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                                                                                                  1         10
MMETSP0253-20130528|6017_1   ..................................................ATGGTGACCCC
MMETSP0253-20130528|95269_1  ............................................................
MMETSP0253-20130528|44304_1  ............................................................
MMETSP0253-20130528|1359_1   ............................................................
MMETSP0253-20130528|15654_1  NNCGCGGTCTTCAGTGTCTTCATGGCTCAAGTGGGTAATTTGGCTCTTTTCAGCGGTCCT

                                     20        30        40        50                60
MMETSP0253-20130528|6017_1   GAGGAAACCGACCGTGTTGAAGCGACCAGCCGCCATAGAGACGGC......CGTGGTGAA
MMETSP0253-20130528|95269_1  ............................................................
MMETSP0253-20130528|44304_1  ....................................NTGATGGCGCTGATGAACAAGAGGCC
MMETSP0253-20130528|1359_1   ............................................................
MMETSP0253-20130528|15654_1  CAAACCGCCAAAAGGCATGAAGAGGCCAGCGGCAGTGGCACCAGTGAAAGCCATGAAGAC

                                70        80        90       100       110       120
MMETSP0253-20130528|6017_1   GCAGTCGACGAAAGTGTCCAGGAAGGTCACGACCAAGACGAAGGTGATGAAGGAGAGCAG
MMETSP0253-20130528|95269_1  ...........................................NNTGCGATGAAGGAACACAA
MMETSP0253-20130528|44304_1  TTCGCGTTCGTCCTTGTCGACTGAGAAGAAGAGTATGAAGGTGAGCATGAAGGTCAAGAG
MMETSP0253-20130528|1359_1   .ATGAAGCCGATGAGAGCGGTTGCAGAAGGAATCATGAAGACGAGCATGAAGGCGAAGAA
MMETSP0253-20130528|15654_1  ACTGAAGACCGCGACGCCGAAGAAAGCTATGAAGGTTGGGAAGGCGATGAAGGCGATGAA

                               130       140       150       160       170       180
MMETSP0253-20130528|6017_1   GGTGAGCAAGATCGCCCGTGGAGTTCGGATGCGTGCCGCCGTTTTCAGAGGCCTAAAGGA
MMETSP0253-20130528|95269_1  GGGAAGCAGTGTCGCTCGCGGAGTGCGTATGCGATATGCCGTGTACAGCGGCCGCAAGGA
MMETSP0253-20130528|44304_1  GCCAAGCGTGGTGGCCCGAGGAAGCCGCATGCGCGTTGCCGTCTTCTTGGGTCGAAAGCA
MMETSP0253-20130528|1359_1   AGCTAGCGCCATTGCACGCGGATCCCGCATGCGGCTTGTTGTGTTCCGGGGTCGCAAGGA
MMETSP0253-20130528|15654_1  AGAGAGCAGGATCGCCAGAGGACGTCTTATGCGGGCAGCGGTCCTTTCTGGCA.GGAAGG

                               190       200       210       220       230       240
MMETSP0253-20130528|6017_1   CAAAACGTATACGGGTTTGAAAAGAACCGACCTCAAAATGAACAGGAAAGGTAAGATCGT
MMETSP0253-20130528|95269_1  CAAGACGTACACGGGTTTGAAAAAAAGCGACCTCAAATATAGCAAGAAAGGCAAGATCGT
MMETSP0253-20130528|44304_1  CAAGACGTACACTGGCTTGACCAAGAACGATTTGATGAAAAATGCGGACGGCAGGATTGT
MMETSP0253-20130528|1359_1   AAAGACATACACAGGACTCTCTCGATCCGATTTGAAGCTGAGCAAAAGTGGAAAATTGT
MMETSP0253-20130528|15654_1  AAAAACAACACTGGCCTAAGGGCGTCCGATTTGAAGAGGAACAAGGTTGGGAAATTGT

                               250       260       270       280       290       300
MMETSP0253-20130528|6017_1   GCCGAAGAAAAGGTCTGCACTGGGCAAAAAAACTTCAAGGAAGCAAGCTTGAGGCGTG
MMETSP0253-20130528|95269_1  CAGCAAAAAGATGTCTATTGCAAGCAAGAAACACTTCAAGGCAGCAAGCTCGAAGCGTG
MMETSP0253-20130528|44304_1  GTCGAAGCGCAGTTCGGCGTTGTCCAGGAAACGTTGGGTTGGCAGCAGGTGTCAGGAATG
MMETSP0253-20130528|1359_1   GACAAAAAAAGTTCTGCAGCAGCCAGGAAGCGATGGGAAGGGAGTCGGCCGCAGCAGTG
MMETSP0253-20130528|15654_1  TACGAAAAGGAAGTCGGCGTTGGCCAAGAAGCGTTATGAAGGAAGCAAGCTGCAACAGTG

                               310       320       330       340       350       360
MMETSP0253-20130528|6017_1   GGGGAAAGCTTGCACAGCCGCGCGCAAGGAGCTCAACATCGTTGGTTTTGTCGCCATGGA
MMETSP0253-20130528|95269_1  GGGAAAGCTTGTACCGCCGCGCGCAAGGAGCTCCATCTTGTCGGTTTTGTTGCCATGGA
MMETSP0253-20130528|44304_1  GAACAAGCCCGTCGTGGCCGCAAGGAAGCAACTCAGCCTCGTTGGTTTCGTGGCAGTAAA
MMETSP0253-20130528|1359_1   GATCAAAGCCGTTGTGTCCGCACGAAAGCAGCTCAGCGTCTCCGGGTTCGTTGCGGTGAA
MMETSP0253-20130528|15654_1  GGTCAAAGCTGTGACTGTCGCGAGAAACAGTTGAGCCTCTCTGGTTATGTAGGGTTGAA

                               370       380       390       400       410       420
MMETSP0253-20130528|6017_1   TTCGACACCCC...AGGGACGTGCTTTGTATGCGAAGACAAAGGCCATCTACACAGNN..
MMETSP0253-20130528|95269_1  TTCCACGCTTC...AAGGACGTGCTTTGTATGCGAAGACAAAGGCCATCTACACAGGTAC
MMETSP0253-20130528|44304_1  TTCGCCCTCACCGCAGGGACGAGCATTGTATGTCAAGGCCCGGTCCATCTTCACTGGCGC
MMETSP0253-20130528|1359_1   N...........................................................
MMETSP0253-20130528|15654_1  CTCGCCAACTCCAGAGGGAAGAGCGCTCTACGCCAAGGCGAGAGCGAACTACAACGCCAT

MMETSP0253-20130528|6017_1   .............
MMETSP0253-20130528|95269_1  AGTTTGA......
MMETSP0253-20130528|44304_1  GGCTTGA......
MMETSP0253-20130528|1359_1   .............
MMETSP0253-20130528|15654_1  TCAGAACGCTTGA
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