	((.((((((((((((((((((((((((((((((
dISFV-NC 016997.1	CAUUAGAAUAGGUUUGAAA <mark>G</mark> AGUUGAAUCAAUGAGAUCAAUGGŪCUU <mark>U</mark> CG.	AGCU 54
$dISFV-NC_{024017.1}$	UGAUAGAGUUGACUAGAUA <mark>A</mark> CGAAU <mark>U</mark> AUCGAUGGUGAC <mark>U</mark> AA	
disfv-NC_017086.1	UGAAUAGGUUUAUUGUGAA <mark>U</mark> CGAAA <mark>U</mark> ACCGAUGGCGAA <mark>U</mark> AA	
dISFV-NC_040610.1 dISFV-NC 027999.1	AAAGAAGUCAAAACUAGAAĀU <mark>U</mark> UAA <mark>U</mark> UUCAAUGG <u>U</u> UAGĀCA. UUGUUUGAGAGAACAGCGA <mark>G</mark> AAGUACUAUGGUUGAGAAAAC <mark>U</mark> AG.	ACUU 45 AGUA 51
TBFV-NC_02/999.1	AGCIJIAGGAGAACAGAGCUGGGAGIGGIJIAIJGAUGAGAAAACUAG.	UAAA 48
TBFV-NC 003690.1	AAAGACAGCUUAGGAGAACAAGAGCUUGGAAGGCCGGGAAGGCCGUUCU.	
TBFV-NC 001809.1	AAAGACAGCUUAGGAGAAC <mark>AAG</mark> AGC <mark>U</mark> GGGGAUGGGCAGGAAGA <mark>C</mark> CAU <mark>C</mark> CU	GAAA 54
TBFV-NC_027709.1	AAAGACAGCUUAGGAGAACAAGAGC <mark>U</mark> GGGGAUGGGCAGGAAGUCCAUCCU	
TBFV-NC_005062.1 TBFV-NC 001672.1	AAAGACAGCUUAGGAGAAC <mark>A</mark> AGAGC <mark>U</mark> GGGGAUGGCCGGGAAGGCCAUUCU AAAGACAGCUUAGGAGAACAAGAGC <mark>U</mark> GGGGAUGGUCAAGAAGGCCAUCCU	GAAA 54
1BFV-NC_001672.1	1020304050	
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	((((((((
dISFV-NC_016997.1	AGGAG <mark>G</mark> CCC <mark>GU</mark> UAAACGGGCCGUCGAUAUAAUAAAACGUAAGCUGC	NN 102
disfv-NC_024017.1	AGAAAGCCCGCCAAAGCGGGCUGUUAAUAUGCUAACACGCAUCACCAAGCGU	
dISFV-NC_017086.1 dISFV-NC 040610.1	AAGAA <mark>G</mark> C CCGGAAGACGGGCUAUCGAUAUAGUGAGACGUGCGUUGCCUCGCGU GGAAAGACC <mark>GU</mark> UGCAAGACGGUCUGUCAAUAUGCG AAAGCGGAUUACCACCCG -	CUNN 102
dISFV-NC 027999.1	GGAUCUCGGGUUAAAACCCGAGGUGUCAAUAUGAUUAGGCGUACUGG.	AANN 102
TBFV-NC 003687.1	GGAAAGGGG <mark>G</mark> CCGCUCCCCCUAGGCGCAAGCUUAAAGUGACCGC-AAAUAA	
TBFV-NC 003690.1	GGAAA <mark>G</mark> GGG <mark>G</mark> GGUCCCCCUCGACGAGCGUCGAAAGUGGCCCC-AA	NN 102
TBFV-NC_001809.1	GGCAAGGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGC-GA	
TBFV-NC_027709.1 TBFV-NC 005062.1	GGUAA <mark>G</mark> GGG <mark>G</mark> CCGUCCCCCUCGACGAGUGUCGAAAGAGACCGC-AA GGAAA <mark>G</mark> GGG <mark>G</mark> CCGUCCCCCUCGACGAGUGUCGAAAGAGACCGC-UA	NN 102
TBFV-NC_001672.1	GGUAAGGGGGCGGUCCCCCUCGACGAGUGUCGAAAGAGACCGC-UA	NN 102 NN 102
1B1 V NC_001072.1	0708090100110	
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dISFV-NC_016997.1	NNNNCAAAAAUGUAUAGGCUAUUAUAUCG <mark>AC</mark> ACC <mark>U</mark> AGGAA-AGACUGGAGAUA	
$dISFV-NC_024017.1$	NNNNC-UGCUGAGAGC-AAAAUAGCAUAU <mark>U</mark> GA <mark>C</mark> ACU <mark>U</mark> GGGAAGAGACCAGAG <mark>A</mark> UA	
disfv-NC_017086.1	NNNNNC-CCAAAAAUGA-AGGGCACUAUAUCGACACU <mark>U</mark> GGGAA-AG <mark>A</mark> CCAGAG <mark>G</mark> UA	
dISFV-NC_040610.1 dISFV-NC_027999.1	NNNNNC-UCAUUAUUUAUAUAACAGCAUAU <mark>UGAC</mark> ACCAGGG-AUAGĀCC <mark>G</mark> GAG <mark>A</mark> AA NNNNNCCAUUGGAAAUUUAGCUUCAUAUU <mark>GAC</mark> ACC <mark>A</mark> GGGAAAAG <mark>A</mark> CCGGAGAAA	
TBFV-NC_003687.1	NNNNNCCAUUGGAAAUUUAGCUUCAUAUUGACACCAGGAAAAGACCGGAGAAAN NNNNN-AUUUUUCCGCCUCCUAUACUAAAUUUCCCCCAGGAAACUGGGGGGGG	
TBFV-NC_003690.1	NNNNGAUAUUUCCUCCUCUAUACCAAA-UUCCCCCUCGUCAGAGGGGGGGGCGGU	
TBFV-NC_001809.1	NNNNNG-UUUUUCCUCCUCCUAUACCAAAU <mark>UUCC</mark> CCC <mark>U</mark> CGACAGAG <mark>G</mark> GGGGGC <mark>G</mark> GU	UCU <mark>U</mark> 161
TBFV-NC_027709.1	NNNNNGAUUUUUCCUCCUAUACCAAA- <mark>UUCC</mark> CCC <mark>U</mark> CGACAGAGGGGGGGGGGU	UCUU 161
TBFV-NC_005062.1 TBFV-NC_001672.1	NNNNNUAUACCAAA- <mark>UUCC</mark> CCC <mark>U</mark> CAUCUGAGGGGGGGC <mark>G</mark> GU NNNNNUAUACAAAA-UUCCCCC <mark>U</mark> CGGUAGAGGGGGGGC <mark>G</mark> GU	UCU <mark>U</mark> 146 UCU U 146
1BF V-NC_001072.1	20130140150160170	1
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dISFV-NC_016997.1	UGGUUCUU 167	
$dISFV-NC_024017.1$	GUUGUUU- 167	
disfv-NC_017086.1	CUGAUUCA 167	
dISFV-NC_040610.1 dISFV-NC_027999.1	C-AAUUUC 167 CUUGCUUA 167	
TBFV-NC 003687.1	GUUCUC 167	
TBFV-NC_003690.1	GUUCUC 167	
TBFV-NC 001809.1	GUUCUC 167	
TBFV-NC_027709.1	GUUCUC 167 GUUCUCCCUGAGCCACCAUCA 167	
TBFV-NC_005062.1 TBFV-NC 001672.1	GUUCUCCCUGAGCCACCAUCA 167 GUUCUCCCUGAGCCACCAUCA 167	
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