

| Accession | Sequence | Position |
|-------------------|--|----------|
| dISFV-NC_017086.1 | AUAACACGUUUGAAUAGGUUUAUUGUGAAUCGAAAUACCGAUG-----GCGAAUAAACCCAAGAAGC---CCGGAAGACGGGCUAUCGAUAUAGUGAGACGUG | 95 |
| dISFV-NC_024017.1 | UCGCGCAUUUUGAUAGAGUUGACUAGAUAAACGAAUUAUCGAUG-----GUGACUAAAUCAAGAAAGCCCGCCGCAAAGCGGGCUGUUAUAUAGCUAACAC--G | 96 |
| dISFV-NC_016997.1 | AAACAUAACGCAUUAAGAAUAGGUUUGAAAGAGUUGAAUCAAUAGAGAUCAAUGGUCUUUCGAGCUAGGAGGC---CCGUUAAACGGGCCGUCGAUAUAUAUAAAACGUA | 104 |
| dISFV-NC_040610.1 | GAACAAUUUUAAAGAAGUCAAAACUAGAAAUUUAAUUUCAUUG-----GUUAGACAACUUGGAAAGACCGUUGCAAGACGGUCUGUCAUAUAGCGAAAGCGG- | 97 |
| dISFV-NC_027999.1 | GGAUUAUUUGUUGUUUGAGAGAACAGCGAGAAGUACUACUAUG---GUUGAGAAAACUAGAGUAGGAUCUCGGGUUAAAACCCGAGGUGUCAUAUAGAUUAGGCGUA | 104 |
| |10.....20.....30.....40.....50.....60.....70.....80.....90.....100..... | |



| | CGUUGCCUCGCGUCUNNNNNNNNAGAGGAGACCCCCCAA | AAAAUGAAGGGCACUAUAUCGACA | CUUGG | GAA-AGA | CCAGAGGU | ACUCGC | UGAUUC | ACCGCC | --- | ACCA | |
|-------------------|---|--------------------------|-------|---------|----------|---------|--------|--------|------|---------|-----|
| dISFV-NC_017086.1 | CGUUGCCUCGCGUCUNNNNNNNNAGAGGAGACCCCCCAA | AAAAUGAAGGGCACUAUAUCGACA | CUUGG | GAA-AGA | CCAGAGGU | ACUCGC | UGAUUC | ACCGCC | --- | ACCA | 198 |
| dISFV-NC_024017.1 | CAUCACCAAGCGU-GNNNNNNN-----CCCCACUGCUGAGAGCA | AAAUAGCAUAUUGACA | CUUGG | GAAGAGA | CCAGAGAU | AUGAUG | UUGUU | UGUC | AAU | GCUGACA | 195 |
| dISFV-NC_016997.1 | AGCUGC-----NNNNNNNAGGAAACCCCCCAA | AAUGUAUAGGCUAUUAUAUCGACA | CUUAG | GAA-AGA | CUGGAGAU | ACCUU | UGUUC | UUGA | --- | CACUAA | 198 |
| dISFV-NC_040610.1 | -AUUACCACCCG-CUNNNNNNNNA-----CCCCUCAUUAUUUAUAUAACAGCAUAUUGACA | | CCAGG | G-AUAGA | CCGAGAA | AUACACA | AAUUC | UGGG | CACU | AAACA | 195 |
| dISFV-NC_027999.1 | C-----UGGAANNNNNNNNUGGCGAACCCCAUUGGAAAUU--UAGCUUCAUAUUGACA | | CCAGG | GAAAAGA | CCGAGAA | ACUUCU | UGC | UUACU | CGC | AACCA | 199 |



(((((...((((.....))))))....)))..)).))).).)))).)))).)))).))....

| | | | | | | |
|-------------------|---|----------------------------|------------|--------------|----------|-----|
| dISFV-NC_017086.1 | GACUACACGGGACAGCGC | GC CG GAAAAGGUAGUUUGG | --AGGUUGUA | AAACAACAAGU | AUCU--- | 258 |
| dISFV-NC_024017.1 | ACAGCAUUGGACAGAGC | GC CUGAAGAGAGCUGUUGUUAUA | AUUUACA | AAAAAUACA | UAUCU--- | 258 |
| dISFV-NC_016997.1 | CUCCUAAUGGACAGAGC | GC CAUAAGAAACGGAGUUAGU | ---AUGG | AAAAACAUAAGU | AUCU--- | 258 |
| dISFV-NC_040610.1 | ACUAC--AAGCACAGAGC | GC UU CGAGA-AGUAGUUGUUUCU | UGACGAG | AGAAUAUGUAU | AUCUUGG | 258 |
| dISFV-NC_027999.1 | CCGA--AGGCACAGUAU | GC CU GAA-ACA UUCGGUGGUUGA | UGU-AGU | AAACAACCAGU | UUUCU--- | 258 |
| |220.....230.....240.....250.....260.....270..... | | | | | |

