCUAGAAAA--CCAAAAAAGAA 51
ISFV-NC_017086.1 UGAAUUGUUUA--UUUGGAAACGAAUACCG-----AUGGCA--UAAAACCAAGAA 50
MBFV-NC_001563.2 -----UGCGAGCUGUUUCUUGGCAC--GAAGA--UCUCGAGUUCU--AAGAAACCAAGGAG 49
MBFV-NC_009942.1 -----UGCGAGCUGUUUCUUGGCAC--GAAGA--UCUCGAGUUCU--AAGAAACCAAGGAG 49
MBFV-NC_006551.1 -----CGCG--CGAUUUUUUUGAGG--GUUGAUUUUCAAUGUCU--AAGAAACCAAGGAG 49
MBFV-NC_000943.1 -----U--GAACAGUUUUUUUGGAGCUUUUGAUUUUCAAUGUCU--AAAAAACCAAGGAG 49
MBFV-NC_009029.2 ACACAGUGUGAACAAGUUUUUUGAACAG-----CAAUGACC--AAAAAACCAAGAA 49
MBFV-NC_032088.1 -----AGUGUGAACAAGUUUUUUUGAAGCAGUGGA-----UAUGACU--AAAAAACCAAGAA 49
MBFV-NC_007580.2 -----GCGAACAGUUUUUUUAGCAGGGAAU--UACCCAAUGUCU--AAAAAACCAAGGAA 49
MBFV-NC_012534.1 -----AGUUUGAACGUUUUUUUGGAGAGAGAUUAU--AUGUUUAACAAAAACCAAGAA 52
MBFV-NC_015843.2 -----AGUUUGAACGUUUUUUUGGAGAGAGACAAU--AUGUUUAACAAAAACCAAGAA 52
MBFV-NC_009028.2 -----UCUGAAUAGUUUUUU--AGCAAGGGA--UUUCCCAUGUCU--AAAAAACCAAGGAA 49
MBFV-NC_040776.1 -----UCUGAAUAGUUUUUU--AGCAGAGGU--ACUACCAUGUCU--AAAAAACCAAGGAG 49
MBFV-NC_034151.1 -----UGAAUAGUUUUUUUGAAGCAGGUGAAAAACCAUGUCU--AAGGUCUAGAACCAAGGAG 49
MBFV-NC_009026.2 UCGAGGAAAGA-----CGAGAAGCGAAUUCUCUGAAUGGCU--AGAAAAACCAAGGAA 49
ISFV-NC_040610.1 A-----AAGAAGUCAAAAACUAGAAAUUAAUUUCGAUGGU--AGACAACUUGAA 49
ISFV-NC_024017.1 -----UGAUAGAGUUAGCUAGAUAACGAAUUAUCGAGUGG--ACUAAAUCAGAA 49
MBFV-NC_001437.1 -----UUAAACAGUUUUUUUAGAACGGAAGAUAAACCAUGACU--AAAAAACCAAGGAG 49
NKV-NC_005039.1 -----CGGAAGAACAACUCUGCCGGAUCAAAGACGAUGAAAAACGUAGACAAAAA 52
NKV-NC_003635.1 ----GUGCGCG--UCACGCCAUUUUUGUGCGGGAAUGUGCGAAAAUAACAGGAAAA 52
NKV-NC_004119.1 ----CCGGCUGGAAACCACGUCGCUCUUCGU--CAAUAGACGAAAGGCAAAAUAGAAA 52



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ISFV-NC_012671.1 AG-----UGGUGGAGAGAGG--ACGUGGAACGCGUCCUCCUGCCAAUCGUGGAGGAG 100
ISFV-NC_024299.2 AU-----UAUGGGAGAAGGG--AAGC--UAUCCCGCCCCUCCAUUAAGGAGAAAA 100
ISFV-NC_027819.1 AG-----AAGAAGGGAAUUU--CCUUGGGAGGAGGAGGAAAAACGGGGCUUUGAG--A 100
ISFV-NC_001564.2 GG-----CGCGAGGAAAAGC--GCCAGGGAGGGAUUUUUUCCACCCCGUUCUGGGGAA 100
ISFV-NC_005064.1 GA-----AGGGCCUCCGGG--CCCAACAUCCAAACCGUCCGUGGAGGGGCGGGCGG 100
ISFV-NC_012932.1 GA-----AAGGGCUUCGGG--CCAAACAUUCGUCGCGCAUCGACCAGACCGCGCG 100
ISFV-NC_034204.1 UU-----CCCAGCCGGGAGG--UGGUCGGGCGCGCCAGAGAGGGGCGUGGGGAAG 100
ISFV-NC_034017.1 UU-----CCGGCCGGGAGG--CGGUCGGGCGGGCCCCAAAGAGGGGCGUGGCGAAGA 100
ISFV-NC_027817.1 UU-----CCCCUCGAGGGA--AGAA--AAAGAAAGCCCCUGUGAUACAUAUCACAGG 100
TBFV-NC_003687.1 AA-----GGAAAGGGGGGCGGUCCCCUAGGCGCAAGCUUAAAGUGACCGCAAAUAAU 100
TBFV-NC_027709.1 AA-----GUAAGGGGGGCGGUCCCCUAGGCGUAGGUGUCGAAAGAGACCGCAA--CUGC 100
TBFV-NC_001809.1 AA-----GGCAAGGGGGGCGGUCCCCUAGGCGUAGGUGUGGAAAGAGACCGCGA-- 100
TBFV-NC_003690.1 AA-----GGAAAGGGGGGGGUGCCCCUAGGCGUAGGUGUGGAAAGAGACCGCGA-- 100
TBFV-NC_001672.1 AA-----GGAAAGGGGGGCGGUCCCCUAGGCGUAGGUGUGGAAAGAGACCGCGA-- 100
TBFV-NC_005062.1 AA-----GGAAAGGGGGGCGGUCCCCUAGGCGUAGGUGUGGAAAGAGACCGCGA-- 100
ISFV-NC_027999.1 GAUCUCGGGUUAAAAACCCGAGGUGUCAAUUAGUUAUAGGCGUAGG-----GAA-- 100
MBFV-NC_012735.1 AGUCUCGGGUCGAUACCCGAGGCGUCAAUUAGGUUAGGUGUAGCGUACAG-----UGA-- 100
MBFV-NC_008719.1 AGUCUCGGGUCACGACCCCGGGGCGUCAAUUAGGUGUAGCGGCAAGG-----CCA-- 100
MBFV-NC_002031.1 AAGCUCAGGGAAAAACCCUGGGCGUCAAUUAGGUUAGGUGUAGCGAGGAGUUCGCUCCUUGU-- 100
ISFV-NC_016997.1 CUGAGGAGGCGGUUAAACCGGGCGGUCGCAAUUAGGUUAGGUGUAGCGGCAAGG----- 100
MBFV-NC_002640.1 G--GU--G--GUUAGACCACCUUUAUUAUAGCUGAAACGCGAGAGAAACCGCGUAUCAA 100
MBFV-NC_001474.2 G--GC--G--AAAAACACCGCUUUAUUAUAGCUGAAACGCGAGAGAAACCGCGUGU-- 100
MBFV-NC_001477.1 G--AC--G--GGUACGCGGCUUUAUUAUAGCUGAAACGCGAGAGAAACCGCGUGU-- 100
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MBFV-NC_012532.1 U--CC--GGAGGA--UCCGGAUUGUCAAUUAGCUGAAACGCGGAGUAGCCCGUG-- 100
MBFV-NC_035889.1 U--CC--GGAGGAUUCGGAUUGUCAAUUAGCUGAAACGCGGAGUAGCCCGUG-- 100
ISFV-NC_017086.1 G--CC--C--GGAAGACGGGCUUUAUUAUAGGAGACGUGCGUAGGCCUUGCCUGCGUCU-- 100
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MBFV-NC_006551.1 GGCC--GGAAGAAACCGGGCGUUAUUAUAGCUGAAACGCGGCAUACCCCGCG-- 100
MBFV-NC_000943.1 GACC--GGGAAGCCCGGGGCGUUAUUAUAGCUGAAACGCGGCAUACCCCGCG-- 100
MBFV-NC_009029.2 GGCC--GGUAGAAAUCCGGGCGUUAUUAUAGCUGAAAGCGCGGUGCGUCCCGCG-- 100
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MBFV-NC_007580.2 AACC--GGUAGAAACCGGGGCGUUAUUAUAGCUGAAACGCGGAGUAGCCCGCG-- 100
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MBFV-NC_009028.2 AGUCC--GCUGCAAGCGGACUGUCAAUUAGCUGAAACGCGGUAUGCGUGU-- 100
MBFV-NC_040776.1 GACC--GCCGGAAGGCGGGUUGUCAAUUAGCUGAAACGCGGAGCAGCGU-- 100
MBFV-NC_034151.1 GACC--GGAAAGAACCGGGGUGUCAAUUAGCUGAAAGCGGCAUACCCCGCGU-- 100
MBFV-NC_009026.2 GACC--GGAGGAACCGGGGUGUCAAUUAGCUGAAACGCGGAGUAGCCCGCG-- 100
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ISFV-NC_024017.1 AGCCC--GCCGCAAGCGGGGUGUCAAUUAGCUGAAACGCGGACCAUACCGCGU-- 100
MBFV-NC_001437.1 GGCCC--GUUAAAAACCGGGGUGUCAAUUAGCUGAAACGCGGUGCAUCCCGCG-- 100
NKV-NC_005039.1 -----CCCGAUUUCGGGUGUCAAUUAAAAAAGCAAAAUCCAAAAAACAAAA-- 100
NKV-NC_003635.1 AGAACCUGGAAAGG-----GUUCAAGAGAAAGAAAGAAUGCUGGAGGAAUG 100
NKV-NC_004119.1 ACGCGAAUAAAAAGA-----AAGCAAAAAACUUAAGACCAAUUCUGGUGCA 100
0.....70.....80.....90.....100.....110.....1



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ISFV-NC_021069.1 NNNNNNN--GUGACGCCACCCCG--AAAUGUGUAUUGUGACAG--GCUUGCUCGCGAGCAGC 157
ISFV-NC_012671.1 NNNNNNN--GUGACGCCACCCCG--AAAUGUGUAUUGUGACAG--GCUUGCUCGCGAGCAGC 157
ISFV-NC_024299.2 NNNNNNN--ACGACACACCCCG--AAAUGUGUAUUAUGCCAUUGAGUUAUACAAAC 155
ISFV-NC_027819.1 NNNNNNN--UUAGCCACACCCCG--AAACGUGAUUUUAUGAUGACAAGAAUACAGUAG 158
ISFV-NC_001564.2 NNNNNNN--AGCGAACCUCCUCCGUUAAUUGUGAAAAAGUAUGGGG--AAAGAUCUAUCUU 157
ISFV-NC_005064.1 NNNNNNN--CGUACCCCUCC--UAAUGUGAAAAAGUAUGGGGAAAAAGAACCCAGCUA 155
ISFV-NC_012932.1 NNNNNNN--UUGCGAACCCCG--AAAUGUGAAAAAUUAGGGGAAAAAGUACCCAUUA 158
ISFV-NC_034204.1 NNNNNNN--ACGACACCCUCUG--AAAUGUUAUUAUAAACUGUGCGGAAAAAUCCAAAGA 156
ISFV-NC_034017.1 NNNNNNN--GCGACACCCUCUG--UAAUGUUUAUUAAGUUGCGGAAAAAUCCAAAGA 156
ISFV-NC_027817.1 NNNNNNN--UAGCCAGCCUCCCG--UAGUUAUUAUUAAGUUGCGGAAAAAUCCAAAGA 156
TBFV-NC_003687.1 NNNNNNN--AUUUUUCGCGCUCCUAUACUAAAUUUCUCCCGAGGAAACUGGGGGGGCGGUUC 159
TBFV-NC_027709.1 NNNNNNN--AUUUUUCGCGCUCCUAUACUAAAUUUCUCCCGAGGAAACUGGGGGGGCGGUUC 159
TBFV-NC_001809.1 NNNNNNN--UUUUUUCGCGCUCCUAUACUAAAUUUCUCCCGAGGAAACUGGGGGGGCGGUUC 159
TBFV-NC_003690.1 NNNNNNN--AUUUUUCGCGCUCCUAUACUAAAUUUCUCCCGAGGAAACUGGGGGGGCGGUUC 159
TBFV-NC_001672.1 NNNNNNN--UUUUUUCGCGCUCCUAUACUAAAUUUCUCCCGAGGAAACUGGGGGGGCGGUUC 144
TBFV-NC_005062.1 NNNNNNN--UUUUUUCGCGCUCCUAUACUAAAUUUCUCCCGAGGAAACUGGGGGGGCGGUUC 144
ISFV-NC_027999.1 NNNNNNN--CC--AUUGGAAAUUAGCUUUAUUAUUGACACCAGGGAA--AAGACC--GGAGAAA 156
MBFV-NC_012735.1 NNNNNNN--CCGAGCACACUAGCGGACCAUUAUUGACCGCCAGG--A--AAGACC--GGAGACA 156
MBFV-NC_008719.1 NNNNNNN--CCGAGCAC--UAUAGCGGACCAUUAUUGACCGCCAGG--A--AAGACC--GGAGACA 155
MBFV-NC_002031.1 NNNNNNN--AGGGAACAAAUAGUGGGACCAUUAUUGACCGCCAGG--A--AAGACC--GGAGUGG 156
ISFV-NC_016997.1 NNNNNNN--CAAAAUGUAUAGGCUUUAUUAUUGACCAUUAUUGAGG--A--AAGACC--GGAGUA 155
MBFV-NC_002640.1 NNNNNNN--CCGAAACCCCG--AACACAAACAGCAUUAUUGACG--CUGGGA--AAGACCAG--AGAU 155
MBFV-NC_001474.2 NNNNNNN--CCGAAACCCCG--AACACAAACAGCAUUAUUGACG--CUGGGA--AAGACCAG--AGAU 155
MBFV-NC_001477.1 NNNNNNN--CCGAAACCCCG--AACACAAACAGCAUUAUUGACG--CUGGGA--AAGACCAG--AGAU 155
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ISFV-NC_017086.1 NNNNNNN--CCGAAACCCCG--AACACAAACAGCAUUAUUGACG--CUGGGA--AAGACCAG--AGAU 155
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MBFV-NC_006551.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_000943.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_009029.2 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_032088.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_007580.2 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_012534.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_015843.2 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_009028.2 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_040776.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_034151.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_009026.2 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
ISFV-NC_040610.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
ISFV-NC_024017.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
MBFV-NC_001437.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
NKV-NC_005039.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
NKV-NC_003635.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
NKV-NC_004119.1 NNNNNNN--CAACAUAUAAACAGCAUUAUUGACACCUGGGA--AAGACCAG--AGAU 155
20.....130.....140.....150.....160.....170.....1



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ISFV-NC_008604.2 AGGUCUCC-----A-- 167
ISFV-NC_021069.1 AAAAUUCC-----AC-- 167
ISFV-NC_012671.1 AAAAUUCC-----AC-- 167
ISFV-NC_024299.2 AGGCUUCC-----ACCG 167
ISFV-NC_027819.1 AGAUUCC-----A-- 167
ISFV-NC_001564.2 AACCUCUCC-----AC-- 167
ISFV-NC_005064.1 AAUUCUCC-----ACCG 167
ISFV-NC_012932.1 AGCCUCC-----A-- 167
ISFV-NC_034204.1 CUGUCC-----ACC 167
ISFV-NC_034017.1 AUGUCC-----ACC 167
ISFV-NC_027817.1 AGAGUC-----C 167
TBFV-NC_003687.1 UUGUUC----- 167
TBFV-NC_027709.1 UUGUUC----- 167
TBFV-NC_001809.1 UUGUUC----- 167
TBFV-NC_003690.1 UUGUUC----- 167
TBFV-NC_001672.1 UUGUUC----- 167
TBFV-NC_005062.1 UUGUUC----- 167
ISFV-NC_027999.1 CUUCUUGAU-----U-A 167
MBFV-NC_012735.1 CUUCUUGAU-----U-C 167
MBFV-NC_008719.1 CUUCUUGAU-----U-C 167
MBFV-NC_002031.1 CUUCUUGAU-----U-C 167
ISFV-NC_016997.1 CUUCUUGAU-----U-C 167
MBFV-NC_002640.1 CUUCUUGAU-----U-C 167
MBFV-NC_001474.2 CUUCUUGAU-----U-C 167
MBFV-NC_001477.1 CUUCUUGAU-----U-C 167
MBFV-NC_001475.2 CUUCUUGAU-----U-C 167
MBFV-NC_012533.1 CUUCUUGAU-----U-C 167
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ISFV-NC_017086.1 CUUCUUGAU-----U-C 167
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MBFV-NC_015843.2 CUUCUUGAU-----U-C 167
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MBFV-NC_040776.1 CUUCUUGAU-----U-C 167
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MBFV-NC_009026.2 CUUCUUGAU-----U-C 167
ISFV-NC_040610.1 CUUCUUGAU-----U-C 167
ISFV-NC_024017.1 CUUCUUGAU-----U-C 167
MBFV-NC_001437.1 CUUCUUGAU-----U-C 167
NKV-NC_005039.1 CUUCUUGAU-----U-C 167
NKV-NC_003635.1 CUUCUUGAU-----U-C 167
NKV-NC_004119.1 CUUCUUGAU-----U-C 167
80.....190.....200.....

