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                                                                  AAACCAGGAAAGUCC-
                                                                                                                                   CUGCAAAG<mark>C</mark>GGACUG<mark>UC</mark>
                                                                                                                                                                                                                     <mark>U</mark>AAAACGCCUGGCUAGC
                                                                                                                                                                                                                                                                                 106
                                                                                                                                  CCGGAAGGCGGGUUGUC
GAAAGAACCGGGUCGUU
GUAAAAACCGGGCUGUC
GCAAGAGCCCGGGCUGUC
GGAAGCCCCGGGUCGUC
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UAAAACGCGGUAUGCCC
UAAAACGCGGAAUGCCC
UAAAACGCGGAAUGCCC
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                                                                                                                                  GGAAGCCCCGGGUCGUC
GAAGAAACCGGGCCAUC
GUAAAAACCGGGUUGUC
GUAGAAACCGGGUUGUC
GGCCAACCCGGGUUGUC
GCUCAGGCCGGGUUGUC
GCUCAGGCCGGGUUGUC
                                                                  AAACCAGGAGGCCC---
                                                                                                                                                                                                                     <mark>U</mark>GAAACGCGGCAUACCC
                                                                                                                                                                                                                                                                                 106
                                                                                                                                                                                                                    UGAAACGCGGCCUACCC
UAAAACGCGGCGUAUCC
                                                                  AAACCAGGAGGCCC
                                                                  AAACCAGGAAAACCC---
                                                                                                                                                                                                                                                                                 106
                                                                                                                                 GGGCAACCCGGGUUGUCAAUA
GGCUCAGGCCGGGUUGUCAAUA
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GCAAUCGUCGGACUGUCAAUA
GAGGAAACCGGGUCGUCAAUA
GGGCAACCCGGGUUGUCAAUA
                                                                                                                                                                                                                    '<mark>U</mark>AAAGCGCGGUGCAUCC
'UAAAGCGCGGAACGUCC
'UAAAGCGCGGUGCGUCC
                                                                  AAACCAGGAAAACCC
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                                                                  AAACCAGGAAGACCC
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                                                                  AAACCAGGAAGGCCC---
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                                                                  AAACCAGGAAAGUCC
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CUAAAGCGCGGCGCAUCC
                                                                  AAACCAGGAAGACCC---
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MBFV-NC_009029.2
MBFV-NC_009026.2
MBFV-NC_018705.3
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MBFV-NC_012533.1

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MBFV-NC_001477.1

MBFV-NC_001477.1

MBFV-NC_002640.1

MBFV-NC_009028.2

MBFV-NC_040776.1

MBFV-NC_034151.1

MBFV-NC_034151.1

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MBFV-NC_009942.1
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                                                                                             CG-CUGGGAA--AGAC AGAGA-UCCUGCUGUCUC---UGCAACAUCAAUCC
CGCCUGGGAA--AGAC CGGGAGAUCCU-CUGCUU---UCCAUCACCAGCCAC
CACCUGGGAA--AGAC CGGGAGACCCU-CUGCUC---UGCAACACCAGCCAC
CACCUGGGAU--AGAC UGGGAGACCCU-CUGCUUA-UUUCAACACCAGCCAC
CACCUGGGA--UAGAC UAGGGGAUCUU-CUGCUCU---GCACAACCAGCCAC
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206
MBFV-NC_001563.2

MBFV-NC_009942.1

MBFV-NC_000943.1

MBFV-NC_006551.1

MBFV-NC_001437.1

MBFV-NC_007580.2

MBFV-NC_012534.1

MBFV-NC_015843.2

MBFV-NC_09029.2

MBFV-NC_032088.1

MBFV-NC_09026.2

MBFV-NC_018705.3
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CACCUGGGA - - UAGAC<mark>U</mark>AGGAGAUCUU - CUGCUCU - - GCACAACCAGCCAC
CACCUGGGAA - AAGAC<mark>U</mark>AGGAGAUCUU - CUGCUCUAUUCCAACAUCAGUCAC
                                                                                                                                                                                                                                                                                 207
                                                                                                                                                                                                                                                                                  205
                                                                                              CACCU<mark>GGGAAUUAGAC<mark>U</mark>AGGAGAUCUU-CUGCUCUAUUCCAACAUCAACCAC</mark>
                                                                                                                                                                                                                                                                                  205
                                                                                             CACCU<mark>G</mark>GGAAU-AGAC<mark>U</mark>GGGAGAUCUU-CUGCUCUAUCUCAACAUCAGCUAC
CACCU<mark>G</mark>GAAAGA----<mark>C</mark>AGGAGAUCCC-CUGCUUU--UUCAACACCAGCCAC
CACCU<mark>G</mark>GGA--GAGAC<mark>U</mark>AGGAGACCCU-CUGUCCUAA--CAACACCAGCCAU
                                                                                                                                                                                                                                                                                  205
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                                                                                             CACCU<mark>G</mark>GGA - UAGACUAGGAGACCCU - CUGUCCUAA - CAACACCAGCCAC
CACCU<mark>G</mark>GGAA - AGACUAGGGGAUUUG - CUGCUC - - - - UGGACUUCCGGCUC
CACCU<mark>G</mark>GGAU - AGAC<mark>U</mark>AGGAGAUCCG - CUGCUC - - - - UGGAACCCCAGUUC
                                                                                                                                                                                                                                                                                  208
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                                                                                      GACACCU<mark>G</mark>GGA - UAGAC<mark>C</mark>GG - AGAAUU - - UGCUGCCUCGCACACCUCCCAC
GUAAACCAGGGGAAGGAC<mark>U</mark>AGAGGUUAGAGGAGACCCCUUGCGAG - UGAACAC
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MBFV-NC_001477.1

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MBFV-NC_002640.1

MBFV-NC_009028.2

MBFV-NC_040776.1
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                                                                  AAAGCACAGAUUGCUU – GAAUGCAGGGCGGGAAGUUGAGAACCAACUAGUGUCU UAAGCACAGAUUGCUU – GAAUGCAGGGUGGGGAAGUUGGGAACCAACUAGUGUCU
                                                                                                                                                                                                                                                                    258
                                                                  A-GGCACAGAUCGCCGA--ACUUCGGCGGCCGGUGUGGG-GAAAUCCAUGGUUUCU
A-GGCACAGAUCGCCGA--AUAGCGGCGGCCGGUGUGGG-GAAAUCCAUGGGUCUU
                                                                                                                                                                                                                                                                    258
                                                                                                                                                                                                                                                                    258
                                                                                                                               -AGAUCGGGGGUUGGUGUGGA-GCAACCCAACAGGACU
                                                                                                                                                                                                                                                                    258
                                                                  U-GGCACAGAGCGCC---
                                                                                                                            --AAAAUGGAAUGGUGCUGU---UGAAUCAACAGGUUCU
--AAAAUGGAAUGGUGCUGU---UGAAUCAACAGGUUCU
                                                                  A-GGCACAGAACGCCAG
                                                                  A-GGCACAGAACGCCAA
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                                                                                                                            --AAGAUGGAUUGGUGUUGU---UGAUCCAACAGGUUCU
                                                                  A-GGCACAGAGCGCCGC
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MBFV-NC_009028.2

MBFV-NC_040776.1

MBFV-NC_034151.1

MBFV-NC_001563.2

MBFV-NC_00942.1

MBFV-NC_000943.1

MBFV-NC_006551.1

MBFV-NC_001437.1

MBFV-NC_007580.2

MBFV-NC_012534.1

MBFV-NC_012534.1

MBFV-NC_015843.2

MBFV-NC_09029.2

MBFV-NC_032088.1

MBFV-NC_032088.1

MBFV-NC_018705.3
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                                                                  UAGGCACAGAUCGCCGC
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                                                                                                                                  AAUCUGUGGCUGGUGCUGCA---AAACACA-GGAUCU
                                                                  UAGGCACAGAUCGCCGA-
                                                                  UCGGCACAGAACGCCGU
                                                                                                                                  AAGU-GUGGCUGGUGGUGAAAAUAAACACA-GAGUCU
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CAAUGGUGGCUGGUGGUGC - GAGAACACA - GGAUCU
                                                                  ACGGCACAGUGCGCCGA
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                                                                  AAGGCACAGAGCGCCGA:
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                                                                  AAGGCACAGAGCGCCGU
                                                                                                                                  AAAGUGUGGCUGGUGGUGAA-AAAAUCACA-GGAUCU
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CCGGCACAGAACGCCGA--CA-UGGUGGGAGGGGUCGUAAGACACCA---GAUUCU
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                                                                  CACACAAAAACAGC---
                                                                                                                            --AUAUUGACACCUGGGAAAGACUAGGAGACCC-UCUGU
                                                                  40......250......260......270......280......290....
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