



610 620 630 640 650 660 670  
GTCAGATCTCTAGAAGCTGGGTACCAGCTGCTAGCAAGCTTGCTAGCGGCCGCTCGAGGCCGCAAGGCCGGATC  
G0184540-1.seq(1>10190) → GTCAGATCTCTAGAAGCTGGGTACCAGCTGCTAGCAAGCTTGCTAGCGGCCGCTCGAGGCCGCAAGGCCGGATC  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → CTCTAGAAGCTGGGTACCAGCTGCTAGCAAGCTTGCTAGCGGCCGCTCGAGGCCGCAAGGCCGGATC

680 690 700 710 720 730 740 750  
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G0184540-1.seq(1>10190) → CAGACATGATAAGATAACATTGATGAGTTTGGACAAACCACAACCTAGAATGCAGTGAAAAAATGCTTTATTTGTG  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → CAGACATGATAAGATAACATTGATGAGTTTGGACAAACCACAACCTAGAATGCAGTGAAAAAATGCTTTATTTGTG

760 770 780 790 800 810 820  
AAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATTC  
G0184540-1.seq(1>10190) → AAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATTC  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → AAATTTGTGATGCTATTGCTTTATTTGTAACCATTATAAGCTGCAATAAACAAGTTAACAACAACAATTGCATTC

830 840 850 860 870 880 890 900  
ATTTTATGTTTCAGGTTTCAGGGGGAGGTGTGGGAGGTTTTTTTAAAGCAAGTAAAACCTCTACAAATGTGGTATGG  
G0184540-1.seq(1>10190) → ATTTTATGTTTCAGGTTTCAGGGGGAGGTGTGGGAGGTTTTTTTAAAGCAAGTAAAACCTCTACAAATGTGGTATGG  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → ATTTTATGTTTCAGGTTTCAGGGGGAGGTGTGGGAGGTTTTTTTAAAGCAAGTAAAACCTCTACAAATGTGGTATGG

910 920 930 940 950 960 970  
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G0184540-1.seq(1>10190) → CTGATTATGATCCGGCTGCCTCGCGCGTTTCGGTGATGACGGTGAAAACCTCTGACACATGCAGCTCCCGGAGAC  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → CTGATTATGATCCGGCTGCCTCGCGCGTTTCGGTGATGACGGTGAAAACCTCTGACACATGCAGCTCCCGGAGAC

980 990 1000 1010 1020 1030 1040 1050  
GGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCCTCAGGGCGCGTCAGCGGGTGTGGCGGGTG  
G0184540-1.seq(1>10190) → GGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCCTCAGGGCGCGTCAGCGGGTGTGGCGGGTG  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → GGTCACAGCTTGTCTGTAAGCGGATGCCGGGAGCAGACAAGCCCCTCAGGGCGCGTCAGCGGGTGTGGCGGGTG

1060 1070 1080 1090 1100 1110 1120  
TCGGGGCGCAGCCATGAGGTCGACTCTAGAGGATCGATCCCCGCCCGGACGAACTAAACCTGACTACGACATCT  
G0184540-1.seq(1>10190) → TCGGGGCGCAGCCATGAGGTCGACTCTAGAGGATCGATCCCCGCCCGGACGAACTAAACCTGACTACGACATCT  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → TCGGGGCGCAGCCATGAGGTCGACTCTAGAGGATCGATCCCCGCCCGGACGAACTAAACCTGACTACGACATCT

1130 1140 1150 1160 1170 1180 1190 1200  
CTGCCCCCTTCTTCGCGGGGCAGTGCATGTAATCCCTTCAGTTGGTTGGTACAACCTTGCCAACCTGGGCCCTGTTCC  
G0184540-1.seq(1>10190) → CTGCCCCCTTCTTCGCGGGGCAGTGCATGTAATCCCTTCAGTTGGTTGGTACAACCTTGCCAACCTGGGCCCTGTTCC  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → CTGCCCCCTTCTTCGCGGGGCAGTGCATGTAATCCCTTCAGTTGGTTGGTACAACCTTGCCAACCTGGGCCCTGTTCC

1210 1220 1230 1240 1250 1260 1270  
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G0184540-1.seq(1>10190) → ACATGTGACACGGGGGGGACCAAACACAAAGGGGTTCTCTGACTGTAGTTGACATCCTTATAAATGGATGTGCA  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → ACATGTGACACGGGGGGGACCAAACACAAAGGGGTTCTCTGACTGTAGTTGACATCCTTATAAATGGATGTGCA

1280 1290 1300 1310 1320 1330 1340 1350  
CATTTGCCAACACTGAGTGGCTTTCATCCTGGAGCAGACTTTGCAGTCTGTGGACTGCAACACAACATTGCCTTT  
G0184540-1.seq(1>10190) → CATTTGCCAACACTGAGTGGCTTTCATCCTGGAGCAGACTTTGCAGTCTGTGGACTGCAACACAACATTGCCTTT  
E09-E04901-G0184540-1\_CMV-F.ab1(1>675) → CATTTGC

1360 1370 1380 1390 1400 1410 1420  
ATGTGTAACCTCTTGGCTGAAGCTCTTACACCAATGCTGGGGGACATGTACCTCCCAGGGGCCAGGAAGACTACG  
G0184540-1.seq(1>10190) → ATGTGTAACCTCTTGGCTGAAGCTCTTACACCAATGCTGGGGGACATGTACCTCCCAGGGGCCAGGAAGACTACG

1430 1440 1450 1460 1470 1480 1490 1500  
GGAGGCTACACCAACGTCAATCAGAGGGGCCTGTGTAGCTACCGATAAGCGGACCCTCAAGAGGGCATTAGCAAT  
G0184540-1.seq(1>10190) → GGAGGCTACACCAACGTCAATCAGAGGGGCCTGTGTAGCTACCGATAAGCGGACCCTCAAGAGGGCATTAGCAAT

1510 1520 1530 1540 1550 1560 1570  
AGTGTTTATAAGGCCCCCTTGTAAACCCTAAACGGGTAGCATATGCTTCCCGGGTAGTAGTATATACTATCCAGA  
G0184540-1.seq(1>10190) → AGTGTTTATAAGGCCCCCTTGTAAACCCTAAACGGGTAGCATATGCTTCCCGGGTAGTAGTATATACTATCCAGA

1580 1590 1600 1610 1620 1630 1640 1650  
CTAACCTAATTCAATAGCATATGTTACCCAACGGGAAGCATATGCTATCGAATTAGGGTTAGTAAAAGGGTCCT  
G0184540-1.seq(1>10190) → CTAACCTAATTCAATAGCATATGTTACCCAACGGGAAGCATATGCTATCGAATTAGGGTTAGTAAAAGGGTCCT

1660 1670 1680 1690 1700 1710 1720  
AAGGAACAGCGATATCTCCACCCCATGAGCTGTCACGGTTTTATTACATGGGGTCAGGATTCCACGAGGGTAG  
G0184540-1.seq(1>10190) → AAGGAACAGCGATATCTCCACCCCATGAGCTGTCACGGTTTTATTACATGGGGTCAGGATTCCACGAGGGTAG





2930 2940 2950 2960 2970 2980 2990 3000  
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G0184540-1.seq(1>10190) → GTAGCATATACTACCCAAATATCTGGATAGCATATGCTATCCTAATCTATATCTGGGTAGCATATGCTATCCTAA

3010 3020 3030 3040 3050 3060 3070  
TCTATATCTGGGTAGCATAGGCTATCCTAATCTATATCTGGGTAGCATATGCTATCCTAATCTATATCTGGGTAG  
G0184540-1.seq(1>10190) → TCTATATCTGGGTAGCATAGGCTATCCTAATCTATATCTGGGTAGCATATGCTATCCTAATCTATATCTGGGTAG

3080 3090 3100 3110 3120 3130 3140 3150  
TATATGCTATCCTAATTTATATCTGGGTAGCATAGGCTATCCTAATCTATATCTGGGTAGCATATGCTATCCTAA  
G0184540-1.seq(1>10190) → TATATGCTATCCTAATTTATATCTGGGTAGCATAGGCTATCCTAATCTATATCTGGGTAGCATATGCTATCCTAA

3160 3170 3180 3190 3200 3210 3220  
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G0184540-1.seq(1>10190) → TCTATATCTGGGTAGTATATGCTATCCTAATCTGTATCCGGGTAGCATATGCTATCCTCATGCATATACAGTCAG

3230 3240 3250 3260 3270 3280 3290 3300  
CATATGATACCCAGTAGTAGAGTGGGAGTGCTATCCTTTGCATATGCCGCCACCTCCCAAGGGGGCGTGAATTTT  
G0184540-1.seq(1>10190) → CATATGATACCCAGTAGTAGAGTGGGAGTGCTATCCTTTGCATATGCCGCCACCTCCCAAGGGGGCGTGAATTTT

3310 3320 3330 3340 3350 3360 3370  
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G0184540-1.seq(1>10190) → CGCTGCTTGTCTTTTCTGCTGGTTGCTCCCATTCTTAGGTGAATTTAAGGAGGCCAGGCTAAAGCCGTCGCAT

3380 3390 3400 3410 3420 3430 3440 3450  
GTCTGATTGCTCACCAGGTAAATGTCGCTAATGTTTTCCAACGCGAGAAGGTGTTGAGCGCGGAGCTGAGTGACG  
G0184540-1.seq(1>10190) → GTCTGATTGCTCACCAGGTAAATGTCGCTAATGTTTTCCAACGCGAGAAGGTGTTGAGCGCGGAGCTGAGTGACG

3460 3470 3480 3490 3500 3510 3520  
TGACAACATGGGTATGCCCAATTGCCCATGTTGGGAGGACGAAAATGGTGACAAGACAGATGGCCAGAAATACA  
G0184540-1.seq(1>10190) → TGACAACATGGGTATGCCCAATTGCCCATGTTGGGAGGACGAAAATGGTGACAAGACAGATGGCCAGAAATACA





4130 4140 4150 4160 4170 4180 4190 4200  
GGAACCTCCTTGACCACGATGCTTTCCAAACCACCCTCCTTTTTTGGCGCTGCCTCCATCACCCCTGACCCCGGGG  
G0184540-1.seq(1>10190) → GGAACCTCCTTGACCACGATGCTTTCCAAACCACCCTCCTTTTTTGGCGCTGCCTCCATCACCCCTGACCCCGGGG

4210 4220 4230 4240 4250 4260 4270  
TCCAGTGCTTGGGCCTTCTCCTGGGTCATCTGCGGGGCCCTGCTCTATCGCTCCCGGGGGCACGTCAGGCTCACC  
G0184540-1.seq(1>10190) → TCCAGTGCTTGGGCCTTCTCCTGGGTCATCTGCGGGGCCCTGCTCTATCGCTCCCGGGGGCACGTCAGGCTCACC

4280 4290 4300 4310 4320 4330 4340 4350  
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G0184540-1.seq(1>10190) → ATCTGGGCCACCTTCTTGGTGGTATTCAAATAATCGGCTTCCCCTACAGGGTGGAAAAATGGCCTTCTACCTGG

4360 4370 4380 4390 4400 4410 4420  
AGGGGGCCTGCGCGGTGGAGACCCGGATGATGATGACTGACTACTGGGACTCCTGGGCCTCTTTTCTCCACGTCC  
G0184540-1.seq(1>10190) → AGGGGGCCTGCGCGGTGGAGACCCGGATGATGATGACTGACTACTGGGACTCCTGGGCCTCTTTTCTCCACGTCC

4430 4440 4450 4460 4470 4480 4490 4500  
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G0184540-1.seq(1>10190) → ACGACCTCTCCCCCTGGCTCTTTTACAGACTTCCCCCCTGGCTCTTTTACAGTCTCTACCCCGGCGGCCTCCACT

4510 4520 4530 4540 4550 4560 4570  
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G0184540-1.seq(1>10190) → ACCTCCTCGACCCCGGCCTCCACTACCTCCTCGACCCCGGCCTCCACTGCCTCCTCGACCCCGGCCTCCACTCC

4580 4590 4600 4610 4620 4630 4640 4650  
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G0184540-1.seq(1>10190) → TGCTCCTGCCCCTCCTGCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCCTGC

4660 4670 4680 4690 4700 4710 4720  
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G0184540-1.seq(1>10190) → CCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTGC



## Project: Untitled.SQD Contig 1

4730 4740 4750 4760 4770 4780 4790 4800  
CCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCCTGC  
G0184540-1.seq(1>10190) → CCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCCTGC

4810 4820 4830 4840 4850 4860 4870  
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G0184540-1.seq(1>10190) → CCCTCCTGCTCCTGCCCTCCTGCTCCTGCCCTCCTGCTCCTGCCCTCCTGCTCCTGCCCTCCTGCCCTCC

4880 4890 4900 4910 4920 4930 4940 4950  
TGCCCCCTCCTCCTGCTCCTGCCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCC  
G0184540-1.seq(1>10190) → TGCCCCCTCCTCCTGCTCCTGCCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCCCTCCTGCTCCTGCCCTCC

4960 4970 4980 4990 5000 5010 5020  
TCCTGCTCCTGCCCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGC  
G0184540-1.seq(1>10190) → TCCTGCTCCTGCCCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGC

5030 5040 5050 5060 5070 5080 5090 5100  
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G0184540-1.seq(1>10190) → CCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTCCTGC

5110 5120 5130 5140 5150 5160 5170  
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G0184540-1.seq(1>10190) → TCCTGCCCTCCTCCTGCTCCTGCCCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTCCTGCTCC

5180 5190 5200 5210 5220 5230 5240 5250  
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G0184540-1.seq(1>10190) → TGCCCCCTCCTGCCCTCCTGCCCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTCCTGCTCCTGCCCTCCTGC

5260 5270 5280 5290 5300 5310 5320  
TCCTGCCCTCCCGCTCCTGCTCCTGCTCCTGTTCCACCGTGGGTCCCTTTGCAGCCAATGCAACTTGGACGTTT  
G0184540-1.seq(1>10190) → TCCTGCCCTCCCGCTCCTGCTCCTGCTCCTGTTCCACCGTGGGTCCCTTTGCAGCCAATGCAACTTGGACGTTT





6530 6540 6550 6560 6570 6580 6590 6600  
AGAATTATGCAGTGCTGCCATAACCATGAGTGATAAACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACC  
G0184540-1.seq(1>10190) → AGAATTATGCAGTGCTGCCATAACCATGAGTGATAAACTGCGGCCAACTTACTTCTGACAACGATCGGAGGACC

6610 6620 6630 6640 6650 6660 6670  
GAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACCTCGCCTTGATCGTTGGGAACCGGAGCTGAA  
G0184540-1.seq(1>10190) → GAAGGAGCTAACCGCTTTTTTGCACAACATGGGGGATCATGTAACCTCGCCTTGATCGTTGGGAACCGGAGCTGAA

6680 6690 6700 6710 6720 6730 6740 6750  
TGAAGCCATAACCAAACGACGAGCGTGACACCACGATGCCTGCAGCAATGGCAACAACGTTGCGCAAACCTATTAAC  
G0184540-1.seq(1>10190) → TGAAGCCATAACCAAACGACGAGCGTGACACCACGATGCCTGCAGCAATGGCAACAACGTTGCGCAAACCTATTAAC

6760 6770 6780 6790 6800 6810 6820  
TGGCGAACTACTTACTCTAGCTTCCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAAGTTGCAGGACCACT  
G0184540-1.seq(1>10190) → TGGCGAACTACTTACTCTAGCTTCCCGGCAACAATTAATAGACTGGATGGAGGCGGATAAAAGTTGCAGGACCACT

6830 6840 6850 6860 6870 6880 6890 6900  
TCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGCGGTAT  
G0184540-1.seq(1>10190) → TCTGCGCTCGGCCCTTCCGGCTGGCTGGTTTATTGCTGATAAATCTGGAGCCGGTGAGCGTGGGTCTCGCGGTAT

6910 6920 6930 6940 6950 6960 6970  
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G0184540-1.seq(1>10190) → CATTGCAGCACTGGGGCCAGATGGTAAGCCCTCCCGTATCGTAGTTATCTACACGACGGGGAGTCAGGCAACTAT

6980 6990 7000 7010 7020 7030 7040 7050  
GGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGACCAAGTTTA  
G0184540-1.seq(1>10190) → GGATGAACGAAATAGACAGATCGCTGAGATAGGTGCCTCACTGATTAAGCATTGGTAACTGTCAGACCAAGTTTA

7060 7070 7080 7090 7100 7110 7120  
CTCATATATACTTTAGATTGATTTAAAACCTTCATTTTTAATTTAAAAGGATCTAGGTGAAGATCCTTTTTGATAA  
G0184540-1.seq(1>10190) → CTCATATATACTTTAGATTGATTTAAAACCTTCATTTTTAATTTAAAAGGATCTAGGTGAAGATCCTTTTTGATAA





8330 8340 8350 8360 8370 8380 8390 8400  
GTGTGGCCTCGAACACCGAGCGACCCTGCAGCGACCCGCTTAACAGCGTCAACAGCGTGCCGCAGATCCCGGGG  
G0184540-1.seq(1>10190) → GTGTGGCCTCGAACACCGAGCGACCCTGCAGCGACCCGCTTAACAGCGTCAACAGCGTGCCGCAGATCCCGGGG

8410 8420 8430 8440 8450 8460 8470  
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G0184540-1.seq(1>10190) → GCAATGAGATATGAAAAGCCTGAACTCACCGCGACGTCTGTGCGAGAAGTTTCTGATCGAAAAGTTCGACAGCGT

8480 8490 8500 8510 8520 8530 8540 8550  
CTCCGACCTGATGCAGCTCTCGGAGGGCGAAGAATCTCGTGCTTTCAGCTTCGATGTAGGAGGGCGTGGATATGT  
G0184540-1.seq(1>10190) → CTCCGACCTGATGCAGCTCTCGGAGGGCGAAGAATCTCGTGCTTTCAGCTTCGATGTAGGAGGGCGTGGATATGT

8560 8570 8580 8590 8600 8610 8620  
CCTGCGGGTAAATAGCTGCGCCGATGGTTTCTACAAAGATCGTTATGTTTATCGGCACTTTGCATCGGCCGCGCT  
G0184540-1.seq(1>10190) → CCTGCGGGTAAATAGCTGCGCCGATGGTTTCTACAAAGATCGTTATGTTTATCGGCACTTTGCATCGGCCGCGCT

8630 8640 8650 8660 8670 8680 8690 8700  
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G0184540-1.seq(1>10190) → CCCGATTCCGGAAGTGCTTGACATTGGGGAATTCAGCGAGAGCCTGACCTATTGCATCTCCCGCCGTGCACAGGG

8710 8720 8730 8740 8750 8760 8770  
TGTCACGTTGCAAGACCTGCCTGAAACCGAACTGCCCGCTGTTCTGCAGCCGGTCGCGGAGGCCATGGATGCGAT  
G0184540-1.seq(1>10190) → TGTCACGTTGCAAGACCTGCCTGAAACCGAACTGCCCGCTGTTCTGCAGCCGGTCGCGGAGGCCATGGATGCGAT

8780 8790 8800 8810 8820 8830 8840 8850  
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G0184540-1.seq(1>10190) → CGCTGCGGCCGATCTTAGCCAGACGAGCGGGTTCGGCCCATTCGGACCGCAAGGAATCGGTCAATACACTACATG

8860 8870 8880 8890 8900 8910 8920  
GCGTGATTTTCATATGCGCGATTGCTGATCCCATGTGTATCACTGGCAAACCTGTGATGGACGACACCGTCAAGTGC  
G0184540-1.seq(1>10190) → GCGTGATTTTCATATGCGCGATTGCTGATCCCATGTGTATCACTGGCAAACCTGTGATGGACGACACCGTCAAGTGC



8930 8940 8950 8960 8970 8980 8990 9000  
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G0184540-1.seq(1>10190) → GTCCGTCGCGCAGGCTCTCGATGAGCTGATGCTTTGGGCCGAGGACTGCCCCGAAGTCCGGCACCTCGTGCACGC

9010 9020 9030 9040 9050 9060 9070  
GGATTTCCGGCTCCAACAATGTCCTGACGGACAATGGCCGCATAACAGCGGTCATTGACTGGAGCGAGGCGATGTT  
G0184540-1.seq(1>10190) → GGATTTCCGGCTCCAACAATGTCCTGACGGACAATGGCCGCATAACAGCGGTCATTGACTGGAGCGAGGCGATGTT

9080 9090 9100 9110 9120 9130 9140 9150  
CGGGGATTCCCAATACGAGGTCGCCAACATCTTCTTCTGGAGGCCGTGGTTGGCTTGTATGGAGCAGCAGACGCG  
G0184540-1.seq(1>10190) → CGGGGATTCCCAATACGAGGTCGCCAACATCTTCTTCTGGAGGCCGTGGTTGGCTTGTATGGAGCAGCAGACGCG

9160 9170 9180 9190 9200 9210 9220  
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G0184540-1.seq(1>10190) → CTACTTCGAGCGGAGGCATCCGGAGCTTGCAGGATCGCCGCGGCTCCGGGCGTATATGCTCCGCATTGGTCTTGA

9230 9240 9250 9260 9270 9280 9290 9300  
CCAACTCTATCAGAGCTTGGTTGACGGCAATTTTCGATGATGCAGCTTGGGCGCAGGGTTCGATGCGACGCAATCGT  
G0184540-1.seq(1>10190) → CCAACTCTATCAGAGCTTGGTTGACGGCAATTTTCGATGATGCAGCTTGGGCGCAGGGTTCGATGCGACGCAATCGT

9310 9320 9330 9340 9350 9360 9370  
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G0184540-1.seq(1>10190) → CCGATCCGGAGCCGGGACTGTCGGGCGTACACAAATCGCCCGCAGAAGCGCGGCCGTCTGGACCGATGGCTGTGT

9380 9390 9400 9410 9420 9430 9440 9450  
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G0184540-1.seq(1>10190) → AGAAGTACTCGCCGATAGTGGA AACCGACGCCCCAGCACTCGTCCGGATCGGGAGATGGGGGAGGCTAACTGAAA

9460 9470 9480 9490 9500 9510 9520  
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G0184540-1.seq(1>10190) → CACGGAAGGAGACAATAACCGGAAGGAACCCGCGCTATGACGGCAATAAAAAGACAGAATAAAACGCACGGGTGTT



10130 10140 10150 10160 10170 10180  
ATACGTTGTATCTATATCATAATATGTACATTTATATTGGCTCATGTCCAATATGACCGCCAT  
G0184540-1.seq(1>10190) → ATACGTTGTATCTATATCATAATATGTACATTTATATTGGCTCATGTCCAATATGACCGCCAT

