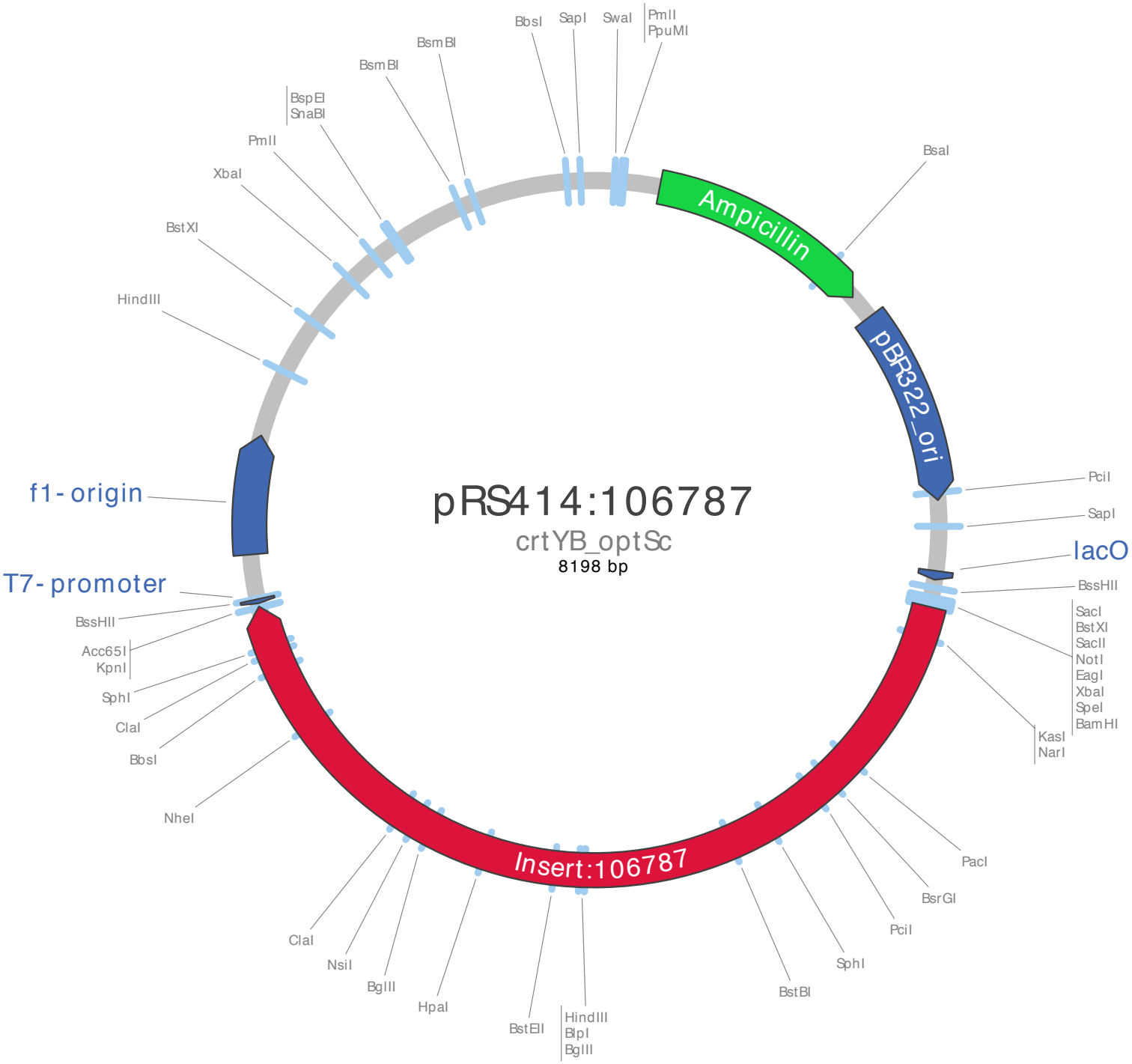
DNA 2.0

Plasmid Map

pRS414:106787

Only single cutters are shown in the map.



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Feature Map

Name	Start	End	
Insert: 106787	2360	5839	
Ampicillin	250	1107	
pBR322_ori	1204	1954	
lacO	2222	2256	
T7-promoter	5867	5851	Reverse
f1-origin	6038	6493	

Restriction Map

Name	Sequence	5' Cut Positions
Acc65I	GGTACC	5835
AccI	GTMKAC	2745, 3196, 3630, 7298
AclI	AACGTT	435, 808, 6481
AlwNI	CAGNNNCTG	1516, 5184, 5772
ApaLI	GTGCAC	365, 1611, 2956, 6551, 7552
AseI	ATTAAT	862, 2097, 2156, 3015, 3714, 4450, 7253, 7367, 8124
BamHI	GGATCC	2359
BbsI	GAAGAC	5595, 8083(C)
BglI	GCCNNNNNGGC	919, 2550, 6023
BglII	AGATCT	4155, 4740
BlpI	GCTNAGC	4136
BsaI	GGTCTC	971
BsmBI	CGTCTC	7684, 7726(C)
BspEI	TCCGGA	7397
BspHI	TCATGA	197, 1205, 5809, 7776
BsrBI	CCGCTC	195, 1996, 2237(C), 2345, 5815(C), 6124
BsrDI	GCAATG	803, 977(C), 3477, 3937(C), 4998(C)
BsrGI	TGTACA	3125
BssHII	GCGCGC	2286, 5869
BstBI	TTCGAA	3569
BstEII	GGTNACC	4250
BstXI	CCANNNNNNTGG	2335, 6972
BtsI	GCAGTG	618, 638(C), 2145, 3285(C), 4914(C)
ClaI	ATCGAT	4873, 5653
EagI	CGGCCG	2340
EcoRV	GATATC	3104, 5121, 5396, 6981
HincII	GTYRAC	4369, 4415, 4522
HindIII	AAGCTT	4133, 6749
HpaI	GTTAAC	4522
KasI	GGCGCC	2478
KpnI	GGTACC	5839
MfeI	CAATTG	2501, 4495, 7122
NarI	GGCGCC	2479

Name	Sequence	5' Cut Positions
NheI	GCTAGC	5346
NotI	GCGGCCGC	2340
NsiI	ATGCAT	4808
PacI	TTAATTAA	3019
PciI	ACATGT	1925, 3204
PmlI	CACGTG	103, 7306
PpuMI	RGGWCCY	114
PvuI	CGATCG	668, 4375, 5994
PvuII	CAGCTG	2105, 4008, 5963
SacI	GAGCTC	2327
SacII	CCGCGG	2334
SapI	GCTCTTC	2042(C), 8136(C)
SnaBI	TACGTA	7414
SpeI	ACTAGT	2353
SphI	GCATGC	3412, 5687
SspI	AATATT	232, 3095, 3778, 6476
SwaI	ATTTAAAT	79
XbaI	TCTAGA	2347, 7178
XmnI	GAANNNNTTC	437, 5035, 7368, 7447, 8114
AfeI	AGCGCT	no cuts
AgeI	ACCGGT	no cuts
ApaI	GGGCCC	no cuts
AscI	GGCGCGCC	no cuts
AsiSI	GCGATCGC	no cuts
AvaI	CYCGRG	no cuts
AvrII	CCTAGG	no cuts
BclI	TGATCA	no cuts
BsiWI	CGTACG	no cuts
EcoRI	GAATTC	no cuts
FseI	GGCCGGCC	no cuts
MluI	ACGCGT	no cuts
MscI	TGGCCA	no cuts
NcoI	CCATGG	no cuts
NdeI	CATATG	no cuts
NruI	TCGCGA	no cuts
PmeI	GTTTAAAC	no cuts
PshAI	GACNNNNGTC	no cuts
PspOMI	GGGCCC	no cuts
PspXI	VCTCGAGB	no cuts
PstI	CTGCAG	no cuts
RsrII	CGGWCCG	no cuts
SalI	GTCGAC	no cuts
SanDI	GGGWCCC	no cuts
SbfI	CCTGCAGG	no cuts
SfiI	GGCCNNNNNGGCC	no cuts
SmaI	CCCGGG	no cuts
XhoI	CTCGAG	no cuts
XmaI	CCCGGG	no cuts

Sequence

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   1 CGGAAAACAA AAACTATTTT TTCTTTAATT TCTTTTTTTA CTTTCTATTT TTAATTTATA
  61 TATTTATATT AAAAAATTTA AATTATAATT ATTTTTATAG CACGTGATGA AAAGGACCCA
 121 GGTGGCACTT TTCGGGGAAA TGTGCGCGGA ACCCCTATTT GTTTATTTTT CTAAATACAT
 181 TCAAATATGT ATCCGCTCAT GAGACAATAA CCCTGATAAA TGCTTCAATA ATATTGAAAA
 241 AGGAAGAGTA TGAGTATTCA ACATTTCCGT GTCGCCCTTA TTCCCTTTTT TGCGGCATTT
 301 TGCCTTCCTG TTTTTGCTCA CCCAGAAACG CTGGTGAAAG TAAAAGATGC TGAAGATCAG
 361 TTGGGTGCAC GAGTGGGTTA CATCGAACTG GATCTCAACA GCGGTAAGAT CCTTGAGAGT
 421 TTTCGCCCCG AAGAACGTTT TCCAATGATG AGCACTTTTA AAGTTCTGCT ATGTGGCGCG
 481 GTATTATCCC GTATTGACGC CGGGCAAGAG CAACTCGGTC GCCGCATACA CTATTCTCAG
 541 AATGACTTGG TTGAGTACTC ACCAGTCACA GAAAAGCATC TTACGGATGG CATGACAGTA
 601 AGAGAATTAT GCAGTGCTGC CATAACCATG AGTGATAACA CTGCGGCCAA CTTACTTCTG
 661 ACAACGATCG GAGGACCGAA GGAGCTAACC GCTTTTTTGC ACAACATGGG GGATCATGTA
 721 ACTCGCCTTG ATCGTTGGGA ACCGGAGCTG AATGAAGCCA TACCAAACGA CGAGCGTGAC
 781 ACCACGATGC CTGTAGCAAT GGCAACAACG TTGCGCAAAC TATTAACTGG CGAACTACTT
 841 ACTCTAGCTT CCCGGCAACA ATTAATAGAC TGGATGGAGG CGGATAAAGT TGCAGGACCA
 901 CTTCTGCGCT CGGCCCTTCC GGCTGGCTGG TTTATTGCTG ATAAATCTGG AGCCGGTGAG
 961 CGTGGGTCTC GCGGTATCAT TGCAGCACTG GGGCCAGATG GTAAGCCCTC CCGTATCGTA
1021 GTTATCTACA CGACGGGGAG TCAGGCAACT ATGGATGAAC GAAATAGACA GATCGCTGAG
1081 ATAGGTGCCT CACTGATTAA GCATTGGTAA CTGTCAGACC AAGTTTACTC ATATATACTT
1141 TAGATTGATT TAAAACTTCA TTTTTAATTT AAAAGGATCT AGGTGAAGAT CCTTTTTGAT
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1381 TTTCCGAAGG TAACTGGCTT CAGCAGAGCG CAGATACCAA ATACTGTCCT TCTAGTGTAG
1441 CCGTAGTTAG GCCACCACTT CAAGAACTCT GTAGCACCGC CTACATACCT CGCTCTGCTA
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1741 ACAGGAGAGC GCACGAGGGA GCTTCCAGGG GGAAACGCCT GGTATCTTTA TAGTCCTGTC
1801 GGGTTTCGCC ACCTCTGACT TGAGCGTCGA TTTTTGTGAT GCTCGTCAGG GGGGCGGAGC
1861 CTATGGAAAA ACGCCAGCAA CGCGGCCTTT TTACGGTTCC TGGCCTTTTG CTGGCCTTTT
1921 GCTCACATGT TCTTTCCTGC GTTATCCCCT GATTCTGTGG ATAACCGTAT TACCGCCTTT
1981 GAGTGAGCTG ATACCGCTCG CCGCAGCCGA ACGACCGAGC GCAGCGAGTC AGTGAGCGAG
2041 GAAGCGGAAG AGCGCCCAAT ACGCAAACCG CCTCTCCCCG CGCGTTGGCC GATTCATTAA
2101 TGCAGCTGGC ACGACAGGTT TCCCGACTGG AAAGCGGGCA GTGAGCGCAA CGCAATTAAT
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2641 GAGAAGCAAT AAGAATGCCG GTTGGGGTTG CGATGATGAC GACCACGACA ACTGGTGTCA

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2761 GCCAAGACTT GCCAGACGCG AGTTTGCCGG TGGTGCGAAC AATAGAGCGA CCATGACCTT

2821 GAAGGTGTGA CGCGCATAAC CGCTAGAGTA CTTTGAAGAG GAAACAGCAA TAGGGTTGCT

2881 ACCAGTATAA ATAGACAGGT ACATACAACA CTGGAAATGG TTGTCTGTTT GAGTACGCTT

2941 TCAATTCATT TGGGTGTGCA CTTTATTATG TTACAATATG GAAGGGAACT TTACACTTCT

3001 CCTATGCACA TATATTAATT AAAGTCCAAT GCTAGTAGAG AAGGGGGGTA ACACCCCTCC

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3121 CGGGTGTACA ATATGGACTT CCTCTTTTCT GGCAACCAAA CCCATACATC GGGATTCCTA

3181 TAATACCTTC GTTGGTCTAC CTAACATGTA GGTGGCGGAG GGGAGATATA CAATAGAACA

3241 GATACCAGAC AAGACATAAT GGGCTAAACA AGACTACACC AATTACACTG CCTCATTGAT

3301 GGTGGTACAT AACGAACTAA TACTGTAGCC CTAGACTTGA TAGCCATCAT CATATCGAAG

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3541 GTTGTTCCAG AGCTGATGAG GGGTATCTTC GAACACACGA AACTTTTTCC TTCCTTCATT

3601 CACGCACACT ACTCTCTAAT GAGCAACGGT ATACGGCCTT CCTTCCAGTT ACTTGAATTT

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3781 TTCAAGCTAT ACCAAGCATA CAATCAACTA AAATGACGGC TCTCGCATAT TACCAGATCC

3841 ATCTGATCTA TACTCTCCCA ATTCTAGGCC TATTAGGACT CTTGACATCT CCTATCTTGA

3901 CGAAATTTGA CATCTACAAG ATAAGTATAT TGGTATTCAT TGCTTTCTCA GCTACAACAC

3961 CTTGGGACAG TTGGATTATT AGAAACGGCG CATGGACTTA CCCATCAGCT GAATCTGGAC

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5101 TACAAGGGTT GATTCCAAGA TATCCATTGG ACGAACTATT GAGAGGTTAC ACTACTGACT

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6781 GTCAGAAATC GAGTTCCAAT CCAAAAGTTC ACCTGTCCCA CCTGCTTCTG AATCAAACAA
6841 GGGAATAAAC GAATGAGGTT TCTGTGAAGC TGCACTGAGT AGTATGTTGC AGTCTTTTGG
6901 AAATACGAGT CTTTTAATAA CTGGCAAACC GAGGAACTCT TGGTATTCTT GCCACGACTC
6961 ATCTCCATGC AGTTGGACGA TATCAATGCC GTAATCATTG ACCAGAGCCA AAACATCCTC
7021 CTTAGGTTGA TTACGAAACA CGCCAACCAA GTATTTCGGA GTGCCTGAAC TATTTTTATA
7081 TGCTTTTACA AGACTTGAAA TTTTCCTTGC AATAACCGGG TCAATTGTTC TCTTTCTATT
7141 GGGCACACAT ATAATACCCA GCAAGTCAGC ATCGGAATCT AGAGCACATT CTGCGGCCTC
7201 TGTGCTCTGC AAGCCGCAAA CTTTCACCAA TGGACCAGAA CTACCTGTGA AATTAATAAC
7261 AGACATACTC CAAGCTGCCT TTGTGTGCTT AATCACGTAT ACTCACGTGC TCAATAGTCA
7321 CCAATGCCCT CCCTCTTGGC CCTCTCCTTT TCTTTTTTCG ACCGAATTAA TTCTTAATCG
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7501 TGCTGTCTAT TAAATGCTTC CTATATTATA TATATAGTAA TGTCGTTTAT GGTGCACTCT
7561 CAGTACAATC TGCTCTGATG CCGCATAGTT AAGCCAGCCC CGACACCCGC CAACACCCGC
7621 TGACGCGCCC TGACGGGCTT GTCTGCTCCC GGCATCCGCT TACAGACAAG CTGTGACCGT
7681 CTCCGGGAGC TGCATGTGTC AGAGGTTTTC ACCGTCATCA CCGAAACGCG CGAGACGAAA
7741 GGGCCTCGTG ATACGCCTAT TTTTATAGGT TAATGTCATG ATAATAATGG TTTCTTAGGA
7801 CGGATCGCTT GCCTGTAACT TACACGCGCC TCGTATCTTT TAATGATGGA ATAATTTGGG

7861 AATTTACTCT GTGTTTATTT ATTTTTATGT TTTGTATTTG GATTTTAGAA AGTAAATAAA

7921 GAAGGTAGAA GAGTTACGGA ATGAAGAAAA AAAAATAAAC AAAGGTTTAA AAAATTTCAA

7981 CAAAAAGCGT ACTTTACATA TATATTTATT AGACAAGAAA AGCAGATTAA ATAGATATAC

8041 ATTCGATTAA CGATAAGTAA AATGTAAAAT CACAGGATTT TCGTGTGTGG TCTTCTACAC

8101 AGACAAGATG AAACAATTCG GCATTAATAC CTGAGAGCAG GAAGAGCAAG ATAAAAGGTA

8161 GTATTTGTTG GCGATCCCCC TAGAGTCTTT TACATCTT

Only the synthesized DNA fragment (in red) has been sequence verified. We do not guarantee the vector sequence.