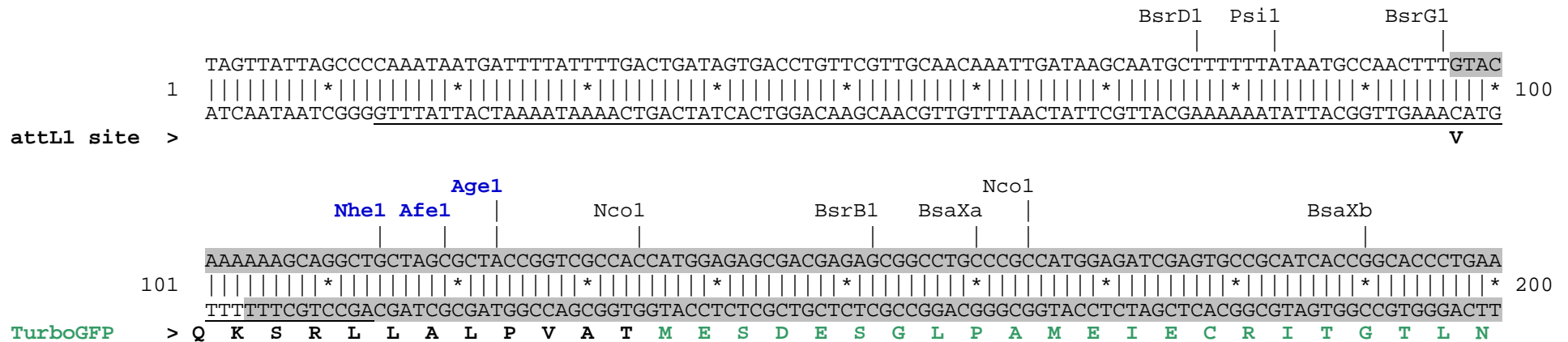
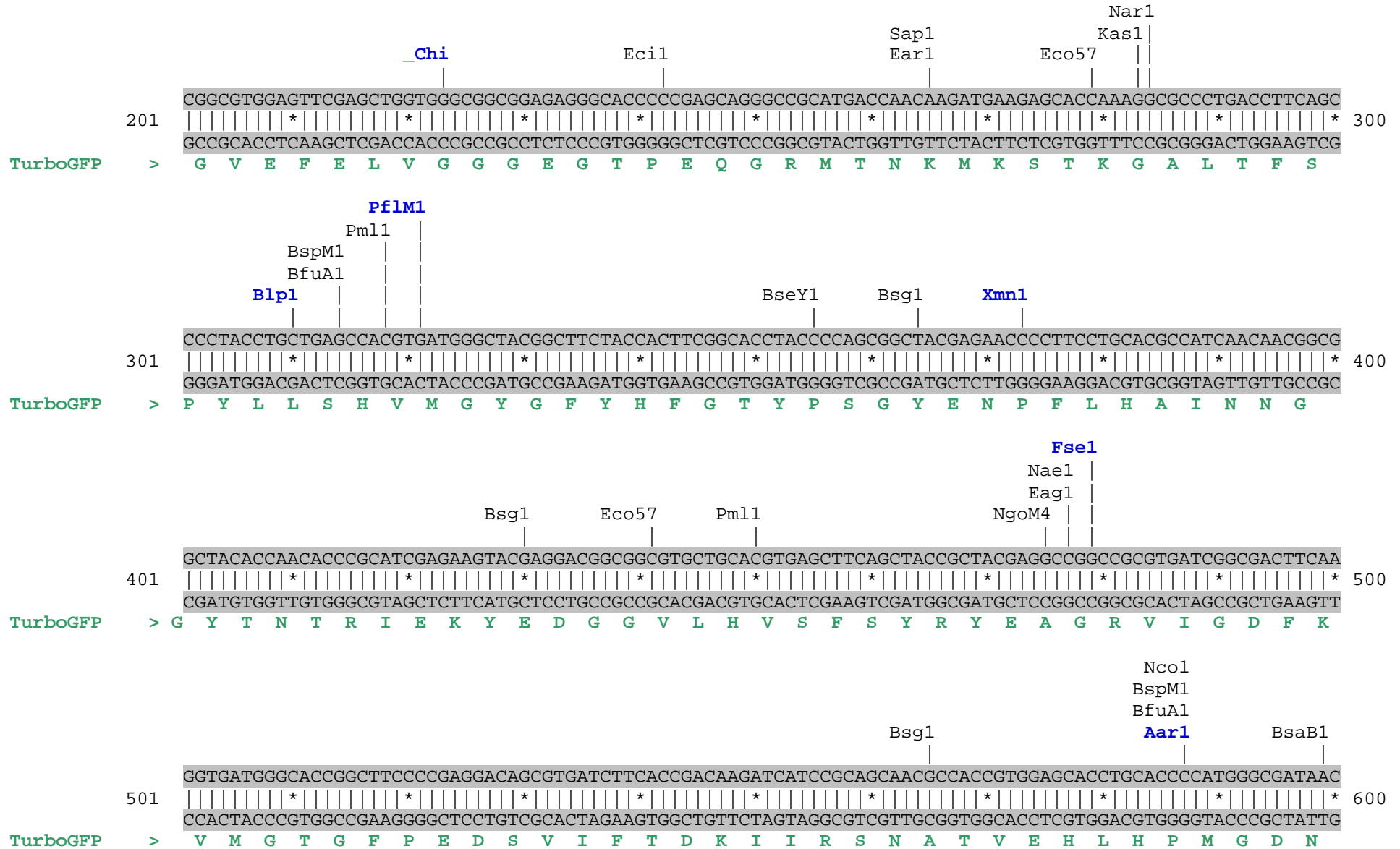


**Gateway® TurboGFP-C entry clone restriction map**

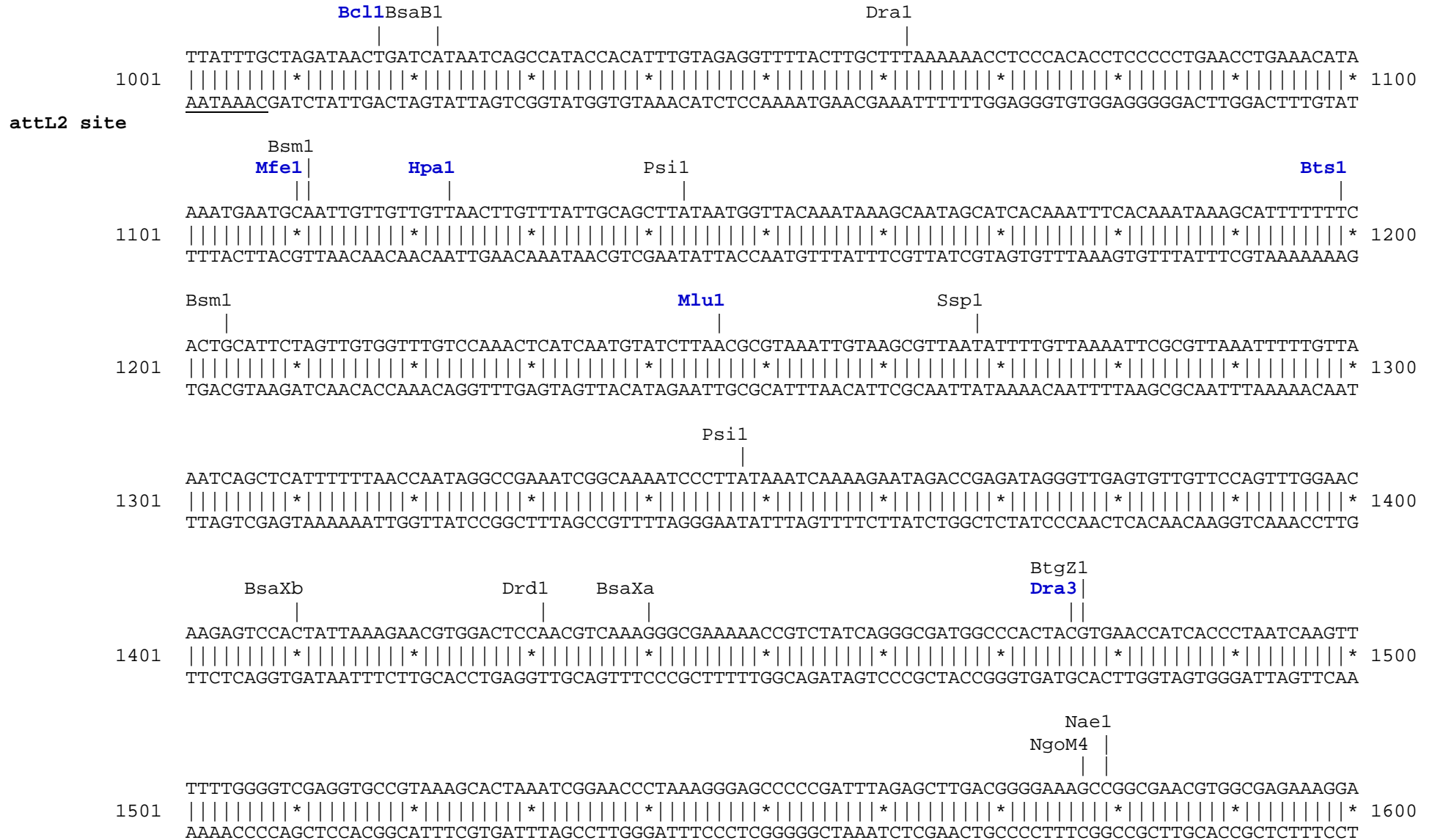
The data has not been verified by restriction digestion with each enzyme listed and does not take into account possible methylation sites. Enzymes that recognize unambiguous sequences less than 6 basepairs long are not included – for the more complete enzyme list please refer to the Table of restriction sites.

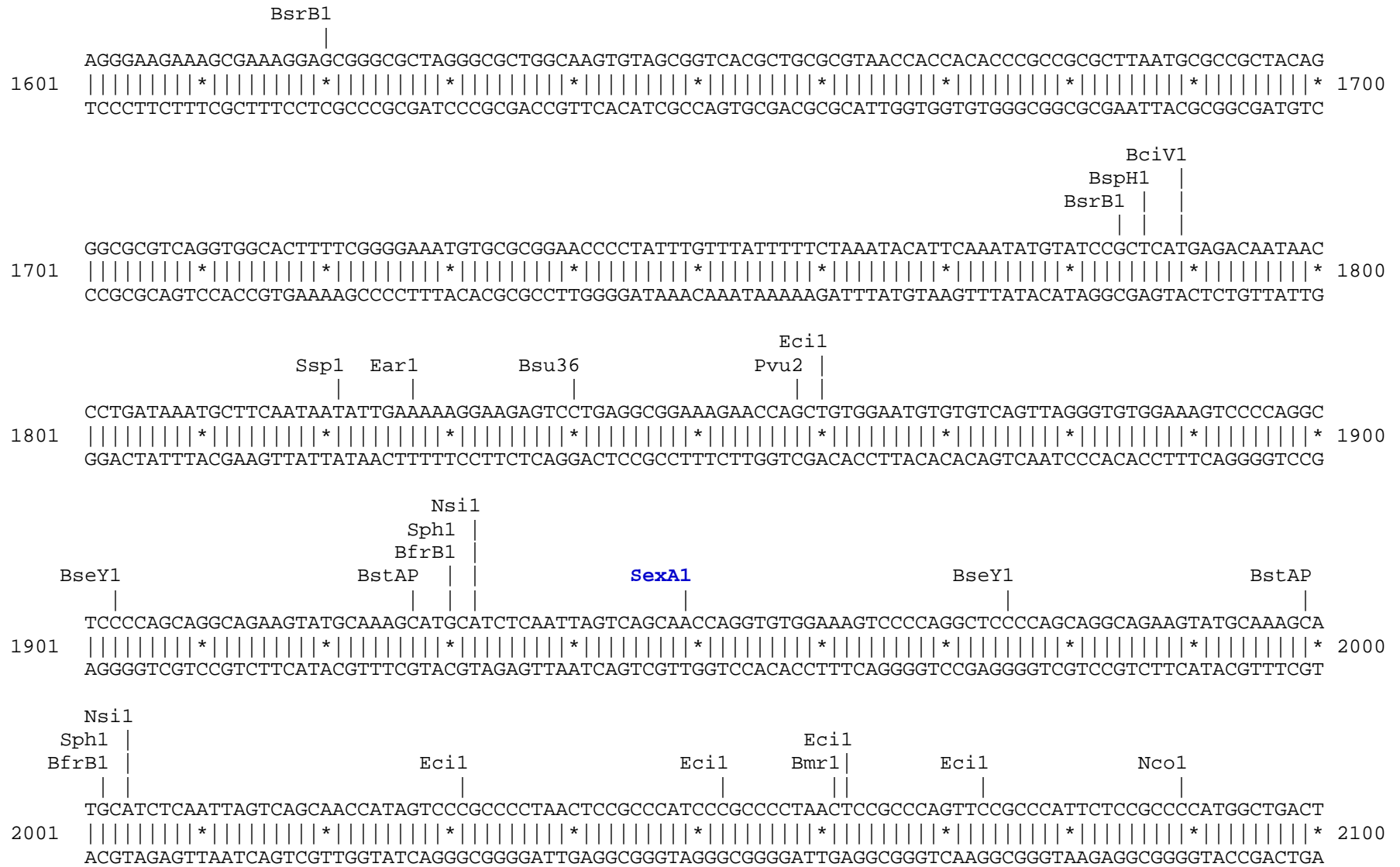
Unique sites are shown in bold blue. The location given specifies the 3' end of the cut DNA (the base to the left of the cut site). MCS sequence is shown in frame, the attL sites are underlined. Shaded regions correspond to DNA sequences transferred from the entry clone into the destination vector following recombination. Non-TurboGFP amino acids coded by those DNA sequences are shown in black, TurboGFP amino acids are shown in green.



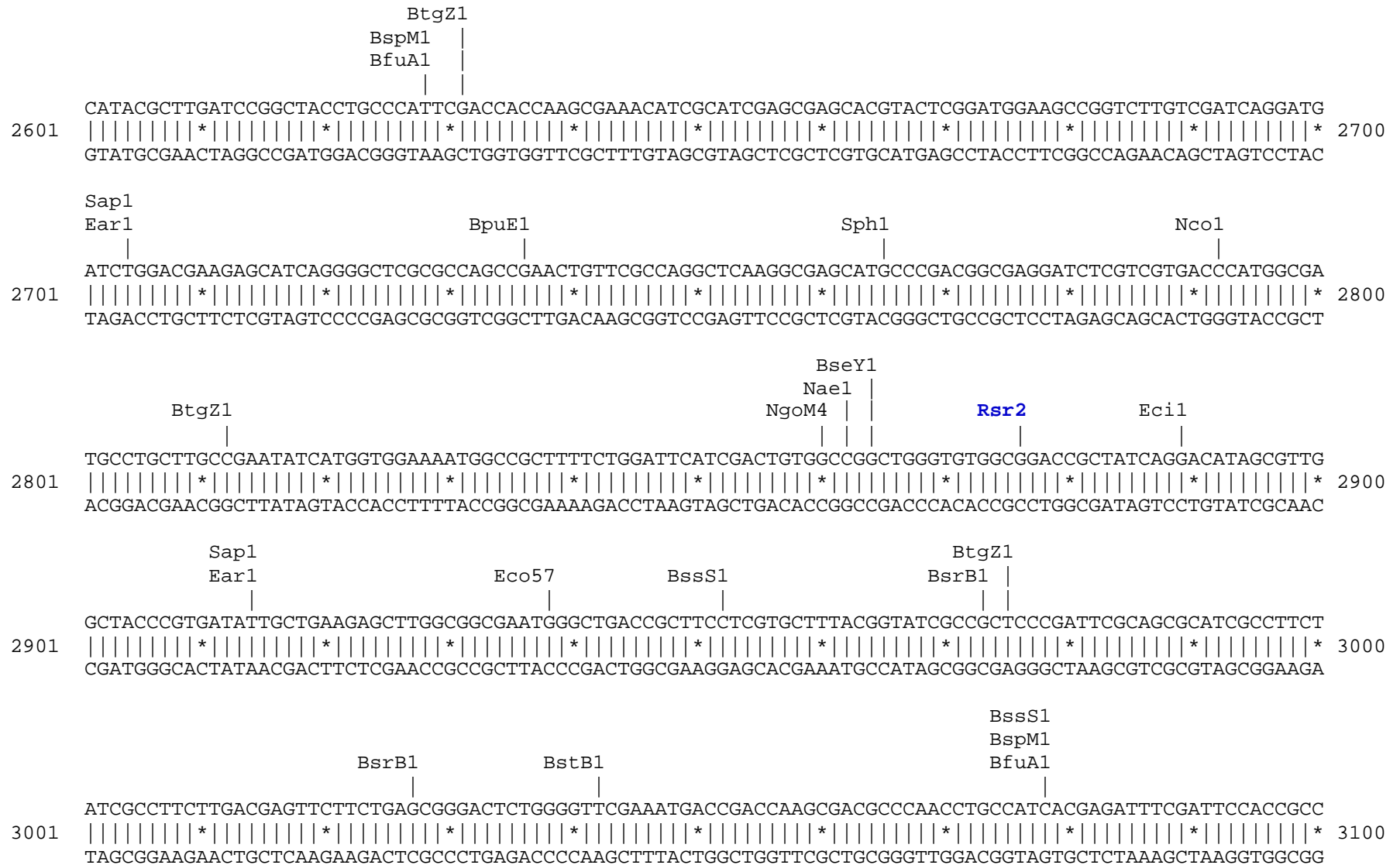












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                                     NaeI
                                     |
                               NgoM4 |
                               Bpm1  |
                                     |
3101 GCCTTCTATGAAAGGTTGGGCTTCGGAATCGTTTTCCGGGACGCCGGCTGGATGATCCTCCAGCGCGGGGATCTCATGCTGGAGTTCTTCGCCCACCCTA
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3200
    CGGAAGATACTTTCCAACCCGAAGCCTTAGCAAAAGGCCCTGCGGCCGACCTACTAGGAGGTCGCGCCCCTAGAGTACGACCTCAAGAAGCGGGTGGGAT

Bpm1
|
3201 GGGGGAGGCTAACTGAAACACGGAAGGAGACAATACCGGAAGGAACCCGCGCTATGACGGCAATAAAAAGACAGAATAAAACGCACGGTGTGGGTGCGTT
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3300
    CCCCCTCCGATTGACTTTGTGCCTTCTCTGTTATGGCCTTCTTGGGCGGATACTGCCGTTATTTTTCTGTCTTATTTTGCCTGCCACAACCCAGCAA

                                     BsaI
                                     |
3301 TGTTTCATAAACCGGGGTTTCGGTCCCAGGGCTGGCACTCTGTGCGATACCCACCGAGACCCCATTGGGGCCAATACGCCCGCGTTTCTTCCTTTTCCCA
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3400
    ACAAGTATTTGCGCCCCAAGCCAGGGTCCCAGCGTGAGACAGCTATGGGGTGGCTCTGGGGTAACCCCGGTTATGCGGGCGCAAAGAAGGAAAAGGGGT

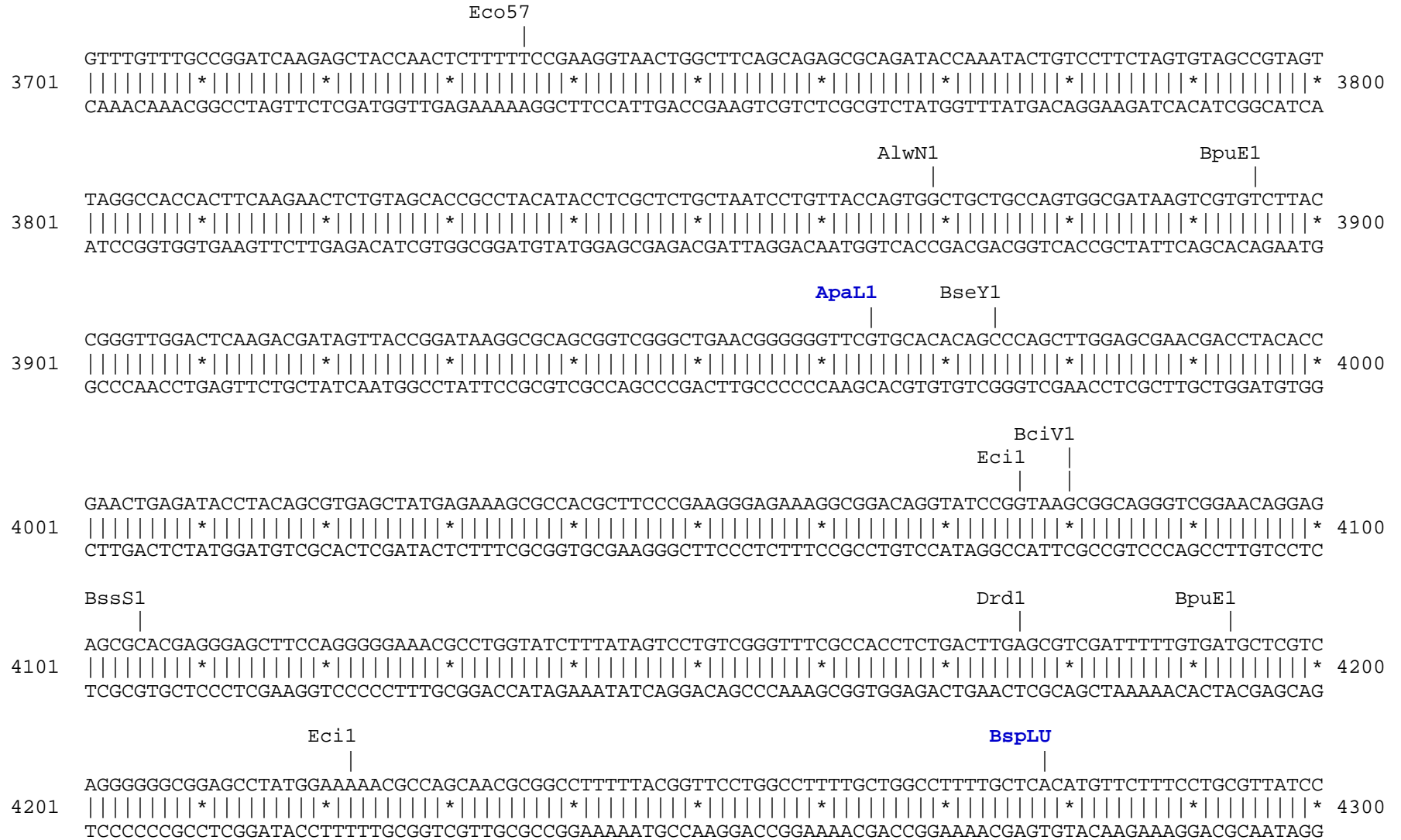
                                     BstAP
                                     |
                               AlwN1 |
                                     |
                               Bsu36 |
                                     |
3401 CCCACCCCCAAGTTCGGGTGAAGGCCAGGGCTCGCAGCCAACGTCGGGGCGGCAGGCCCTGCCATAGCCTCAGGTTACTCATATATACTTTAGATTG
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3500
    GGGGTGGGGGTTCAAGCCACTTCCGGGTCCCGAGCGTCGGTTGCAGCCCCGCCGTCCGGGACGGTATCGGAGTCCAATGAGTATATATGAAATCTAAC

DraI           DraI           BspH1
|             |             |
3501 ATTTAAAACCTTCATTTTTAATTTAAAAGGATCTAGGTGAAGATCCTTTTTGATAATCTCATGACCAAAATCCCTTAACGTGAGTTTTCGTTCCACTGAGC
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3600
    TAAATTTTGAAGTAAAAATTAATTTTCTAGATCCACTTCTAGGAAAACTATTAGAGTACTGGTTTTAGGGAATTGCACTCAAAGCAAGGTGACTCG

                                     BpuE1
                                     |
3601 GTCAGACCCCGTAGAAAAGATCAAAGGATCTTCTTGAGATCCTTTTTTTCTGCGGTAATCTGCTGCTTGCAAACAAAAAAACCACCGCTACCAGCGGTG
    |||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||*|||||* 3700
    CAGTCTGGGGCATCTTTTCTAGTTTCTTAGAAGAACTCTAGGAAAAAAGACGCGCATTAGACGACGAACGTTTGTTTTTTTGGTGGCGATGGTCGCCAC

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                                NsiI
                                BfrB1 |
                                | |
4301 CCTGATTCTGTGGATAACCGTATTACCGCCATGCAT 4336
      |||||*|||||*|||||*|||||
      GGACTAAGACACCTATTGGCATAATGGCGGTACGTA

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Found:

<a href="#">Aar1</a>	<a href="#">Acc65</a>	<a href="#">Afe1</a>	<a href="#">Age1</a>	AlwN1	Apa1	<a href="#">ApaL1</a>	Avr2	<a href="#">BamH1</a>	BciV1	<a href="#">Bcl1</a>	BfrB1	BfuA1	<a href="#">Bgl1</a>
<a href="#">Bgl2</a>	<a href="#">Blp1</a>	Bmr1	Bpm1	BpuE1	<a href="#">Bsa1</a>	BsaB1	BsaXa	BsaXb	BseR1	BseY1	Bsg1	Bsm1	BspH1
<a href="#">BspLU</a>	BspM1	BsrB1	BsrD1	BsrG1	BssS1	BstAP	BstB1	Bsu36	BtgZ1	<a href="#">Bts1</a>	<a href="#">_Chi</a>	<a href="#">Cla1</a>	Dra1
<a href="#">Dra3</a>	Drd1	Eag1	Ear1	Eci1	Eco57	<a href="#">EcoR1</a>	<a href="#">Fse1</a>	<a href="#">Fsp1</a>	<a href="#">Hind3</a>	<a href="#">Hpa1</a>	Kas1	<a href="#">Kpn1</a>	<a href="#">Mfe1</a>
<a href="#">Mlu1</a>	<a href="#">Msc1</a>	Nae1	Nar1	Nco1	NgoM4	<a href="#">Nhe1</a>	Nsi1	<a href="#">PflF1</a>	<a href="#">PflM1</a>	Pml1	Psi1	PspOM	Pst1
Pvu2	<a href="#">Rsr2</a>	<a href="#">Sac1</a>	<a href="#">Sac2</a>	<a href="#">Sal1</a>	Sap1	<a href="#">SexA1</a>	<a href="#">Sfi1</a>	<a href="#">Sma1</a>	Sph1	Ssp1	<a href="#">Stu1</a>	<a href="#">Xho1</a>	<a href="#">Xmn1</a>

Unique:

Aar1	Acc65	Afe1	Age1	ApaL1	BamH1	Bcl1	Bgl1	Bgl2	Blp1	Bsa1	BspLU	Bts1	_Chi
Cla1	Dra3	EcoR1	Fse1	Fsp1	Hind3	Hpa1	Kpn1	Mfe1	Mlu1	Msc1	Nhe1	PflF1	PflM1
Rsr2	Sac1	Sac2	Sall	SexA1	Sfi1	Sma1	Stu1	Xho1	Xmn1				

Not found:

Aat2	Acl1	Afl2	Ahd1	Ale1	Asc1	Ase1	AsiS1	Baela	Baelb	Bbs1	BbvC1	BcglA	Bcglb
BmgB1	Bpu10	BsiW1	BsmB1	BspE1	BssH2	BstE2	BstX1	BstZ1	EcoK	EcoN1	EcoRV	ScFRT	FspA1
I_Ceu	loxP	Nde1	Not1	Nru1	Pac1	Pme1	PshA1	Pvu1	SanD1	Sbf1	Sca1	Sgf1	SgrA1
SnaB1	Spe1	Srf1	Swal	PISce	Xba1	Xcm1							

Excluded by site complexity:

Acc1	Ac1	Afl3	Alu1	Alw1	Apo1	Ava1	Ava2	Ban1	Ban2	Bbv1	BceA1	Bfa1	Bme15
BsaA1	BsaH1	BsaJ1	BsaW1	BseM2	BsiE1	BsiH1	Bsl1	BsmA1	BsmF1	Bsp12	BspCa	BspCb	Bsr1
BsrF1	BssK1	BstF5	BstN1	BstU1	BstY1	Btg1	Cac8	CviJ1	Dde1	Eae1	EcoO1	Fau1	Fnu4H
Fok1	Hae2	Hae3	Hga1	Hha1	Hinc2	Hinf1	HinP1	Hpa2	Hph1	Hpy99	Hpy1	Hpy3	HpyC3
HpyC4	HpyC5	Mae3	Mbo2	Mnl1	Mse1	Msl1	MspA1	Mwo1	Nci1	Nla3	Nla4	Nsp1	Ple1
PpuM1	Rsa1	Sau3A	Sau96	SfaN1	Sfc1	Sml1	Sty1	Taq1	Tat1	Tfi1	Tse1	Tsp45	Tsp50
TspR1													