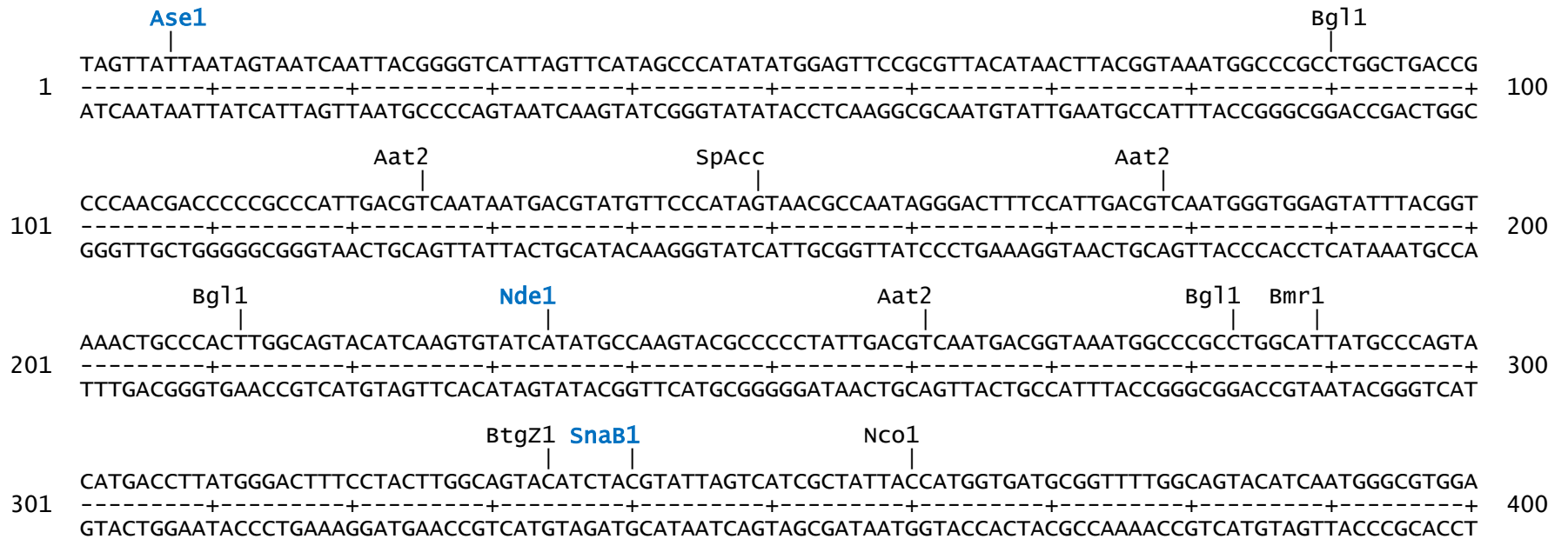
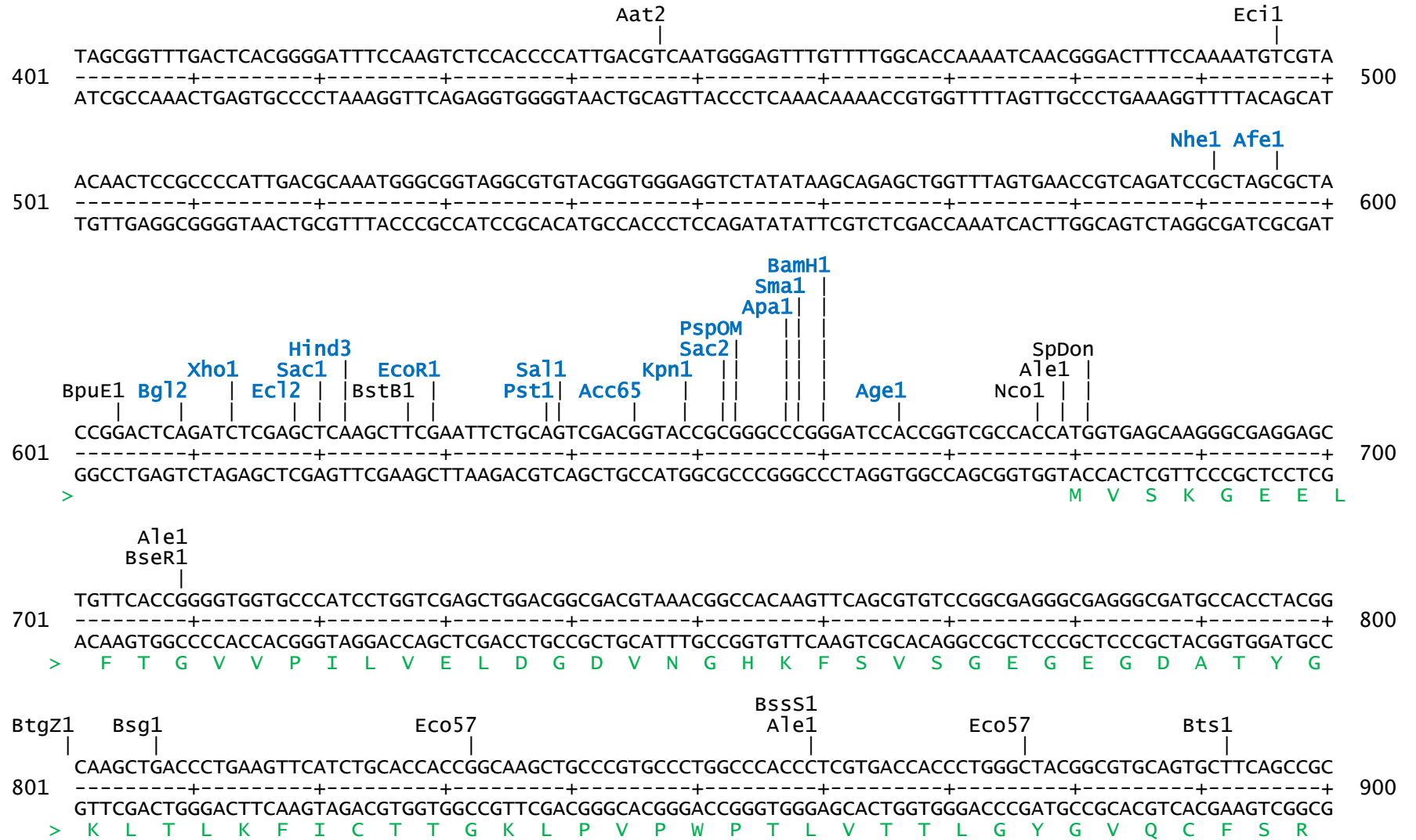


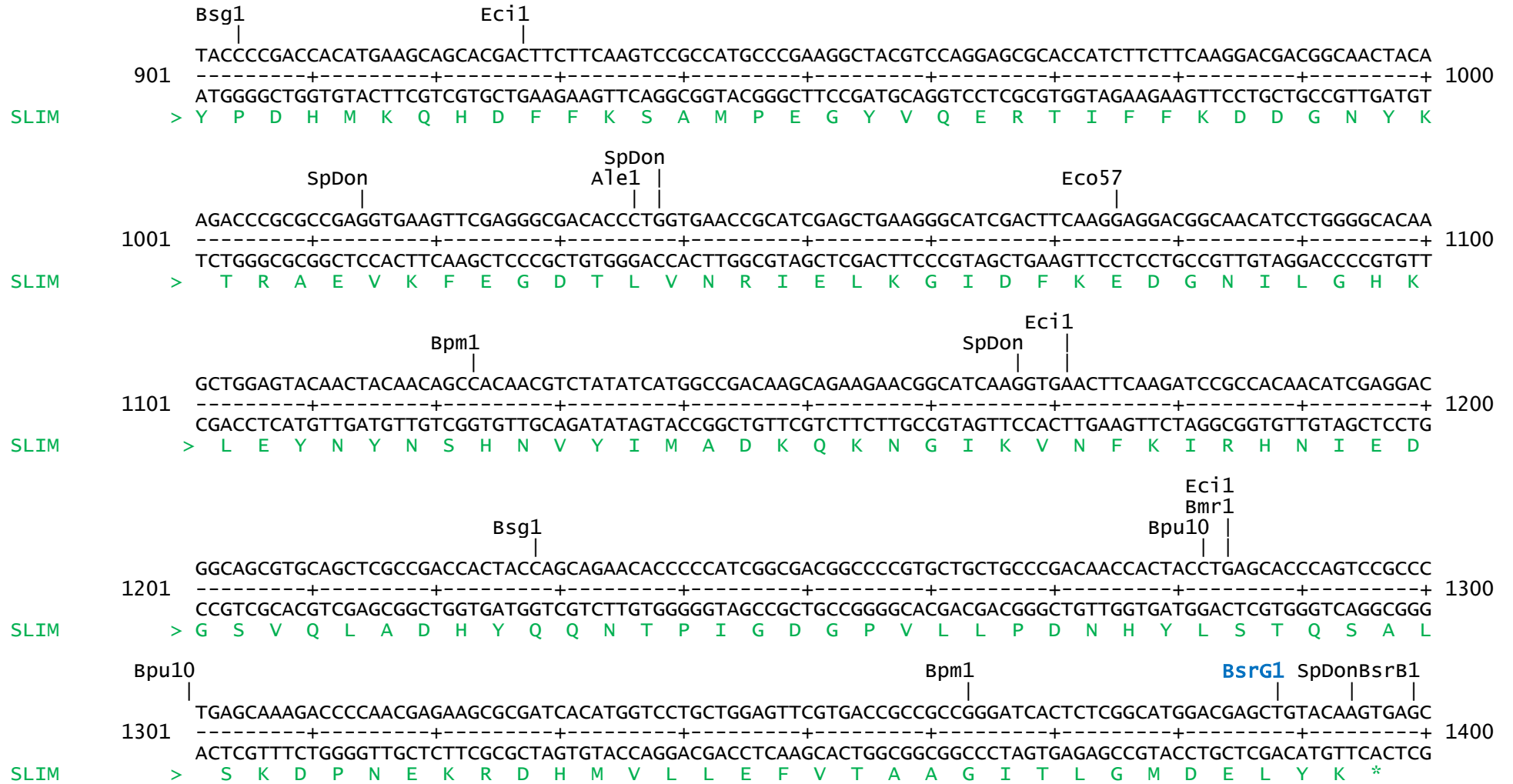
**pSLIM-N** vector restriction map

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.

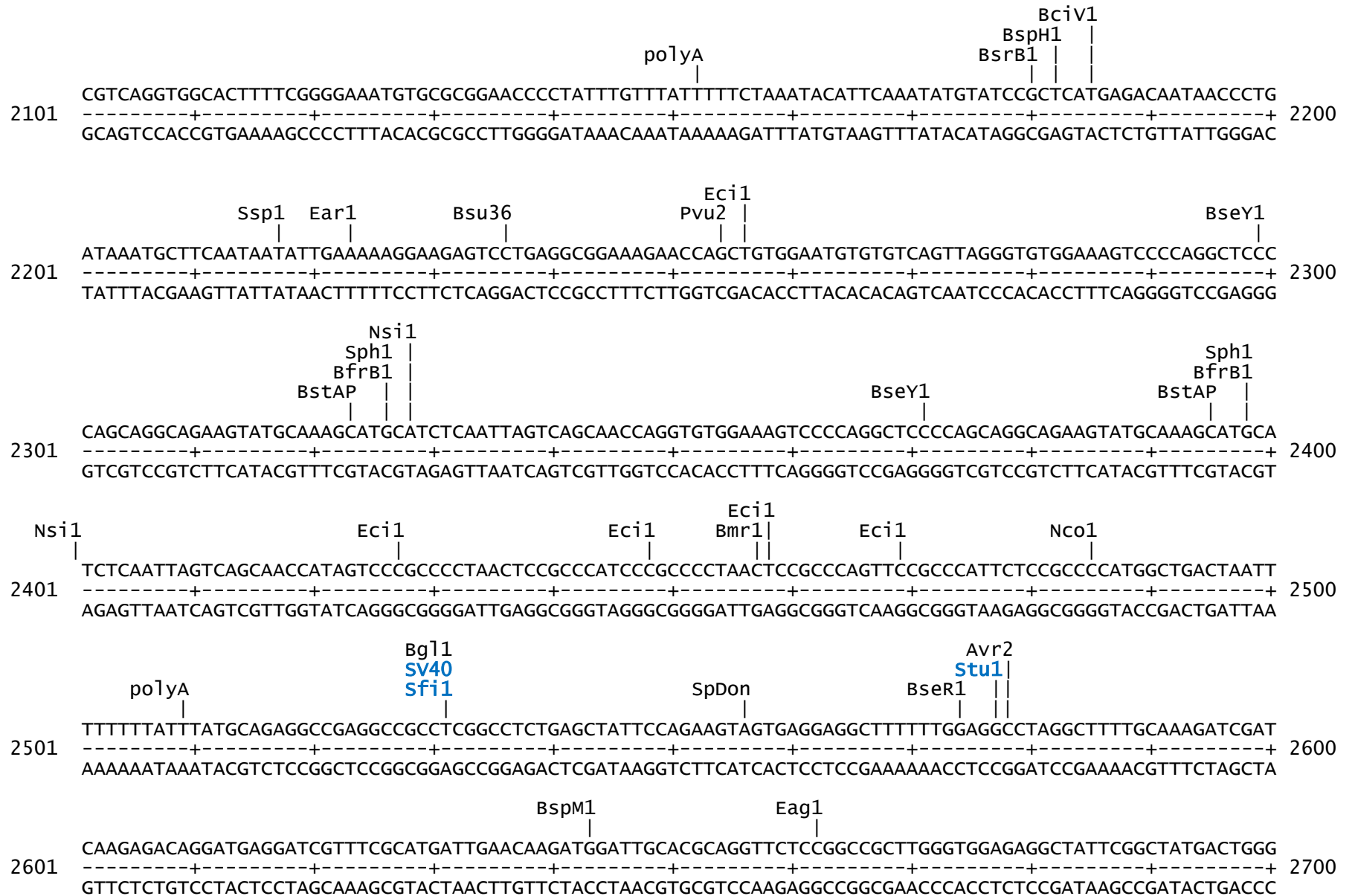
Unique sites are shown in bold blue. The location given specifies the 3' end of the cut DNA (the base to the left of the cut site).

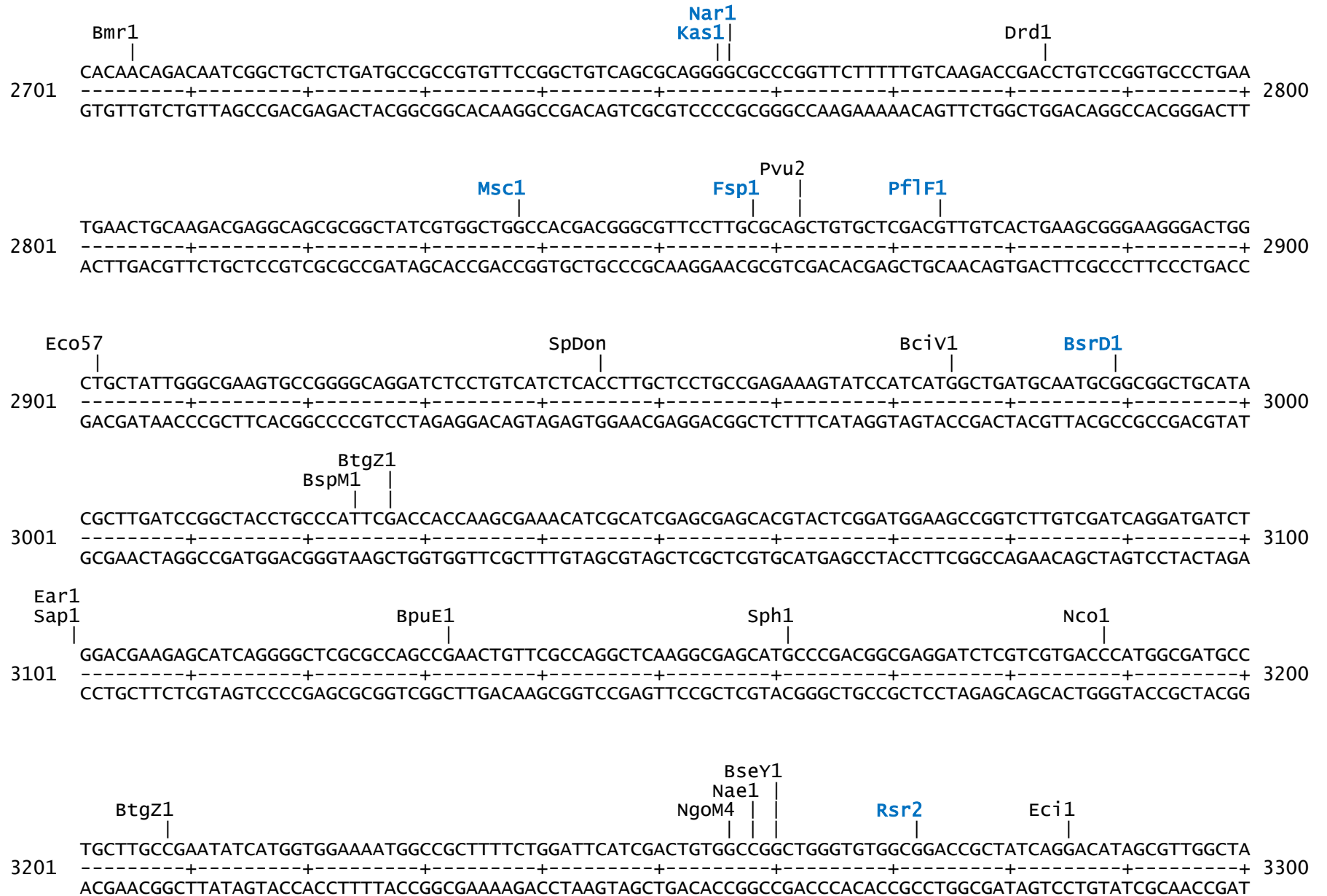


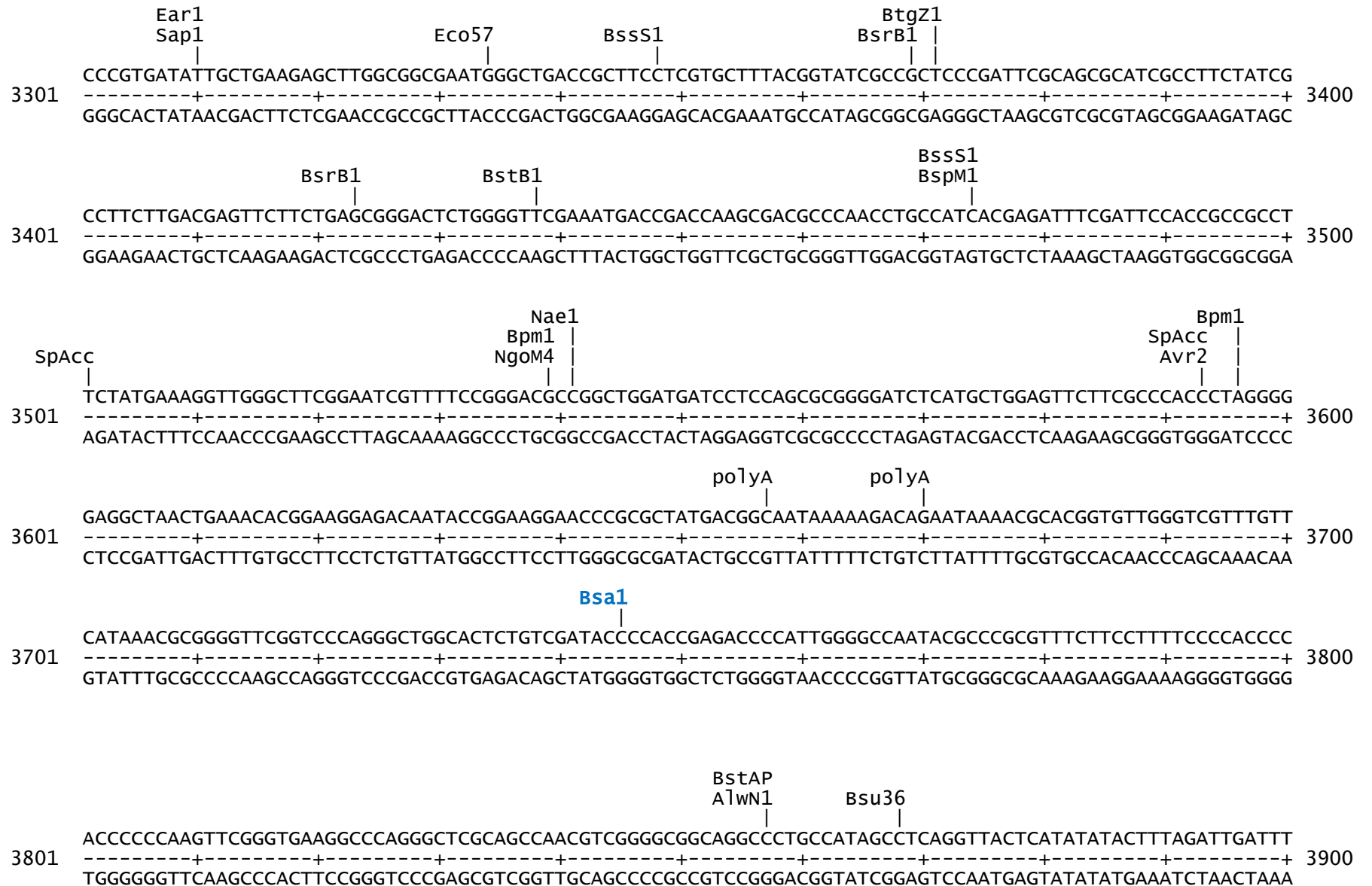


















Found

Aat2	<b>Acc65</b>	<b>Afe1</b>	<b>Afl2</b>	<b>Age1</b>	Ale1	AlwN1	<b>Apa1</b>	<b>ApaL1</b>	<b>Ase1</b>	Avr2	<b>BamH1</b>	BciV1	BfrB1
Bgl1	<b>Bgl2</b>	Bmr1	Bpm1	Bpu10	BpuE1	<b>Bsa1</b>	BseR1	BseY1	Bsg1	Bsm1	BspH1	<b>BspLU</b>	BspM1
BsrB1	<b>BsrD1</b>	<b>BsrG1</b>	BssS1	BstAP	BstB1	Bsu36	BtgZ1	Bts1	Dra1	<b>Dra3</b>	Drd1	Eag1	Ear1
Ec11	<b>Ec12</b>	Eco57	<b>EcoR1</b>	<b>Fsp1</b>	<b>Hind3</b>	<b>Hpa1</b>	<b>Kas1</b>	<b>Kpn1</b>	<b>Mfe1</b>	<b>Msc1</b>	Nae1	<b>Nar1</b>	Nco1
<b>Nde1</b>	Ngom4	<b>Nhe1</b>	<b>Not1</b>	Nsi1	<b>Pf1F1</b>	polyA	Psi1	<b>PspOM</b>	<b>Pst1</b>	Pvu2	<b>Rsr2</b>	<b>Sac1</b>	<b>Sac2</b>
<b>Sal1</b>	Sap1	<b>Sfi1</b>	<b>Sma1</b>	<b>SnaB1</b>	SpAcc	SpDon	Sph1	Ssp1	<b>Stu1</b>	<b>SV40</b>	<b>Xho1</b>		

Unique

<b>Acc65</b>	<b>Afe1</b>	<b>Afl2</b>	<b>Age1</b>	<b>Apa1</b>	<b>ApaL1</b>	<b>Ase1</b>	<b>BamH1</b>	<b>Bgl2</b>	<b>Bsa1</b>	<b>BspLU</b>	<b>BsrD1</b>	<b>BsrG1</b>	<b>Dra3</b>
<b>Ec12</b>	<b>EcoR1</b>	<b>Fsp1</b>	<b>Hind3</b>	<b>Hpa1</b>	<b>Kas1</b>	<b>Kpn1</b>	<b>Mfe1</b>	<b>Msc1</b>	<b>Nar1</b>	<b>Nde1</b>	<b>Nhe1</b>	<b>Not1</b>	<b>Pf1F1</b>
<b>PspOM</b>	<b>Pst1</b>	<b>Rsr2</b>	<b>Sac1</b>	<b>Sac2</b>	<b>Sal1</b>	<b>Sfi1</b>	<b>Sma1</b>	<b>SnaB1</b>	<b>Stu1</b>	<b>SV40</b>	<b>Xho1</b>		

Not found

Aar1	Ac11	Ahd1	Asc1	AsiS1	Bbs1	BbvC1	Bcl1	B1p1	BmgB1	BsaB1	BsiW1	BsmB1	BspE1
BssH2	BstE2	BstX1	BstZ1	BxatB	BxatL	BxatR	BxatP	_Chi	ClA1	EcoN1	EcoRV	FCatB	FCatL
FCatR	FCatP	ScFRT	Fse1	FspA1	I_Ceu	loxP	Mlu1	Nru1	Pac1	Pf1M1	Pme1	Pml1	PshA1
Pvu1	R4atB	R4atL	R4atP	R4atR	SanD1	Sbf1	Sca1	SexA1	Sgf1	SgrA1	SPRNA	Spe1	Srf1
Swa1	T3RNA	T7RNA	T7Ter	PISce	Xba1	Xcm1	Xmn1						

Excluded by site complexity

Acc1	Ac11	Afl3	Alu1	Alw1	Apo1	Ava1	Ava2	Ban1	Ban2	Bbv1	BceA1	Bfa1	Bme15
BsaA1	BsaH1	BsaJ1	Bsaw1	Bsem2	BsiE1	BsiH1	Bsl1	BsmA1	BsmF1	Bsp12	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	