

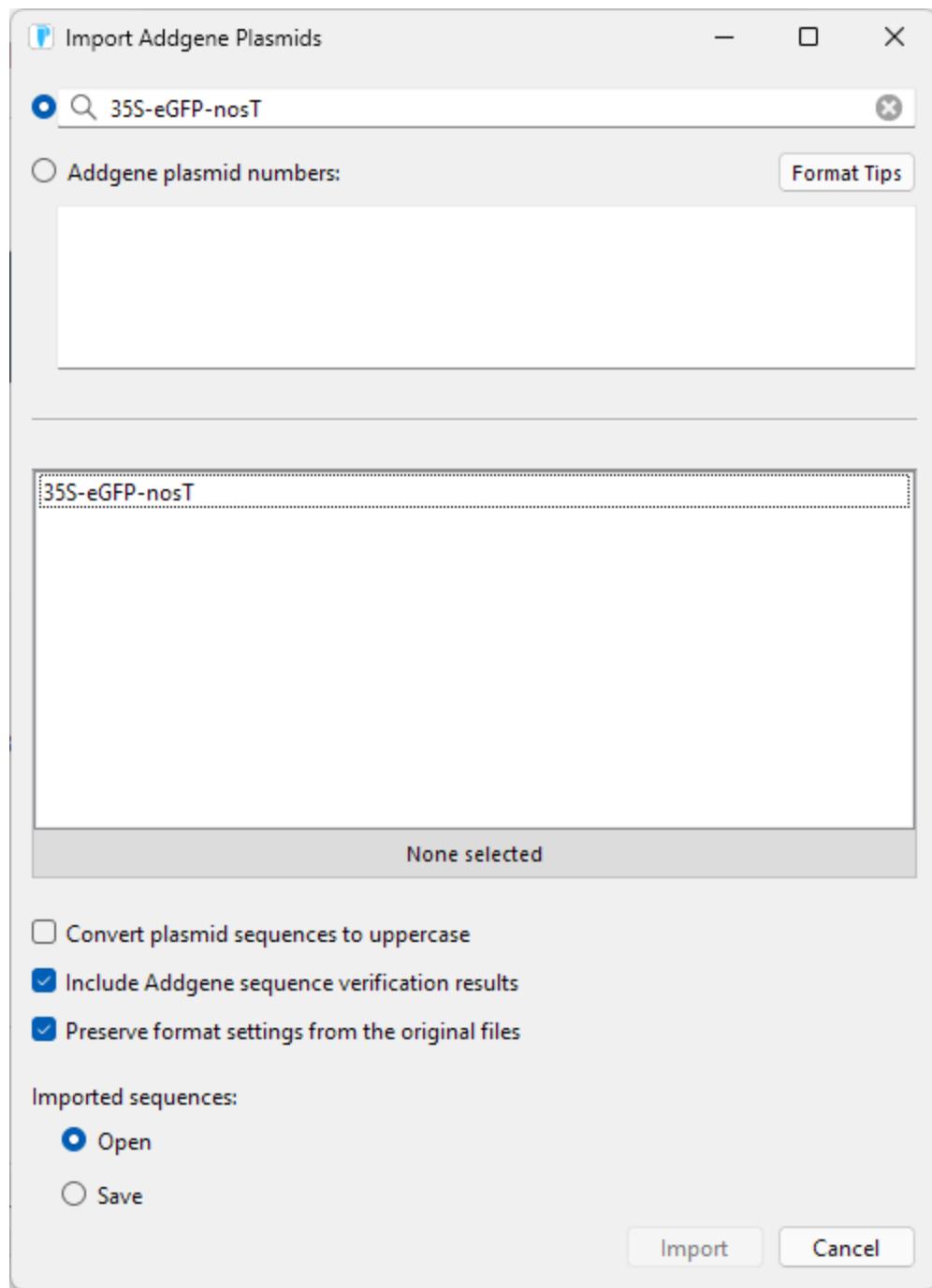
## 022-practicals

### **Insertion of gene for cry protein in a plasmid with the sequence of GFP protein, called 35S-eGFP-nosT, and used for transient expression**

To complete this task follow the steps described in 2.1 section, except two important things:

- sequences for the restriction enzymes should be replaced. Which restriction enzymes can be used to open the plasmid between CaMV 35S promoter and eGFP gene? You have to select restriction enzymes which don't have recognition sites in the cry gene.
  - when you will design reverse primer you should skip the last three nucleotides (represent stop codon) in order to keep ORF open
- 

1. Importam plasmid



Da vstavim cry gen moram izbrat restrikcijska mesta ki niso v njemu, torej.. če grem na Enzymes tab od cry gena in izberem print lahko dobim seznam:

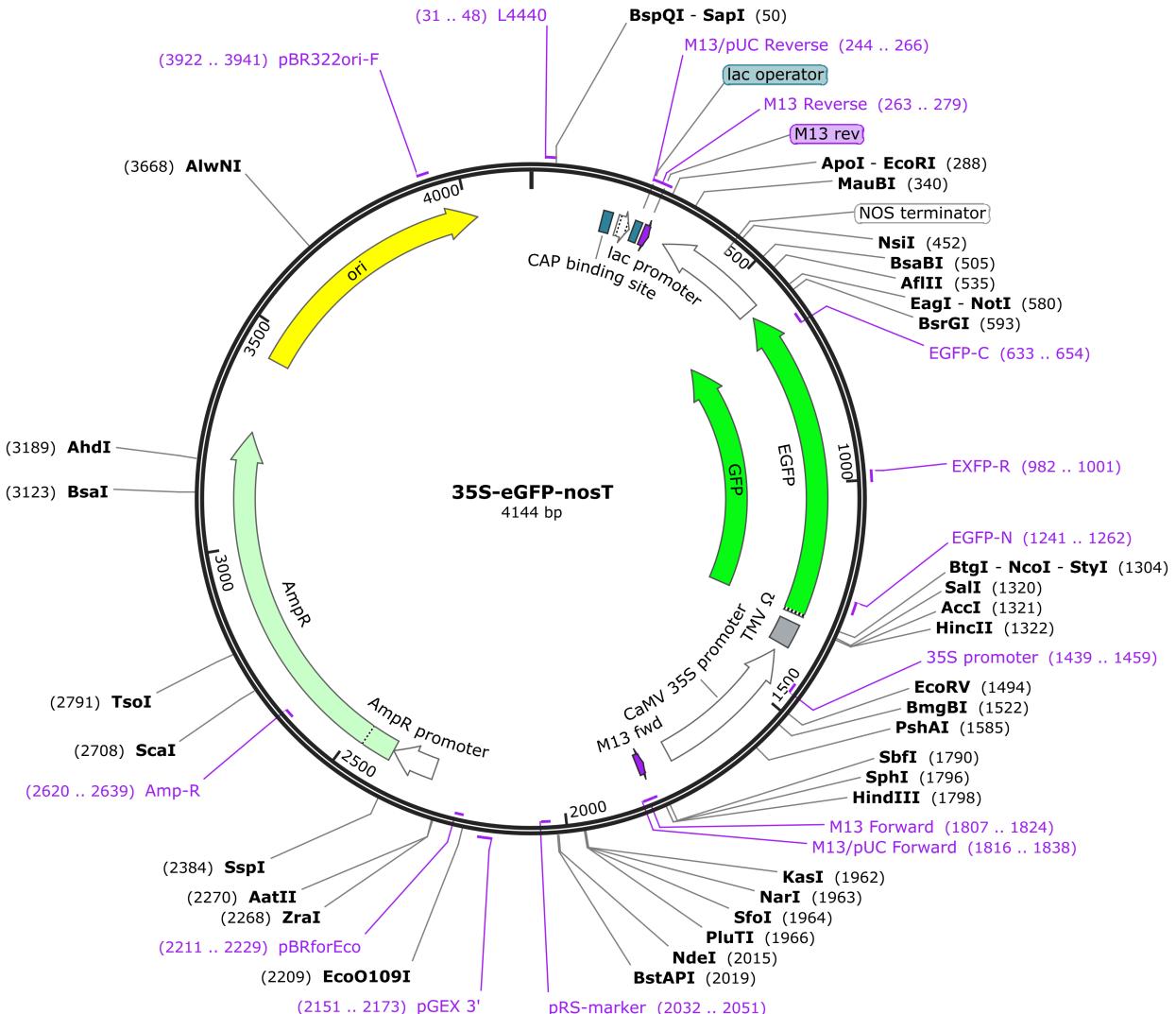
Sequence: Amplified\_from021.dna (Linear / 1794 bp)

Enzymes: < Unsaved Enzyme Set > (36 of 686 total)

Enzyme	Sites	Location
<b>AccI</b>	1	1352
<b>AflIII</b>	1	889
<b>ApaI</b>	1	1605
<b>ApaLI</b>	1	1476
<b>BbsI</b>	1	751
<b>BIpI</b>	1	1702
<b>BpuEI</b>	1	1054
<b>BsaI</b>	1	1228
<b>BseRI</b>	1	520
<b>BsmBI</b>	1	327
<b>BstXI</b>	1	1437
<b>BtgZI</b>	1	516
<b>DraIII</b>	1	588
<b>Esp3I</b>	1	327
<b>HincII</b>	1	1353
<b>MluI</b>	1	889
<b>MreI</b>	1	1744
<b>MscI</b>	1*	1678*
<b>NaeI</b>	1	1746
<b>NgoMIV</b>	1	1744
<b>NspI</b>	1	402
<b>PaeR7I</b>	1	324
<b>PaqCI</b>	1	183
<b>PpuMI</b>	1	168
<b>PspOMI</b>	1	1601
<b>PstI</b>	1	1101
<b>SalI</b>	1	1351
<b>Sau3AI</b>	8	53 218 695 1367 1382 1574 1721 1769
<b>SphI</b>	1	402
<b>TsoI</b>	1	1057
<b>XbaI</b>	1	609
<b>XhoI</b>	1	324
<b>XmnI</b>	1	181

**3** of 36 enzymes in the chosen set do not cut

Moram pa opret med CaMV 35S promoter in eGFP gene ...



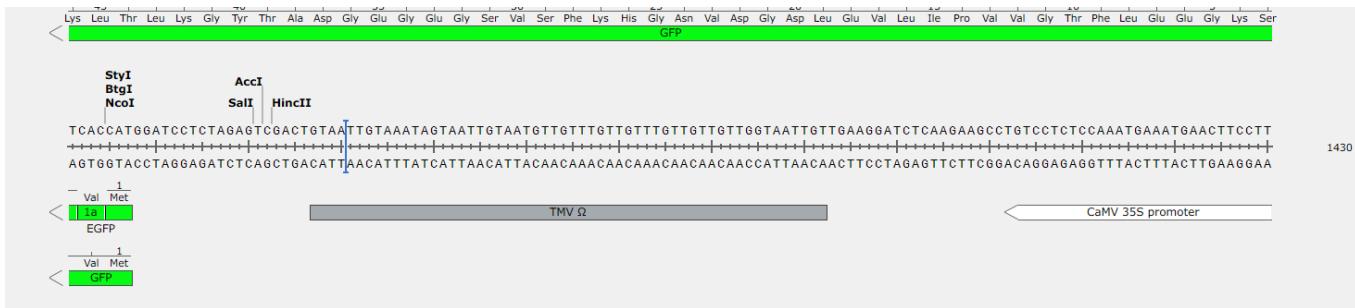
Okej, sem kopiral seznam encimov od obeh v svoje datoteke, jih odpr z R-jem in preveril kateri so unikatni za gen (glej `candidate_REs.html`):

Encimi, ki jih lahko izberem za vstavljanje (prisotni sami v `genu`):

`uncommon_f`

```
## AfIIII ApaI ApaII BbsI BlpI BpuEI BseRI BsmBI BstXI BtgZI DraIII
## 2 3 4 5 6 7 9 10 11 12 13
## EspIII MluI MreI MscI NaeI NgoMIV NspI PaeR7I PaqCI PpuMI PspOMI
## 14 16 17 18 19 20 21 22 23 24 25
## PstI Sau3AI XbaI XhoI XmnI
## 26 28 31 32 33
```

In zdaj lahko bi sporogramiral da najde še kateri so med promotorjem in GFPjem, ampak bom naoko pogledal ker jih ni precej. Mislim da moram te gledati:



In dobim:

Kateri od njih ni prisoten v fragmentu?

```
plasmid_candidates[which(!(plasmid_candidates %in% fvec2))]
```

```
## [1] "BtgI" "NcoI"
```

Te lahko zdaj uporabim za PCR in vstavljanje... to pomeni, da moram prilagoditi svoj gen - mu spremeniti primerje, narediti PCR in vstaviti. Hmm... a naj dam vsakega na en konec ali uporabim samo enega...? Glede na to, da smo prej tudi izbrali različna za isto regijo, se mi zdi prav da tudi zdaj naredim tako.

`BtgI` na začetek in `NcoI` na konec. Pazi, da dodaš encima za primerji (primerji so samo za PCR). Rekel je, da naj dodamo še en random site pred tem, da ni takoj za promotorjem.. ampak verjetno ne sme biti site prisoten v fragmentu..? Idk, prej sem dal za `SauAI3` pa je bilo okej... idk. Bom zdaj tudi.

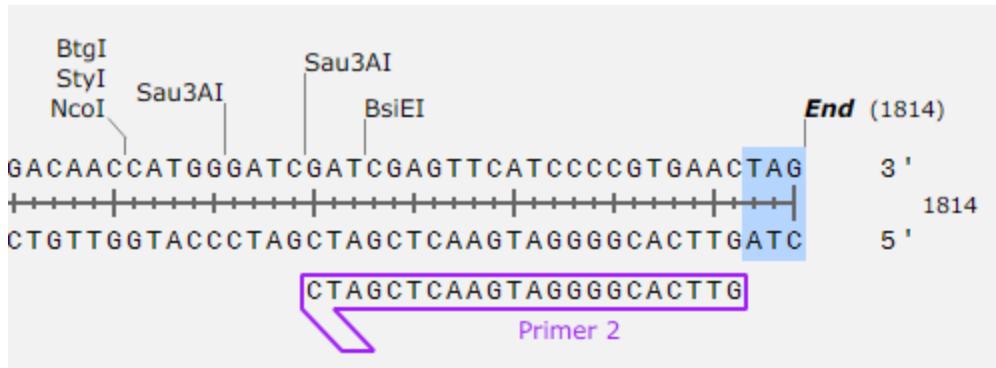
No in tudi `BtgI` je med encimi ampak ga ni označilo v export... I dont fucking know.

`NcoI` se osveti odbeljeno, `BtgI` pa ne.... bom `NcoI` uporabu...

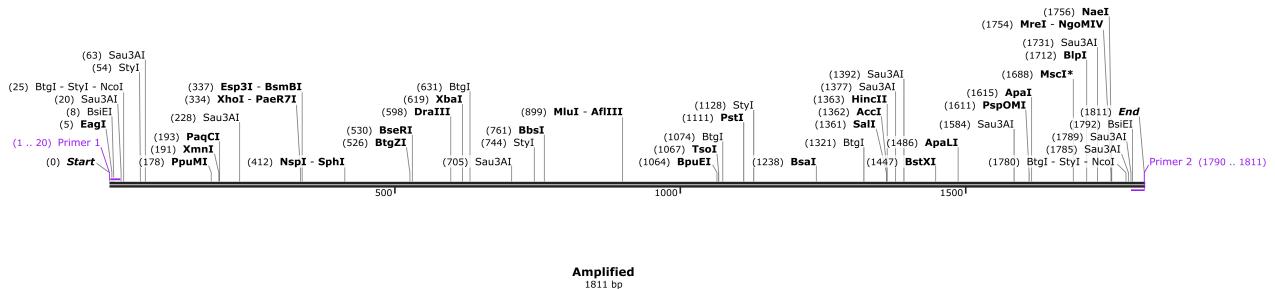
No itak mi kaže da je isti site...



Pazil, da sem preskočil stop kodon.

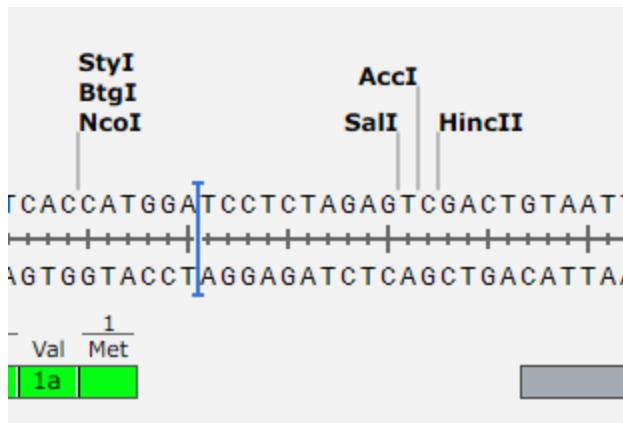


In dobim tole



In zdaj to vstavim v plazmid...

Hold up...



Pač kaže mi da ne morem izbrat dva na istem baznem paru... ampak vsi te ki so na desni, so prisotni v fragmentu in jih ne smem izbrat...

Siva regija pa je enhancer in tega ne smem izrezat, kot smo imeli pri [2.1 MCS](#)... I don't know what to do.

Detailed description: This is a screenshot of a bioinformatics interface. It shows a sequence region from 1327 to 1382 bp, which is 56 bp long. A note below states: "translational enhancer from the tobacco mosaic virus 5'-leader sequence (Gallie et al., 1988)".

Pusti mi če izberem samo eno regijo pri plazmidu in encimu... sem izbral NcoI . Anyways,

**Restriction and Insertion Cloning: Insert Fragment**

**Vector:** 35S-eGFP-nosT.dna

**Cut** with **NcoI**  
and fill in 5' overhangs with dNTPs

**Cut** with  
and then blunt

**Site of insertion:** NcoI (1304)

**Orientation of Vector:** ↗ ↘

**Create product:** Cloned.dna  and close this window

**Source of Fragment:** Amplified.dna

**Cut** with **NcoI**  
and fill in 5' overhangs with dNTPs

**Cut** with  
and then blunt

**Fragment to insert:** Start (0) — NcoI (25)  
NcoI (25) — NcoI (1780)  
NcoI (1780) — End (1811) 1755 bp

**Oriantation of Fragment:** ↗ ↘

**Create product:** Cloned.dna  and close this window

**Clone** | **Cancel**

Restriction and Insertion Cloning: Insert Fragment

File Edit View Enzymes Features Primers Actions Tools Window Help

Vector Fragment Product

Map Sequence Enzymes

Sequence

Enzymes

Map

10 20 30 40 50 60 70 80 90

2970

3060

3150

Bacterial Transformation Strain  
Unspecified

Lineared Vector  
Source: 35S-eGFP-nosT.dna  
 Make file:  
Linearized Vector.dna

Fragment  
Source: Amplified.dna  
 Make file:  
Fragment.dna

Alternative Product with Flipped Fragment  
 Make file:  
Alternative Product.dna

Unique 6+ Cutters | Nonredundant

Ready to clone  
Product: 399 bp  
1 warning

Create product:  
Cloned.dna

and close this window

Clone Cancel