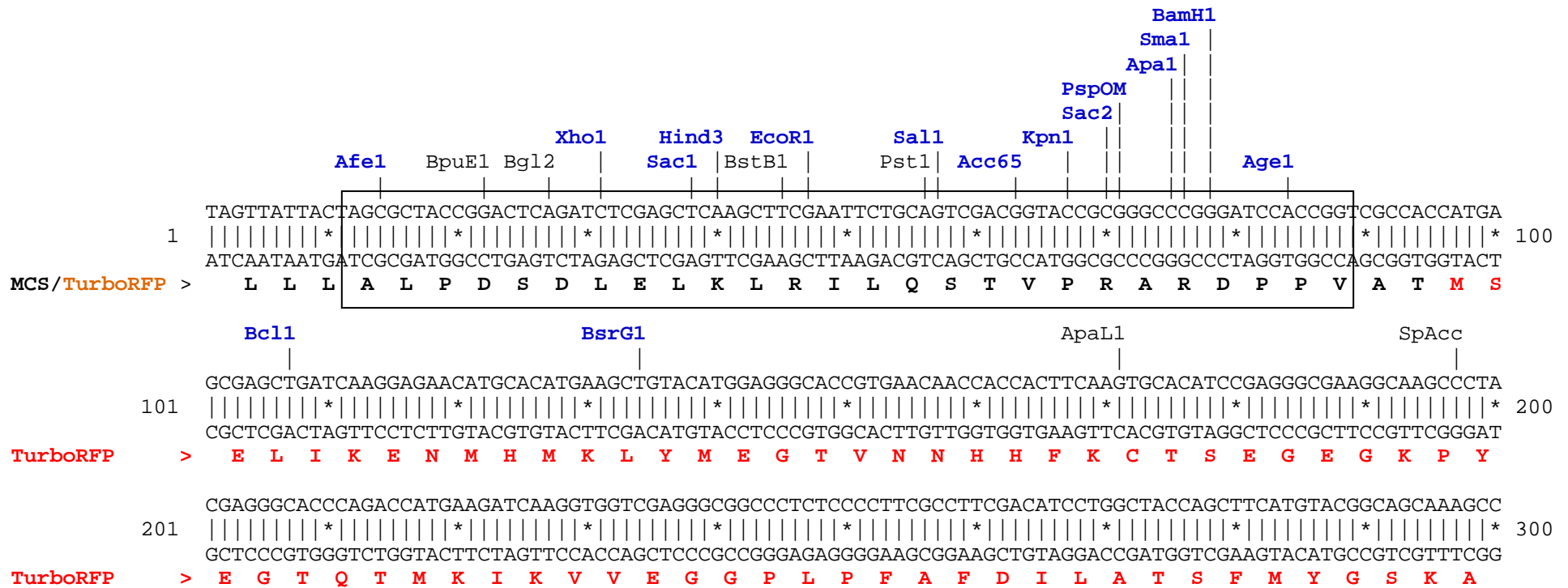


**pTurboRFP-PRL-dest1 vector restriction map**

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).

MCS sequence shown in frame, amino acids coded by MCS and MODC amino acids are shown in bold black.





















Found:

<b>Acc65</b>	<b>Afe1</b>	<b>Afl2</b>	<b>Age1</b>	AlwN1	<b>Apa1</b>	ApaL1	Avr2	<b>BamH1</b>	<b>Bbs1</b>	BciV1	<b>Bcl1</b>	BfrB1	BfuA1
Bgl1	Bgl2	Bmr1	Bpm1	BpuE1	Bsa1	BsaB1	<b>BsaXa</b>	<b>BsaXb</b>	<b>BseR1</b>	BseY1	<b>Bsg1</b>	Bsm1	<b>BspE1</b>
BspH1	<b>BspLU</b>	BspM1	BsrB1	<b>BsrD1</b>	<b>BsrG1</b>	BssS1	BstAP	BstB1	Bsu36	BtgZ1	Bts1	<b>Clal</b>	Dra1
Dra3	Drd1	Eag1	Ear1	Eci1	Eco57	<b>EcoR1</b>	<b>Fsp1</b>	<b>Hind3</b>	<b>Hpa1</b>	<b>Kas1</b>	<b>Kpn1</b>	<b>Mfe1</b>	Msc1
Nae1	<b>Nar1</b>	Nco1	NgoM4	Nsi1	<b>PflF1</b>	PflM1	polyA	<b>PshA1</b>	Psi1	<b>PspOM</b>	Pst1	Pvu2	<b>Rsr2</b>
<b>Sac1</b>	<b>Sac2</b>	<b>Sall</b>	Sap1	<b>Scal</b>	<b>SexA1</b>	Sfi1	<b>SgrA1</b>	<b>Smal</b>	SpAcc	SpDon	Sph1	Ssp1	<b>Stu1</b>
<b>Xba1</b>	<b>Xho1</b>												

Unique:

<b>Acc65</b>	<b>Afe1</b>	<b>Afl2</b>	<b>Age1</b>	<b>Apa1</b>	<b>BamH1</b>	<b>Bbs1</b>	<b>Bcl1</b>	<b>BsaXa</b>	<b>BsaXb</b>	<b>BseR1</b>	<b>Bsg1</b>	<b>BspE1</b>	<b>BspLU</b>
<b>BsrD1</b>	<b>BsrG1</b>	<b>Clal</b>	<b>EcoR1</b>	<b>Fsp1</b>	<b>Hind3</b>	<b>Hpa1</b>	<b>Kas1</b>	<b>Kpn1</b>	<b>Mfe1</b>	<b>Nar1</b>	<b>PflF1</b>	<b>PshA1</b>	<b>PspOM</b>
<b>Rsr2</b>	<b>Sac1</b>	<b>Sac2</b>	<b>Sall</b>	<b>Scal</b>	<b>SexA1</b>	<b>SgrA1</b>	<b>Smal</b>	<b>Stu1</b>	<b>Xba1</b>	<b>Xho1</b>			

Not found:

Aar1	Aat2	Acl1	Ahd1	Ale1	Asc1	Ase1	AsiS1	Baela	Baelb	BbvC1	Bcg1a	Bcg1b	Blp1
BmgB1	Bpu10	BsiW1	BsmB1	BssH2	BstE2	BstX1	BstZ1	_Chi	EcoK	EcoN1	EcoRV	ScFRT	Fse1
FspA1	I_Ceu	loxP	Mlu1	Nde1	Nhe1	Not1	Nru1	Pac1	Pme1	Pml1	Pvu1	SanD1	Sbf1
Sgf1	SnaB1	Spe1	Srf1	Swal	T3RNA	T7RNA	T7Ter	PISce	Xcm1	Xmn1			

Excluded by site complexity:

Acc1	Acil	Afl3	Alu1	Alw1	Apo1	Ava1	Ava2	Ban1	Ban2	Bbv1	BceA1	Bfal	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	Faul	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													