

Aligned using an external algorithm

Consensus

1. First analysis after plaque purification
2. Analysis after seven cycles of amplification

1	1	CATCATCAATAATACCTTATTTGGATTGAAGCCAATATGATAATGAGGGGTGGAGT	60
2	1	CATCATCAATAATACCTTATTTGGATTGAAGCCAATATGATAATGAGGGGTGGAGT	60
1	61	TTGTGACGTGGCGCGGGCGTGGAACGGGCAGGGTGACGTAGTAGTGTGGCGGAAGTGT	120
2	61	TTGTGACGTGGCGCGGGCGTGGAACGGGCAGGGTGACGTAGTAGTGTGGCGGAAGTGT	120
1	121	GATGTTGCAAGTGTGGCGAACACATGTAAGCGACGGATGTGGCAAAAGTGACGTTTG	180
2	121	GATGTTGCAAGTGTGGCGAACACATGTAAGCGACGGATGTGGCAAAAGTGACGTTTG	180
1	181	GTGTGCGCCGGTGTACACAGGAAGTGACAATTTCGCGCGGTTAGGCAGTGTGAG	240
2	181	GTGTGCGCCGGTGTACACAGGAAGTGACAATTTCGCGCGGTTAGGCAGTGTGAG	240
1	241	TAAATTGGCGTAACCGAGTAAGATTGCCATTTCGCGGGAAACTGAATAAGAGGA	300
2	241	TAAATTGGCGTAACCGAGTAAGATTGCCATTTCGCGGGAAACTGAATAAGAGGA	300
1	301	AGTGAAATCTGAATAATTGTGTTACTCATAGCGCGTAATATTGCTAGGGCCGCGGG	360
2	301	AGTGAAATCTGAATAATTGTGTTACTCATAGCGCGTAATATTGCTAGGGCCGCGGG	360
1	361	GACTTTGACCGTTACGTGGAGACTCGCCCAGGTGTTTCAGGTGTTCCCGCGTTC	420
2	361	GACTTTGACCGTTACGTGGAGACTCGCCCAGGTGTTTCAGGTGTTCCCGCGTTC	420
1	421	CGGGTCAAAGTGGCGTTATTATAGTCAGCTGACGTAGTAGTGTATTACCCGG	480
2	421	CGGGTCAAAGTGGCGTTATTATAGTCAGCTGACGTAGTAGTGTATTACCCGG	480
1	481	TGAGTTCTCAAGAGGCCACTCTTGAGTGCCAGCGAGTAGAGTTCTCCTCCGAGCCGC	540
2	481	TGAGTTCTCAAGAGGCCACTCTTGAGTGCCAGCGAGTAGAGTTCTCCTCCGAGCCGC	540

1 541 TCCGACACCAGGGACTGAAAATGAGACATATTATGCCACGGAGGTGTTATTACCGAAGA 600
2 541 TCCGACACCAGGGACTGAAAATGAGACATATTATGCCACGGAGGTGTTATTACCGAAGA 600

1 601 AATGGCCGCCAGTCTTGGACCAGCTGATCGAAGAGGTACTGGCTGATAATCTTCCACC 660
2 601 AATGGCCGCCAGTCTTGGACCAGCTGATCGAAGAGGTACTGGCTGATAATCTTCCACC 660

1 661 TCCTAGCCATTTGAACCACCTACCCCTCACGAACGTATGATTAGACGTGACGGCCCC 720
2 661 TCCTAGCCATTTGAACCACCTACCCCTCACGAACGTATGATTAGACGTGACGGCCCC 720

1 721 CGAAGATCCAACGAGGAGGC GGTT CGCAGATTTCCC ACTCTGTAATGTTGGCGGT 780
2 721 CGAAGATCCAACGAGGAGGC GGTT CGCAGATTTCCC ACTCTGTAATGTTGGCGGT 780

1 781 GCAGGAAGGGATTGACTTACTCACTTTCCGCCGGCGCCGGTTCTCCGGAGCCGCTCA 840
2 781 GCAGGAAGGGATTGACTTACTCACTTTCCGCCGGCGCCGGTTCTCCGGAGCCGCTCA 840

1 841 CCTTCCCGCAGCCCAGCAGCCGGAGCAGAGAGCCTTGGTCCGGTTCTATGCCAAA 900
2 841 CCTTCCCGCAGCCCAGCAGCCGGAGCAGAGAGCCTTGGTCCGGTTCTATGCCAAA 900

1 901 CCTTGTACCGGAGGTGATCGATCCACCCAGTGACGACGAGGATGAAGAGGGTGAGGAGTT 960
2 901 CCTTGTACCGGAGGTGATCGATCCACCCAGTGACGACGAGGATGAAGAGGGTGAGGAGTT 960

1 961 TGTGTTAGATTATGTGGAGCACCCGGCACGGTTGCAGGTCTTGTCAATTACCGGAG 1020
2 961 TGTGTTAGATTATGTGGAGCACCCGGCACGGTTGCAGGTCTTGTCAATTACCGGAG 1020

1 1021 GAATACGGGGACCCAGATATTATGTGTTCGCTTGCTATATGAGGACCTGTGGCATGTT 1080
2 1021 GAATACGGGGACCCAGATATTATGTGTTCGCTTGCTATATGAGGACCTGTGGCATGTT 1080

1 1081 TGTCTACAGTAAGTGAAAATTATGGCAGTGGTGATAGAGTGGTGGTTGGTGTGGTA 1140
2 1081 TGTCTACAGTAAGTGAAAATTATGGCAGTGGTGATAGAGTGGTGGTTGGTGTGGTA 1140

1	1141	ATTTTTTTTAATTTACAGTTGTGGTTAAAGAATTGTATTGTGATTTTTA	1200
2	1141	ATTTTTTTTAATTTACAGTTGTGGTTAAAGAATTGTATTGTGATTTTTA	1200
1	1201	AAAGGTCTGTCTGAACCTGAGCCTGAGCCCAGCCAGAACCGGAGCCTGCAAGACCT	1260
2	1201	AAAGGTCTGTCTGAACCTGAGCCTGAGCCCAGCCAGAACCGGAGCCTGCAAGACCT	1260
1	1261	ACCCGCCGTCTAAAATGGCGCCTGCTATCCTGAGACGCCGACATCACCTGTGTCTAGA	1320
2	1261	ACCCGCCGTCTAAAATGGCGCCTGCTATCCTGAGACGCCGACATCACCTGTGTCTAGA	1320
1	1321	GAATGCAATAGTAGTACGGATAGCTGTGACTCCGGTCCTTCTAACACACCTCCTGAGATA	1380
2	1321	GAATGCAATAGTAGTACGGATAGCTGTGACTCCGGTCCTTCTAACACACCTCCTGAGATA	1380
1	1381	CACCCGGTGGTCCCGCTGTGCCCATTAACCAACAGTTGCCGTGAGAGTTGGTGGCGTCGC	1440
2	1381	CACCCGGTGGTCCCGCTGTGCCCATTAACCAACAGTTGCCGTGAGAGTTGGTGGCGTCGC	1440
1	1441	CAGGCTGTGGAATGTATCGAGGACTTGCTTAACGAGCCTGGCAACCTTGGACTTGAGC	1500
2	1441	CAGGCTGTGGAATGTATCGAGGACTTGCTTAACGAGCCTGGCAACCTTGGACTTGAGC	1500
1	1501	TGTAAACGCCAGGCCATAAGGTGTAAACCTGTGATTGCGTGTGGTTACGCCTTG	1560
2	1501	TGTAAACGCCAGGCCATAAGGTGTAAACCTGTGATTGCGTGTGGTTACGCCTTG	1560
1	1561	TTTGCTGAATGAGTTGATGTAAGTTAATAAAGGGTAGAGATAATGTTAACGGCATGGC	1620
2	1561	TTTGCTGAATGAGTTGATGTAAGTTAATAAAGGGTAGAGATAATGTTAACGGCATGGC	1620
1	1621	GTGTTAAATGGGGCGGGGCTTAAGGGTATATAATGCGCCGTGGCTAATCTGGTTACA	1680
2	1621	GTGTTAAATGGGGCGGGGCTTAAGGGTATATAATGCGCCGTGGCTAATCTGGTTACA	1680
1	1681	TCTGACCTCATGGAGGCTTGGAGTGTGTTGGAAGATTTCTGCTGTGCGTAACCTGCTG	1740
2	1681	TCTGACCTCATGGAGGCTTGGAGTGTGTTGGAAGATTTCTGCTGTGCGTAACCTGCTG	1740

1	1741	GAACAGAGCTAACAGTACCTCTGGTTGGAGGTTCTGTGGGCTCATCCCAGGCA	1800
2	1741	GAACAGAGCTAACAGTACCTCTGGTTGGAGGTTCTGTGGGCTCATCCCAGGCA	1800
1	1801	AAGTTAGTCTGCAGAATTAAAGGAGGATTACAAGTGGAAATTGAAGAGCTTTGAAATCC	1860
2	1801	AAGTTAGTCTGCAGAATTAAAGGAGGATTACAAGTGGAAATTGAAGAGCTTTGAAATCC	1860
1	1861	TGTGGTGAGCTGTTGATTCTTGAATCTGGTCACCAGGCCTTTCCAAGAGAAGGTC	1920
2	1861	TGTGGTGAGCTGTTGATTCTTGAATCTGGTCACCAGGCCTTTCCAAGAGAAGGTC	1920
1	1921	ATCAAGACTTGGATTTCCACACCGGGCGCGCTGCGGCTGCTGTTGCTTTTGAGT	1980
2	1921	ATCAAGACTTGGATTTCCACACCGGGCGCGCTGCGGCTGCTGTTGCTTTTGAGT	1980
1	1981	TTTATAAAGGATAATGGAGCGAAGAAACCCATCTGAGCGGGGGTACCTGCTGGATTT	2040
2	1981	TTTATAAAGGATAATGGAGCGAAGAAACCCATCTGAGCGGGGGTACCTGCTGGATTT	2040
1	2041	CTGGCCATGCATCTGTGGAGAGCGGTTGTGAGACACAAGAACGCTGCTACTGTTGTCT	2100
2	2041	CTGGCCATGCATCTGTGGAGAGCGGTTGTGAGACACAAGAACGCTGCTACTGTTGTCT	2100
1	2101	TCCGTCGCCGGCGATAATACCGACGGAGGAGCAGCAGCAGCAGGAGGAAGCCAGG	2160
2	2101	TCCGTCGCCGGCGATAATACCGACGGAGGAGCAGCAGCAGCAGGAGGAAGCCAGG	2160
1	2161	CGGC GGCGGCAGGAGCAGAGCCATGGAACCGAGAGCCGGCTGGACCCCTCGGAATGA	2220
2	2161	CGGC GGCGGCAGGAGCAGAGCCATGGAACCGAGAGCCGGCTGGACCCCTCGGAATGA	2220
1	2221	ATGTTGTACAGGTGGCTGAACGTATCCAGAACTGAGACGCATTTGACAATTACAGAGG	2280
2	2221	ATGTTGTACAGGTGGCTGAACGTATCCAGAACTGAGACGCATTTGACAATTACAGAGG	2280
1	2281	ATGGGCAGGGGCTAAAGGGGTAAGAGGGAGCGGGGGCTTGTGAGGCTACAGAGGAGG	2340
2	2281	ATGGGCAGGGGCTAAAGGGGTAAGAGGGAGCGGGGGCTTGTGAGGCTACAGAGGAGG	2340

1	2341	CTAGGAATCTAGCTTTAGCTTAATGACCAGACACCGTCCTGAGTGTATTACTTTCAAC	2400
2	2341	CTAGGAATCTAGCTTTAGCTTAATGACCAGACACCGTCCTGAGTGTATTACTTTCAAC	2400
1	2401	AGATCAAGGATAATTGCCTAATGAGCTTGATCTGCTGGCGAGAAGTATTCCATAGAGC	2460
2	2401	AGATCAAGGATAATTGCCTAATGAGCTTGATCTGCTGGCGAGAAGTATTCCATAGAGC	2460
1	2461	AGCTGACCACTTACTGGCTGCAGCCAGGGATGATTTGAGGAGGCTATTAGGGTATATG	2520
2	2461	AGCTGACCACTTACTGGCTGCAGCCAGGGATGATTTGAGGAGGCTATTAGGGTATATG	2520
1	2521	CAAAGGTGGCACTTAGGCCAGATTGCAAGTACAAGATCAGCAAACCTTGTAAATATCAGGA	2580
2	2521	CAAAGGTGGCACTTAGGCCAGATTGCAAGTACAAGATCAGCAAACCTTGTAAATATCAGGA	2580
1	2581	ATTGTTGCTACATTCGGAACGGGCCGAGGTGGAGATAGATACTGGAGGATAGGGTGG	2640
2	2581	ATTGTTGCTACATTCGGAACGGGCCGAGGTGGAGATAGATACTGGAGGATAGGGTGG	2640
1	2641	CCTTAGATGTAGCATGATAAATATGTGGCGGGGTGCTGGCATGGACGGGTGGTTA	2700
2	2641	CCTTAGATGTAGCATGATAAATATGTGGCGGGGTGCTGGCATGGACGGGTGGTTA	2700
1	2701	TTATGAATGTAAGGTTACTGGCCCCAATTTAGCGGTACGGTTCTGGCAATACCA	2760
2	2701	TTATGAATGTAAGGTTACTGGCCCCAATTTAGCGGTACGGTTCTGGCAATACCA	2760
1	2761	ACCTTATCCTACACGGTGTAAAGCTTCTATGGTTAACAAACCTGTGTGGAAAGCCTGG	2820
2	2761	ACCTTATCCTACACGGTGTAAAGCTTCTATGGTTAACAAACCTGTGTGGAAAGCCTGG	2820
1	2821	CCGATGTAAGGGTTCGGGCTGTGCCTTTACTGCTGCTGGAAAGGGTGGTGTGCGCC	2880
2	2821	CCGATGTAAGGGTTCGGGCTGTGCCTTTACTGCTGCTGGAAAGGGTGGTGTGCGCC	2880
1	2881	CCAAAAGCAGGGCTTCAATTAAGAAATGCCTCTTGAAGGTGTACCTGGTATCCTGT	2940
2	2881	CCAAAAGCAGGGCTTCAATTAAGAAATGCCTCTTGAAGGTGTACCTGGTATCCTGT	2940

1	2941	CTGAGGGTAACTCCAGGGTGC GCCACAATGTGGCTCCGACTGTGGTTGCTTCATGCTAG	3000
2	2941	CTGAGGGTAACTCCAGGGTGC GCCACAATGTGGCTCCGACTGTGGTTGCTTCATGCTAG	3000
1	3001	TGAAAAGCGTGGCTGTGATTAAGCATAACATGGTATGTGGCAACTGC GAGGACAGGGCCT	3060
2	3001	TGAAAAGCGTGGCTGTGATTAAGCATAACATGGTATGTGGCAACTGC GAGGACAGGGCCT	3060
1	3061	CTCAGATGCTGACCTGCTCGGACGGCAACTGTCACCTGCTGAAGACCATTACGTAGCCA	3120
2	3061	CTCAGATGCTGACCTGCTCGGACGGCAACTGTCACCTGCTGAAGACCATTACGTAGCCA	3120
1	3121	GCCACTCTCGCAAGGCCTGGCCAGTGTGTTGAGCATAACATACTGACCCGCTGTTCTTGC	3180
2	3121	GCCACTCTCGCAAGGCCTGGCCAGTGTGTTGAGCATAACATACTGACCCGCTGTTCTTGC	3180
1	3181	ATTTGGGTAACAGGAGGGGGTGTTCCTACCTTACCAATGCAATTGAGTCACACTAAGA	3240
2	3181	ATTTGGGTAACAGGAGGGGGTGTTCCTACCTTACCAATGCAATTGAGTCACACTAAGA	3240
1	3241	TATTGCTTGAGCCCGAGAGCATGTCCAAGGTGAACCTGAACGGGTGTTGACATGACCA	3300
2	3241	TATTGCTTGAGCCCGAGAGCATGTCCAAGGTGAACCTGAACGGGTGTTGACATGACCA	3300
1	3301	TGAAGATCTGGAAGGTGCTGAGGTACGATGAGACCCGACCAGGTGCAGACCTGCGAGT	3360
2	3301	TGAAGATCTGGAAGGTGCTGAGGTACGATGAGACCCGACCAGGTGCAGACCTGCGAGT	3360
1	3361	GTGGCGGTAAACATATTAGAACCCAGCCTGTGATGCTGGATGTGACCGAGGAGCTGAGGC	3420
2	3361	GTGGCGGTAAACATATTAGAACCCAGCCTGTGATGCTGGATGTGACCGAGGAGCTGAGGC	3420
1	3421	CCGATCACTTGGTGCCTGCCCTGCACCCCGCCTGAGTTGGCTCTAGCGATGAAGATAACAG	3480
2	3421	CCGATCACTTGGTGCCTGCCCTGCACCCCGCCTGAGTTGGCTCTAGCGATGAAGATAACAG	3480
1	3481	ATTGAGGTACTGAAATGTGTGGCGTGGCTTAAGGGTGGAAAGAATATATAAGGTGGGG	3540
2	3481	ATTGAGGTACTGAAATGTGTGGCGTGGCTTAAGGGTGGAAAGAATATATAAGGTGGGG	3540

1	3541	GTCTTATGTTAGTTGTATCTGTTGCAGCAGCCGCCATGAGCACCAACTCGT	3600
2	3541		
1	3601	TTGATGGAAGCATTGTGAGCTCATATTGACAACGCGCATGCCCATGGGCCGGGTGC	3660
2	3601		
1	3661	GTCAGAACATGTGATGGGCTCCAGCATTGATGGTCGCCCGTCCTGCCGAAACTCTACTA	3720
2	3661		
1	3721	CCTTGACCTACGAGACCGTGTCTGGAACGCCGTTGGAGACTGCAGCCTCGGCCGCCGCTT	3780
2	3721		
1	3781	CAGCCGCTGCAGCCACCGCCGCCGGATTGTGACTGACTTTGCTTCCTGAGCCCCTTG	3840
2	3781		
1	3841	CAAGCAGTGCAGCTTCCGTTATCCGCCCGCATGACAAGTTGACGGCTTTGGCAC	3900
2	3841		
1	3901	AATTGGATTCTTGACCCGGAACTTAATGTCGTTCTCAGCAGCTTGGATCTGCGCC	3960
2	3901		
1	3961	AGCAGGTTCTGCCCTGAAGGCTTCCCTCCCAATGCGGTTAAACATAAATAAAA	4020
2	3961		
1	4021	AACCAGACTCTGTTGGATTTGGATCAAGCAAGTGTCTTGCTGTCTTATTTAGGGTTT	4080
2	4021		
1	4081	TGCGCGCGCGTAGGCCCGGGACCAGCGGTCTGGTCGTTGAGGGCCTGTGTATTTTT	4140
2	4081		
1	4081	TGCGCGCGCGTAGGCCCGGGACCAGCGGTCTGGTCGTTGAGGGCCTGTGTATTTTT	4140
2	4081		

1	4141	CCAGGACGTGGTAAAGGTGACTCTGGATGTTAGATACTGGCATAAGCCCCTCTGG	4200
2	4141	CCAGGACGTGGTAAAGGTGACTCTGGATGTTAGATACTGGCATAAGCCCCTCTGG	4200
1	4201	GGTGGAGGTAGCACCCTGCAGAGCTTCATGCTGCCGGTGGTGTAGATGATCCAGT	4260
2	4201	GGTGGAGGTAGCACCCTGCAGAGCTTCATGCTGCCGGTGGTGTAGATGATCCAGT	4260
1	4261	CGTAGCAGGAGCGCTGGCGTGGTGCCTAAAAATGTCTTCAGTAGCAAGCTGATTGCCA	4320
2	4261	CGTAGCAGGAGCGCTGGCGTGGTGCCTAAAAATGTCTTCAGTAGCAAGCTGATTGCCA	4320
1	4321	GGGGCAGGCCCTGGTGTAAAGTGTACAAAGCGGTTAACAGCTGGATGGTGCAACGTG	4380
2	4321	GGGGCAGGCCCTGGTGTAAAGTGTACAAAGCGGTTAACAGCTGGATGGTGCAACGTG	4380
1	4381	GGGATATGAGATGCATCTGGACTGTATTTAGGTTGGCTATGTTCCAGCCATATCCC	4440
2	4381	GGGATATGAGATGCATCTGGACTGTATTTAGGTTGGCTATGTTCCAGCCATATCCC	4440
1	4441	TCCGGGGATTCATGTTGTGCAGAACCAACCAGCACAGTGTATCCGGTGCACGGAAATT	4500
2	4441	TCCGGGGATTCATGTTGTGCAGAACCAACCAGCACAGTGTATCCGGTGCACGGAAATT	4500
1	4501	TGTCATGTAGCTAGAAGGAAATGCGTGGAAAGAACCTGGAGACGCCCTGTGACCTCAA	4560
2	4501	TGTCATGTAGCTAGAAGGAAATGCGTGGAAAGAACCTGGAGACGCCCTGTGACCTCAA	4560
1	4561	GATTTCCATGCATCGTCCATAATGATGGCAATGGGCCACGGCGGCCCTGGCGA	4620
2	4561	GATTTCCATGCATCGTCCATAATGATGGCAATGGGCCACGGCGGCCCTGGCGA	4620
1	4621	AGATATTCTGGATCACTAACGTCAAGTTGTGTCAGGATGAGATCGTCATAGGCCA	4680
2	4621	AGATATTCTGGATCACTAACGTCAAGTTGTGTCAGGATGAGATCGTCATAGGCCA	4680
1	4681	TTTTTACAAAGCGCGGGCGGAGGGTGCAGACTGCGGTATAATGGTCCATCCGGCCAG	4740
2	4681	TTTTTACAAAGCGCGGGCGGAGGGTGCAGACTGCGGTATAATGGTCCATCCGGCCAG	4740

1	4741	GGCGTAGTTACCCCTCACAGATTGCATTTCCCACGCTTGAGTCAGATGGGGGATCA	4800
2	4741	GGCGTAGTTACCCCTCACAGATTGCATTTCCCACGCTTGAGTCAGATGGGGGATCA	4800
1	4801	TGTCTACCTGCGGGGCATGAAGAAAACGGTTCCGGGTAGGGAGATCAGCTGGAAAG	4860
2	4801	TGTCTACCTGCGGGGCATGAAGAAAACGGTTCCGGGTAGGGAGATCAGCTGGAAAG	4860
1	4861	AAAGCAGGTTCTGAGCAGCTGCAGACTTACCGCAGCCGGTGGGCCGTAAATCACACCTA	4920
2	4861	AAAGCAGGTTCTGAGCAGCTGCAGACTTACCGCAGCCGGTGGGCCGTAAATCACACCTA	4920
1	4921	TTACCGGCTGCAACTGGTAGTTAAGAGAGCTGCAGCTGCCGTATCCCTGAGCAGGGGG	4980
2	4921	TTACCGGCTGCAACTGGTAGTTAAGAGAGCTGCAGCTGCCGTATCCCTGAGCAGGGGG	4980
1	4981	CCACTTCGTTAACGATGTCCTGACTCGCATGTTTCCCTGACCAAATCCGCCAGAAGGC	5040
2	4981	CCACTTCGTTAACGATGTCCTGACTCGCATGTTTCCCTGACCAAATCCGCCAGAAGGC	5040
1	5041	GCTCGCCGCCAGCGATAGCAGTTCTTGCAAGGAAGCAAAGTTTCAACGGTTGAGAC	5100
2	5041	GCTCGCCGCCAGCGATAGCAGTTCTTGCAAGGAAGCAAAGTTTCAACGGTTGAGAC	5100
1	5101	CGTCCGCCGTAGGCATGCTTTGAGCGTTGACCAAGCAGCTCCAGGCGTCCACAGCT	5160
2	5101	CGTCCGCCGTAGGCATGCTTTGAGCGTTGACCAAGCAGCTCCAGGCGTCCACAGCT	5160
1	5161	CGGTCACCTGCTCTACGGCATCTCGATCCAGCATATCTCCTCGTTCGCGGGTTGGGCG	5220
2	5161	CGGTCACCTGCTCTACGGCATCTCGATCCAGCATATCTCCTCGTTCGCGGGTTGGGCG	5220
1	5221	GCTTCGCTGTACGGCAGTAGTCGGTGCTCGTCCAGACGGGCCAGGGTCATGTCTTCCA	5280
2	5221	GCTTCGCTGTACGGCAGTAGTCGGTGCTCGTCCAGACGGGCCAGGGTCATGTCTTCCA	5280
1	5281	CGGGCGCAGGGCCTCGTCAGCGTAGTCTGGTCACGGTAAGGGTGCCTCCGGCTG	5340
2	5281	CGGGCGCAGGGCCTCGTCAGCGTAGTCTGGTCACGGTAAGGGTGCCTCCGGCTG	5340

1	5341	CGCGCTGGCCAGGGTGCCTGAGGCTGGTCTGCTGGTGCTGAAGCGCTGCCGTCTTC	5400
2	5341	CGCGCTGGCCAGGGTGCCTGAGGCTGGTCTGCTGGTGCTGAAGCGCTGCCGTCTTC	5400
1	5401	GCCCTGCGCGTCGCCAGGTAGCATTGACCATGGTGTATAGTCCAGCCCCCTCCGCGGC	5460
2	5401	GCCCTGCGCGTCGCCAGGTAGCATTGACCATGGTGTATAGTCCAGCCCCCTCCGCGGC	5460
1	5461	GTGGCCCTTGGCGCGCAGCTTGCCTTGAGGAGGCCGCACGAGGGCAGTGCAGACT	5520
2	5461	GTGGCCCTTGGCGCGCAGCTTGCCTTGAGGAGGCCGCACGAGGGCAGTGCAGACT	5520
1	5521	TTTGAGGGCGTAGAGCTGGCGAGAAATACCGATTCCGGGAGTAGGCATCCGCGCC	5580
2	5521	TTTGAGGGCGTAGAGCTGGCGAGAAATACCGATTCCGGGAGTAGGCATCCGCGCC	5580
1	5581	GCAGGGCCCCGCAGACGGTCTCGCATTCCACGCCAGGTGAGCTCTGCCGTTGGGT	5640
2	5581	GCAGGGCCCCGCAGACGGTCTCGCATTCCACGCCAGGTGAGCTCTGCCGTTGGGT	5640
1	5641	AAAAACCAGGTTCCCCATGCTTTGATGCGTTCTTACCTCTGGTTCCATGAGCCG	5700
2	5641	AAAAACCAGGTTCCCCATGCTTTGATGCGTTCTTACCTCTGGTTCCATGAGCCG	5700
1	5701	GTGTCCACGCTCGGTGACGAAAAGGCTGTCCTGTCCTCGTATACTGAGAGGCCT	5760
2	5701	GTGTCCACGCTCGGTGACGAAAAGGCTGTCCTGTCCTCGTATACTGAGAGGCCT	5760
1	5761	GTCCTCGAGCGGTGTTCCCGGGCTCTCGTATACTGAGACACTCTGAGACAAA	5820
2	5761	GTCCTCGAGCGGTGTTCCCGGGCTCTCGTATACTGAGACACTCTGAGACAAA	5820
1	5821	GGCTCGCGTCCAGGCCAGCACGAAGGAGGCTAAGTGGAGGGTAGCGGTGTTGTCCAC	5880
2	5821	GGCTCGCGTCCAGGCCAGCACGAAGGAGGCTAAGTGGAGGGTAGCGGTGTTGTCCAC	5880
1	5881	TAGGGGGTCCACTCGCTCCAGGGTGTGAAGACACATGTCGCCCTTCCGCACTAAGGAA	5940
2	5881	TAGGGGGTCCACTCGCTCCAGGGTGTGAAGACACATGTCGCCCTTCCGCACTAAGGAA	5940

1 5941 GGTGATTGGTTGTAGGTGTAGGCCACGTGACCGGGTGTTCCTGAAGGGGGCTATAAAA 6000
2 5941 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||

1 6001 GGGGGTGGGGCGCGTCCTCACTCTTCCGCATCGCTGTCTGCAGGGCCAGCTG 6060
2 6001 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6061 TTGGGGTGAGTACTCCCTCTGAAAAGCGGGCATGACTTCTGCCTAAGATTGTCAGTTTC 6120
2 6061 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6121 CAAAAACGAGGAGGATTGATATTCACCTGGCCC CGGTGATGCCTTGAGGGTGGCCGC 6180
2 6121 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6181 ATCCATCTGGTCAGAAAGACAATCTTTGTTGTCAAGCTTGGTGGCAAACGACCCGTA 6240
2 6181 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6241 GAGGGCGTTGGACAGCAACTTGGCGATGGAGCGCAGGGTTGGTTTGTCGCGATCGGC 6300
2 6241 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6301 GCGCTCCTTGGCCCGATGTTAGCTGCACGTATT CGCGCAACGCAACGCCATT CGGG 6360
2 6301 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6361 AAAGACGGTGGTGCCTCGTCGGCACCAAGGTGCACGCCAACCGCGTTGTGCAGGGT 6420
2 6361 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6421 GACAAGGTCAACGCTGGTGGCTACCTCTCCCGTAGGCGCTCGTTGGCCAGCAGAGGCG 6480
2 6421 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1 6481 GCCGCCCTTGCAGAGCAGAATGGCGTAGGGGGCTAGCTGCCTCGTCCGGGGTC 6540
2 6481 ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| ||||||| |||||||

1	6541	TGC GT CCACGGTAAAGACCCCGGGCAGCAGGCGCGCTCGAAGTAGTCTATCTTGCATCC	6600
2	6541	TGC GT CCACGGTAAAGACCCCGGGCAGCAGGCGCGCTCGAAGTAGTCTATCTTGCATCC	6600
1	6601	TTGCAAGTCTAGCGCCTGCTGCCATGCGCGGGCGGAAGCGCGCGCTCGTATGGTTGAG	6660
2	6601	TTGCAAGTCTAGCGCCTGCTGCCATGCGCGGGCGGAAGCGCGCGCTCGTATGGTTGAG	6660
1	6661	TGGGGGACCCC ATGGCATGGGTGGGTGAGCGCGGAGGCGTACATGCCGCAAATGTCGTA	6720
2	6661	TGGGGGACCCC ATGGCATGGGTGGGTGAGCGCGGAGGCGTACATGCCGCAAATGTCGTA	6720
1	6721	AACGTAGAGGGCTCTTGAGTATTCCAAGATATTAGGGTAGCATCTTCCACCGCGGAT	6780
2	6721	AACGTAGAGGGCTCTTGAGTATTCCAAGATATTAGGGTAGCATCTTCCACCGCGGAT	6780
1	6781	GCTGGCGCGCACGTAATCGTATAGTTCGTGCAGGGAGCGAGGAGGTGGGACCGAGGTT	6840
2	6781	GCTGGCGCGCACGTAATCGTATAGTTCGTGCAGGGAGCGAGGAGGTGGGACCGAGGTT	6840
1	6841	GCTACGGGCGGGCTGCTCTGCTCGGAAGACTATCTGCCTGAAGATGGCATGTGAGTTGGA	6900
2	6841	GCTACGGGCGGGCTGCTCTGCTCGGAAGACTATCTGCCTGAAGATGGCATGTGAGTTGGA	6900
1	6901	TGATATGGTTGGACGCTGGAAAGACGTTGAAGCTGGCGTCTGTGAGACCTACCGCGTCACG	6960
2	6901	TGATATGGTTGGACGCTGGAAAGACGTTGAAGCTGGCGTCTGTGAGACCTACCGCGTCACG	6960
1	6961	CACGAAGGAGGCGTAGGAGTCGCGCAGCTTGTGACCAGCTCGCGGTGACCTGCACGTC	7020
2	6961	CACGAAGGAGGCGTAGGAGTCGCGCAGCTTGTGACCAGCTCGCGGTGACCTGCACGTC	7020
1	7021	TAGGGCGCAGTAGTCCAGGGTTCCCTGATGATGTCATACTTATCCTGTCCTTTTTTT	7080
2	7021	TAGGGCGCAGTAGTCCAGGGTTCCCTGATGATGTCATACTTATCCTGTCCTTTTTTT	7080
1	7081	CCACAGCTCGCGGTTGAGGAACAAACTCTTCGGGTCTTCCAGTACTCTGGATCGGAAA	7140
2	7081	CCACAGCTCGCGGTTGAGGAACAAACTCTTCGGGTCTTCCAGTACTCTGGATCGGAAA	7140

1	7141	CCCGTCGGCCTCCGAACGGTAAGAGCCTAGCATGTAGAACTGGTTGACGGCCTGGTAGGC	7200
2	7141	CCCGTCGGCCTCCGAACGGTAAGAGCCTAGCATGTAGAACTGGTTGACGGCCTGGTAGGC	7200
1	7201	GCAGCATCCCTTTCTACGGTAGCGCGTATGCCTGCGCGGCCCTCCGGAGCGAGGTGTG	7260
2	7201	GCAGCATCCCTTTCTACGGTAGCGCGTATGCCTGCGCGGCCCTCCGGAGCGAGGTGTG	7260
1	7261	GGTGAGCGCAAAGGTGTCCTGACCATGACTTGAGGTACTGGTATTGAAGTCAGTGTG	7320
2	7261	GGTGAGCGCAAAGGTGTCCTGACCATGACTTGAGGTACTGGTATTGAAGTCAGTGTG	7320
1	7321	GTCGCATCCGCCCTGCTCCCAGAGCAAAAGTCCGTGCGCTTTGGAACGCGGATTGG	7380
2	7321	GTCGCATCCGCCCTGCTCCCAGAGCAAAAGTCCGTGCGCTTTGGAACGCGGATTGG	7380
1	7381	CAGGGCGAAGGTGACATCGTTGAAGAGTATCTTCCCGCGAGGCATAAGTTGCGTGT	7440
2	7381	CAGGGCGAAGGTGACATCGTTGAAGAGTATCTTCCCGCGAGGCATAAGTTGCGTGT	7440
1	7441	GATGCGGAAGGGTCCGGCACCTCGAACGGTTGTTAATTACCTGGCGGCGAGCACGAT	7500
2	7441	GATGCGGAAGGGTCCGGCACCTCGAACGGTTGTTAATTACCTGGCGGCGAGCACGAT	7500
1	7501	CTCGTCAAAGCCGTTGATGTTGGCCCACAATGTAAGTTCCAAGAACGCGGGATGCC	7560
2	7501	CTCGTCAAAGCCGTTGATGTTGGCCCACAATGTAAGTTCCAAGAACGCGGGATGCC	7560
1	7561	CTTGATGGAAGGCAATTAAAGTTCTCGTAGGTGAGCTTCAGGGAGCTGAGCCC	7620
2	7561	CTTGATGGAAGGCAATTAAAGTTCTCGTAGGTGAGCTTCAGGGAGCTGAGCCC	7620
1	7621	GTGCTCTGAAAGGGCCAGTCTGCAAGATGAGGGTTGGAAGCGACGAATGAGCTCCACAG	7680
2	7621	GTGCTCTGAAAGGGCCAGTCTGCAAGATGAGGGTTGGAAGCGACGAATGAGCTCCACAG	7680
1	7681	GTCACGGGCCATTAGCATTGCAAGGTGGTCGAAAGGTCTAAACTGGCGACCTATGGC	7740
2	7681	GTCACGGGCCATTAGCATTGCAAGGTGGTCGAAAGGTCTAAACTGGCGACCTATGGC	7740

1	7741	CATTTTCTGGGTGATGCAGTAGAAGGTAAAGCGGGCTTGTCCAGCGGCCATCC	7800
2	7741	CATTTTCTGGGTGATGCAGTAGAAGGTAAAGCGGGCTTGTCCAGCGGCCATCC	7800
1	7801	AAGGTTCGCGCTAGGTCTCGCGGGCAGTCACTAGAGGCTCATCTCCGCCGAACCTCAT	7860
2	7801	AAGGTTCGCGCTAGGTCTCGCGGGCAGTCACTAGAGGCTCATCTCCGCCGAACCTCAT	7860
1	7861	GACCAGCATGAAGGGCACGAGCTGCTTCCAAAGGCCCCATCCAAGTATAAGGTCTCTAC	7920
2	7861	GACCAGCATGAAGGGCACGAGCTGCTTCCAAAGGCCCCATCCAAGTATAAGGTCTCTAC	7920
1	7921	ATCGTAGGTGACAAAGAGACGCTCGGTGCGAGGATGCGAGCCGATCGGAAGAACGGAT	7980
2	7921	ATCGTAGGTGACAAAGAGACGCTCGGTGCGAGGATGCGAGCCGATCGGAAGAACGGAT	7980
1	7981	CTCCCGCCACCAATTGGAGGAGTGGCTATTGATGTGGTCAAAGTAGAACGTCCTGCGACG	8040
2	7981	CTCCCGCCACCAATTGGAGGAGTGGCTATTGATGTGGTCAAAGTAGAACGTCCTGCGACG	8040
1	8041	GGCCGAACACTCGTGCTGGCTTTGTAAAAACGTGCGCAGTACTGGCAGCGGTGCACGGG	8100
2	8041	GGCCGAACACTCGTGCTGGCTTTGTAAAAACGTGCGCAGTACTGGCAGCGGTGCACGGG	8100
1	8101	CTGTACATCCTGCACGAGGTTGACCTGACGACCGCGCACAGGAAGCAGAGTGGAAATT	8160
2	8101	CTGTACATCCTGCACGAGGTTGACCTGACGACCGCGCACAGGAAGCAGAGTGGAAATT	8160
1	8161	GAGCCCTCGCCTGGCGGGTTGGCTGGTGGCTTCTACTTCGGCTGTTGTCCTTGACC	8220
2	8161	GAGCCCTCGCCTGGCGGGTTGGCTGGTGGCTTCTACTTCGGCTGTTGTCCTTGACC	8220
1	8221	GTCTGGCTGCTCGAGGGAGTTACGGTGGATCGGACCAACGCCGCCGAGCCAAAGT	8280
2	8221	GTCTGGCTGCTCGAGGGAGTTACGGTGGATCGGACCAACGCCGCCGAGCCAAAGT	8280
1	8281	CCAGATGTCCCGCGCGCGGGTGGAGCTTGATGACAACATCGCGCAGATGGAGCTGTC	8340
2	8281	CCAGATGTCCCGCGCGCGGGTGGAGCTTGATGACAACATCGCGCAGATGGAGCTGTC	8340

1	8341	CATGGTCTGGAGCTCCCGCGCGTCAGGTCAAGGCGGGAGCTCCTGCAGGTTACCTCGCA	8400
2	8341	CATGGTCTGGAGCTCCCGCGCGTCAGGTCAAGGCGGGAGCTCCTGCAGGTTACCTCGCA	8400
1	8401	TAGACGGGTCAAGGGCGCGGGCTAGATCCAGGTGATAACCTAATTCCAGGGCTGGTTGGT	8460
2	8401	TAGACGGGTCAAGGGCGCGGGCTAGATCCAGGTGATAACCTAATTCCAGGGCTGGTTGGT	8460
1	8461	GGCGGCGTCGATGGCTTGCAAGAGGCCGCATCCCCGCGCGACTACGGTACCGCGCGG	8520
2	8461	GGCGGCGTCGATGGCTTGCAAGAGGCCGCATCCCCGCGCGACTACGGTACCGCGCGG	8520
1	8521	CGGGCGGTGGGCCGCGGGGGTGTCCCTGGATGATGCATCTAAAGCGGTGACCGCGGGCGA	8580
2	8521	CGGGCGGTGGGCCGCGGGGGTGTCCCTGGATGATGCATCTAAAGCGGTGACCGCGGGCGA	8580
1	8581	GCCCCCGGAGGTAGGGGGGCTCCGGACCCGCCGGAGAGGGGGCAGGGGACGTGGCG	8640
2	8581	GCCCCCGGAGGTAGGGGGGCTCCGGACCCGCCGGAGAGGGGGCAGGGGACGTGGCG	8640
1	8641	CCGCGCGCGGGCAGGAGCTGGTGCTGCGCGTAGGTTGCTGGGAACCGCACGACGCGG	8700
2	8641	CCGCGCGCGGGCAGGAGCTGGTGCTGCGCGTAGGTTGCTGGGAACCGCACGACGCGG	8700
1	8701	CGGTTGATCTCCTGAATCTGGCGCCTCTCGCTGAAGACGACGGGCCCGGTGAGCTTGAAC	8760
2	8701	CGGTTGATCTCCTGAATCTGGCGCCTCTCGCTGAAGACGACGGGCCCGGTGAGCTTGAAC	8760
1	8761	CTGAAAGAGAGTTCGACAGAACATCAATTCCGGTGTGCGTTGACGGCGGCCCTGGCGAAAATC	8820
2	8761	CTGAAAGAGAGTTCGACAGAACATCAATTCCGGTGTGCGTTGACGGCGGCCCTGGCGAAAATC	8820
1	8821	TCCTGCACGTCTCCTGAGTTGTCTGATAGGCGATCTCGGCCATGAAC TGCTCGATCTCT	8880
2	8821	TCCTGCACGTCTCCTGAGTTGTCTGATAGGCGATCTCGGCCATGAAC TGCTCGATCTCT	8880
1	8881	TCCTCCTGGAGATCTCCCGCGTCCGGCTCGCTCCACGGTGGCGGGGAGGTCGTTGGAAATG	8940
2	8881	TCCTCCTGGAGATCTCCCGCGTCCGGCTCGCTCCACGGTGGCGGGGAGGTCGTTGGAAATG	8940

1	8941	CGGGCCATGAGCTGCGAGAAGGC GTT GAGGCCTCCCTCGTTCCAGACGCCGGCTGTAGACC	9000
2	8941	CGGGCCATGAGCTGCGAGAAGGC GTT GAGGCCTCCCTCGTTCCAGACGCCGGCTGTAGACC	9000
1	9001	ACGCCCCCTCGGCATCGCGGGCGCGCATGACCACCTGCGCGAGATTGAGCTCCACGTGC	9060
2	9001	ACGCCCCCTCGGCATCGCGGGCGCGCATGACCACCTGCGCGAGATTGAGCTCCACGTGC	9060
1	9061	CGGGCGAAGACGGCGTAGTTCGCAGGC GCTGAAAGAGGTAGTTGAGGGTGGTGGCGGTG	9120
2	9061	CGGGCGAAGACGGCGTAGTTCGCAGGC GCTGAAAGAGGTAGTTGAGGGTGGTGGCGGTG	9120
1	9121	TGTTCTGCCACGAAGAAGTACATAACCCAGCGTCGCAACGTGGATTGTTGATATCCCCC	9180
2	9121	TGTTCTGCCACGAAGAAGTACATAACCCAGCGTCGCAACGTGGATTGTTGATATCCCCC	9180
1	9181	AAGGCCTCAAGGC GCTCCATGGCCTCGTAGAAGTCCACGGCGAAGTTGAAAAACTGGGAG	9240
2	9181	AAGGCCTCAAGGC GCTCCATGGCCTCGTAGAAGTCCACGGCGAAGTTGAAAAACTGGGAG	9240
1	9241	TTGCGCGCCGACACGGTTA ACT CCT CCT CCAGAACAGGGATGAGCTGGCGACAGTGTG	9300
2	9241	TTGCGCGCCGACACGGTTA ACT CCT CCT CCAGAACAGGGATGAGCTGGCGACAGTGTG	9300
1	9301	CGCACCTCGCGCTAAAGGCTACAGGGGCCTTCTTCTTCAATCTCCCTTTCCATA	9360
2	9301	CGCACCTCGCGCTAAAGGCTACAGGGGCCTTCTTCTTCAATCTCCCTTTCCATA	9360
1	9361	AGGGCCTCCCCTTCTTCTTCTGGCGGCGGTGGGGAGGGGGACACGGCGGCGACGA	9420
2	9361	AGGGCCTCCCCTTCTTCTTCTGGCGGCGGTGGGGAGGGGGACACGGCGGCGACGA	9420
1	9421	CGGCGCACCGGGAGGC GGTCGACAAAGCGCTCGATCATCTCCCCGCGGCACGGCGCATG	9480
2	9421	CGGCGCACCGGGAGGC GGTCGACAAAGCGCTCGATCATCTCCCCGCGGCACGGCGCATG	9480
1	9481	GTCTCGGTGACGGCGCGGCCGTTCTCGCGGGGGCGCAGTTGGAAGACGCCGCCGTCA	9540
2	9481	GTCTCGGTGACGGCGCGGCCGTTCTCGCGGGGGCGCAGTTGGAAGACGCCGCCGTCA	9540

1	9541	TCCCGGTTATGGTTGGCGGGGGCTGCCATGCAGGGATAACGGCTAACGATGCAT	9600
2	9541	TCCCGGTTATGGTTGGCGGGGGCTGCCATGCAGGGATAACGGCTAACGATGCAT	9600
1	9601	CTCAACAATTGTTGTAGGTACTCCGCCGAGGGACCTGAGCGAGTCCGCATCGACC	9660
2	9601	CTCAACAATTGTTGTAGGTACTCCGCCGAGGGACCTGAGCGAGTCCGCATCGACC	9660
1	9661	GGATCGGAAAACCTCTCGAGAAAGGCGTCAACCAGTCACAGTCGAAGGTAGGCTGAGC	9720
2	9661	GGATCGGAAAACCTCTCGAGAAAGGCGTCAACCAGTCACAGTCGAAGGTAGGCTGAGC	9720
1	9721	ACCGTGGCGGGCGGCAGCGGGCGGTGGGTTGTTCTGGCGGAGGTGCTGCTGATG	9780
2	9721	ACCGTGGCGGGCGGCAGCGGGCGGTGGGTTGTTCTGGCGGAGGTGCTGCTGATG	9780
1	9781	ATGTAATTAAAGTAGGCGGTCTTGAGACGGCGGATGGTCGACAGAACCATGTCCTTG	9840
2	9781	ATGTAATTAAAGTAGGCGGTCTTGAGACGGCGGATGGTCGACAGAACCATGTCCTTG	9840
1	9841	GGTCGGCCTGCTGAATGCGCAGGCAGGTGGCCATGCCCAAGGCTTCGTTTGACATCGG	9900
2	9841	GGTCGGCCTGCTGAATGCGCAGGCAGGTGGCCATGCCCAAGGCTTCGTTTGACATCGG	9900
1	9901	CGCAGGTCTTGTAGTAGTCTTGCATGAGCCTTCTACCGGCACCTTCTTCTCCTTCC	9960
2	9901	CGCAGGTCTTGTAGTAGTCTTGCATGAGCCTTCTACCGGCACCTTCTTCTCCTTCC	9960
1	9961	TCTTGTCTGCATCTTGCATCTATCGCTGGCGGGCGGAGTTGGCGTAGGTGG	10 020
2	9961	TCTTGTCTGCATCTTGCATCTATCGCTGGCGGGCGGAGTTGGCGTAGGTGG	10 020
1	10 021	CGCCCTCTTCCCAGCGTGTGACCCGAAGCCCTCATCGCTGAAGCAGGGCTAGG	10 080
2	10 021	CGCCCTCTTCCCAGCGTGTGACCCGAAGCCCTCATCGCTGAAGCAGGGCTAGG	10 080
1	10 081	TCGGCGACAACCGCGCTGGCTAATATGGCTGCTGCACCTCGTGAGGGTAGACTGGAAG	10 140
2	10 081	TCGGCGACAACCGCGCTGGCTAATATGGCTGCTGCACCTCGTGAGGGTAGACTGGAAG	10 140

1 10 141 TCATCCATGTCCACAAAGCGGTGGTATGCGCCCGTGTGATGGTGTAAAGTGCAGTTGCC 10 200
2 |||||||
1 10 141 TCATCCATGTCCACAAAGCGGTGGTATGCGCCCGTGTGATGGTGTAAAGTGCAGTTGCC 10 200

1 10 201 ATAACGGACCAGTTAACGGTCTGGTGACCCGGCTGCAGAGCTCGGTGTACCTGAGACGC 10 260
2 |||||||
1 10 201 ATAACGGACCAGTTAACGGTCTGGTGACCCGGCTGCAGAGCTCGGTGTACCTGAGACGC 10 260

1 10 261 GAGTAAGCCCTCGAGTCAAATACGTAGTCGTTGCAAGTCCGCACCAGGTACTGGTATCCC 10 320
2 |||||||
1 10 261 GAGTAAGCCCTCGAGTCAAATACGTAGTCGTTGCAAGTCCGCACCAGGTACTGGTATCCC 10 320

1 10 321 ACCAAAAAGTGC GGCGGGCTGGCGTAGAGGGGCCAGCGTAGGGTGGCCGGGCTCCG 10 380
2 |||||||
1 10 321 ACCAAAAAGTGC GGCGGGCTGGCGTAGAGGGGCCAGCGTAGGGTGGCCGGGCTCCG 10 380

1 10 381 GGGCGAGATCTTCAACATAAGGCGATGATATCCGTAGATGTACCTGGACATCCAGGTG 10 440
2 |||||||
1 10 381 GGGCGAGATCTTCAACATAAGGCGATGATATCCGTAGATGTACCTGGACATCCAGGTG 10 440

1 10 441 ATGCCGGCGGCGGTGGTGGAGGCGCGCGAAAGTCGCGGACGC GGTTCCAGATGTTGCC 10 500
2 |||||||
1 10 441 ATGCCGGCGGCGGTGGTGGAGGCGCGCGAAAGTCGCGGACGC GGTTCCAGATGTTGCC 10 500

1 10 501 AGCGGGAAAAAGTGCCTCATGGTCGGGACGCTCTGCCGGTCAGGCGCGCAATCGTTG 10 560
2 |||||||
1 10 501 AGCGGGAAAAAGTGCCTCATGGTCGGGACGCTCTGCCGGTCAGGCGCGCAATCGTTG 10 560

1 10 561 ACGCTCTAGACCGTGCAAAAGGAGAGCCTGTAAGCGGGCACTCTTCCGTGGTCTGGTGG 10 620
2 |||||||
1 10 561 ACGCTCTAGACCGTGCAAAAGGAGAGCCTGTAAGCGGGCACTCTTCCGTGGTCTGGTGG 10 620

1 10 621 TAAATTGCAAGGGTATCATGGCGGACGACC GGTTCGAGCCCCGTATCCGGCCGTCCG 10 680
2 |||||||
1 10 621 TAAATTGCAAGGGTATCATGGCGGACGACC GGTTCGAGCCCCGTATCCGGCCGTCCG 10 680

1 10 681 CCGTGATCCATGCGGTTACCGCCCCGCGTGTGAAACCCAGGTGTGCGACGTCAGACAACGG 10 740
2 |||||||
1 10 681 CCGTGATCCATGCGGTTACCGCCCCGCGTGTGAAACCCAGGTGTGCGACGTCAGACAACGG 10 740

1	10 741	GGGAGTGCTCCTTGGCTTCAGGCAGGGCTGCTGCGCTAGCTTTTGCC	10 800
2	10 741	GGGAGTGCTCCTTGGCTTCAGGCAGGGCTGCTGCGCTAGCTTTTGCC	10 800
1	10 801	ACTGGCCGCGCAGCGTAAGCGGTTAGGCTGGAAAGCGAAAGCATTAAGTGGCTCGCTC	10 860
2	10 801	ACTGGCCGCGCAGCGTAAGCGGTTAGGCTGGAAAGCGAAAGCATTAAGTGGCTCGCTC	10 860
1	10 861	CCTGTAGCCGGAGGGTTATTTCCAAGGGTTGAGTCGCGGGACCCCCGGTTCGAGTCTCG	10 920
2	10 861	CCTGTAGCCGGAGGGTTATTTCCAAGGGTTGAGTCGCGGGACCCCCGGTTCGAGTCTCG	10 920
1	10 921	GACCGGCCGGACTGCAGCGAACGGGGTTGCCTCCCCGTATGCAAGACCCGCTTGCA	10 980
2	10 921	GACCGGCCGGACTGCAGCGAACGGGGTTGCCTCCCCGTATGCAAGACCCGCTTGCA	10 980
1	10 981	AATTCTCCGGAAACAGGGACGAGCCCTTTTGCCTTCCAGATGCATCCGGTGCTG	11 040
2	10 981	AATTCTCCGGAAACAGGGACGAGCCCTTTTGCCTTCCAGATGCATCCGGTGCTG	11 040
1	11 041	CGGCAGATGCAGCCCCCTCCTCAGCAGCGCAAGAGCAAGAGCAGCGGCAGACATGCAGG	11 100
2	11 041	CGGCAGATGCAGCCCCCTCCTCAGCAGCGCAAGAGCAAGAGCAGCGGCAGACATGCAGG	11 100
1	11 101	GCACCCCTCCCTCCTACCGCGTCAGGAGGGCGACATCCGGTTGACGCGGCAGCA	11 160
2	11 101	GCACCCCTCCCTCCTACCGCGTCAGGAGGGCGACATCCGGTTGACGCGGCAGCA	11 160
1	11 161	GATGGTGATTACGAACCCCCCGCGCGCCGGCCGGCACTACCTGGACTTGGAGGGAGGGC	11 220
2	11 161	GATGGTGATTACGAACCCCCCGCGCGCCGGCCGGCACTACCTGGACTTGGAGGGAGGGC	11 220
1	11 221	GAGGGCCTGGCGCGTAGGAGCGCCCTCCTGAGCGGCACCCAAGGGTGCAGCTGAAG	11 280
2	11 221	GAGGGCCTGGCGCGTAGGAGCGCCCTCCTGAGCGGCACCCAAGGGTGCAGCTGAAG	11 280
1	11 281	CGTGATACCGCGTAGGCGTACGTGCCCGGCAGAACCTGTTCGCACCGCGAGGGAGAG	11 340
2	11 281	CGTGATACCGCGTAGGCGTACGTGCCCGGCAGAACCTGTTCGCACCGCGAGGGAGAG	11 340

1 11 341 GAGCCCGAGGAGATGCAGGGATCGAAAGTTCCACGCAGGGCGCAGCTGCAGGCATGGCTG 11 400
2 11 341 GAGCCCGAGGAGATGCAGGGATCGAAAGTTCCACGCAGGGCGCAGCTGCAGGCATGGCTG 11 400

1 11 401 AATCGCGAGCGGTTGCTGCGCGAGGAGGACTTGTAGCCCCACGCACGCCAACCGGGATTAGT 11 460
2 11 401 AATCGCGAGCGGTTGCTGCGCGAGGAGGACTTGTAGCCCCACGCACGCCAACCGGGATTAGT 11 460

1 11 461 CCCGCGCGCGCACACGTGGCGGCCGCCACCTGGTAACCGCATACGAGCAGACGGTGAAC 11 520
2 11 461 CCCGCGCGCGCACACGTGGCGGCCGCCACCTGGTAACCGCATACGAGCAGACGGTGAAC 11 520

1 11 521 CAGGAGATTAACTTCAAAAAAGCTTAACAAACCACGTGCGTACGCTTGTGGCGCGAG 11 580
2 11 521 CAGGAGATTAACTTCAAAAAAGCTTAACAAACCACGTGCGTACGCTTGTGGCGCGAG 11 580

1 11 581 GAGGTGGCTATAAGGACTGATGCATCTGTGGACTTTGTAAGCGCGCTGGAGCAAAACCCA 11 640
2 11 581 GAGGTGGCTATAAGGACTGATGCATCTGTGGACTTTGTAAGCGCGCTGGAGCAAAACCCA 11 640

1 11 641 AATAGCAAGCCGCTCATGGCGCAGCTGTTCTTATAGTCAGCACAGCAGGGACAACGAG 11 700
2 11 641 AATAGCAAGCCGCTCATGGCGCAGCTGTTCTTATAGTCAGCACAGCAGGGACAACGAG 11 700

1 11 701 GCATTCAAGGGATGCGCTGCTAACATAGTAGAGCCGAGGGCCGCTGGCTGCTCGATTTG 11 760
2 11 701 GCATTCAAGGGATGCGCTGCTAACATAGTAGAGCCGAGGGCCGCTGGCTGCTCGATTTG 11 760

1 11 761 ATAAACATCCTGCAGAGCATAGTGGTGCAGGAGCGCAGCTTGAGCCTGGCTGACAAGGTG 11 820
2 11 761 ATAAACATCCTGCAGAGCATAGTGGTGCAGGAGCGCAGCTTGAGCCTGGCTGACAAGGTG 11 820

1 11 821 GCCGCCATCAACTATTCCATGCTTAGCCTGGCAAGTTTACGCCCAAGATATACCAT 11 880
2 11 821 GCCGCCATCAACTATTCCATGCTTAGCCTGGCAAGTTTACGCCCAAGATATACCAT 11 880

1 11 881 ACCCCTTACGTTCCCATAAGACAAGGAGGTAAAGATCGAGGGGTTCTACATGCGCATGGCG 11 940
2 11 881 ACCCCTTACGTTCCCATAAGACAAGGAGGTAAAGATCGAGGGGTTCTACATGCGCATGGCG 11 940

1	11 941	CTGAAGGTGCTTACCTTGAGCGACGACCTGGCGTTATCGCAACGAGCGCATCCACAAG	12 000
2	11 941	CTGAAGGTGCTTACCTTGAGCGACGACCTGGCGTTATCGCAACGAGCGCATCCACAAG	12 000
1	12 001	GCCGTGAGCGTGAGCCGGCGCGAGCTCAGCGACC CGAGCTGATGCACAGCCTGCAA	12 060
2	12 001	GCCGTGAGCGTGAGCCGGCGCGAGCTCAGCGACC CGAGCTGATGCACAGCCTGCAA	12 060
1	12 061	AGGGCCCTGGCTGGCACGGCAGCGCGATAGAGAGGCCGAGTCCTACTTTGACGCGGGC	12 120
2	12 061	AGGGCCCTGGCTGGCACGGCAGCGCGATAGAGAGGCCGAGTCCTACTTTGACGCGGGC	12 120
1	12 121	GCTGACCTGCGCTGGGCCCAAGCCGACGCGCCCTGGAGGCAGCTGGGGCCGGACCTGGG	12 180
2	12 121	GCTGACCTGCGCTGGGCCCAAGCCGACGCGCCCTGGAGGCAGCTGGGGCCGGACCTGGG	12 180
1	12 181	CTGGCGGTGGCACCCGCGCGCTGGCAACGTCGGCGGTGGAGGAATATGACGAGGAC	12 240
2	12 181	CTGGCGGTGGCACCCGCGCGCTGGCAACGTCGGCGGTGGAGGAATATGACGAGGAC	12 240
1	12 241	GATGAGTACGAGCCAGAGGACGGCGAGTACTAAGCGGTGATGTTCTGATCAGATGATGC	12 300
2	12 241	GATGAGTACGAGCCAGAGGACGGCGAGTACTAAGCGGTGATGTTCTGATCAGATGATGC	12 300
1	12 301	AAGACGCAACGGACCCGGCGGTGCGGGCGCGCTGCAGAGCCAGCCGTCCGGCTTA ACT	12 360
2	12 301	AAGACGCAACGGACCCGGCGGTGCGGGCGCGCTGCAGAGCCAGCCGTCCGGCTTA ACT	12 360
1	12 361	CCACGGACGACTGGCGCCAGGT CATGGACCGCATCATGTCGCTGACTGCGCGCAATCCTG	12 420
2	12 361	CCACGGACGACTGGCGCCAGGT CATGGACCGCATCATGTCGCTGACTGCGCGCAATCCTG	12 420
1	12 421	ACGCGTTCCGGCAGCAGCCGAGGCCAACCGGCTCTCGCAATTCTGGAAGCGGTGGTCC	12 480
2	12 421	ACGCGTTCCGGCAGCAGCCGAGGCCAACCGGCTCTCGCAATTCTGGAAGCGGTGGTCC	12 480
1	12 481	CGGCGCGCGCAAACCCCACGCACGAGAAGGTGCTGGCGATCGTAAACGCGCTGGCGAAA	12 540
2	12 481	CGGCGCGCGCAAACCCCACGCACGAGAAGGTGCTGGCGATCGTAAACGCGCTGGCGAAA	12 540

1	12 541	ACAGGGCCATCCGGCCCACGAGGCCGGCTGGTCTACGACGCGCTGTTCAGCGCGTGG	12 600
2	12 541	ACAGGGCCATCCGGCCCACGAGGCCGGCTGGTCTACGACGCGCTGTTCAGCGCGTGG	12 600
1	12 601	CTCGTTACAACAGCGGCAACGTGCAGACCAACCTGGACCGGCTGGTGGGGATGTGCGCG	12 660
2	12 601	CTCGTTACAACAGCGGCAACGTGCAGACCAACCTGGACCGGCTGGTGGGGATGTGCGCG	12 660
1	12 661	AGGCCGTGGCGCAGCGTGAGCGCGCAGCAGCAGGGCAACCTGGCTCCATGGTTGCAC	12 720
2	12 661	AGGCCGTGGCGCAGCGTGAGCGCGCAGCAGCAGGGCAACCTGGCTCCATGGTTGCAC	12 720
1	12 721	TAAACGCCCTCCTGAGTACACAGCCGCCAACGTGCCGGGGACAGGAGGACTACACCA	12 780
2	12 721	TAAACGCCCTCCTGAGTACACAGCCGCCAACGTGCCGGGGACAGGAGGACTACACCA	12 780
1	12 781	ACTTTGTGAGCGCACTGCGGCTAATGGTACTGAGACACCGCAAAGTGAGGTGTACCA	12 840
2	12 781	ACTTTGTGAGCGCACTGCGGCTAATGGTACTGAGACACCGCAAAGTGAGGTGTACCA	12 840
1	12 841	CTGGGCCAGACTATTTTCCAGACCAAGTAGACAAGGCCCTGCAGACCGTAAACCTGAGCC	12 900
2	12 841	CTGGGCCAGACTATTTTCCAGACCAAGTAGACAAGGCCCTGCAGACCGTAAACCTGAGCC	12 900
1	12 901	AGGCTTCAAAAACTTGCAGGGCTGTGGGGGTGCAGGGCTCCACAGGCGACCGCGCGA	12 960
2	12 901	AGGCTTCAAAAACTTGCAGGGCTGTGGGGGTGCAGGGCTCCACAGGCGACCGCGCGA	12 960
1	12 961	CCGTGTCTAGCTTGCTGACGCCAACTCGCGCCTGTTGCTGCTGCTAATAGGCCCTCA	13 020
2	12 961	CCGTGTCTAGCTTGCTGACGCCAACTCGCGCCTGTTGCTGCTGCTAATAGGCCCTCA	13 020
1	13 021	CGGACAGTGGCAGCGTGTCCGGGACACATACTAGGTCACTTGCTGACACTGTACCGCG	13 080
2	13 021	CGGACAGTGGCAGCGTGTCCGGGACACATACTAGGTCACTTGCTGACACTGTACCGCG	13 080
1	13 081	AGGCCATAGGTAGGCATGTGGACGAGCATACTTCCAGGAGATTACAAGTGTAGCC	13 140
2	13 081	AGGCCATAGGTAGGCATGTGGACGAGCATACTTCCAGGAGATTACAAGTGTAGCC	13 140

1	13 141	GCGCGCTGGGCAGGAGGACACGGCAGCCTGGAGGAACCTAAACTACCTGCTGACCA	13 200
2	13 141	GCGCGCTGGGCAGGAGGACACGGCAGCCTGGAGGAACCTAAACTACCTGCTGACCA	13 200
1	13 201	ACCGGCCAGAAGATCCCCTCGTTGCACAGTTAACAGCGAGGAGGAGCGCATTTGC	13 260
2	13 201	ACCGGCCAGAAGATCCCCTCGTTGCACAGTTAACAGCGAGGAGGAGCGCATTTGC	13 260
1	13 261	GCTACGTGCAGCAGAGCGTGAGCCTAACCTGATGCGCACGGGTAACGCCAGCGTGG	13 320
2	13 261	GCTACGTGCAGCAGAGCGTGAGCCTAACCTGATGCGCACGGGTAACGCCAGCGTGG	13 320
1	13 321	CGCTGGACATGACCGCGCAACATGGAACCAGGGATGTATGCCTCAAACCGGCCGTTA	13 380
2	13 321	CGCTGGACATGACCGCGCAACATGGAACCAGGGATGTATGCCTCAAACCGGCCGTTA	13 380
1	13 381	TCAACCGCTAATGGACTACTTGCATCGCGGGCCCGTGAACCCGAGTATTACCCA	13 440
2	13 381	TCAACCGCTAATGGACTACTTGCATCGCGGGCCCGTGAACCCGAGTATTACCCA	13 440
1	13 441	ATGCCATCTGAACCCGCACTGGCTACCGCCCCCTGGTTCTACACCGGGGATTGAGG	13 500
2	13 441	ATGCCATCTGAACCCGCACTGGCTACCGCCCCCTGGTTCTACACCGGGGATTGAGG	13 500
1	13 501	TGCCCAGGGTAACGATGGATTCTCTGGGACGACATAGACGACAGCGTGTTCGGC	13 560
2	13 501	TGCCCAGGGTAACGATGGATTCTCTGGGACGACATAGACGACAGCGTGTTCGGC	13 560
1	13 561	AACCGCAGACCTGCTAGAGTTGCAACAGCGCGAGCAGGCAGAGCGGCGCTGCGAAAGG	13 620
2	13 561	AACCGCAGACCTGCTAGAGTTGCAACAGCGCGAGCAGGCAGAGCGGCGCTGCGAAAGG	13 620
1	13 621	AAAGCTTCCGCAGGCCAAGCAGCTTGTCCGATCTAGGCCTGGGCCCCCGCGTCAGATG	13 680
2	13 621	AAAGCTTCCGCAGGCCAAGCAGCTTGTCCGATCTAGGCCTGGGCCCCCGCGTCAGATG	13 680
1	13 681	CTAGTAGCCCATTCCAAGCTTGTAGGGTCTTACCAAGCACTCGCACCAACCGCCC	13 740
2	13 681	CTAGTAGCCCATTCCAAGCTTGTAGGGTCTTACCAAGCACTCGCACCAACCGCCC	13 740

1	13 741	GCCTGCTGGCGAGGAGGTACCTAAACAACCGCTGCTGCAGCCGAGCGCGAAAAAA	13 800
2	13 741	GCCTGCTGGCGAGGAGGTACCTAAACAACCGCTGCTGCAGCCGAGCGCGAAAAAA	13 800
1	13 801	ACCTGCCTCCGGCATTCCCAACAACGGATAGAGAGCCTAGTGGACAAGATGAGTAGAT	13 860
2	13 801	ACCTGCCTCCGGCATTCCCAACAACGGATAGAGAGCCTAGTGGACAAGATGAGTAGAT	13 860
1	13 861	GGAAGACGTACCGCAGGGAGCACAGGGACGTGCCAGGCCGCGCCACCCGTGTC	13 920
2	13 861	GGAAGACGTACCGCAGGGAGCACAGGGACGTGCCAGGCCGCGCCACCCGTGTC	13 920
1	13 921	AAAGGCACGACCGTCAGCGGGCTGGTGTGGAGGGACGATGACTCGCAGACGACAGCA	13 980
2	13 921	AAAGGCACGACCGTCAGCGGGCTGGTGTGGAGGGACGATGACTCGCAGACGACAGCA	13 980
1	13 981	GCGTCCTGGATTGGAGGGAGTGGCAACCCGTTGCGCACCTCGCCCCAGGCTGGGA	14 040
2	13 981	GCGTCCTGGATTGGAGGGAGTGGCAACCCGTTGCGCACCTCGCCCCAGGCTGGGA	14 040
1	14 041	GAATGTTTAAAAAAAAAAGCATGATGCAAAATAAAACTCACCAAGGCCATGGCA	14 100
2	14 041	GAATGTTTAAAAAAAAAAGCATGATGCAAAATAAAACTCACCAAGGCCATGGCA	14 100
1	14 101	CCGAGCGTTGGTTTCTTGTATTCCCTTAGTATGCGCGCGCGATGTATGAGGAAG	14 160
2	14 101	CCGAGCGTTGGTTTCTTGTATTCCCTTAGTATGCGCGCGCGATGTATGAGGAAG	14 160
1	14 161	GTCCTCCCTCCCTACGAGAGTGTGGTGAGCGCGCCAGTGGCGCGCTGGTT	14 220
2	14 161	GTCCTCCCTCCCTACGAGAGTGTGGTGAGCGCGCCAGTGGCGCGCTGGTT	14 220
1	14 221	CTCCCTCGATGCTCCCTGGACCCGCCGTTGTGCCTCCGCGGTACCTGCGGCCAACCG	14 280
2	14 221	CTCCCTCGATGCTCCCTGGACCCGCCGTTGTGCCTCCGCGGTACCTGCGGCCAACCG	14 280
1	14 281	GGGGGAGAACAGCATCCGTTACTCTGAGTTGGCACCCCTATTGACACCACCCGTGT	14 340
2	14 281	GGGGGAGAACAGCATCCGTTACTCTGAGTTGGCACCCCTATTGACACCACCCGTGT	14 340

1	14 341	ACCTGGTGGACAACAAGTCACGGATGTGGCATCCCTGAACCTACCAAGAACGACCAAGCA	14 400
2	14 341	ACCTGGTGGACAACAAGTCACGGATGTGGCATCCCTGAACCTACCAAGAACGACCAAGCA	14 400
1	14 401	ACTTTCTGACCACGGTCATTAAAACAATGACTACAGCCCCGGGGAGGCAAGCACACAGA	14 460
2	14 401	ACTTTCTGACCACGGTCATTAAAACAATGACTACAGCCCCGGGGAGGCAAGCACACAGA	14 460
1	14 461	CCATCAATCTTGACGACCGGTCGCACCTGGGGCGGCACCTGAAACCATCCTGCATACCA	14 520
2	14 461	CCATCAATCTTGACGACCGGTCGCACCTGGGGCGGCACCTGAAACCATCCTGCATACCA	14 520
1	14 521	ACATGCCAAATGTGAACGAGTTCATGTTACCAATAAGTTAAGGCGCGGGTGTGGTGT	14 580
2	14 521	ACATGCCAAATGTGAACGAGTTCATGTTACCAATAAGTTAAGGCGCGGGTGTGGTGT	14 580
1	14 581	CGCGCTTGCCTACTAAGGACAATCAGGTGGAGCTGAAATACGAGTGGGTGGAGTTCACGC	14 640
2	14 581	CGCGCTTGCCTACTAAGGACAATCAGGTGGAGCTGAAATACGAGTGGGTGGAGTTCACGC	14 640
1	14 641	TGCCCAGGGCAACTACTCCGAGACCATGACCATAGACCTTATGAACAAACGCGATCGTGG	14 700
2	14 641	TGCCCAGGGCAACTACTCCGAGACCATGACCATAGACCTTATGAACAAACGCGATCGTGG	14 700
1	14 701	AGCACTACTTGAAAGTGGCAGACAGAACGGGTTCTGGAAAGCGACATCGGGTAAAGT	14 760
2	14 701	AGCACTACTTGAAAGTGGCAGACAGAACGGGTTCTGGAAAGCGACATCGGGTAAAGT	14 760
1	14 761	TTGACACCCGCAACTTCAGACTGGGTTTACCGTCACTGGCTTGTATGCCTGGGG	14 820
2	14 761	TTGACACCCGCAACTTCAGACTGGGTTTACCGTCACTGGCTTGTATGCCTGGGG	14 820
1	14 821	TATATACAAACGAAGCCTTCCATCCAGACATCATTGCTGCCAGGATGCGGGTGGACT	14 880
2	14 821	TATATACAAACGAAGCCTTCCATCCAGACATCATTGCTGCCAGGATGCGGGTGGACT	14 880
1	14 881	TCACCCACAGCCGCCTGAGCAACTTGTGGCATCCGCAAGCGGCAACCCTCCAGGAGG	14 940
2	14 881	TCACCCACAGCCGCCTGAGCAACTTGTGGCATCCGCAAGCGGCAACCCTCCAGGAGG	14 940

1	14 941	GCTTTAGGATCACCTACGATGATCTGGAGGGTGGTAACATTCCGCACGTGTTGGATGTGG	15 000
2	14 941	GCTTTAGGATCACCTACGATGATCTGGAGGGTGGTAACATTCCGCACGTGTTGGATGTGG	15 000
1	15 001	ACGCCTACCAGGCAGCTTGAAAGATGACACCGAACAGGGCGGGGTGGCGCAGGCCGGCA	15 060
2	15 001	ACGCCTACCAGGCAGCTTGAAAGATGACACCGAACAGGGCGGGGTGGCGCAGGCCGGCA	15 060
1	15 061	GCAACAGCAGTGGCAGCGCGCGAACAGAGAACCTCAACCGCAGCCCGGGCAATGCAGC	15 120
2	15 061	GCAACAGCAGTGGCAGCGCGCGAACAGAGAACCTCAACCGCAGCCCGGGCAATGCAGC	15 120
1	15 121	CGGTGGAGGACATGAACGATCATGCCATTGGCGAACACCTTGGCACACGGGCTGAGG	15 180
2	15 121	CGGTGGAGGACATGAACGATCATGCCATTGGCGAACACCTTGGCACACGGGCTGAGG	15 180
1	15 181	AGAAGCGCGCTGAGGCCGAAGCAGCGGCCGAAGCTGCCGCCCCGCTGCGCAACCCGAGG	15 240
2	15 181	AGAAGCGCGCTGAGGCCGAAGCAGCGGCCGAAGCTGCCGCCCCGCTGCGCAACCCGAGG	15 240
1	15 241	TCGAGAACGCTCAGAAGAAACCGGTGATCAAACCCCTGACAGAGGACAGCAAGAAACGCA	15 300
2	15 241	TCGAGAACGCTCAGAAGAAACCGGTGATCAAACCCCTGACAGAGGACAGCAAGAAACGCA	15 300
1	15 301	GTTACAACCTAATAAGCAATGACAGCACCTCACCCAGTACCGCAGCTGGTACCTTGCAT	15 360
2	15 301	GTTACAACCTAATAAGCAATGACAGCACCTCACCCAGTACCGCAGCTGGTACCTTGCAT	15 360
1	15 361	ACAACCTACGGCGACCCCTCAGACCGGAATCCGCTCATGGACCTGCTTGCACTCCTGACG	15 420
2	15 361	ACAACCTACGGCGACCCCTCAGACCGGAATCCGCTCATGGACCTGCTTGCACTCCTGACG	15 420
1	15 421	TAACCTGCGGCTCGGAGCAGGTCTACTGGTCGTTGCCAGACATGATGCAAGACCCGTGA	15 480
2	15 421	TAACCTGCGGCTCGGAGCAGGTCTACTGGTCGTTGCCAGACATGATGCAAGACCCGTGA	15 480
1	15 481	CCTTCCGCTCCACGCGCCAGATCAGCAACTTCCGGTGGTGGCGCCGAGCTGTTGCCCG	15 540
2	15 481	CCTTCCGCTCCACGCGCCAGATCAGCAACTTCCGGTGGTGGCGCCGAGCTGTTGCCCG	15 540

1	15 541	TGC ACT CCA AGAG GCT TCA ACG ACC AGG CGT CTA CT CCC AACT CAT CC GCA GTT TA	15 600
2	15 541	TGC ACT CCA AGAG GCT TCA ACG ACC AGG CGT CTA CT CCC AACT CAT CC GCA GTT TA	15 600
1	15 601	CCT CT CTG ACC CAC GT GT CAAT CG CT TCCC GAG AACC AG AT TT GG CG CG CC GC AG	15 660
2	15 601	CCT CT CTG ACC CAC GT GT CAAT CG CT TCCC GAG AACC AG AT TT GG CG CG CC GC AG	15 660
1	15 661	CCCC CACCAT CACC ACC CGT CAG T GAAA AC GT TT CCT GCT CT CA CAG AT CA CG GG AC GCT AC	15 720
2	15 661	CCCC CACCAT CACC ACC CGT CAG T GAAA AC GT TT CCT GCT CT CA CAG AT CA CG GG AC GCT AC	15 720
1	15 721	CG CT GCG CAAC AG CAT CG GAGG AGT CC AG CG AG T GAC CATT ACT GAC GCC AG AC GCG GCA	15 780
2	15 721	CG CT GCG CAAC AG CAT CG GAGG AGT CC AG CG AG T GAC CATT ACT GAC GCC AG AC GCG GCA	15 780
1	15 781	CCT GCCC CTAC GT TT ACA AGG CCT GGG CA TAG T CTG CC CG CG TC CT ATC GAG CC GCA	15 840
2	15 781	CCT GCCC CTAC GT TT ACA AGG CCT GGG CA TAG T CTG CC CG CG TC CT ATC GAG CC GCA	15 840
1	15 841	CTTT TGAG CAAG CAT GT CC AT CTT AT AT CG CC AG CA AT AA CA AC AGG CT GGG GCT GC	15 900
2	15 841	CTTT TGAG CAAG CAT GT CC AT CTT AT AT CG CC AG CA AT AA CA AC AGG CT GGG GCT GC	15 900
1	15 901	GCT TCCC AAG CAAG AT GT TT GG CG GGG CC AAG AAG CG CT CC GAC CA AC ACC AG T GCG CG	15 960
2	15 901	GCT TCCC AAG CAAG AT GT TT GG CG GGG CC AAG AAG CG CT CC GAC CA AC ACC AG T GCG CG	15 960
1	15 961	TG CG CG GG CACT ACC CG CG CC CT GG GG CG CG CA CA AC CG CG CC GCA CT GG CG CA CC	16 020
2	15 961	TG CG CG GG CACT ACC CG CG CC CT GG GG CG CG CA CA AC CG CG CC GCA CT GG CG CA CC	16 020
1	16 021	CCG TCG AT GAC GCC AT CG AC GCG GT GG TG AGG AGG CG CG CA ACT AC AC GCC AC GCG C	16 080
2	16 021	CCG TCG AT GAC GCC AT CG AC GCG GT GG TG AGG AGG CG CG CA ACT AC AC GCC AC GCG C	16 080
1	16 081	CAC CAG TGT CC AC AGT GG AC GCG GG CC ATT CAG ACC GT GG TG CG CG GG AG CC CG CG CT AT G	16 140
2	16 081	CAC CAG TGT CC AC AGT GG AC GCG GG CC ATT CAG ACC GT GG TG CG CG GG AG CC CG CG CT AT G	16 140

1	16 141	CTAAAATGAAGAGACGGCGGAGGCACGTAGCACGTCGCCACCGCCGCCGACCCGGCACTG	16 200
2	16 141	CTAAAATGAAGAGACGGCGGAGGCACGTAGCACGTCGCCACCGCCGCCGACCCGGCACTG	16 200
1	16 201	CCGCCAACGCGCGCGCGGCCCTGCTTAACCGCGCACGTCGACCGGCCGACGGCGG	16 260
2	16 201	CCGCCAACGCGCGCGGCCCTGCTTAACCGCGCACGTCGACCGGCCGACGGCGG	16 260
1	16 261	CCATGCGGGCCGCTCGAAGGCTGGCCGCCGGTATTGTCACTGTGCCCCCCAGGTCCAGGC	16 320
2	16 261	CCATGCGGGCCGCTCGAAGGCTGGCCGCCGGTATTGTCACTGTGCCCCCCAGGTCCAGGC	16 320
1	16 321	GACGAGCGGCCGCCGCAGCAGCCGCCATTAGTGCTATGACTCAGGGTCGCAGGGCA	16 380
2	16 321	GACGAGCGGCCGCCGCAGCAGCCGCCATTAGTGCTATGACTCAGGGTCGCAGGGCA	16 380
1	16 381	ACGTGTATTGGGTGCGCAGCTCGTTAGCGGCCCTGCGCGTGCCGTGCGCACCCGCCCC	16 440
2	16 381	ACGTGTATTGGGTGCGCAGCTCGTTAGCGGCCCTGCGCGTGCCGTGCGCACCCGCCCC	16 440
1	16 441	CGCGCAACTAGATTGCAAGAAAAACTACTTAGACTCGTACTGTTGTATGTATCCAGCGG	16 500
2	16 441	CGCGCAACTAGATTGCAAGAAAAACTACTTAGACTCGTACTGTTGTATGTATCCAGCGG	16 500
1	16 501	CGGCGCGCGAACGAAGCTATGTCCAAGCGAAATCAAAGAAGAGATGCTCCAGGTCA	16 560
2	16 501	CGGCGCGCGAACGAAGCTATGTCCAAGCGAAATCAAAGAAGAGATGCTCCAGGTCA	16 560
1	16 561	TCGCGCCGGAGATCTATGGCCCCCGAAGAAGGAAGAGCAGGATTACAAGCCCCGAAAGC	16 620
2	16 561	TCGCGCCGGAGATCTATGGCCCCCGAAGAAGGAAGAGCAGGATTACAAGCCCCGAAAGC	16 620
1	16 621	TAAAGCGGGTCAAAAGAAAAAGAAAGATGATGATGATGAACTTGACGACGAGGTGGAAC	16 680
2	16 621	TAAAGCGGGTCAAAAGAAAAAGAAAGATGATGATGATGAACTTGACGACGAGGTGGAAC	16 680
1	16 681	TGCTGCACGCTACCGCGCCCAGGCGACGGTACAGTGGAAAGGTGACGCGTAAACGTG	16 740
2	16 681	TGCTGCACGCTACCGCGCCCAGGCGACGGTACAGTGGAAAGGTGACGCGTAAACGTG	16 740

1 16 741 TTTTGCAGCCGGCACCAACCGTAGTCTTACGCCGGTGAGCGCTCCACCGCACCTACA 16 800
2 16 741 TTTTGCAGCCGGCACCAACCGTAGTCTTACGCCGGTGAGCGCTCCACCGCACCTACA 16 800

1 16 801 AGCGCGTGTATGAGGTGTACGGCAGCAGGACCTGCTTGAGCAGGCCAACGAGCGCC 16 860
2 16 801 AGCGCGTGTATGAGGTGTACGGCAGCAGGACCTGCTTGAGCAGGCCAACGAGCGCC 16 860

1 16 861 TCGGGGAGTTGCCTACGGAAAGCGGCATAAGGACATGCTGGCGTTGCCGCTGGACGAGG 16 920
2 16 861 TCGGGGAGTTGCCTACGGAAAGCGGCATAAGGACATGCTGGCGTTGCCGCTGGACGAGG 16 920

1 16 921 GCAACCCAACACCTAGCCTAAAGCCC GTAACACTGCAGCAGGTGCTGCCGCGCTTCAC 16 980
2 16 921 GCAACCCAACACCTAGCCTAAAGCCC GTAACACTGCAGCAGGTGCTGCCGCGCTTCAC 16 980

1 16 981 CGTCCGAAGAAAAGCGCGGCCCTAAAGCGCGAGTCTGGTACTTGGCACCCACCGTGCAGC 17 040
2 16 981 CGTCCGAAGAAAAGCGCGGCCCTAAAGCGCGAGTCTGGTACTTGGCACCCACCGTGCAGC 17 040

1 17 041 TGATGGTACCCAAGCGCCAGCGACTGGAAAGATGTCTTGGAAAAAATGACCGTGGAACCTG 17 100
2 17 041 TGATGGTACCCAAGCGCCAGCGACTGGAAAGATGTCTTGGAAAAAATGACCGTGGAACCTG 17 100

1 17 101 GGCTGGAGCCGAGGTCCCGGTGCGGCCAATCAAGCAGGTGGCGCCGGACTGGCGTGC 17 160
2 17 101 GGCTGGAGCCGAGGTCCCGGTGCGGCCAATCAAGCAGGTGGCGCCGGACTGGCGTGC 17 160

1 17 161 AGACCGTGGACGTTAGATACCCACTACCAGTAGCACCAAGTATTGCCACCGGCCACAGAGG 17 220
2 17 161 AGACCGTGGACGTTAGATACCCACTACCAGTAGCACCAAGTATTGCCACCGGCCACAGAGG 17 220

1 17 221 GCATGGAGACACAAACGTCCCCGGTTGCCTCAGCGGTGGCGGATGCCCGGGTGCAGGC 17 280
2 17 221 GCATGGAGACACAAACGTCCCCGGTTGCCTCAGCGGTGGCGGATGCCCGGGTGCAGGC 17 280

1 17 281 TCGCTCGGGCCCGTCCAAGACCTCTACGGAGGTGCAAACGGACCCGTGGATTTCGCG 17 340
2 17 281 TCGCTCGGGCCCGTCCAAGACCTCTACGGAGGTGCAAACGGACCCGTGGATTTCGCG 17 340

1	17 341	TTTCAGCCCCCGGCGCCCGGCCGTTCGAGGAAGTACGGCGCCAGCGCGCTACTGC	17 400
2	17 341	TTTCAGCCCCCGGCGCCCGGCCGTTCGAGGAAGTACGGCGCCAGCGCGCTACTGC	17 400
1	17 401	CCGAATATGCCCTACATCCTTCCATTGCGCCTACCCCCGGCTATCGTGGCTACACCTACC	17 460
2	17 401	CCGAATATGCCCTACATCCTTCCATTGCGCCTACCCCCGGCTATCGTGGCTACACCTACC	17 460
1	17 461	GCCCCAGAACGAGCAACTACCCGACGCCAACCACTGGAACCCGCCGCCGT	17 520
2	17 461	GCCCCAGAACGAGCAACTACCCGACGCCAACCACTGGAACCCGCCGCCGT	17 520
1	17 521	GCCGTCGCCAGCCCCGTGCTGGCCCCGATTCGTGCGCAGGGTGGCTCGCGAAGGAGGCA	17 580
2	17 521	GCCGTCGCCAGCCCCGTGCTGGCCCCGATTCGTGCGCAGGGTGGCTCGCGAAGGAGGCA	17 580
1	17 581	GGACCTGGTGCTGCCAACAGCGCGCTACCAACCCAGCATCGTTAAAAGCCGGTCTTG	17 640
2	17 581	GGACCTGGTGCTGCCAACAGCGCGCTACCAACCCAGCATCGTTAAAAGCCGGTCTTG	17 640
1	17 641	TGGTTCTTGCAGATATGCCCTCACCTGCCGCTCCGTTCCGGTGCCGGATTCCGAG	17 700
2	17 641	TGGTTCTTGCAGATATGCCCTCACCTGCCGCTCCGTTCCGGTGCCGGATTCCGAG	17 700
1	17 701	GAAGAATGCACCGTAGGAGGGCATGGCCGCCACGGCCTGACGGGGCATGCGTCGTG	17 760
2	17 701	GAAGAATGCACCGTAGGAGGGCATGGCCGCCACGGCCTGACGGGGCATGCGTCGTG	17 760
1	17 761	CGCACCAACGGCGGGCGCGCGTCGCACCGTCGCATGCGGGCGGTATCCTGCCCTCC	17 820
2	17 761	CGCACCAACGGCGGGCGCGCGTCGCACCGTCGCATGCGGGCGGTATCCTGCCCTCC	17 820
1	17 821	TTATTCCACTGATGCCGCGGCGATTGGCGCCGTGCCCGGAATTGCATCCGTGGCCTTGC	17 880
2	17 821	TTATTCCACTGATGCCGCGGCGATTGGCGCCGTGCCCGGAATTGCATCCGTGGCCTTGC	17 880
1	17 881	AGGCAGAGACACTGATTAAAAACAAGTTGCATGTGGAAAATCAAATAAAAGTCTG	17 940
2	17 881	AGGCAGAGACACTGATTAAAAACAAGTTGCATGTGGAAAATCAAATAAAAGTCTG	17 940

1	17 941	GA C T C T C A C G C T C G C T T G G C C T G T A A C T A T T T G T A G A A T G G A A G A C A T C A A C T T T G C G	18 000
2	17 941	GA C T C T C A C G C T C G C T T G G C C T G T A A C T A T T T G T A G A A T G G A A G A C A T C A A C T T T G C G	18 000
1	18 001	T C T C T G G C C C C G C G A C A C G G C T C G C G C C C G T T C A T G G G A A C T G G C A A G A T A T C G G C A C C	18 060
2	18 001	T C T C T G G C C C C G C G A C A C G G C T C G C G C C C G T T C A T G G G A A C T G G C A A G A T A T C G G C A C C	18 060
1	18 061	A G C A A T A T G A G C G G T G G C G C C T T C A G C T G G G G C T C G C T G T G G A G C G G C A T T A A A A A T T C	18 120
2	18 061	A G C A A T A T G A G C G G T G G C G C C T T C A G C T G G G G C T C G C T G T G G A G C G G C A T T A A A A A T T C	18 120
1	18 121	G G T T C C A C C G T T A A G A A C T A T G G C A G C A A G G C C T G G A A C A G C A G C A C A G G C C A G A T G C T G	18 180
2	18 121	G G T T C C A C C G T T A A G A A C T A T G G C A G C A A G G C C T G G A A C A G C A G C A C A G G C C A G A T G C T G	18 180
1	18 181	A G G G A T A A G T T G A A A G A G C A A A A T T C C A A C A A A A G G T G G T A G A T G G C C T G G C C T C T G G C	18 240
2	18 181	A G G G A T A A G T T G A A A G A G C A A A A T T C C A A C A A A A G G T G G T A G A T G G C C T G G C C T C T G G C	18 240
1	18 241	A T T A G C G G G G T G G T G G A C C T G G C C A A C C A G G C A G T G C A A A A T A A G A T T A A C A G T A A G C T T	18 300
2	18 241	A T T A G C G G G G T G G T G G A C C T G G C C A A C C A G G C A G T G C A A A A T A A G A T T A A C A G T A A G C T T	18 300
1	18 301	G A T C C C C G C C C T C C C G T A G A G G A G C C T C C A C C G G C C G T G G A G A C A G T G T C T C C A G A G G G G	18 360
2	18 301	G A T C C C C G C C C T C C C G T A G A G G A G C C T C C A C C G G C C G T G G A G A C A G T G T C T C C A G A G G G G	18 360
1	18 361	C G T G G C G A A A A G C G T C C G C G C C C G A C A G G G A A G A A A C T C T G G T G A C G C A A A T A G A C G A G	18 420
2	18 361	C G T G G C G A A A A G C G T C C G C G C C C G A C A G G G A A G A A A C T C T G G T G A C G C A A A T A G A C G A G	18 420
1	18 421	C C T C C C T C G T A C G A G G A G G C A C T A A A G C A A G G C C T G C C C A C C A C C C G T C C C A T C G C G C C C	18 480
2	18 421	C C T C C C T C G T A C G A G G A G G C A C T A A A G C A A G G C C T G C C C A C C A C C C G T C C C A T C G C G C C C	18 480
1	18 481	A T G G C T A C C G G A G T G C T G G G C C A G C A C A C A C C C G T A A C G C T G G A C C T G C C T C C C C C G C C	18 540
2	18 481	A T G G C T A C C G G A G T G C T G G G C C A G C A C A C A C C C G T A A C G C T G G A C C T G C C T C C C C C G C C	18 540

1	18 541	GACACCCAGCAGAACCTGTGCTGCCAGGCCGACCGCCGTTGTTGTAACCGTCCTAGC	18 600
2	18 541	GACACCCAGCAGAACCTGTGCTGCCAGGCCGACCGCCGTTGTTGTAACCGTCCTAGC	18 600
1	18 601	CGCGCGTCCCTGCGCCGCGCCGCCAGCGGTCCCGCATCGTTGC GGCCGTAGCCAGTGGC	18 660
2	18 601	CGCGCGTCCCTGCGCCGCGCCGCCAGCGGTCCCGCATCGTTGC GGCCGTAGCCAGTGGC	18 660
1	18 661	AACTGGCAAAGCACACTGAACAGCATCGTGGGTCTGGGGTGCAATCCCTGAAGCGCCGA	18 720
2	18 661	AACTGGCAAAGCACACTGAACAGCATCGTGGGTCTGGGGTGCAATCCCTGAAGCGCCGA	18 720
1	18 721	CGATGCTTCTGATA GCTAACGTGTCGTATGTGTGTCATGTATCGTCCATGTCGCCGCCA	18 780
2	18 721	CGATGCTTCTGATA GCTAACGTGTCGTATGTGTGTCATGTATCGTCCATGTCGCCGCCA	18 780
1	18 781	GAGGAGCTGCTGAGCCGCCGCGGCCGCTTCCAAGATGGCTACCCCTTCGATGATGCC	18 840
2	18 781	GAGGAGCTGCTGAGCCGCCGCGGCCGCTTCCAAGATGGCTACCCCTTCGATGATGCC	18 840
1	18 841	GCAGTGGTCTTACATGCACATCTCGGGCCAGGACGCCCTGGAGTACCTGAGCCCCGGGCT	18 900
2	18 841	GCAGTGGTCTTACATGCACATCTCGGGCCAGGACGCCCTGGAGTACCTGAGCCCCGGGCT	18 900
1	18 901	GGTGCAGTTGCCGCCACCGAGACGTACTTCAGCCTGAATAACAAGTTAGAAACCC	18 960
2	18 901	GGTGCAGTTGCCGCCACCGAGACGTACTTCAGCCTGAATAACAAGTTAGAAACCC	18 960
1	18 961	CACGGTGGCGCCTACGCACGACGTGACCACAGACCGGTCCCAGCGTTGACGCTGCGGTT	19 020
2	18 961	CACGGTGGCGCCTACGCACGACGTGACCACAGACCGGTCCCAGCGTTGACGCTGCGGTT	19 020
1	19 021	CATCCCTGTGGACCGTGAGGATACTGCGTACTCGTACAAGGCGCGGTTACCCTAGCTGT	19 080
2	19 021	CATCCCTGTGGACCGTGAGGATACTGCGTACTCGTACAAGGCGCGGTTACCCTAGCTGT	19 080
1	19 081	GGGTGATAACCGTGTGGACATGGCTTCCACGTACTTGTACATCCGGCGGTGCTGGA	19 140
2	19 081	GGGTGATAACCGTGTGGACATGGCTTCCACGTACTTGTACATCCGGCGGTGCTGGA	19 140

1	19 141	CAGGGGCCCTACTTTAAGCCCTACTCTGGCACTGCCTACAAACGCCCTGGCTCCCAAGGG	19 200
2	19 141	CAGGGGCCCTACTTTAAGCCCTACTCTGGCACTGCCTACAAACGCCCTGGCTCCCAAGGG	19 200
1	19 201	TGCCCCAAATCCTTGCATGGATGAAGCTGCTACTGCTCTGAAATAAACCTAGAAGA	19 260
2	19 201	TGCCCCAAATCCTTGCATGGATGAAGCTGCTACTGCTCTGAAATAAACCTAGAAGA	19 260
1	19 261	AGAGGACGATGACAACGAAGACGAAGTAGACGAGCAAGCTGAGCAGCAAAAAACTCACGT	19 320
2	19 261	AGAGGACGATGACAACGAAGACGAAGTAGACGAGCAAGCTGAGCAGCAAAAAACTCACGT	19 320
1	19 321	ATTTGGGCAGGCGCCTTATTCTGGTATAAATATTACAAAGGAGGGTATTCAAATAGGTGT	19 380
2	19 321	ATTTGGGCAGGCGCCTTATTCTGGTATAAATATTACAAAGGAGGGTATTCAAATAGGTGT	19 380
1	19 381	CGAAGGTCAAACACCTAAATATGCCGATAAAACATTCAACCTGAACCTCAAATAGGAGA	19 440
2	19 381	CGAAGGTCAAACACCTAAATATGCCGATAAAACATTCAACCTGAACCTCAAATAGGAGA	19 440
1	19 441	ATCTCAGTGGTACGAAACAGAAATTAATCATGCAGCTGGAGAGTCCTAAAAAGACTAC	19 500
2	19 441	ATCTCAGTGGTACGAAACAGAAATTAATCATGCAGCTGGAGAGTCCTAAAAAGACTAC	19 500
1	19 501	CCCAATGAAACCATGTTACGGTTCATATGCAAAACCCACAAATGAAAATGGAGGGCAAGG	19 560
2	19 501	CCCAATGAAACCATGTTACGGTTCATATGCAAAACCCACAAATGAAAATGGAGGGCAAGG	19 560
1	19 561	CATTCTTGTAAAGCAACAAAATGGAAAGCTAGAAAGTCAAGTGGAAATGCAATTTCCTC	19 620
2	19 561	CATTCTTGTAAAGCAACAAAATGGAAAGCTAGAAAGTCAAGTGGAAATGCAATTTCCTC	19 620
1	19 621	AACTACTGAGGCAGCCGCAGGCAATGGTATACTTGACTCCTAAAGTGGTATTGTACAG	19 680
2	19 621	AACTACTGAGGCAGCCGCAGGCAATGGTATACTTGACTCCTAAAGTGGTATTGTACAG	19 680
1	19 681	TGAAGATGTAGATATAGAAACCCAGACACTCATATTCTTACATGCCCACTATTAAGGA	19 740
2	19 681	TGAAGATGTAGATATAGAAACCCAGACACTCATATTCTTACATGCCCACTATTAAGGA	19 740

1	19 741	AGGTAAC T CAC GAG AACT AAT GGGCC AACA AT CT AT GCCA ACAG GCCT AATT AC ATT GC	19 800
2	19 741	AGGTAAC T CAC GAG AACT AAT GGGCC AACA AT CT AT GCCA ACAG GCCT AATT AC ATT GC	19 800
1	19 801	TTT TAGGGACA ATT TATT GGTCT AAT GTATT ACAC AGC AC GGG TAAT AT GGGT GTT CT	19 860
2	19 801	TTT TAGGGACA ATT TATT GGTCT AAT GTATT ACAC AGC AC GGG TAAT AT GGGT GTT CT	19 860
1	19 861	GGCGGGCCAAGCATCGCAGTTGAATGCTGTTGAGATTGCAAGACAGAAACACAGAGCT	19 920
2	19 861	GGCGGGCCAAGCATCGCAGTTGAATGCTGTTGAGATTGCAAGACAGAAACACAGAGCT	19 920
1	19 921	TTCATACCAGCTTGCTTGATTCCATTGGT GATA GA ACCAGGT ACT TTTCTATGTGGAA	19 980
2	19 921	TTCATACCAGCTTGCTTGATTCCATTGGT GATA GA ACCAGGT ACT TTTCTATGTGGAA	19 980
1	19 981	TCAGGGCTGTTGACAGCTATGATCCAGATGTTAGAATTATTGAAAATCATGGAAC TGAA GA	20 040
2	19 981	TCAGGGCTGTTGACAGCTATGATCCAGATGTTAGAATTATTGAAAATCATGGAAC TGAA GA	20 040
1	20 041	TGAAC TTCCA AATTACTGCTTCCACTGGGAGGTGTGATTAATACAGAGACTCTTACCAA	20 100
2	20 041	TGAAC TTCCA AATTACTGCTTCCACTGGGAGGTGTGATTAATACAGAGACTCTTACCAA	20 100
1	20 101	GGTAAAACCTAACAGGT CAGGAAAATGGATGGAAAAAGATGCTACAGAATTTCAGA	20 160
2	20 101	GGTAAAACCTAACAGGT CAGGAAAATGGATGGAAAAAGATGCTACAGAATTTCAGA	20 160
1	20 161	TAAAAATGAAATAAGAGTTGGAAATAATTTGCCATGGAAATCAATCTAAATGCCAACCT	20 220
2	20 161	TAAAAATGAAATAAGAGTTGGAAATAATTTGCCATGGAAATCAATCTAAATGCCAACCT	20 220
1	20 221	GTGGAGAAATT CCTGTACTCC AACATAGCGCTGTAT TGC CCGACAAGCTAAAGTACAG	20 280
2	20 221	GTGGAGAAATT CCTGTACTCC AACATAGCGCTGTAT TGC CCGACAAGCTAAAGTACAG	20 280
1	20 281	TCCTTCCAACGTAAAATTCTGATAACCCAAACACCTACGACTACATGAACAGCGAGT	20 340
2	20 281	TCCTTCCAACGTAAAATTCTGATAACCCAAACACCTACGACTACATGAACAGCGAGT	20 340

1 20 341 GGTGGCTCCGGCTAGTGGACTGCTACATTAACCTGGAGCACGCTGGTCCCTGACTA 20 400
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 341 GGTGGCTCCGGCTAGTGGACTGCTACATTAACCTGGAGCACGCTGGTCCCTGACTA 20 400

1 20 401 TATGGACAACGTCAACCCATTAAACCACCAACCGCAATGCTGGCCTGCCTACCGCTCAAT 20 460
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 401 TATGGACAACGTCAACCCATTAAACCACCAACCGCAATGCTGGCCTGCCTACCGCTCAAT 20 460

1 20 461 GTTGCTGGCAATGGTCGCTATGTGCCCTCCACATCCAGGTGCCTCAGAAGTTCTTG 20 520
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 461 GTTGCTGGCAATGGTCGCTATGTGCCCTCCACATCCAGGTGCCTCAGAAGTTCTTG 20 520

1 20 521 CATTAAAAACCTCCTCTCCTGCCGGCTCATACACCTACGAGTGGAACTTCAGGAAGGA 20 580
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 521 CATTAAAAACCTCCTCTCCTGCCGGCTCATACACCTACGAGTGGAACTTCAGGAAGGA 20 580

1 20 581 TGTTAACATGGTTCTGCAGAGCTCCCTAGGAAATGACCTAAGGGTTACGGAGCCAGCAT 20 640
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 581 TGTTAACATGGTTCTGCAGAGCTCCCTAGGAAATGACCTAAGGGTTACGGAGCCAGCAT 20 640

1 20 641 TAAGTTGATAGCATTGCCCTTACGCCACCTTCTCCCCATGGCCACAAACACCGCCTC 20 700
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 641 TAAGTTGATAGCATTGCCCTTACGCCACCTTCTCCCCATGGCCACAAACACCGCCTC 20 700

1 20 701 CACGCTTGAGGCCATGCTTAGAAACGACACCAACGACCAAGTCCTTAACGACTATCTCTC 20 760
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 701 CACGCTTGAGGCCATGCTTAGAAACGACACCAACGACCAAGTCCTTAACGACTATCTCTC 20 760

1 20 761 CGCCGCCAACATGCTCTACCCATACCGCCAACGCTACCAACGTGCCATATCCATCCC 20 820
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 761 CGCCGCCAACATGCTCTACCCATACCGCCAACGCTACCAACGTGCCATATCCATCCC 20 820

1 20 821 CTCCCGCAACTGGCGGCTTCCGCGGCTGGCCTTCACGCGCCTTAAGACTAAGGAAAC 20 880
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 821 CTCCCGCAACTGGCGGCTTCCGCGGCTGGCCTTCACGCGCCTTAAGACTAAGGAAAC 20 880

1 20 881 CCCATCACTGGCGTGGCTACGACCCATTACACCTACTCTGGCTCTATAACCTACCT 20 940
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 20 881 CCCATCACTGGCGTGGCTACGACCCATTACACCTACTCTGGCTCTATAACCTACCT 20 940

1	20 941	AGATGGAACCTTTACCTCAACCACACCTTAAGAAGGTGGCCATTACCTTGACTCTTC	21 000
2	20 941	AGATGGAACCTTTACCTCAACCACACCTTAAGAAGGTGGCCATTACCTTGACTCTTC	21 000
1	21 001	TGTCAGCTGGCCTGGCAATGACCGCCTGCTTACCCCCAACGAGTTGAAATTAAGCGCTC	21 060
2	21 001	TGTCAGCTGGCCTGGCAATGACCGCCTGCTTACCCCCAACGAGTTGAAATTAAGCGCTC	21 060
1	21 061	AGTTGACGGGGAGGGTTACAACGTTGCCAGTGTAAACATGACCAAAGACTGGTTCCTGGT	21 120
2	21 061	AGTTGACGGGGAGGGTTACAACGTTGCCAGTGTAAACATGACCAAAGACTGGTTCCTGGT	21 120
1	21 121	ACAAATGCTAGCTAACTATAACATTGGCTACCAGGGCTTCTATATCCCAGAGAGCTACAA	21 180
2	21 121	ACAAATGCTAGCTAACTATAACATTGGCTACCAGGGCTTCTATATCCCAGAGAGCTACAA	21 180
1	21 181	GGACCGCATGTACTCCTTCTTAGAAACTTCCAGGCCATGAGCCGTAGGTGGTGGATGA	21 240
2	21 181	GGACCGCATGTACTCCTTCTTAGAAACTTCCAGGCCATGAGCCGTAGGTGGTGGATGA	21 240
1	21 241	TACTAAATACAAGGACTACCAACAGGTGGCATCCTACACCAACACAACACTCTGGATT	21 300
2	21 241	TACTAAATACAAGGACTACCAACAGGTGGCATCCTACACCAACACAACACTCTGGATT	21 300
1	21 301	TGTTGGCTACCTTGCACCATGCGCGAAGGACAGGCCTACCCGTAACTTCCCTA	21 360
2	21 301	TGTTGGCTACCTTGCACCATGCGCGAAGGACAGGCCTACCCGTAACTTCCCTA	21 360
1	21 361	TCCGCTTATAGGCAAGACCGCAGTTGACAGCATTACCCAGAAAAAGTTCTTGCATCG	21 420
2	21 361	TCCGCTTATAGGCAAGACCGCAGTTGACAGCATTACCCAGAAAAAGTTCTTGCATCG	21 420
1	21 421	CACCCCTTGGCGCATCCATTCTCCAGTAACTTATGTCATGGCGCACTCACAGACCT	21 480
2	21 421	CACCCCTTGGCGCATCCATTCTCCAGTAACTTATGTCATGGCGCACTCACAGACCT	21 480
1	21 481	GGGCCAAACCTCTACGCCAACCTCGCCCCACGCGCTAGACATGACTTTGAGGTGGA	21 540
2	21 481	GGGCCAAACCTCTACGCCAACCTCGCCCCACGCGCTAGACATGACTTTGAGGTGGA	21 540

1	21 541	TCCCATGGACGAGCCCACCCCTTCTTATGTTTGTTGAAGTC TTGACGTGGTCCGTGT	21 600
2	21 541	TCCCATGGACGAGCCCACCCCTTCTTATGTTTGTTGAAGTC TTGACGTGGTCCGTGT	21 600
1	21 601	GCACCAGCGCACCGCGCGTACATGAAACCGTGTACCTGCGCACGCCCTCTCGGCCGG	21 660
2	21 601	GCACCAGCGCACCGCGCGTACATGAAACCGTGTACCTGCGCACGCCCTCTCGGCCGG	21 660
1	21 661	CAACGCCACAACATAAAGAACAGCAACATCAACAAACAGCTGCCGCCATGGGCTCCAGT	21 720
2	21 661	CAACGCCACAACATAAAGAACAGCAACATCAACAAACAGCTGCCGCCATGGGCTCCAGT	21 720
1	21 721	GAGCAGGAACTGAAAGCCATTGTCAAAGATCTTGGTTGTGGGCCATATTTTGGCACC	21 780
2	21 721	GAGCAGGAACTGAAAGCCATTGTCAAAGATCTTGGTTGTGGGCCATATTTTGGCACC	21 780
1	21 781	TATGACAAGCGCTTCCAGGCTTGTCCACACAAGCTCGCCTGCGCCATAGTCAT	21 840
2	21 781	TATGACAAGCGCTTCCAGGCTTGTCCACACAAGCTCGCCTGCGCCATAGTCAT	21 840
1	21 841	ACGGCCGGTCGCGAGACTGGGGCGTACACTGGATGGCCTTGCCTGGAACCCGCACTCA	21 900
2	21 841	ACGGCCGGTCGCGAGACTGGGGCGTACACTGGATGGCCTTGCCTGGAACCCGCACTCA	21 900
1	21 901	AAAACATGCTACCTCTTGAGCCCTTGGCTTTCTGACCAGCGACTCAAGCAGGTTAC	21 960
2	21 901	AAAACATGCTACCTCTTGAGCCCTTGGCTTTCTGACCAGCGACTCAAGCAGGTTAC	21 960
1	21 961	CAGTTGAGTACGAGTCACTCCTGCGCCGTAGCGCATTGCTTCTTCCCCGACCGCTGT	22 020
2	21 961	CAGTTGAGTACGAGTCACTCCTGCGCCGTAGCGCATTGCTTCTTCCCCGACCGCTGT	22 020
1	22 021	ATAACGCTGGAAAAGTCCACCCAAAGCGTACAGGGGCCAACTCGGCCGCTGTGGACTA	22 080
2	22 021	ATAACGCTGGAAAAGTCCACCCAAAGCGTACAGGGGCCAACTCGGCCGCTGTGGACTA	22 080
1	22 081	TTCTGCTGCATGTTCTCACGCCCTTGCCAAGTGGCCCCAACTCCCATGGATCACAAAC	22 140
2	22 081	TTCTGCTGCATGTTCTCACGCCCTTGCCAAGTGGCCCCAACTCCCATGGATCACAAAC	22 140

1	22 141 CCCACCATGAAACCTTATTACCGGGGTACCCAACTCCATGCTAACAGTCCCCAGGTACAG	22 200
2	22 141 CCCACCATGAAACCTTATTACCGGGGTACCCAACTCCATGCTAACAGTCCCCAGGTACAG	22 200
1	22 201 CCCACCCTGCGTCGCAACCAGGAACAGCTCTACAGCTTGGAGCGCCACTGCCCTAC	22 260
2	22 201 CCCACCCTGCGTCGCAACCAGGAACAGCTCTACAGCTTGGAGCGCCACTGCCCTAC	22 260
1	22 261 TTCCGCAGCCACAGTGCGCAGATTAGGAGCGCCACTTCTTTGTCACTTGAAAAACATG	22 320
2	22 261 TTCCGCAGCCACAGTGCGCAGATTAGGAGCGCCACTTCTTTGTCACTTGAAAAACATG	22 320
1	22 321 TAAAAATAATGTACTAGAGACACTTCAATAAGGCAAATGCTTTATTTGTACACTCTC	22 380
2	22 321 TAAAAATAATGTACTAGAGACACTTCAATAAGGCAAATGCTTTATTTGTACACTCTC	22 380
1	22 381 GGGTGATTATTTACCCCCACCCCTGCCGTCTGCCGTTAAAAATCAAAGGGTTCTGC	22 440
2	22 381 GGGTGATTATTTACCCCCACCCCTGCCGTCTGCCGTTAAAAATCAAAGGGTTCTGC	22 440
1	22 441 CGCGCATCGCTATGCCCACTGGCAGGGACACGTTGCGATACTGGTGTAGTGCTCCAC	22 500
2	22 441 CGCGCATCGCTATGCCCACTGGCAGGGACACGTTGCGATACTGGTGTAGTGCTCCAC	22 500
1	22 501 TTAAACTCAGGCACAACCATCCGGCAGCTCGGTGAAGTTTCACTCCACAGGCTGCGC	22 560
2	22 501 TTAAACTCAGGCACAACCATCCGGCAGCTCGGTGAAGTTTCACTCCACAGGCTGCGC	22 560
1	22 561 ACCATCACCAACCGCGTTAGCAGGTGGCGCCGATATCTTGAAGTCGCAGTTGGGCCT	22 620
2	22 561 ACCATCACCAACCGCGTTAGCAGGTGGCGCCGATATCTTGAAGTCGCAGTTGGGCCT	22 620
1	22 621 CCGCCCTGCGCGCGAGTTGCGATAACACAGGGTTGCAGCACTGGAACACTATCAGCGCC	22 680
2	22 621 CCGCCCTGCGCGCGAGTTGCGATAACACAGGGTTGCAGCACTGGAACACTATCAGCGCC	22 680
1	22 681 GGGTGGTGCACGCTGGCCAGCACGCTTGTGGAGATCAGATCCGCGTCCAGGTCTCC	22 740
2	22 681 GGGTGGTGCACGCTGGCCAGCACGCTTGTGGAGATCAGATCCGCGTCCAGGTCTCC	22 740

1	22 741	GCGTTGCTCAGGGCGAACGGAGTCACACTTGGTAGCTGCCTCCAAAAAGGGCGCGTGC	22 800
2	22 741	GCGTTGCTCAGGGCGAACGGAGTCACACTTGGTAGCTGCCTCCAAAAAGGGCGCGTGC	22 800
1	22 801	CCAGGGCTTGAGTTGCACTCGCACCGTAGTGGCATAAAAGGTGACCGTGCCGGTCTGG	22 860
2	22 801	CCAGGGCTTGAGTTGCACTCGCACCGTAGTGGCATAAAAGGTGACCGTGCCGGTCTGG	22 860
1	22 861	GCGTTAGGATACAGCGCCTGCATAAAAGCCTTGATCTGCTTAAAGCCACCTGAGCCTTT	22 920
2	22 861	GCGTTAGGATACAGCGCCTGCATAAAAGCCTTGATCTGCTTAAAGCCACCTGAGCCTTT	22 920
1	22 921	GCGCCTTCAGAGAAGAACATGCCGCAAGACTTGCCGGAAACTGATTGGCCGGACAGGCC	22 980
2	22 921	GCGCCTTCAGAGAAGAACATGCCGCAAGACTTGCCGGAAACTGATTGGCCGGACAGGCC	22 980
1	22 981	GCGTCGTGCACGCAGCACCTTGCCTCGGTGTTGGAGATCTGCACCACATTCGGCCCCAC	23 040
2	22 981	GCGTCGTGCACGCAGCACCTTGCCTCGGTGTTGGAGATCTGCACCACATTCGGCCCCAC	23 040
1	23 041	CGGTTCTTCACGATCTTGGCCTTGCTAGACTGCTCCTTCAGCGCGCGCTGCCGTTTCG	23 100
2	23 041	CGGTTCTTCACGATCTTGGCCTTGCTAGACTGCTCCTTCAGCGCGCGCTGCCGTTTCG	23 100
1	23 101	CTCGTCACATCCATTCAATCACGTGCTCCTTATTATCATAATGCTTCCGTGTAGACAC	23 160
2	23 101	CTCGTCACATCCATTCAATCACGTGCTCCTTATTATCATAATGCTTCCGTGTAGACAC	23 160
1	23 161	TTAAGCTCGCCTTCGATCTCAGCGCAGCGGTGCAGCCACAACGCGCAGCCGTGGCTCG	23 220
2	23 161	TTAAGCTCGCCTTCGATCTCAGCGCAGCGGTGCAGCCACAACGCGCAGCCGTGGCTCG	23 220
1	23 221	TGATGCTTGTAGGTACCCCTGCAAACGACTGCAGGTACGCCCTGCAGGAATGCCCATC	23 280
2	23 221	TGATGCTTGTAGGTACCCCTGCAAACGACTGCAGGTACGCCCTGCAGGAATGCCCATC	23 280
1	23 281	ATCGTCACAAAGGTCTTGGTAGCTGGTAAGGTCAAGCTGCAACCCGCGGTGCTCCTCGTTC	23 340
2	23 281	ATCGTCACAAAGGTCTTGGTAGCTGGTAAGGTCAAGCTGCAACCCGCGGTGCTCCTCGTTC	23 340

1 23 341 AGCCAGGTCTTGCATACGGCCGCCAGAGCTTCCACTTGGTCAGGCAGTAGTTGAAGTTC 23 400
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 341 AGCCAGGTCTTGCATACGGCCGCCAGAGCTTCCACTTGGTCAGGCAGTAGTTGAAGTTC 23 400

1 23 401 GCCTTAGATCGTTATCCACGTGGTACTTGTCCATCAGCGCGCGCAGCCTCCATGCC 23 460
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 401 GCCTTAGATCGTTATCCACGTGGTACTTGTCCATCAGCGCGCGCAGCCTCCATGCC 23 460

1 23 461 TTCTCCCACGCAGACACGATCGGCACACTCAGCGGGTTCATCACCGTAATTCACTTCC 23 520
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 461 TTCTCCCACGCAGACACGATCGGCACACTCAGCGGGTTCATCACCGTAATTCACTTCC 23 520

1 23 521 GCTTCGCTGGGCTTTCTCTTGCCTCGGCATACCACGCGCCACTGGGTCGTCT 23 580
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 521 GCTTCGCTGGGCTTTCTCTTGCCTCGGCATACCACGCGCCACTGGGTCGTCT 23 580

1 23 581 TCATTCAGCCGCCGCACTGTGCGCTTACCTCCTTGCATGCTTGATTAGCACCGGTGG 23 640
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 581 TCATTCAGCCGCCGCACTGTGCGCTTACCTCCTTGCATGCTTGATTAGCACCGGTGG 23 640

1 23 641 TTGCTGAAACCCACCATTGTAGCGCCACATCTCTCTTCTCGCTGTCCACGATT 23 700
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 641 TTGCTGAAACCCACCATTGTAGCGCCACATCTCTCTTCTCGCTGTCCACGATT 23 700

1 23 701 ACCTCTGGTGATGGCGGGCGCTCGGGCTTGGAGAAGGGCGCTTCTTTCTTGGC 23 760
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 701 ACCTCTGGTGATGGCGGGCGCTCGGGCTTGGAGAAGGGCGCTTCTTTCTTGGC 23 760

1 23 761 GCAATGGCAAATCCGCCGCCGAGGTCGATGGCGGGCTGGGTGTGCGCGGACCA 23 820
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 761 GCAATGGCAAATCCGCCGCCGAGGTCGATGGCGGGCTGGGTGTGCGCGGACCA 23 820

1 23 821 GCGTCTTGTGATGAGTCTTCTCGTCCTCGGACTCGATAACGCCGCCTCATCCGCT 23 880
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 821 GCGTCTTGTGATGAGTCTTCTCGTCCTCGGACTCGATAACGCCGCCTCATCCGCT 23 880

1 23 881 GGGGGCGCCGGGAGGCGGCGACGGGACGGGACGACACGTCCATGGTTGGG 23 940
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 23 881 GGGGGCGCCGGGAGGCGGCGACGGGACGGGACGACACGTCCATGGTTGGG 23 940

1 23 941 GGACGTCGCCGCACCGCGTCCGCGCTCGGGGTGGTTCGCGCTGCTCCTCTTCCC 24 000
2 23 941 GGACGTCGCCGCACCGCGTCCGCGCTCGGGGTGGTTCGCGCTGCTCCTCTTCCC 24 000

1 24 001 CTGGCCATTCCTCTCCTATAGGCAGAAAAAGATCATGGAGTCAGTCGAGAAGAAGGAC 24 060
2 24 001 CTGGCCATTCCTCTCCTATAGGCAGAAAAAGATCATGGAGTCAGTCGAGAAGAAGGAC 24 060

1 24 061 AGCCTAACCGCCCCCTTGAGTTGCCACCAACCGCCTCCACCGATGCCGCCAACGCGCCT 24 120
2 24 061 AGCCTAACCGCCCCCTTGAGTTGCCACCAACCGCCTCCACCGATGCCGCCAACGCGCCT 24 120

1 24 121 ACCACCTTCCCCGTCGAGGCACCCCGCTTGAGGAGGAAGTGATTATCGAGCAGGAC 24 180
2 24 121 ACCACCTTCCCCGTCGAGGCACCCCGCTTGAGGAGGAAGTGATTATCGAGCAGGAC 24 180

1 24 181 CCAGGTTTGTAAGCGAACGACGAGGACCGCTCAGTACCAACAGAGGATAAAAAGCAA 24 240
2 24 181 CCAGGTTTGTAAGCGAACGACGAGGACCGCTCAGTACCAACAGAGGATAAAAAGCAA 24 240

1 24 241 GACCAGGACAACGCAGAGGCAAACGAGGAACAAGTCGGCGGGGGACGAAAGGCATGGC 24 300
2 24 241 GACCAGGACAACGCAGAGGCAAACGAGGAACAAGTCGGCGGGGGACGAAAGGCATGGC 24 300

1 24 301 GACTACCTAGATGTGGGAGACGACGTGCTGTTGAAGCATTGCAGCGCCAGTGC 24 360
2 24 301 GACTACCTAGATGTGGGAGACGACGTGCTGTTGAAGCATTGCAGCGCCAGTGC 24 360

1 24 361 ATCTCGACCGCTTGCAAGAGCGCAGCGATGTGCCCTGCCATAGCGGATGTCAGC 24 420
2 24 361 ATCTCGACCGCTTGCAAGAGCGCAGCGATGTGCCCTGCCATAGCGGATGTCAGC 24 420

1 24 421 GCCTACGAACGCCACCTATTCTACCGCGCGTACCCCCAAACGCCAAGAAAACGGCACA 24 480
2 24 421 GCCTACGAACGCCACCTATTCTACCGCGCGTACCCCCAAACGCCAAGAAAACGGCACA 24 480

1 24 481 TCGAGCCAAACCGCGCCTCAACTTCTACCCGTATTCGGCGTGCAGAGGTGCTTG 24 540
2 24 481 TCGAGCCAAACCGCGCCTCAACTTCTACCCGTATTCGGCGTGCAGAGGTGCTTG 24 540

1	24 541	ACCTATCACATTTTCCAAAAGTCAAGATAACCCCTATCCTGCCGTGCCAACCGCAGC	24 600
2	24 541	ACCTATCACATTTTCCAAAAGTCAAGATAACCCCTATCCTGCCGTGCCAACCGCAGC	24 600
1	24 601	CGAGCGGACAAGCAGCTGGCTTGCGGCAGGGCGCTGTCATAACCTGATATGCCCTCGCTC	24 660
2	24 601	CGAGCGGACAAGCAGCTGGCTTGCGGCAGGGCGCTGTCATAACCTGATATGCCCTCGCTC	24 660
1	24 661	AACGAAGTGCCAAAAATCTTGAGGGCTTGGACGCGACGAGAACGCGCGGCAAACGCT	24 720
2	24 661	AACGAAGTGCCAAAAATCTTGAGGGCTTGGACGCGACGAGAACGCGCGGCAAACGCT	24 720
1	24 721	CTGCAACAGGAAACAGCGAAATGAAAGTCACTCTGGAGTGGTGGAACTCGAGGGT	24 780
2	24 721	CTGCAACAGGAAACAGCGAAATGAAAGTCACTCTGGAGTGGTGGAACTCGAGGGT	24 780
1	24 781	GACAACGCGCGCCTAGCCGTACTAAAACGCAGCATCGAGGTACCCACTTGCCTACCCG	24 840
2	24 781	GACAACGCGCGCCTAGCCGTACTAAAACGCAGCATCGAGGTACCCACTTGCCTACCCG	24 840
1	24 841	GCACCTAACCTACCCCCCAAGGTATGAGCACAGTCATGAGTGAGCTGATCGTGC CGCGT	24 900
2	24 841	GCACCTAACCTACCCCCCAAGGTATGAGCACAGTCATGAGTGAGCTGATCGTGC CGCGT	24 900
1	24 901	GCGCAGCCCTGGAGAGGGATGCAAATTGCAAGAACAAACAGAGGAGGGCTACCGCA	24 960
2	24 901	GCGCAGCCCTGGAGAGGGATGCAAATTGCAAGAACAAACAGAGGAGGGCTACCGCA	24 960
1	24 961	GTTGGCGACGAGCAGCTAGCGCGTGGCTTCAACCGCGAGCCTGCCGACTGGAGGAG	25 020
2	24 961	GTTGGCGACGAGCAGCTAGCGCGTGGCTTCAACCGCGAGCCTGCCGACTGGAGGAG	25 020
1	25 021	CGACGCAAACATAATGATGGCCCGAGTGCAGCGAAGCTAGAGGAAACATTGCACTACACCTT CGA	25 080
2	25 021	CGACGCAAACATAATGATGGCCCGAGTGCAGCGAAGCTAGAGGAAACATTGCACTACACCTT CGA	25 080
1	25 081	TTCTTGCTGACCCGGAGATGCAGCGAAGCTAGAGGAAACATTGCACTACACCTT CGA	25 140
2	25 081	TTCTTGCTGACCCGGAGATGCAGCGAAGCTAGAGGAAACATTGCACTACACCTT CGA	25 140

1	25 141	CAGGGCTACGTACGCCAGGCCTGCAAGATCTCCAACGTGGAGCTCTGCAACCTGGTCTCC	25 200
2	25 141	CAGGGCTACGTACGCCAGGCCTGCAAGATCTCCAACGTGGAGCTCTGCAACCTGGTCTCC	25 200
1	25 201	TACCTTGAATTTCACGAAAACCGCCTGGCAAAACGTGCTTCATTCCACGCTCAAG	25 260
2	25 201	TACCTTGAATTTCACGAAAACCGCCTGGCAAAACGTGCTTCATTCCACGCTCAAG	25 260
1	25 261	GGCGAGGCCGCGCCGCGACTACGTCCCGACTGCCTTACTTATTCTATGCTACACCTGG	25 320
2	25 261	GGCGAGGCCGCGCCGCGACTACGTCCCGACTGCCTTACTTATTCTATGCTACACCTGG	25 320
1	25 321	CAGACGGCCATGGCGTTGGCAGCAGTGCTTGGAGGAGTGCAACCTCAAGGAGCTGCAG	25 380
2	25 321	CAGACGGCCATGGCGTTGGCAGCAGTGCTTGGAGGAGTGCAACCTCAAGGAGCTGCAG	25 380
1	25 381	AAACTGCTAAAGCAAAACTTGAGGACCTATGGACGGCCTTCAACGAGCGCTCCGTGGCC	25 440
2	25 381	AAACTGCTAAAGCAAAACTTGAGGACCTATGGACGGCCTTCAACGAGCGCTCCGTGGCC	25 440
1	25 441	GCGCACCTGGCGGACATCATTTCGGACGGCTGCTTAAACCCCTGCAACAGGGTCTG	25 500
2	25 441	GCGCACCTGGCGGACATCATTTCGGACGGCTGCTTAAACCCCTGCAACAGGGTCTG	25 500
1	25 501	CCAGACTTCACCAGTCAAAGCATGTTGAGAACCTTAGGAACCTTATCCTAGAGCGCTCA	25 560
2	25 501	CCAGACTTCACCAGTCAAAGCATGTTGAGAACCTTAGGAACCTTATCCTAGAGCGCTCA	25 560
1	25 561	GGAATCTTGCACCTGCTGTGACTTCCTAGCGACTTGTGCCATTAGTACCGC	25 620
2	25 561	GGAATCTTGCACCTGCTGTGACTTCCTAGCGACTTGTGCCATTAGTACCGC	25 620
1	25 621	GAATGCCCTCCGCCGCTTGGGCCACTGCTACCTCTGCAGCTAGCCAACCTTGCC	25 680
2	25 621	GAATGCCCTCCGCCGCTTGGGCCACTGCTACCTCTGCAGCTAGCCAACCTTGCC	25 680
1	25 681	TACCACTCTGACATAATGGAAGACGTGAGCGGTGACGGTCACTGGAGTGTCACTGTCGC	25 740
2	25 681	TACCACTCTGACATAATGGAAGACGTGAGCGGTGACGGTCACTGGAGTGTCACTGTCGC	25 740

1	25 741	TGCAACCTATGCACCCCGCACCGCTCCCTGGTTGCAATTGAGCTGCTTAACGAAAGT	25 800
2	25 741	TGCAACCTATGCACCCCGCACCGCTCCCTGGTTGCAATTGAGCTGCTTAACGAAAGT	25 800
1	25 801	CAAATTATCGGTACCTTGAGCTGCAGGGTCCCTGCCCTGACGAAAAGTCCGCGGCTCCG	25 860
2	25 801	CAAATTATCGGTACCTTGAGCTGCAGGGTCCCTGCCCTGACGAAAAGTCCGCGGCTCCG	25 860
1	25 861	GGGTTGAAACTCACTCCGGGGCTGTGGACGTCGGCTTACCTTCGCAAATTGTACCTGAG	25 920
2	25 861	GGGTTGAAACTCACTCCGGGGCTGTGGACGTCGGCTTACCTTCGCAAATTGTACCTGAG	25 920
1	25 921	GACTACCACGCCACGAGATTAGGTTCTACGAAGACCAATCCGCCGCTAATGCGGAG	25 980
2	25 921	GACTACCACGCCACGAGATTAGGTTCTACGAAGACCAATCCGCCGCTAATGCGGAG	25 980
1	25 981	CTTACCGCCTGCGTCATTACCCAGGGCACATTCTGGCAATTGCAAGCCATCAACAAA	26 040
2	25 981	CTTACCGCCTGCGTCATTACCCAGGGCACATTCTGGCAATTGCAAGCCATCAACAAA	26 040
1	26 041	GCCC GCCAAGAGTTCTGCTACGAAAGGGACGGGGGTTACTTGGACCCCCAGTCCGGC	26 100
2	26 041	GCCC GCCAAGAGTTCTGCTACGAAAGGGACGGGGGTTACTTGGACCCCCAGTCCGGC	26 100
1	26 101	GAGGAGCTAACCCAACTCCCCCGCCGCCAGCCCTATCAGCAGCAGCCGCGGGCCCTT	26 160
2	26 101	GAGGAGCTAACCCAACTCCCCCGCCGCCAGCCCTATCAGCAGCAGCCGCGGGCCCTT	26 160
1	26 161	GCTTCCCAGGATGGCACCCAAAAAGAAGCTGCAGCTGCCGCCACCCACGGACGAGGA	26 220
2	26 161	GCTTCCCAGGATGGCACCCAAAAAGAAGCTGCAGCTGCCGCCACCCACGGACGAGGA	26 220
1	26 221	GGAATACTGGGACAGTCAGGCAGAGGAGTTTGACGAGGAGGAGGACATGATGGA	26 280
2	26 221	GGAATACTGGGACAGTCAGGCAGAGGAGTTTGACGAGGAGGAGGACATGATGGA	26 280
1	26 281	AGACTGGGAGAGCCTAGACGAGGAAGCTTCCGAGGTCGAAGAGGTGTCAGACGAAACACC	26 340
2	26 281	AGACTGGGAGAGCCTAGACGAGGAAGCTTCCGAGGTCGAAGAGGTGTCAGACGAAACACC	26 340

1	26 341	GTCACCCCTGGTCGCATTCCTCGCCGGCGCCAGAAATCGGCAACC GGTTCCAGCAT	26 400
2	26 341	GTCACCCCTGGTCGCATTCCTCGCCGGCGCCAGAAATCGGCAACC GGTTCCAGCAT	26 400
1	26 401	GGCTACAACCTCCGCTCCTCAGGCGCCGCCGGCACTGCCGTTGCCGACCCAACCGTAG	26 460
2	26 401	GGCTACAACCTCCGCTCCTCAGGCGCCGCCGGCACTGCCGTTGCCGACCCAACCGTAG	26 460
1	26 461	ATGGGACACCACTGGAACCAGGGCCGGTAAGTCCAAGCAGCCGCCGCGTTAGCCCAAGA	26 520
2	26 461	ATGGGACACCACTGGAACCAGGGCCGGTAAGTCCAAGCAGCCGCCGCGTTAGCCCAAGA	26 520
1	26 521	GCAACAACAGCGCCAAGGCTACCGCTCATGGCGCGGGCACAGAACGCCATAGTTGCTTG	26 580
2	26 521	GCAACAACAGCGCCAAGGCTACCGCTCATGGCGCGGGCACAGAACGCCATAGTTGCTTG	26 580
1	26 581	CTTGCAAGACTGTGGGGCAACATCTCCTCGCCGCCGCTTCTTCTACCATCACGG	26 640
2	26 581	CTTGCAAGACTGTGGGGCAACATCTCCTCGCCGCCGCTTCTTCTACCATCACGG	26 640
1	26 641	CGTGGCCTTCCCCGTAACATCCTGCATTACTACCGTCATCTCTACAGCCCATACTGCAC	26 700
2	26 641	CGTGGCCTTCCCCGTAACATCCTGCATTACTACCGTCATCTCTACAGCCCATACTGCAC	26 700
1	26 701	CGGCGGCAGCGGCAGCAACAGCAGCGGCCACACAGAACAGCAAGGCAGCGGATAGCAAGA	26 760
2	26 701	CGGCGGCAGCGGCAGCAACAGCAGCGGCCACACAGAACAGCAAGGCAGCGGATAGCAAGA	26 760
1	26 761	CTCTGACAAAGCCAAGAAATCCACAGCGCGGCAGCAGCAGGAGGAGGAGCGCTCGTC	26 820
2	26 761	CTCTGACAAAGCCAAGAAATCCACAGCGCGGCAGCAGCAGGAGGAGGAGCGCTCGTC	26 820
1	26 821	TGGCGCCCAACGAACCCGTATCGACCCGCGAGCTTAGAACAGGATTTCCTCACTCTGT	26 880
2	26 821	TGGCGCCCAACGAACCCGTATCGACCCGCGAGCTTAGAACAGGATTTCCTCACTCTGT	26 880
1	26 881	ATGCTATATTCAACAGAGCAGGGCCAAGAACAGAGCTGAAAATAAAACAGGTCTC	26 940
2	26 881	ATGCTATATTCAACAGAGCAGGGCCAAGAACAGAGCTGAAAATAAAACAGGTCTC	26 940

1 26 941 T GCGATCCCTCACCGCAGCTGCCGTATCACAAAGCGAAGATCAGCTTCGGCGCACGC 27 000
2 26 941 T GCGATCCCTCACCGCAGCTGCCGTATCACAAAGCGAAGATCAGCTTCGGCGCACGC 27 000

1 27 001 T GGAAGACGCGGAGGCTCTTCAGTAATACTGCGCGCTGACTCTTAAGGACTAGTTTC 27 060
2 27 001 T GGAAGACGCGGAGGCTCTTCAGTAATACTGCGCGCTGACTCTTAAGGACTAGTTTC 27 060

1 27 061 G CGCCCTTCTCAAATTAAAGCGCGAAAACGTACATCTCCAGCGGCCACACCCGGCGC 27 120
2 27 061 G CGCCCTTCTCAAATTAAAGCGCGAAAACGTACATCTCCAGCGGCCACACCCGGCGC 27 120

1 27 121 C AGCACCTGTTGTCAGCGCCATTATGAGCAAGGAAATTCCCACGCCACATGTGGAGTT 27 180
2 27 121 C AGCACCTGTTGTCAGCGCCATTATGAGCAAGGAAATTCCCACGCCACATGTGGAGTT 27 180

1 27 181 A CCAGCCACAAATGGGACTTGCGGCTGGAGCTGCCAAGACTACTCAACCGAATAAACT 27 240
2 27 181 A CCAGCCACAAATGGGACTTGCGGCTGGAGCTGCCAAGACTACTCAACCGAATAAACT 27 240

1 27 241 A CATGAGCGCGGGACCCACATGATATCCGGTCAACGGAATACGCGCCCACCGAAACC 27 300
2 27 241 A CATGAGCGCGGGACCCACATGATATCCGGTCAACGGAATACGCGCCCACCGAAACC 27 300

1 27 301 G AATTCTCCTGGAACAGGCAGCTATTACCAACACCTCGTAATAACCTTAATCCCCGTA 27 360
2 27 301 G AATTCTCCTGGAACAGGCAGCTATTACCAACACCTCGTAATAACCTTAATCCCCGTA 27 360

1 27 361 G TTGGCCCGCTGCCCTGGTGTACCAAGGAAAGTCCCGCTCCACCACTGTGGTACTTCCA 27 420
2 27 361 G TTGGCCCGCTGCCCTGGTGTACCAAGGAAAGTCCCGCTCCACCACTGTGGTACTTCCA 27 420

1 27 421 G AGACGCCAGGCCAGGTTCACTGACTAACCTGAGGGCAGCTTGCAGGGCGGCTTTC 27 480
2 27 421 G AGACGCCAGGCCAGGTTCACTGACTAACCTGAGGGCAGCTTGCAGGGCGGCTTTC 27 480

1 27 481 G TCAACAGGGTGCAGCTGCCCGGGCAGGGTATAACTCACCTGACAATCAGAGGGCGAGGTA 27 540
2 27 481 G TCAACAGGGTGCAGCTGCCCGGGCAGGGTATAACTCACCTGACAATCAGAGGGCGAGGTA 27 540

1	27 541	TTCAGCTAACGACGAGTCGGTGAGCTCCTCGCTTGGTCTCCGTCCGGACGGGACATTTC	27 600
2	27 541	TTCAGCTAACGACGAGTCGGTGAGCTCCTCGCTTGGTCTCCGTCCGGACGGGACATTTC	27 600
1	27 601	AGATCGGC GGCGCCGGCGCTTT CATT CAC GCCT CGT CAG GCA AT CCT AACT CT GC AGA	27 660
2	27 601	AGATCGGC GGCGCCGGCGCTTT CATT CAC GCCT CGT CAG GCA AT CCT AACT CT GC AGA	27 660
1	27 661	CCTCGT CCTCTGAGCCGCGCTCTGGAGGCATTGGA ACTCTGCA ATT TATTGAGGAGTTG	27 720
2	27 661	CCTCGT CCTCTGAGCCGCGCTCTGGAGGCATTGGA ACTCTGCA ATT TATTGAGGAGTTG	27 720
1	27 721	TGCCATCGGTCTACTTTAACCCCTTCTCGGGACCTCCCGGCCACTATCGGATCAATTAA	27 780
2	27 721	TGCCATCGGTCTACTTTAACCCCTTCTCGGGACCTCCCGGCCACTATCGGATCAATTAA	27 780
1	27 781	TTCCTAACCTTGACGCGGTAAAGGACTCGGC GGACGGCTACGACTGAATGTTAAGTGGAG	27 840
2	27 781	TTCCTAACCTTGACGCGGTAAAGGACTCGGC GGACGGCTACGACTGAATGTTAAGTGGAG	27 840
1	27 841	AGGCAGAGCAACTGCGCCTGAAACACCTGGTCCACTGTCGCCGCCACAAGTGCTTGC	27 900
2	27 841	AGGCAGAGCAACTGCGCCTGAAACACCTGGTCCACTGTCGCCGCCACAAGTGCTTGC	27 900
1	27 901	GCGACTCCGGTGAGTTTGCTACTTGAATTGCCGAGGATCATATCGAGGGCCGGCGC	27 960
2	27 901	GCGACTCCGGTGAGTTTGCTACTTGAATTGCCGAGGATCATATCGAGGGCCGGCGC	27 960
1	27 961	ACGGCGTCCGGCTTACCGCCCAGGGAGAGCTTGCCGTAGCCTGATTGGAGTTACCC	28 020
2	27 961	ACGGCGTCCGGCTTACCGCCCAGGGAGAGCTTGCCGTAGCCTGATTGGAGTTACCC	28 020
1	28 021	AGCGCCCCCTGCTAGTTGAGCGGGACAGGGGACCCCTGTGTTCTCACTGTGATTGCAACT	28 080
2	28 021	AGCGCCCCCTGCTAGTTGAGCGGGACAGGGGACCCCTGTGTTCTCACTGTGATTGCAACT	28 080
1	28 081	GTCCTAACCCCTGGATTACATCAAGATCTTGTTGCCATCTCTGTGCTGAGTATAATAAAT	28 140
2	28 081	GTCCTAACCCCTGGATTACATCAAGATCTTGTTGCCATCTCTGTGCTGAGTATAATAAAT	28 140

1	28 141	ACAGAAATTAAAATATACTGGGGCTCCTATGCCATCCTGTAACGCCACCGTCTTCACC	28 200
2	28 141	ACAGAAATTAAAATATACTGGGGCTCCTATGCCATCCTGTAACGCCACCGTCTTCACC	28 200
1	28 201	CGCCCAGCAAACCAAGGCGAACCTTACCTGGTACTTTAACATCTCTCCCTGTGATT	28 260
2	28 201	CGCCCAGCAAACCAAGGCGAACCTTACCTGGTACTTTAACATCTCTCCCTGTGATT	28 260
1	28 261	TACAACAGTTCAACCCAGACGGAGTGAGTCTACGAGAGAACCTCTCCGAGCTCAGCTAC	28 320
2	28 261	TACAACAGTTCAACCCAGACGGAGTGAGTCTACGAGAGAACCTCTCCGAGCTCAGCTAC	28 320
1	28 321	TCCATCAGAAAAAACACCACCCCTCCTTACCTGCCGGAACGTACGAGTGCACCGGCC	28 380
2	28 321	TCCATCAGAAAAAACACCACCCCTCCTTACCTGCCGGAACGTACGAGTGCACCGGCC	28 380
1	28 381	GCTGCACCAACACCTACCGCCTGACCGTAAACCAGACTTTCCGGACAGACCTCAATAAC	28 440
2	28 381	GCTGCACCAACACCTACCGCCTGACCGTAAACCAGACTTTCCGGACAGACCTCAATAAC	28 440
1	28 441	TCTGTTACCAGAACAGGAGGTGAGCTTAGAAAACCTTAGGGTATTAGGCCAAGGCGC	28 500
2	28 441	TCTGTTACCAGAACAGGAGGTGAGCTTAGAAAACCTTAGGGTATTAGGCCAAGGCGC	28 500
1	28 501	AGCTACTGTGGGTTTACCATGGAAAGGGTCCAACCCCTGGAAAGAGAACATGTGGAAATGC	28 560
2	28 501	AGCTACTGTGGGTTTACCATGGAAAGGGTCCAACCCCTGGAAAGAGAACATGTGGAAATGC	28 560
1	28 561	AGCCAGGCCAAGATTGAGAGGAACAAGCTATTGCTGGTGGCCTCTGTAATTCAAGGACT	28 620
2	28 561	AGCCAGGCCAAGATTGAGAGGAACAAGCTATTGCTGGTGGCCTCTGTAATTCAAGGACT	28 620
1	28 621	GGGGCTGCTCCTGTGCTTCACCTACATCTGCCTGCACTTCTGCTCTTCAGGTATCACA	28 680
2	28 621	GGGGCTGCTCCTGTGCTTCACCTACATCTGCCTGCACTTCTGCTCTTCAGGTATCACA	28 680
1	28 681	TCGGTATCCTCGAATTCAAAGTATCAAAGTACAATTACCGAATATAAGAAGGAGAAAGG	28 740
2	28 681	TCGGTATCCTCGAATTCAAAGTATCAAAGTACAATTACCGAATATAAGAAGGAGAAAGG	28 740

1	28 741	TTTCATCCTCACTTCCAAAAGGAGGATGAAATCATGAAGGTGCAGAACAACTCAGTCAT	28 800
2	28 741	TTTCATCCTCACTTCCAAAAGGAGGATGAAATCATGAAGGTGCAGAACAACTCAGTCAT	28 800
1	28 801	CATCAACTGTGATGGGTTTATCTCATCTCCCTGAAGGGCTACTTCTCCCAGGAAGTCAA	28 860
2	28 801	CATCAACTGTGATGGGTTTATCTCATCTCCCTGAAGGGCTACTTCTCCCAGGAAGTCAA	28 860
1	28 861	CATTAGCCTTCATTACCAAGAAGGATGAGGAGCCCCCTTTCCAAGTGAAGAACGGTCAGGTC	28 920
2	28 861	CATTAGCCTTCATTACCAAGAAGGATGAGGAGCCCCCTTTCCAAGTGAAGAACGGTCAGGTC	28 920
1	28 921	TGTCAACTCCTTGATGGTGGCCTCTCTGACTTACAAAGACAAAGTCTACTTGAATGTGAC	28 980
2	28 921	TGTCAACTCCTTGATGGTGGCCTCTCTGACTTACAAAGACAAAGTCTACTTGAATGTGAC	28 980
1	28 981	CACTGACAATAACCTCCCTGGATGACTTCCATGTGAATGGCGGAGAACGTGATTCTTATCCA	29 040
2	28 981	CACTGACAATAACCTCCCTGGATGACTTCCATGTGAATGGCGGAGAACGTGATTCTTATCCA	29 040
1	29 041	TCAAAATCCTGGTGAATTCTGTGTCCTTAGGAAGAGGAGGAGCAGCAGCGAGGCCAG	29 100
2	29 041	TCAAAATCCTGGTGAATTCTGTGTCCTTAGGAAGAGGAGGAGCAGCAGCGAGGCCAG	29 100
1	29 101	GCACAAAGCAGAAGATCGTGGCCCCGTGAAGCAGACCCCTGAACCTGACCTGCTGAAGCT	29 160
2	29 101	GCACAAAGCAGAAGATCGTGGCCCCGTGAAGCAGACCCCTGAACCTGACCTGCTGAAGCT	29 160
1	29 161	GGCCGGCGACGTGGAGAGCAACCCGGCCCCATGATGAAACATACAACCAAACCTCTCC	29 220
2	29 161	GGCCGGCGACGTGGAGAGCAACCCGGCCCCATGATGAAACATACAACCAAACCTCTCC	29 220
1	29 221	CCGATCTGCAGGCCACTGGACTGCCATCAGCATGAAAATTGTATTTACTTACTGT	29 280
2	29 221	CCGATCTGCAGGCCACTGGACTGCCATCAGCATGAAAATTGTATTTACTTACTGT	29 280
1	29 281	TTTTCTTATCACCCAGATGATTGGTCAGCACTTTGCTGTATCTCATAGAACGGTT	29 340
2	29 281	TTTTCTTATCACCCAGATGATTGGTCAGCACTTTGCTGTATCTCATAGAACGGTT	29 340

1 29 341 GGACAAGATAGAAGATGAAAGGAATCTTCATGAAGATTTGTATTCATGAAAACGATACA 29 400
2 29 341 GGACAAGATAGAAGATGAAAGGAATCTTCATGAAGATTTGTATTCATGAAAACGATACA 29 400

1 29 401 GAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACGTGAGGAGATAAAAGCCA 29 460
2 29 401 GAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACGTGAGGAGATAAAAGCCA 29 460

1 29 461 GTTTGAAGGCTTGTAAGGATATAATGTTAACAAAGAGGGAGACGAAGAAAGAAAACAG 29 520
2 29 461 GTTTGAAGGCTTGTAAGGATATAATGTTAACAAAGAGGGAGACGAAGAAAGAAAACAG 29 520

1 29 521 CTTTGAATGCAAAAGGTGATCAGAACATCCTCAAATTGCGGCACATGTCATAAGTGAGGC 29 580
2 29 521 CTTTGAATGCAAAAGGTGATCAGAACATCCTCAAATTGCGGCACATGTCATAAGTGAGGC 29 580

1 29 581 CAGCAGTAAACAAACATCTGTGTTACAGTGGCTGAAAAGGATACTACACCATGAGCAA 29 640
2 29 581 CAGCAGTAAACAAACATCTGTGTTACAGTGGCTGAAAAGGATACTACACCATGAGCAA 29 640

1 29 641 CAACTTGGTAACCCCTGGAAAATGGAAACAGCTGACCGTTAAAGACAAGGACTCTATT 29 700
2 29 641 CAACTTGGTAACCCCTGGAAAATGGAAACAGCTGACCGTTAAAGACAAGGACTCTATT 29 700

1 29 701 TATCTATGCCAAGTCACCTTCTGTTCAATCGGAAGCTCGAGTCAGCTCCATTAT 29 760
2 29 701 TATCTATGCCAAGTCACCTTCTGTTCAATCGGAAGCTCGAGTCAGCTCCATTAT 29 760

1 29 761 AGCCAGCCTCTGCCTAAAGTCCCCGGTAGATTGAGAGAATCTTACTCAGAGCTGCAA 29 820
2 29 761 AGCCAGCCTCTGCCTAAAGTCCCCGGTAGATTGAGAGAATCTTACTCAGAGCTGCAA 29 820

1 29 821 TACCCACAGTCCGCCAACCTTGCAGGGCAACAATCCATTCACTTGGGAGGAGTATTG 29 880
2 29 821 TACCCACAGTCCGCCAACCTTGCAGGGCAACAATCCATTCACTTGGGAGGAGTATTG 29 880

1 29 881 ATTGCAACCAGGTGCTTCGGTGTCAATGTGACTGATCCAAGCCAAGTGAGCCATGG 29 940
2 29 881 ATTGCAACCAGGTGCTTCGGTGTCAATGTGACTGATCCAAGCCAAGTGAGCCATGG 29 940

1 29 941 CACTGGCTTCACGTCCCTTGGCTTACTCAAACCTCTGATAATTACTAAGTTACAAAGCTA 30 000
2 29 941 CACTGGCTTCACGTCCCTTGGCTTACTCAAACCTCTGATAATTACTAAGTTACAAAGCTA 30 000

1 30 001 ATGTCACCACTAACTGCTTACTCGCTGCTGAAAACAAATTCAAAAAGTTAGCATTAT 30 060
2 30 001 ATGTCACCACTAACTGCTTACTCGCTGCTGAAAACAAATTCAAAAAGTTAGCATTAT 30 060

1 30 061 AATTAGAACAGGATTAAACCCCCCGGTCAATTCTGCTCAATACCATTCCCCTGAACAA 30 120
2 30 061 AATTAGAACAGGATTAAACCCCCCGGTCAATTCTGCTCAATACCATTCCCCTGAACAA 30 120

1 30 121 TTGACTCTATGTGGATATGCTCCAGCGTACAACCTTGAAGTCAGGCTTCTGGATGTC 30 180
2 30 121 TTGACTCTATGTGGATATGCTCCAGCGTACAACCTTGAAGTCAGGCTTCTGGATGTC 30 180

1 30 181 AGCATCTGACTTGGCCAGCACCTGTCCCGCGGATTGTTCCAGTCCAACATACAGCGACC 30 240
2 30 181 AGCATCTGACTTGGCCAGCACCTGTCCCGCGGATTGTTCCAGTCCAACATACAGCGACC 30 240

1 30 241 CACCCCTAACAGAGATGACCAACACAACCAACGCGGCCGCCGTACCGGACTTACATCTAC 30 300
2 30 241 CACCCCTAACAGAGATGACCAACACAACCAACGCGGCCGCCGTACCGGACTTACATCTAC 30 300

1 30 301 CACAAATACACCCCAAGTTCTGCCTTGTCAATAACTGGATAACTTGGCATGTGGTG 30 360
2 30 301 CACAAATACACCCCAAGTTCTGCCTTGTCAATAACTGGATAACTTGGCATGTGGTG 30 360

1 30 361 GTTCTCCATAGCGCTTATGTTGTATGCCTTATTATTATGTGGCTCATCTGCTGCCTAAA 30 420
2 30 361 GTTCTCCATAGCGCTTATGTTGTATGCCTTATTATTATGTGGCTCATCTGCTGCCTAAA 30 420

1 30 421 GCGCAAACGCGCCGACCCACCATCTATAGTCCCATCATTGTGCTACACCCAAACAATGA 30 480
2 30 421 GCGCAAACGCGCCGACCCACCATCTATAGTCCCATCATTGTGCTACACCCAAACAATGA 30 480

1 30 481 TGGAAATCCATAGATTGGACGGACTGAAACACATGTTCTTTCTTACAGTATGATTAAA 30 540
2 30 481 TGGAAATCCATAGATTGGACGGACTGAAACACATGTTCTTTCTTACAGTATGATTAAA 30 540

1	30 541	TGAGACATGATTCCCTCGAGTTTATATTACTGACCCCTGTTGCGCTTTTGTCGTGC	30 600
2	30 541	TGAGACATGATTCCCTCGAGTTTATATTACTGACCCCTGTTGCGCTTTTGTCGTGC	30 600
1	30 601	TCCACATTGGCTCGGGTTCTCACATCGAAGTAGACTGCATTCCAGCCTCACAGTCTAT	30 660
2	30 601	TCCACATTGGCTCGGGTTCTCACATCGAAGTAGACTGCATTCCAGCCTCACAGTCTAT	30 660
1	30 661	TTGCTTTACGGATTTCGTACCCCTCACGCTCATCTGCAGCCTCATCACTGTGGTCATGCC	30 720
2	30 661	TTGCTTTACGGATTTCGTACCCCTCACGCTCATCTGCAGCCTCATCACTGTGGTCATGCC	30 720
1	30 721	TTTATCCAGTGCATTGACTGGGCTGTGCGCTTGCATATCTCAGACACCATCCCCAG	30 780
2	30 721	TTTATCCAGTGCATTGACTGGGCTGTGCGCTTGCATATCTCAGACACCATCCCCAG	30 780
1	30 781	TACAGGGACAGGACTATAGCTGAGCTTCTAGAATTCTTAATTATGAAATTACTGTGA	30 840
2	30 781	TACAGGGACAGGACTATAGCTGAGCTTCTAGAATTCTTAATTATGAAATTACTGTGA	30 840
1	30 841	CTTTCTGCTGATTATTCGACCCATCTCGTTTGTGTTCCCCGACCTCCAAGCCTCAA	30 900
2	30 841	CTTTCTGCTGATTATTCGACCCATCTCGTTTGTGTTCCCCGACCTCCAAGCCTCAA	30 900
1	30 901	GACATATATCATGCAGATTCACTCGTATATGGAATTCCAAAGTTGCTACAATGAAAAAA	30 960
2	30 901	GACATATATCATGCAGATTCACTCGTATATGGAATTCCAAAGTTGCTACAATGAAAAAA	30 960
1	30 961	GCGATTTCCGAAGCCTGGTTATGCAATCATCTCTGTTATGGTGTCTGCAGTACCA	31 020
2	30 961	GCGATTTCCGAAGCCTGGTTATGCAATCATCTCTGTTATGGTGTCTGCAGTACCA	31 020
1	31 021	TCTTAGCCCTAGCTATATCCCTACCTTGACATTGGCTGGAACGCAATAGATGCCATGA	31 080
2	31 021	TCTTAGCCCTAGCTATATCCCTACCTTGACATTGGCTGGAACGCAATAGATGCCATGA	31 080
1	31 081	ACCACCCAACCTTCCCCGCGCCCGCTATGCTTCACTGCAACAAAGTTGTTGCCGGCGGCT	31 140
2	31 081	ACCACCCAACCTTCCCCGCGCCCGCTATGCTTCACTGCAACAAAGTTGTTGCCGGCGGCT	31 140

1 31 141 TTGTCCCAGCCAATCAGCCTGCCACCTTCTCCACCCCCACTGAAATCAGCTACTTTA 31 200
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 141 TTGTCCCAGCCAATCAGCCTGCCACCTTCTCCACCCCCACTGAAATCAGCTACTTTA 31 200

1 31 201 ATCTAACAGGAGGAGACGACTGATAAAAAAAAATAATAAGCATCACTTACTTAAATCA 31 260
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 201 ATCTAACAGGAGGAGACGACTGATAAAAAAAAATAATAAGCATCACTTACTTAAATCA 31 260

1 31 261 GTTAGCAAATTCTGTCCAGTTATTCAAGCAGCACCTCCTGCCCTCCTCCAGCTCTGG 31 320
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 261 GTTAGCAAATTCTGTCCAGTTATTCAAGCAGCACCTCCTGCCCTCCTCCAGCTCTGG 31 320

1 31 321 TATTGCAGCTTCCTCCTGGCTGCAAACCTTCTCCACAATCTAAATGGAATGTCAGTTCC 31 380
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 321 TATTGCAGCTTCCTCCTGGCTGCAAACCTTCTCCACAATCTAAATGGAATGTCAGTTCC 31 380

1 31 381 TCCTGTTCTGTCCATCCGCACCCACTATCTCATGTTGTTGCAGATGAAGCGCGCAAGA 31 440
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 381 TCCTGTTCTGTCCATCCGCACCCACTATCTCATGTTGTTGCAGATGAAGCGCGCAAGA 31 440

1 31 441 CCGTCTGAAGATACTTCAACCCGTGTATCCATATGACACGGAAACCGGTCTCCAACT 31 500
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 441 CCGTCTGAAGATACTTCAACCCGTGTATCCATATGACACGGAAACCGGTCTCCAACT 31 500

1 31 501 GTGCCTTTCTTACTCCTCCCTTGTATCCCCAATGGGTTCAAGAGAGTCCCCCTGGG 31 560
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 501 GTGCCTTTCTTACTCCTCCCTTGTATCCCCAATGGGTTCAAGAGAGTCCCCCTGGG 31 560

1 31 561 GTACTCTTTGCGCCTATCGAACCTCTAGTTACCTCAATGGCATGCTTGCCTCAAA 31 620
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 561 GTACTCTTTGCGCCTATCGAACCTCTAGTTACCTCAATGGCATGCTTGCCTCAAA 31 620

1 31 621 ATGGGCAACGGCCTCTCTGGACGAGGCCGGAACCTTACCTCCAAAATGTAACCACT 31 680
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 621 ATGGGCAACGGCCTCTCTGGACGAGGCCGGAACCTTACCTCCAAAATGTAACCACT 31 680

1 31 681 GTGAGCCCACCTCTCAAAAAAACCAAGTAAACATAAACCTGGAAATATCTGCACCCCTC 31 740
| | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
2 31 681 GTGAGCCCACCTCTCAAAAAAACCAAGTAAACATAAACCTGGAAATATCTGCACCCCTC 31 740

1 31 741 ACAGTTACCTCAGAAGCCCTAACGTGGCTGCCGCCGACCTCTAATGGTCGGGGCAAC 31 800
2 |||||||
31 741 ACAGTTACCTCAGAAGCCCTAACGTGGCTGCCGCCGACCTCTAATGGTCGGGGCAAC 31 800

1 31 801 ACACTCACCATGCAATCACAGGCCCGCTAACCGTGCACGACTCCAAACTTAGCATTGCC 31 860
2 |||||||
31 801 ACACTCACCATGCAATCACAGGCCCGCTAACCGTGCACGACTCCAAACTTAGCATTGCC 31 860

1 31 861 ACCCAAGGACCCCTCACAGTGTAGAAGGAAAGCTAGCCCTGCAAACATCAGGCCCTC 31 920
2 |||||||
31 861 ACCCAAGGACCCCTCACAGTGTAGAAGGAAAGCTAGCCCTGCAAACATCAGGCCCTC 31 920

1 31 921 ACCACCACCGATAGCAGTACCCCTACTATCACTGCCTCACCCCTCTAACTACTGCCACT 31 980
2 |||||||
31 921 ACCACCACCGATAGCAGTACCCCTACTATCACTGCCTCACCCCTCTAACTACTGCCACT 31 980

1 31 981 GGTAGCTTGGCATTGACTTGAAAGAGGCCATTATACACAAAATGGAAAAGTAGGACTA 32 040
2 |||||||
31 981 GGTAGCTTGGCATTGACTTGAAAGAGGCCATTATACACAAAATGGAAAAGTAGGACTA 32 040

1 32 041 AAGTACGGGCTCCTTGCATGTAACAGACGACCTAACACACTTGTACCGTAGCAACTGGT 32 100
2 |||||||
32 041 AAGTACGGGCTCCTTGCATGTAACAGACGACCTAACACACTTGTACCGTAGCAACTGGT 32 100

1 32 101 CCAGGTGTGACTATTAATAACTTCCTTGCAAACTAAAGTTACTGGAGCCTGGTTTT 32 160
2 |||||||
32 101 CCAGGTGTGACTATTAATAACTTCCTTGCAAACTAAAGTTACTGGAGCCTGGTTTT 32 160

1 32 161 GATTACAAGGAATATGCAACTTAATGTAGCAGGAGGACTAAGGATTGATTCTCAAAAC 32 220
2 |||||||
32 161 GATTACAAGGAATATGCAACTTAATGTAGCAGGAGGACTAAGGATTGATTCTCAAAAC 32 220

1 32 221 AGACGCCTTATACCGATGTTAGTTATCCGTTGATGCTCAAAACCAACTAAATCTAAGA 32 280
2 |||||||
32 221 AGACGCCTTATACCGATGTTAGTTATCCGTTGATGCTCAAAACCAACTAAATCTAAGA 32 280

1 32 281 CTAGGACAGGGCCCTTTTATAAACTCAGCCCACAACTGGATATTAACACTACAACAA 32 340
2 |||||||
32 281 CTAGGACAGGGCCCTTTTATAAACTCAGCCCACAACTGGATATTAACACTACAACAA 32 340

1	32 341	GGCCTTACTTACAGCTCAAACAATTCCAAAAAGCTTGAGGTTAACCTAAGCACT	32 400
2	32 341	GGCCTTACTTACAGCTCAAACAATTCCAAAAAGCTTGAGGTTAACCTAAGCACT	32 400
1	32 401	GCCAAGGGGTTGATGTTGACGCTACAGCCATAGCATTAAATGCAGGAGATGGCTTGAA	32 460
2	32 401	GCCAAGGGGTTGATGTTGACGCTACAGCCATAGCATTAAATGCAGGAGATGGCTTGAA	32 460
1	32 461	TTTGGTTCACCTAATGCACCAAACACAATCCCCTCAAAACAAAAATTGCCATGGCTA	32 520
2	32 461	TTTGGTTCACCTAATGCACCAAACACAATCCCCTCAAAACAAAAATTGCCATGGCTA	32 520
1	32 521	GAATTTGATTCAAACAAGGCTATGGTCTAAACTAGGAACCTGGCTTAGTTGACAGC	32 580
2	32 521	GAATTTGATTCAAACAAGGCTATGGTCTAAACTAGGAACCTGGCTTAGTTGACAGC	32 580
1	32 581	ACAGGTGCCATTACAGTAGGAAACAAAAATAATGATAAGCTAACCTATGGACAGGTCCA	32 640
2	32 581	ACAGGTGCCATTACAGTAGGAAACAAAAATAATGATAAGCTAACCTATGGACAGGTCCA	32 640
1	32 641	AAACCAGAAGCCAAC TG CATA ATT G A AT AC GGG AA AC AAA ACC CAG AT AG CAA ACT AACT	32 700
2	32 641	AAACCAGAAGCCAAC TG CATA ATT G A AT AC GGG AA AC AAA ACC CAG AT AG CAA ACT AACT	32 700
1	32 701	TTAATCCTTGTAAAAATGGAGGAATTGTTAATGGATATGTAACGCTAATGGAGCCTCA	32 760
2	32 701	TTAATCCTTGTAAAAATGGAGGAATTGTTAATGGATATGTAACGCTAATGGAGCCTCA	32 760
1	32 761	GACTACGTTAACACCTTATTTAAAAACAAAAATGTCTCCATTAATGTAGAACTATACTTT	32 820
2	32 761	GACTACGTTAACACCTTATTTAAAAACAAAAATGTCTCCATTAATGTAGAACTATACTTT	32 820
1	32 821	GATGCCACTGGTCATATATTACCA GACTCATCTTCTCTTAAACAGATCTAGAACTAAAA	32 880
2	32 821	GATGCCACTGGTCATATATTACCA GACTCATCTTCTCTTAAACAGATCTAGAACTAAAA	32 880
1	32 881	TACAAGCAAACCGCTGACTTTAGTGCAAGAGGTTTATGCCAAGTACTACAGCGTATCCA	32 940
2	32 881	TACAAGCAAACCGCTGACTTTAGTGCAAGAGGTTTATGCCAAGTACTACAGCGTATCCA	32 940

1	32 941	TTTGTCCCTCCTAATGCAGGGAAACACATAATGAAAATTATATTTTGGTCAATGCTACTAC	33 000
2	32 941	TTTGTCCCTCCTAATGCAGGGAAACACATAATGAAAATTATATTTTGGTCAATGCTACTAC	33 000
1	33 001	AAAGCAAGCGATGGTGCCCTTTCCGTTGGAAGTTACTGTTATGCTTAATAAACGCCTG	33 060
2	33 001	AAAGCAAGCGATGGTGCCCTTTCCGTTGGAAGTTACTGTTATGCTTAATAAACGCCTG	33 060
1	33 061	CCAGATAGTCGACATCCTATGTTATGACTTTTATGGCCTTGAATGCTGGTAGCT	33 120
2	33 061	CCAGATAGTCGACATCCTATGTTATGACTTTTATGGCCTTGAATGCTGGTAGCT	33 120
1	33 121	CCAGAAACTACTCAGGCAACCCTCATAACCTCCCCATTTACCTTTCTATATTAGAGAA	33 180
2	33 121	CCAGAAACTACTCAGGCAACCCTCATAACCTCCCCATTTACCTTTCTATATTAGAGAA	33 180
1	33 181	GATGACTAATAAACTCTAAAGAACATCGTTGTGTTATGTTAACGTGTTATTTCAAT	33 240
2	33 181	GATGACTAATAAACTCTAAAGAACATCGTTGTGTTATGTTAACGTGTTATTTCAAT	33 240
1	33 241	TGCAGAAAATTCAAGTCATTTTCATTCAAGTAGTATAGCCCCACCAACATAGCTTAT	33 300
2	33 241	TGCAGAAAATTCAAGTCATTTTCATTCAAGTAGTATAGCCCCACCAACATAGCTTAT	33 300
1	33 301	ACAGATCACCGTACCTTAATCAAACTCACAGAACCTAGTATTCAACCTGCCACCTCCCT	33 360
2	33 301	ACAGATCACCGTACCTTAATCAAACTCACAGAACCTAGTATTCAACCTGCCACCTCCCT	33 360
1	33 361	CCCAACACACAGAGTACACAGTCCTTCTCCCCGGCTGGCCTTAAAAGCATCATATCAT	33 420
2	33 361	CCCAACACACAGAGTACACAGTCCTTCTCCCCGGCTGGCCTTAAAAGCATCATATCAT	33 420
1	33 421	GGGTAACAGACATATTCTAGGTGTTATTCACACGGTTCTGTCGAGCCAAACGCT	33 480
2	33 421	GGGTAACAGACATATTCTAGGTGTTATTCACACGGTTCTGTCGAGCCAAACGCT	33 480
1	33 481	CATCAGTGATATTAATAAACTCCCCGGGCAGCTCACTTAAGTTCATGTCGCTGTCCAGCT	33 540
2	33 481	CATCAGTGATATTAATAAACTCCCCGGGCAGCTCACTTAAGTTCATGTCGCTGTCCAGCT	33 540

1 33 541 GCTGAGCCACAGGCTGCTGTCCAACTTGC GGTTGCTTAACGGGCGGAAGGAGAAGTCC 33 600
2 33 541 GCTGAGCCACAGGCTGCTGTCCAACTTGC GGTTGCTTAACGGGCGGAAGGAGAAGTCC 33 600

1 33 601 ACGCCTACATGGGGTAGAGTCATAATCGTCATCAGGATAGGGCGGTGGTGCTGCAGCA 33 660
2 33 601 ACGCCTACATGGGGTAGAGTCATAATCGTCATCAGGATAGGGCGGTGGTGCTGCAGCA 33 660

1 33 661 GCGCGCGAATAAACTGCTGCCGCCGCTCCGT CCTGCAGGAATACAACATGGCAGTGG 33 720
2 33 661 GCGCGCGAATAAACTGCTGCCGCCGCTCCGT CCTGCAGGAATACAACATGGCAGTGG 33 720

1 33 721 TCTCCTCAGCGATGATTGCACCGCCCCG CAGCATAAGGCGC TTGT CCTCCGGCACAGC 33 780
2 33 721 TCTCCTCAGCGATGATTGCACCGCCCCG CAGCATAAGGCGC TTGT CCTCCGGCACAGC 33 780

1 33 781 AGCGCACCCCTGATCTCACTTAAATCAGCACAGTA ACTGCAGCACAGCACCAATATTGT 33 840
2 33 781 AGCGCACCCCTGATCTCACTTAAATCAGCACAGTA ACTGCAGCACAGCACCAATATTGT 33 840

1 33 841 TCAAAATCCCACAGTGCAAGGCGCTGTATCAAAGCTCATGGCGGGACCACAGAACCCA 33 900
2 33 841 TCAAAATCCCACAGTGCAAGGCGCTGTATCAAAGCTCATGGCGGGACCACAGAACCCA 33 900

1 33 901 CGTGGCCATCATACCACAAGCGCAGGTAGATTAAGTGGCGACCCCTCATAAACACGCTGG 33 960
2 33 901 CGTGGCCATCATACCACAAGCGCAGGTAGATTAAGTGGCGACCCCTCATAAACACGCTGG 33 960

1 33 961 ACATAAACATTACCTTTGGCATGTTGTAATTCA CCACCTCCGGTACCATATAAACCC 34 020
2 33 961 ACATAAACATTACCTTTGGCATGTTGTAATTCA CCACCTCCGGTACCATATAAACCC 34 020

1 34 021 TCTGATTAAACATGGCGCCATCCACCA CCTAAACCAAGCTGGCCAAAACCTGCCCGC 34 080
2 34 021 TCTGATTAAACATGGCGCCATCCACCA CCTAAACCAAGCTGGCCAAAACCTGCCCGC 34 080

1 34 081 CGGCTATAACTGCAGGGAACCGGGACTGGAA CAATGACAGTGGAGAGCCCAGGACTCGT 34 140
2 34 081 CGGCTATAACTGCAGGGAACCGGGACTGGAA CAATGACAGTGGAGAGCCCAGGACTCGT 34 140

1 34 141 AACCATGGATCATCATGCTCGTCATGATATCAATTTGGCACAAACACAGGCACACGTGCA 34 200
2 |||||||
34 141 AACCATGGATCATCATGCTCGTCATGATATCAATTTGGCACAAACACAGGCACACGTGCA 34 200

1 34 201 TACACTTCCTCAGGATTACAAGCTCCCTCCCGCGTTAGAACCATATCCCAGGGAACAACCC 34 260
2 |||||||
34 201 TACACTTCCTCAGGATTACAAGCTCCCTCCCGCGTTAGAACCATATCCCAGGGAACAACCC 34 260

1 34 261 ATT CCT GAAT CAG CGT AAAT CCC AC ACT GC AG GG AAG AC CT CG CA CG TA ACT CAC GT T GT 34 320
2 |||||||
34 261 ATT CCT GAAT CAG CGT AAAT CCC AC ACT GC AG GG AAG AC CT CG CA CG TA ACT CAC GT T GT 34 320

1 34 321 GC ATT GT CAA AGT GT T AC ATT CG GG C AG C AG CG G AT G AT CC T CC AG T AT GG T AG CG CG GG 34 380
2 |||||||
34 321 GC ATT GT CAA AGT GT T AC ATT CG GG C AG C AG CG G AT G AT CC T CC AG T AT GG T AG CG CG GG 34 380

1 34 381 TTT CT GT CT CAA AAA AGG AGG T AG AC G AT CC CT ACT GT AC GG AG T GC G CC G AG AC A ACC G AG 34 440
2 |||||||
34 381 TTT CT GT CT CAA AAA AGG AGG T AG AC G AT CC CT ACT GT AC GG AG T GC G CC G AG AC A ACC G AG 34 440

1 34 441 AT CG T GT TT GG T CG T AG T GT CAT GC CAA AT GG A AC G CC GG AC GT AG T CAT AT TT C CT GA AG 34 500
2 |||||||
34 441 AT CG T GT TT GG T CG T AG T GT CAT GC CAA AT GG A AC G CC GG AC GT AG T CAT AT TT C CT GA AG 34 500

1 34 501 CAAAACCAGGTGCAGGGCGTGACAAACAGATCTGCGTCTCCGGTCTGCCGCTTAGATCGC 34 560
2 |||||||
34 501 CAAAACCAGGTGCAGGGCGTGACAAACAGATCTGCGTCTCCGGTCTGCCGCTTAGATCGC 34 560

1 34 561 TCT GT GT TAG T GT TAG T AT AT CC ACT CT CT CAA AG C AT CC AG GG G C C C C T GG C TT CG 34 620
2 |||||||
34 561 TCT GT GT TAG T GT TAG T AT AT CC ACT CT CT CAA AG C AT CC AG GG G C C C C T GG C TT CG 34 620

1 34 621 GG TT CT AT GT TAA ACT C CTT CAT GC GCC G CT G CC CT GATA AC AT CC ACC ACC G C AG A AT AA 34 680
2 |||||||
34 621 GG TT CT AT GT TAA ACT C CTT CAT GC GCC G CT G CC CT GATA AC AT CC ACC ACC G C AG A AT AA 34 680

1 34 681 GCC AC ACC CAG CC A AC CT AC AC AT TC GT T CT G CG AG T CAC AC AC G GG AG G AG C GG G A AG A 34 740
2 |||||||
34 681 GCC AC ACC CAG CC A AC CT AC AC AT TC GT T CT G CG AG T CAC AC AC G GG AG G AG C GG G A AG A 34 740

1 34 741 GCTGGAAAGAACCATGTTTTTTTTTATTCCAAAAGATTATCCAAAACCTCAAAATGAAG 34 800
2 34 741 GCTGGAAAGAACCATGTTTTTTTATTCCAAAAGATTATCCAAAACCTCAAAATGAAG 34 800

1 34 801 ATCTATTAAGTGAACCGCGCTCCCCCTCCGGTGGCGTGGTCAAACCTCTACAGCCAAAGAAC 34 860
2 34 801 ATCTATTAAGTGAACCGCGCTCCCCCTCCGGTGGCGTGGTCAAACCTCTACAGCCAAAGAAC 34 860

1 34 861 GATAATGGCATTGTAAGATGTTGCACAATGGCTTCCAAAAGGCCAACGGCCCTCACGTC 34 920
2 34 861 GATAATGGCATTGTAAGATGTTGCACAATGGCTTCCAAAAGGCCAACGGCCCTCACGTC 34 920

1 34 921 CAAGTGGACGTAAAGGCTAAACCCTTCAGGGTGAATCTCCTCTATAAACATTCCAGCACC 34 980
2 34 921 CAAGTGGACGTAAAGGCTAAACCCTTCAGGGTGAATCTCCTCTATAAACATTCCAGCACC 34 980

1 34 981 TTCAACCATGCCAAATAATTCTCATCTGCCACCTTCTCAATATATCTTAAGCAAATC 35 040
2 34 981 TTCAACCATGCCAAATAATTCTCATCTGCCACCTTCTCAATATATCTTAAGCAAATC 35 040

1 35 041 CCGAATATTAAGTCCGGCCATTGTAAAAATCTGCTCCAGAGCGCCCTCCACCTTCAGCCT 35 100
2 35 041 CCGAATATTAAGTCCGGCCATTGTAAAAATCTGCTCCAGAGCGCCCTCCACCTTCAGCCT 35 100

1 35 101 CAAGCAGCGAATCATGATTGAAAAATTCAAGGTTCTCACAGACCTGTATAAGATTCAA 35 160
2 35 101 CAAGCAGCGAATCATGATTGAAAAATTCAAGGTTCTCACAGACCTGTATAAGATTCAA 35 160

1 35 161 AGCGGAACATTAACAAAAATACCGCGATCCGTAGGTCCTCGCAGGGCCAGCTGAACA 35 220
2 35 161 AGCGGAACATTAACAAAAATACCGCGATCCGTAGGTCCTCGCAGGGCCAGCTGAACA 35 220

1 35 221 TAATCGTGCAGGTCTGCACGGACCAGCGCGGCCACTTCCCCGCCAGGAACCATGACAAA 35 280
2 35 221 TAATCGTGCAGGTCTGCACGGACCAGCGCGGCCACTTCCCCGCCAGGAACCATGACAAA 35 280

1 35 281 GAACCCACACTGATTATGACACGCATACTCGGAGCTATGCTAACCAAGCGTAGCCCCGATG 35 340
2 35 281 GAACCCACACTGATTATGACACGCATACTCGGAGCTATGCTAACCAAGCGTAGCCCCGATG 35 340

1 35 341 TAAGCTTGTGCATGGCGCGATATAAAATGCAAGGTGCTGCTCAAAAAATCAGGCAA 35 400
2 |||||||
1 35 341 TAAGCTTGTGCATGGCGCGATATAAAATGCAAGGTGCTGCTCAAAAAATCAGGCAA 35 400

1 35 401 GCCTCGCGCAAAAAGAAAGCACATCGTAGTCATGCTCATGCAGATAAAGGCAGGTAAGC 35 460
2 |||||||
1 35 401 GCCTCGCGCAAAAAGAAAGCACATCGTAGTCATGCTCATGCAGATAAAGGCAGGTAAGC 35 460

1 35 461 TCCGGAACCACCCACAGAAAAAGACACCATTTCCTCAAACATGTCCTGCCGGTTCTGC 35 520
2 |||||||
1 35 461 TCCGGAACCACCCACAGAAAAAGACACCATTTCCTCAAACATGTCCTGCCGGTTCTGC 35 520

1 35 521 ATAAACACAAAATAAACAAAAACATTAAACATTAGAACGCTGTCTTACAACAG 35 580
2 |||||||
1 35 521 ATAAACACAAAATAAACAAAAACATTAAACATTAGAACGCTGTCTTACAACAG 35 580

1 35 581 GAAAAACAAACCTTATAAGCATAAGACGGACTACGCCATGCCGGCGTACCGTAAAAAA 35 640
2 |||||||
1 35 581 GAAAAACAAACCTTATAAGCATAAGACGGACTACGCCATGCCGGCGTACCGTAAAAAA 35 640

1 35 641 ACTGGTCACCGTGATTAAGCACCACCGACAGCTCCTCGGTATGTCGGAGTCATAA 35 700
2 |||||||
1 35 641 ACTGGTCACCGTGATTAAGCACCACCGACAGCTCCTCGGTATGTCGGAGTCATAA 35 700

1 35 701 TGTAAGACTCGTAAACACATCAGGTTGATTCACATCGTCAGTGCTAAAAAGCGACCGA 35 760
2 |||||||
1 35 701 TGTAAGACTCGTAAACACATCAGGTTGATTCACATCGTCAGTGCTAAAAAGCGACCGA 35 760

1 35 761 AATAGCCC GGGAATACATACCCGCAGGGTAGAGACAAACATTACAGCCCCATAGGAG 35 820
2 |||||||
1 35 761 AATAGCCC GGGAATACATACCCGCAGGGTAGAGACAAACATTACAGCCCCATAGGAG 35 820

1 35 821 GTATAACAAAATTAATAGGAGAGAAAAACACATAAACACCTGAAAAACCCCTCCTGCCTAG 35 880
2 |||||||
1 35 821 GTATAACAAAATTAATAGGAGAGAAAAACACATAAACACCTGAAAAACCCCTCCTGCCTAG 35 880

1 35 881 GCAAAATAGCACCCCTCCCGCTCCAGAACACATACAGCGCTTCCACAGCGGCAGCCATAA 35 940
2 |||||||
1 35 881 GCAAAATAGCACCCCTCCCGCTCCAGAACACATACAGCGCTTCCACAGCGGCAGCCATAA 35 940

1 35 941 CAGTCAGCCTTACCAAGTAAAAAGAAAACCTATTAAAAAACACCACTCGACACGGCACC 36 000
2 35 941 CAGTCAGCCTTACCAAGTAAAAAGAAAACCTATTAAAAAACACCACTCGACACGGCACC 36 000

1 36 001 AGCTCAATCAGTCACAGTGTAAGGAGGCCAAGTGCAGAGCGAGTATATAGGACTAA 36 060
2 36 001 AGCTCAATCAGTCACAGTGTAAGGAGGCCAAGTGCAGAGCGAGTATATAGGACTAA 36 060

1 36 061 AAAATGACGTAACGGTTAAAGTCCACAAAAACACCCAGAAAACCGCACGCGAACCTACG 36 120
2 36 061 AAAATGACGTAACGGTTAAAGTCCACAAAAACACCCAGAAAACCGCACGCGAACCTACG 36 120

1 36 121 CCCAGAACGAAAGCCAAAAACCCACAACCTCCTCAAATCGTCACTCCGTTTCCCAC 36 180
2 36 121 CCCAGAACGAAAGCCAAAAACCCACAACCTCCTCAAATCGTCACTCCGTTTCCCAC 36 180

1 36 181 GTTACGTCACTTCCCATTAAAGAAAACATACAATTCCAACACATACAAGTTACTCCGCC 36 240
2 36 181 GTTACGTCACTTCCCATTAAAGAAAACATACAATTCCAACACATACAAGTTACTCCGCC 36 240

1 36 241 CTAAAACCTACGTCACCCGCCCCGTTCCACGCCACGTACAAACTCCACCC 36 300
2 36 241 CTAAAACCTACGTCACCCGCCCCGTTCCACGCCACGTACAAACTCCACCC 36 300

1 36 301 CTCATTATCATATTGGCTTCAATCCAAAATAAGGTATATTATTGATGATG 36 350
2 36 301 CTCATTATCATATTGGCTTCAATCCAAAATAAGGTATATTATTGATGATG 36 350

Sequence Logo: 50% GC base composition

Consensus Threshold: >50%

Compare to: the consensus

Bases that match the reference are marked with yellow highlighting.

Created: 22. tammik. 2021

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