



























# Part domestication report

The table below shows how the different parts have been domesticated. The small icons are unique identifiers for the sequences. If two sequences have the same icon, they are very probably the same !

## Summary Table

Record	Order ID	Domesticator	Domesticated Record	Added bp	Edited bp
 p1	p1	p1	 p1	22	0
 p12	p12	p12	 p12	22	0
 p13	p13	p13	 p13	22	0
 p14	p14	p14	 p14	22	0
 p15	p15	p15	 p15	22	0
 p16	p16	p16	 p16	22	0
 p17	p17	p17	 p17	22	0
 p18	p18	p18	 p18	22	0
 p2	p2	p2	 p2	22	0
 p22	p22	p22	 p22	22	0
 pCL	pCL	pCL	 pCL	22	0
 pSV	pSV	pSV	 pSV	22	0
 pWZ	pWZ	pWZ	 pWZ	22	0

## Domesticators

p1

**Name:** p1

**Description:** For slot p1 of the EMMA standard

**Left addition:** CGTCTCATAGG

**Right addition:** ATGGTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** TAGG

**Right overhang:** ATGG

p12

**Name:** p12

**Description:** For slot p12 of the EMMA standard

**Left addition:** CGTCTCACGTC

**Right addition:** TCACTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** CGTC

**Right overhang:** TCAC

p13

**Name:** p13

**Description:** For slot p13 of the EMMA standard

**Left addition:** CGTCTCATCAC

**Right addition:** CTACTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** TCAC

**Right overhang:** CTAC

p14

**Name:** p14

**Description:** For slot p14 of the EMMA standard

**Left addition:** CGTCTCACTAC

**Right addition:** GCAATGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** CTAC

**Right overhang:** GCAA

p15

**Name:** p15

**Description:** For slot p15 of the EMMA standard

**Left addition:** CGTCTCAGCAA

**Right addition:** CCCTTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** GCAA

**Right overhang:** CCCT

p16

**Name:** p16

**Description:** For slot p16 of the EMMA standard

**Left addition:** CGTCTCACCT

**Right addition:** GCTCTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** CCCT

**Right overhang:** GCTC

p17

**Name:** p17

**Description:** For slot p17 of the EMMA standard

**Left addition:** CGTCTCAGCTC

**Right addition:** CGGTTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** GCTC

**Right overhang:** CGGT

p18

**Name:** p18

**Description:** For slot p18 of the EMMA standard

**Left addition:** CGTCTCACGGT

**Right addition:** GTGCTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))  
**Left overhang:** CGGT  
**Right overhang:** GTGC

p2

**Name:** p2  
**Description:** For slot p2 of the EMMA standard  
**Left addition:** CGTCTCAATGG  
**Right addition:** GACTTGAGACG  
**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1  
Seq('CGTCTC', DNAAlphabet()))  
**Left overhang:** ATGG  
**Right overhang:** GACT

p22

**Name:** p22  
**Description:** For slot p22 of the EMMA standard  
**Left addition:** CGTCTCAGTTG  
**Right addition:** CGAATGAGACG  
**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1  
Seq('CGTCTC', DNAAlphabet()))  
**Left overhang:** GTTG  
**Right overhang:** CGAA

pCL

**Name:** pCL  
**Description:** pCL  
**Left addition:** CGTCTCAGACT  
**Right addition:** CGTCTGAGACG  
**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1  
Seq('CGTCTC', DNAAlphabet()))  
**Left overhang:** GACT  
**Right overhang:** CGTC

pSV

**Name:** pSV

**Description:** pSV

**Left addition:** CGTCTCAGTGC

**Right addition:** GTTGTGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** GTGC

**Right overhang:** GTTG

pWZ

**Name:** pWZ

**Description:** pWZ

**Left addition:** CGTCTCACGAA

**Right addition:** ACGATGAGACG

**Enzyme:** BsmBI (ID: Name: Description: Number of features: 1

Seq('CGTCTC', DNAAlphabet()))

**Left overhang:** CGAA

**Right overhang:** ACGA