

Selected Sequences: 1 /Selected Residues: 1679
Deleted Sequences: 0 /Deleted Residues: 0

Gaps Scores: =0= <.001 <.050 <.100 <.150 <.200 <.250 <.350 <.500 <.750 <1.00 =1=
Similarity Scores: =0= <1e-6 <1e-5 <1e-4 <.001 <.010 <.100 <.250 <.500 <.750 <1.00 =1=

10 20 30 40 50 60 70 80 90 100 110 120
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 cctgagccaaggttccgagcggaggagacgcggggggaggatcagcaccgcgagggggatgtcacggtaaaagagcatgggaacgtcggagaaactactcccaagaagcaaaaggaggcttta

Selected Cols:

Gaps Scores:
Similarity Scores:

130 140 150 160 170 180 190 200 210 220 230 240
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 ggaagcggacgagatccccacaacgccggagaaatctctggaaggggaaaagggaagggtggaagaaaaaggggcgggcctcccgatccgaggggcccgaatctccagatctggagagcactc

Selected Cols:

Gaps Scores:
Similarity Scores:

250 260 270 280 290 300 310 320 330 340 350 360
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 cggcccgaagggttgagttagcactcagagggaggaaatccactcggagatgagcagagaaatcacctccagaggacccttcagcgaacaagaggccttcgagcggtaggagtaagacca

Selected Cols:

Gaps Scores:
Similarity Scores:

370 380 390 400 410 420 430 440 450 460 470 480
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 tagcgataggaggagatgctaggagttagggggagaccgaagcaggagggaagcaaaagaaagcaacggggctagccggtgggtgtttccgccccccgagaggggacgagtgaggcttatcc

Selected Cols:

Gaps Scores:
Similarity Scores:

490 500 510 520 530 540 550 560 570 580 590 600
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 cggggaactcgacttatcgtccccatctagcgggacccccggaccctctcgaaagtaccggaaggggtgtctgggaacaccggggaccagtggaagccaagggatgccccctcccgatgctc

Selected Cols:

Gaps Scores:
Similarity Scores:

610 620 630 640 650 660 670 680 690 700 710 720
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 gattccgactccccccccaaagggtcgcccggaatggcgggacccactctgcagggtccggttccatcctttcttacctgatggccggcatggtcccagccctcctcgctgggcggcg

Selected Cols:

Gaps Scores:
Similarity Scores:

730 740 750 760 770 780 790 800 810 820 830 840
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 ctgggcaacattccgaggggaccgtccctcggtaatggcgaaatgggaccacaaaatctctctagattccgatagagaatcgagagaaaagtggctctcccttagccatccgagtggaac

Selected Cols:

Gaps Scores:
Similarity Scores:

850 860 870 880 890 900 910 920 930 940 950 960
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 tggctcctccttcggatgcccagggtcggaccgcgaggaggtggagatgccatgccgaccggaagaggaaaagaggacgcgagacgcaaacctgtgagtggaaacccgctttattcactgg

Selected Cols:

Gaps Scores:
Similarity Scores:

970 980 990 1000 1010 1020 1030 1040 1050 1060 1070 1080
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 ggtcgacaactctggggagaaaaaggcgatcggtcgggaagagtatatccctatggaaatccctgggtttccctgatgtccagccccctccccgggtccgagagaagggggactccgggact

Selected Cols:

Gaps Scores:
Similarity Scores:

1090 1100 1110 1120 1130 1140 1150 1160 1170 1180 1190 1200
=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+=====+
M21012.1 cccctgcagattggggacgaagccgccccgggcgtccccctcgatccaccttcgaaggggttcacaccccccaaccggcgggccggctactcttctttcccttctctcgtcttctcggctc

Selected Cols:

Gaps Scores:
Similarity Scores:

