

cISFV-NC_027817.1 ----UUGAAAGGUUCAAAACUCUGGUGCCAACGUUUGUAUUAAAAUGAGUGGAUUAGGAG 56
cISFV-NC_034204.1 ----AUUUUCUUUAGCGACAGAAGGAACCCAAAAAGGGAACCAUGGAGCAAAA--CGG 54
cISFV-NC_034017.1 ----CUAGUGUUUUAAAAAUUUUUUGGCGACGUACGGAAAGUAUGGAGCAAUUC--GAG 54
cISFV-NC_008604.2 AAGGCUUGAGAUUGUUU--AUAAUAGUCGUUUUUUCUCGCAGAA--AUGGGAAAGGACGACG 56
cISFV-NC_021069.1 AA--UUUCGCUACCUU--AUUACCUUGGUACUAAAGUGGCACUAUGAAGAAGAAGGA-- 54
cISFV-NC_012671.1 UA--AUUUGCUACUUU--AUUACCUUGGUUACUAACGUGGCGCUAUGAAGAAGAAGGA-- 54
cISFV-NC_024299.2 ----CUUUGGUUGUUUGAAAAACUUUUGCGACGUUGUUGAGAAGAUGGAGCGAAAGAAAG 56
cISFV-NC_027819.1 UAGACGUUUUGUCACGG----AAUAGCUUGAGAAGAAGCUUACUUAUGUCAAGUAAGGAAG 56
cISFV-NC_001564.2 AAUUUUUGCCGAGAGACCGUGA----GGUUGAACCCGGCAAGGAAUGAAGAGGAAG-- 53
cISFV-NC_005064.1 AAAUCAAAAGGAGUUGUUCGGAA----AAGUGACCUUGGUUUGUUAUGAAUAGAGGAA-- 53
cISFV-NC_012932.1 AAAAUCUAGAAGUUGUUUGGAA----AAGAGAUUUCUGGUUCGUUAGAAUAGGAUGC-- 53
.....10.....20.....30.....40.....50.....6



cISFV-NC_027817.1 GCCUUCUUCCCCUCGAGGGAAGAAAAAGAAAGCCCCUGUGA--UACAAUCACAGGNNN 113
cISFV-NC_034204.1 AAAGUUUUUCCCGACCGGGAGGUGGUCGGGCCCGGCCAGAGA--GGGGCUGGGGAAGANNN 113
cISFV-NC_034017.1 GAAGUUUUUCCCGGCCGGGAGGCGGUCGGGCGGGCCCCAAAGA--GGGGCUGGCGAAGANNN 113
cISFV-NC_008604.2 GUAAGAAGAAGAGGGGGCGGGCUCUCCGGAUGGU--UGUUAACCA--CCGGGGCGCGNNN 113
cISFV-NC_021069.1 GUCGGGAGUAGUGGAGAGAGAGGACGCGGUACGCGCCC--UCCGACAACCGUGGAGGAANNN 113
cISFV-NC_012671.1 GUCUGGAGUGGUGGAGAGAGGACGUGGAACGCGUCC--UCCUGCCAUCGUGGAGGAGNNN 113
cISFV-NC_024299.2 GAGUAAAUAUUGGGGAGAAGGGAAGC--UAUCCCGCC--CCUCCCAUUAAGGAGAAAANNN 113
cISFV-NC_027819.1 CACUGAAGAAGAAGGGAAUUAUCCUGGGAGGAGGAG--GAAAACGGGGCUUUGAG--ANNN 113
cISFV-NC_001564.2 AUUUGGAGGCGCGAGGAAAAAGCGCCAGGGGAGGGAUUUUUCCACCCCGUUCUGGGGAANNN 113
cISFV-NC_005064.1 GCUUGAAGAAGGGGCCUCCGGGCCCAACAUCCAACCGUCCGUCGUGGACCGCCCCGNNN 113
cISFV-NC_012932.1 UCUUGAAGAAAGGGCCUUCGGGCCAAACAUCUGGUCGGCCAUCGACCAGACCGCCCGNNN 113
0.....70.....80.....90.....100.....110.....1



cISFV-NC_027817.1 NNNNAUGUG-----CAGGGCAACAAAGUUCUAACGAACUAGGGUGAGUAG--CGUCACCCC 167
cISFV-NC_034204.1 NNNN--CUC-----CAGAGCAAACAAGGGAUUUAUCCUUAAGGUGAUUUGACGACCACCC 166
cISFV-NC_034017.1 NNNNACCUC-----CAGAGCAAACGAGGGAGAAAUCCCU--AGGUGAGUUGGCGACCACCC 167
cISFV-NC_008604.2 NNNNUCACCGCAAGGUGAGGGCGCACAGGAUCGAAUAUCCGGGUGAGU--GACGCCACCCU 170
cISFV-NC_021069.1 NNNNUCACCGUAAGGUGAGGGCGCACAGGAUCGAAUAUCCGGGUGAGU--GACGCCACCCU 172
cISFV-NC_012671.1 NNNN--CACCGCAAGGUGAGGGCGCACAGGAUCGAGUAUCCGGGUGAGU--GACGCCACCCU 171
cISFV-NC_024299.2 NNNN--ACCGCAAGGUGAGGGCGCACAGGACGAAAGUCCAGGGUGAGU--GACGACACCCU 170
cISFV-NC_027819.1 NNNNAAAGGCCAUCCACGCAAGGCAACAGGAAGAAAUCCGGGUGAUU--AGCCACACCCU 172
cISFV-NC_001564.2 NNNNCUUUCAAACUGGACGGGGCAACAGGGAGAAAUCCCGGGG--UA--GCGAACCUCU 170
cISFV-NC_005064.1 NNNN--UUCAAAACCGGACGGAGCAACAGGGA--AUUUCCGGGGA--C--GCGUACCCCU 167
cISFV-NC_012932.1 NNNN--A--AAAAACGGGCGGCAACAGGGAGAAAUCCCGGGGAUU--GCGAACCUCU 168
20.....130.....140.....150.....160.....170.....1



cISFV-NC_027817.1 CCGGU--UGUGAAACGAUUGCGACUAGAACUAAAGUCGAGAGU 209
cISFV-NC_034204.1 UCUGAAAUGUUAAUAAAAACUGUGCGGAAAAAUCCAAA--GACUGU 209
cISFV-NC_034017.1 UCUGUAAUGUUUAAUAAAGUUGUGCGGAAAAAUCCAAA--GAUG-- 209
cISFV-NC_008604.2 CCG--AAAUGUGUAUAGUAACA--GAGCA---UGCCUGCAGCAGCA 209
cISFV-NC_021069.1 CCG--AAAUGUGUAUUGUGACA--G--GCU---UGCUCGCAGCAGC-- 209
cISFV-NC_012671.1 CCG--AAAUGUGUAUUGUGACA--G--GCU---UGCUCGCAGCAGCA 209
cISFV-NC_024299.2 CCG--AAAUGUGUAUUAAGCCC--AUGUU---GAGUCUAUACAACA 209
cISFV-NC_027819.1 CCG--AAACGUGAUUUUAUGA--UGACA---AGAAUCAGAUGG-- 209
cISFV-NC_001564.2 CCGUUAAGUGUAAAAAGUAUG--GGG--AAAGAACUCAUCUUA--- 209
cISFV-NC_005064.1 CCG--UAAUGUGAAAAAGUAUG--GGGAAAAGAACCAGCUAAAU 209
cISFV-NC_012932.1 CCG--AAAUGUGAAAAAUUAUG--GGGAAAAGUACCCAUCU--AAGC 209
80.....190.....200.....210.....220..

