

NsI I  
SphI  
Ppu10 I

EcoNI |  
CTCGATTAGGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAAAGCATGCATCTCAATT  
GAGCTAACCCACACCTTCAGGGTCCGAGGGTCTCGTCCGTCTTCATACGTTCTGTACGTAGAGTTAAT  
10 20 30 40 50 60 70

NsI I  
SphI  
Ppu10 I

SexAI |  
GTCAGCAACCAGGTGTGGAAAGTCCCCAGGCTCCCCAGCAGGCAGAAGTATGCAAAGCATGCATCTCAAT  
CAGTCGTTGGTCCACACCTTCAGGGTCCGAGGGTCTCGTCCGTCTTCATACGTTCTGTACGTAGAGTTA  
80 90 100 110 120 130 140

TAGTCAGCAACCATAGTCCC GCCC CTA ACTCCGCC ATCCGCC CTA ACTCCGCC AGTTCCGCC ATT  
ATCAGTCGTTGGTATCAGGGCGGGGATTGAGGC GG GTAGGGCGGGATTGAGGC GG GTCAAGGC GG GTAA  
150 160 170 180 190 200 210

SfiI  
Bgl I

NcoI |  
CTCCGCC C ATGGCTGACTAATTTTTTATT TATGCAGAGGCC GAGGCC G C C T C G G C C T C T G A G C T A T T  
GAGGCC GGGTACCGACTGATTAAAAAAATAACGTCTCCGGCTCCGGAGCCGGAGACTCGATAA  
220 230 240 250 260 270 280

AvrII  
StuI  
BseRI |||  
ClaI  
BsABl

CCAGAAGTAGTGAGGAGGCTTTTGAGGCC TAGGCTTTGCAAAGATCGATCAAGAGAGACAGGATGAGG  
GGTCTTCATCACTCCTCCGAAAAAACCTCCGATCCGAAAACGTTCTAGCTAGTTCTGTCCCTACTCC  
290 300 310 320 330 340 350

BspMI  
EagI

ATCGTTCGCATGATTGAACAAGATGGATTGCACGCAGGTTCTCCGGCGCTGGGTGGAGAGGGTATT  
TAGCAAAGCGTACTAACTGTTCTACCTAACGTGCGTCAAGAGGCCGAAACCCACCTCCGATAAG  
360 370 380 390 400 410 420

NarI  
KasI ||

GGCTATGACTGGGCACAAACAGACAATCGGCTGCTCTGATGCCGCCGTGTTCCGGCTGTCAGCGCAGGGC  
CCGATACTGACCCGTGTTGTCTGTTAGCCGACGAGACTACGGCGGCACAAGGCCGACAGTCGCGTCCCCG  
430 440 450 460 470 480 490

BbeI  
EheI  
DrdI

GCCCCGGTTCTTTGTCAAGACCGACCTGTCCGGTGCCTGAATGAACTGCAAGACGAGGCAGCGCG  
CGGGCCAAGAAAAACAGTTCTGGCTGGACAGGCCACGGGACTTACTTGACGTTCTGCTCCGTGCGCCGA  
500 510 520 530 540 550 560

PvuII

MscI	FspI	Tth111 I
ATCGTGGCTGGCCACGACGGCGTCCTGCGCAGCTGTGCTCGACGTTGTCACTGAAGCGGGAAAGGGAC		
TAGCACCGACCGGTGCTGCCGCAAGGAACGCGTCGACACGAGCTGCAACAGTGAATTGCCCTCCCTG		
570      580      590      600      610      620      630		

Eco57 I

BsrDI	BspMI
GGTAGTACCGACTACGTTACGCCGACGTATGCGAACTAGGCCGATGGACGGTAAGCTGGTGGTCG	
710      720      730      740      750      760      770	

SapI  
EarI

NcoI	
GAGCATCAGGGCTCGGCCAGCCGAAGTGGATGGAAGCCGGTCTGTCGATCAGGATGATCTGGACGAA	
CTTTGTAGCGTAGCTCGCTCGTGCATGAGCCTACCTTCCGCCAGAACAGCTAGTCCTACTAGACCTGCTT	
780      790      800      810      820      830      840	

SphI

NaeI	RsrII	SapI EarI
TCGTCGTGACCCATGGCGATGCCCTGCTTGCGAATATCATGGTGGAAAATGCCGCTTTCTGGATTCA		
AGCAGCACTGGGTACCGCTACGGACGAACGGCTTATAGTACCACTTTACCGCGAAAAGACCTAAGTA		
920      930      940      950      960      970      980		

NgoMI

Eco57 I	BssSI	BsrBI
GAGCTGGCGCGAATGGGCTGACCGCTTCCTCGTGCCTTACGGTATGCCGCTCCGATTGAGCGCA		
CTCGAACCGCCGCTTACCGACTGGGAAGGAGCACGAAATGCCATAGCGCGAGGGCTAAGCGTCGCGT		
1060      1070      1080      1090      1100      1110      1120		

BsrBI

BstBI	
TCGCCTTCTATGCCCTTCTGACGAGTTCTCTGAGCGGGACTCTGGGTTGAAATGACCGACCAAGCG	
AGCGGAAGATAGCGGAAGAACTGCTCAAGAACGACTGCCCTGAGACCCCCAAGCTTACTGGCTGGTCG	
1130      1140      1150      1160      1170      1180      1190	

BssSI  
BspMI  
|  
ACGCCAACCTGCCATCACGAGATTGATTCCACCGCCGCCTCTATGAAAGGTTGGCTTCGGAATCG  
TGC GGTTGGACGGTAGTGCTCTAAAGCTAAGGTGGCGCGGAAGATACTTCCAACCCGAAGCCTTAGC  
1200 1210 1220 1230 1240 1250 1260

NaeI  
NgoMI  
BpmI  
| |  
TTTCCGGGACGCCGGCTGGATGATCCTCCAGCGCGGGGATCTCATGCTGGAGTTCTCGCCCACCC TAG  
AAAAGGCCCTGCGGCCGACCTACTAGGAGGTGCGGCCCTAGAGTACGACCTCAAGAACGGTGGGATC  
1270 1280 1290 1300 1310 1320 1330

BpmI  
AvrII  
| |  
GGGGAGGCTAACTGAAACACGGAAGGAGACAATACCGGAAGGAACCCCGCGCTATGACGGCAATAAAAAGA  
CCCCTCCGATTGACTTGTGCCTTCCTCTGTTATGGCCTTCCTGGCGCGATACTGCCGTATTTTCT  
1340 1350 1360 1370 1380 1390 1400

BseRI  
BsrDI  
| |  
CAGAATAAAACGCACGGTGGCTTGGCTTGGCTGAGGAGCTTGGCCCATGGCATACGGTATCCATA  
GTCTTATTTGCGTGCCACAACCCAGCAAACAGCTCCTCGAACCGGGTAACGTATGCAACATAGGTAT  
1410 1420 1430 1440 1450 1460 1470

BsrGI  
|  
TCATAATATGTACATTATATTGGCTCATGTCAAACATTACCGCCATGTTGACATTGATTATTGACTAGT  
AGTATTATACATGTAAATATAACCGAGTACAGGTTGTAATGGCGGTACAACGTAACTAAACTGATCA  
1480 1490 1500 1510 1520 1530 1540

SpeI  
|  
TATTAATAGTAATCAATTACGGGTCAATTAGTCATAGCCATATATGGAGTTCCGCGTTACATAACTTA  
ATAATTATCATTAGTTAATGCCCAAGTAATCAAGTATCGGGTATATACCTCAAGGCATGTATTGAAT  
AseI  
|  
CGGTAAATGGCCCGCCTGGCTGACCGCCAAACGACCCCCGCCATTGACGTCAATAATGACGTATGTTCC  
GCCATTACCGGGCGGACCGACTGGCGGGTTGCTGGGGCGGGTAACTCAGTTATTACTGCATACAAGG  
1620 1630 1640 1650 1660 1670 1680

Bgl I AatII  
| |  
CATAGTAACGCCAATAGGACTTTCCATTGACGTCAATGGGGAGTATTACGCTAAACTGCCACTTG  
GTATCATTGCGGTTATCCCTGAAAGGTAACTCAGTTACCCACCTCATAATGCGATTGACGGGTGAAC  
1690 1700 1710 1720 1730 1740 1750

AatII Bgl I  
| |  
GCAGTACATCAAGTGTATCATATGCCAAGTACGCCCTATTGACGTCAATGACGGTAAATGGCCCGCCT  
CGTCATGTAGTTCACATAGTACGGTCATGCGGGGATAACTGCAGTTACTGCCATTACCGGGCGGA  
1760 1770 1780 1790 1800 1810 1820



The figure displays a DNA sequence with various restriction enzyme cleavage sites indicated by vertical lines and labels. The sequence is numbered from 2460 to 3080. Key features include:

- BstXI**: Cleavage site at position 2490.
- PpuMI**: Cleavage site at position 2520.
- BcgI**: Cleavage site at position 2570.
- BsgI**: Cleavage site at position 2660.
- BseRI**: Cleavage site at position 2600.
- SspI**: Cleavage site at position 2730.
- AseI**: Cleavage site at position 2780.
- EarI**: Cleavage site at position 2850.
- NheI**: Cleavage site at position 2860.
- BpmI**: Cleavage site at position 2990.
- Bgl I**: Cleavage site at position 2990.
- Clai**: Cleavage site at position 3060.
- BamHI**: Cleavage site at position 3050.
- Eco57 I**: Cleavage site at position 3060.
- NcoI**: Cleavage site at position 3010.

The sequence itself consists of several blocks of DNA, each starting with a different nucleotide (A, T, C, G) and containing various restriction sites. The labels above the sequence indicate the positions of these sites.

PpuMI  
 AGTTGGGGACCTTGATTGTTCTTCTTCGCTATTGAAAATTCATGTTATATGGAGGGGCAAA  
 TCAAACCCCTGGGAACATAACAAGAAGAAAAAGCGATAACATTAAAGTACAATATACCTCCCCGTT  
 3090 3100 3110 3120 3130 3140 3150

NcoI BspHI  
 GTTTCAGGGTGTGTTAGAATGGGAAGATGCCCTGTATCACCATGGACCCATGATAATTTGTT  
 CAAAAGTCCCACAACAAACTTACCCCTCTACAGGGAACATAGGGTACCTGGGAGTACTATTAAAACAA  
 3160 3170 3180 3190 3200 3210 3220

BseRI  
 TCTTCACTTCTACTCTGTTGACAACCATTGTCCTCTTATTTCTTCATTTCTGTAACCTTTC  
 AGAAAAGTGAAGATGAGACAACTGTTGGTAACAGAGGAGATAAAAGAAAAGTAAAAGACATTGAAAAG  
 3230 3240 3250 3260 3270 3280 3290

DraI ScaI  
 GTTAAACTTAGCTGCATTGTAACGAATTAAATTCACTTCGTTATTGTCAGATTGTAAGTAC  
 CAATTGAAATCGAACGTAACATTGCTTAAAATTAAAGTGAAGCAAATAACAGTCTAACATTCATG  
 3300 3310 3320 3330 3340 3350 3360

Eco57 I  
 TTTCTCTAATCACTTTTCAAGGCAATCAGGGTATTATTATTGACTTCAGCACAGTTAGAGAA  
 AAAGAGATTAGTGAACAAAGTTCCGTTAGTCCCATTAAACATGAAGTCGTGTCAAATCTCTT  
 3370 3380 3390 3400 3410 3420 3430

MfeI SspI SspI  
 CAATTGTTATAATTAAATGATAAGGTAGAATTCTGCATATAAAATTCTGGCTGGCGTGGAAATATTCT  
 GTTAACAATATTAATTACTATTCCATCTTATAAAGACGTATATTAAAGACCGACCGCACCTTATAAGA  
 3440 3450 3460 3470 3480 3490 3500

ATAACCATCTTGTGATGGGACCAAGTAGTAGGACGGAAAGAGAAATACCAATGTTACTATATGTGNC  
 3510 3520 3530 3540 3550 3560 3570

ApaI Bsp120 I  
 TTTGAGATGAGGATAAAACTCTGAGTCCAACCGGGCCCTCTGCTAACCATGTTCATGCCTTCTTCT  
 AAACTCTACTCCTATTGAGACTCAGGTTGGCCGGGAGACGATTGGTACAAGTACGGAAAGAAGA  
 3580 3590 3600 3610 3620 3630 3640

AlwNI EcoRI BseRI  
 TTTCCCTACAGCTCCTGGCAACGTGCTGGTTGTGCTGTCTCATCATTGGCAAAGAATTCACTCC  
 AAAAGGATGTCGAGGACCCGTTGCACGACCAACACGACAGAGTAGTAAACCGTTCTTAAGTGAGG  
 3650 3660 3670 3680 3690 3700 3710



SapI  
EarI  
|  
GGAGAGGCGGTTGCGTATTGGGCCTCTTCGCTTCCTCGCTCACTGACTCGCTGCCTCGTCGTTCG  
CCTCTCCGCCAACGCATAACCGCGAGAAGGCGAAGGAGCGAGTGACTGAGCGACGCCAGCAAGC  
4420 4430 4440 4450 4460 4470 4480

BsrBI  
|  
GCTGCGCGAGCGGTATCAGCTCACTCAAAGGCGTAATACGGTTATCCACAGAATCAGGGATAACGCA  
CGACGCCGCTGCCATAGTCGAGTGAGTTCCGCCATTATGCCAATAGGTGCTTAGTCCCCTATTGCGT  
4490 4500 4510 4520 4530 4540 4550

BspLU11 I  
|  
GGAAAGAACATGTGAGCAAAAGGCCAGCAAAAGGCCAGGAACCGTAAAAAGGCCGCGTTGCTGGCGTTTT  
CCTTCTTGTACACTCGTTCCGGTCGTTCCGGCCTTGGCATTTCGGCGAACGACCGCAAAA  
4560 4570 4580 4590 4600 4610 4620

DrdI  
|  
TCCATAGGCTCCGCCCCCTGACGAGCATCACAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGAC  
AGGTATCCGAGGCGGGGGGACTGCTCGTAGTGTAGCTGCGAGTCAGTCTCCACCGCTTGGCTG  
4630 4640 4650 4660 4670 4680 4690

BssSI  
|  
AGGACTATAAGATAACCAGGCCTTCCCCCTGGAAGCTCCCTCGTGCCTCTCCTGTTCCGACCCCTGCCG  
TCCTGATATTCTATGGTCCGAAAGGGGACCTTCGAGGGAGCACGCGAGAGGACAAGGCTGGGACGGC  
4700 4710 4720 4730 4740 4750 4760

CTTACCGGATACTGTCCGCCTTCTCCCTCGGGAAAGCGTGGCGCTTCTCAATGCTCACGCTGTAGGT  
GAATGGCCTATGGACAGGGCGAAAGAGGGAAAGCCCTCGCACCGCAAAGAGATTACGAGTGCGACATCCA  
4770 4780 4790 4800 4810 4820 4830

ApaLI  
|  
ATCTCAGTTCGGTAGGTCGTCGCTCCAAGCTGGCTGTGACGAACCCCCCGTTCAGCCGACCG  
TAGAGTCAGCACATCCAGCAAGCGAGGTTCGACCCACACACGTTGCTGGGGGCAAGTCGGCTGGC  
4840 4850 4860 4870 4880 4890 4900

CTGCGCCTATCCGTAACTATCGTCTTGAGTCCAACCCGGTAAGACACGACTTATGCCACTGGCAGCA  
GACGCCGAAAGGCCATTGATAGCAGAACTCAGGTTGGCCATTCTGTGCTGAATAGCGGTGACCGTCGT  
4910 4920 4930 4940 4950 4960 4970

AlwNI  
|  
GCCACTGGTAACAGGATTAGCAGAGCGAGGTATGTAGGCGGTGCTACAGAGTTCTGAAGTGGTGGCATA  
CGGTGACCATTGTCCTAAATCGTCTCGCTCCATACATCCGCCACGATGTCTCAAGAACATTCAACCACCGGAT  
4980 4990 5000 5010 5020 5030 5040

Eco57 I  
|  
ACTACGGCTACACTAGAAGGACAGTATTGGTATCTCGCTCTGCTGAAGCCAGTTACCTTCGGAAAAAG  
TGATGCCGATGTGATCTCCTGTCAAACCATAGACGCGAGACGACTTCGGTCAATGGAAGCCTTTTC  
5050 5060 5070 5080 5090 5100 5110

AGTTGGTAGCTCTGATCCGGCAAACAAACCACCGCTGGTAGCGGTGGTTTTTGTTGCAAGCAGCAG  
TCAACCATCGAGAACTAGGCCGTTGTTGGTGGCGACCATGCCACCAAAAAAACAGTCGTCGTC  
5120 5130 5140 5150 5160 5170 5180

ATTACGCGCAGAAAAAAAGATCTAAGAAGATCCTTGATCTTCTACGGGGTCTGACGCTCAGTGGAA  
TAATGCGCGTCTTTCTAGAGTCTCTAGGAAACTAGAAAAGATGCCAGACTGCGAGTCACCT  
5190 5200 5210 5220 5230 5240 5250

BspHI | DraI

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ACGAAAACCTACGTTAAGGGATTTGGTCATGAGATTATCAAAAGGATCTCACCTAGATCCTTTAAA
TGCTTTGAGTGCATTCCCTAAACCAGTACTCTAATAGTTTCCTAGAAGTGGATCTAGGAAAATT
      5260      5270      5280      5290      5300      5310      5320

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DraI  
 |  
 TTAATAATGAAGTTTAAATCAATCTAAAGTATATATGAGTAAACTGGTCTGACAGTTACCAATGCTTA  
 AATTTTACTTCAAAATTAGTTAGATTTCATATATACTCATTGAACCAGACTGTCAATGGTTACGAAT  
 5330 5340 5350 5360 5370 5380 5390

ATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTCGTTCATCCATAGTTGCCTGACTCCCCGTCGTGT  
 TAGTCACTCCGGATAGAGTCGCTAGACAGATAAAGCAAGTAGGTATCAACGGACTGAGGGGCAGCACA  
           5400          5410          5420          5430          5440          5450          5460

AGATAACTACGATACGGGAGGGCTTACCATCTGGCCCCAGTGCTGCAATGATAACCGCGAGACCCACGCTC  
 TCTATTGATGCTATGCCCTCCCGAATGGTAGACCGGGGTCACGACGTTACTATGGCGCTCTGGGTGCGAG  
 5479 5489 5499 5509 5519 5529 5539

TGGCCGAGGTCTAAATAGTCGTTATTGGTCGGTCGGCCTTCCCGGCTCGCGTCTTCACCAGGACGTTGA  
5540 5550 5560 5570 5580 5590 5600

AseI

```

TTATCCGCCTCCATCCAGTCTATTATTGTTGCCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTT
AATAGGCGGAGGTAGGTCAAGATAATTAAACAACGGCCCTTCGATCTCATTCAAGCGGTCAATTATCAA
      5610      5620      5630      5640      5650      5660      5670

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Psp1406 I  
FspI BsrDI  
TGCGAACGTTGTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTCGTTGGTATGGCTTCATTCA  
ACCGCTTGCAACAACGGTAACGATGTCCGTAGCACCACAGTGCAGCAGCAAACCATACCGAAGTAAGTC  
5680 5690 5700 5710 5720 5730 5740

CTCCGGTTCCCAACGATCAAGGCAGGTACATGATCCCCCATGTTGCAAAAAAGCGGTTAGCTCCTTC  
GAGGCCAACGGGTTGCTAGTCCGCTCAATGTACTAGGGGGTACAACACGTTTTCGCCAATCGAGGAAG  
5750 5760 5770 5780 5790 5800 5810

PvuI

GGTCCTCCGATCGTTGTCAGAAGTAAGTTGGCCGCAGTGTATCACTCATGGTTATGGCAGCACCGATA  
CCAGGAGGCTAGAACAGTCTTCATTCAACCGGCGTACAATAGTGAGTACCAATACCGTCGTACGTAT  
5820 5830 5840 5850 5860 5870 5880

ScaI

ATTCTCTTACTGTCATGCCATCCGTAAGATGCTTTCTGTGACTGGTGAGTACTCAACCAAGTCATTCTG  
TAAGAGAAATGACAGTACGGTAGGCATTCTACGAAAAGACACTGACCAACTCATGAGTTGGTCAGTAAGAC  
5890 5900 5910 5920 5930 5940 5950

BcgI

AGAATAGTGTATCGGGCACCGAGTTGCTCTGCCCGGCGTCAATACGGGATAATACCGGCCACATAGC  
TCTTATCACATACGCCGCTGGCTAACGAGAACGGGCCAGTTATGCCCTATTATGGCGCGGTATCG  
5960 5970 5980 5990 6000 6010 6020

BcgI

XbaI

DraI

Psp1406 I

AGAACCTTAAAAGTGCTCATCATTGGAAACGTTCTCGGGCGAAAACCTCTCAAGGATCTTACCGCTGT  
TCTTGAAATTTCACGAGTAGTAAACCTTTGCAAGAACGGCCGCTTGAGAGTTCCTAGAATGGCGACA  
6030 6040 6050 6060 6070 6080 6090

ApalI

Eco57 I

BssSI

TGAGATCCAGTCGATGTAACCCACTCGTGCACCCAAC TGATCTTCAGCATCTTACTTCACCAGCGT  
ACTCTAGGTCAAGCTACATTGGGTGAGCACGTGGTTGACTAGAAGTCGTAGAAAATGAAAGTGGTCGCA  
6100 6110 6120 6130 6140 6150 6160

TTCTGGGTGAGCAAAACAGGAAGGCAAAATGCCGCAAAAAGGGATAAGGGCGACACGGAAATGTTGA  
AAGACCCACTCGTTTGTCCCTCCGTTTACGGCGTTTCCCTTATTCCCGCTGTCCTTACAAC  
6170 6180 6190 6200 6210 6220 6230

EarI SspI

BspHI BsrBI

ATACTCATAACTCTCCTTTCAATATTATTGAAGCATTATCAGGGTTATTGTCTCATGAGCGGATACA  
TATGAGTATGAGAAGGAAAAGTTATAATAACTCGTAAATAGTCCAATAACAGAGTACTCGCCTATGT  
6240 6250 6260 6270 6280 6290 6300

TATTTGAATGTATTAGAAAAATAACAAATAGGGTCCGCGCACATTCCCCGAAAAGTGCCACCTGA  
ATAAAACCTACATAATCTTTTATTGTATCCCAAGGCGCGTGTAAAGGGCTTTCACGGTGGACT  
6310 6320 6330 6340 6350 6360 6370

AatII BspHI

BssSI

CGTCTAAGAAACCATTATTATCATGACATTAACCTATAAAATAGGCGTATCACGAGGCCCTTCGTC  
GCAGATTCTTGGTAATAATAGTACTGTAATTGGATATTTCATCCGCATAGTGCTCCGGAAAGCAG  
6380 6390 6400 6410 6420 6430 6440