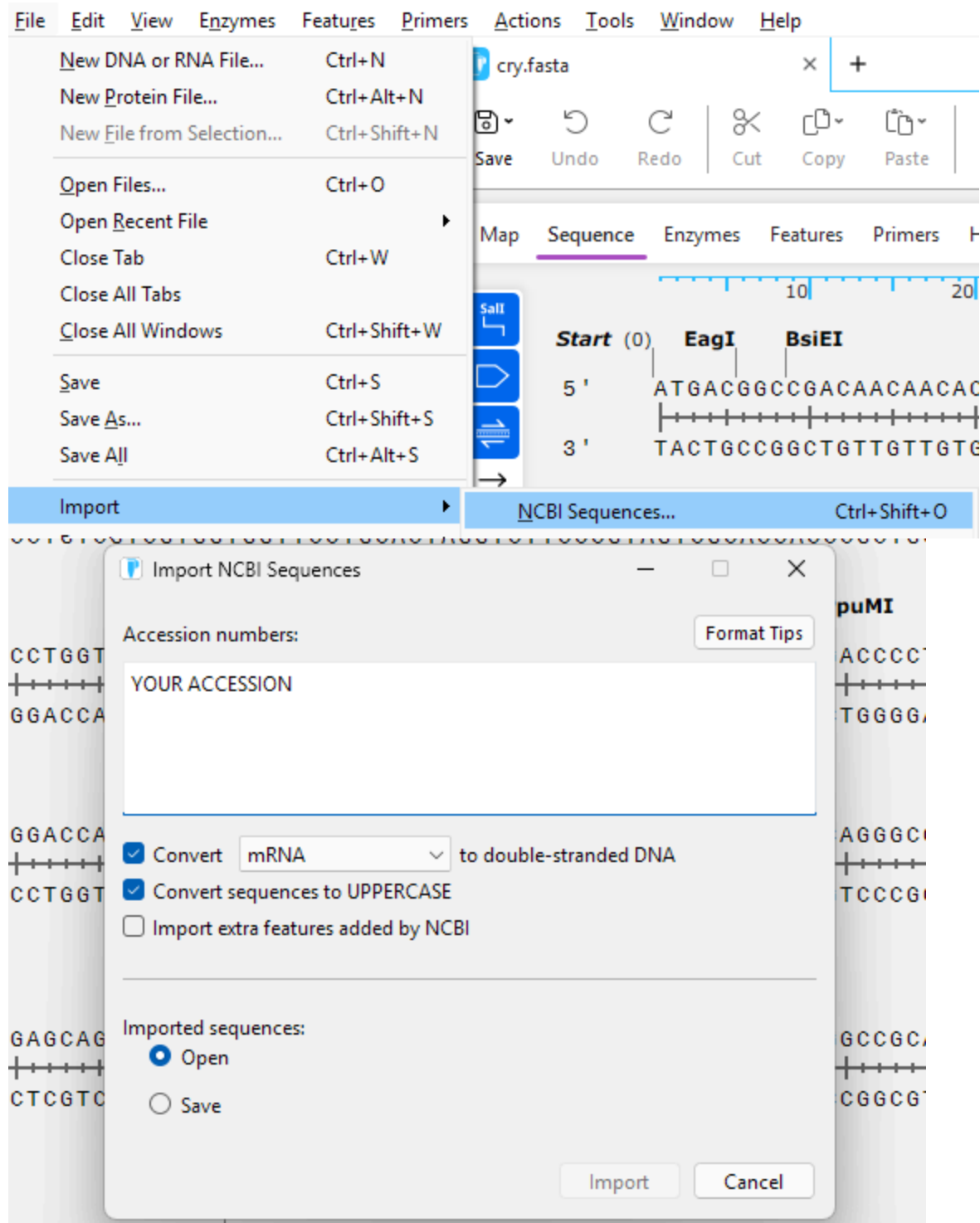


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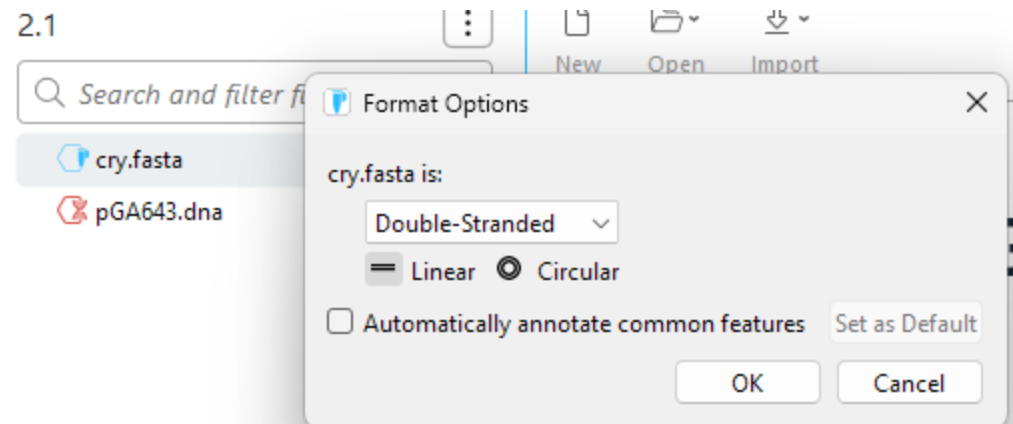
Insertion of gene for cry protein into a pGA643 plasmid.

1. *Import fasta sequence of cry protein in SnapGene.*

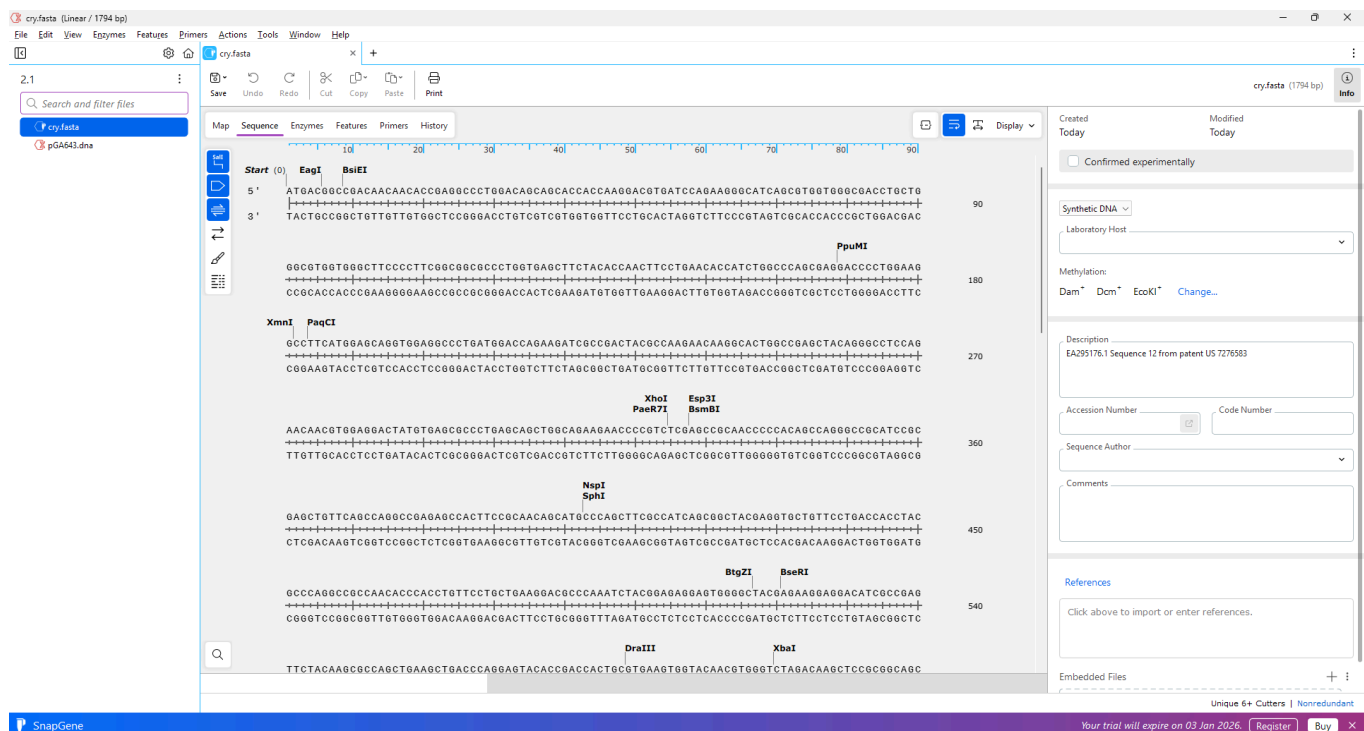
Either like this:



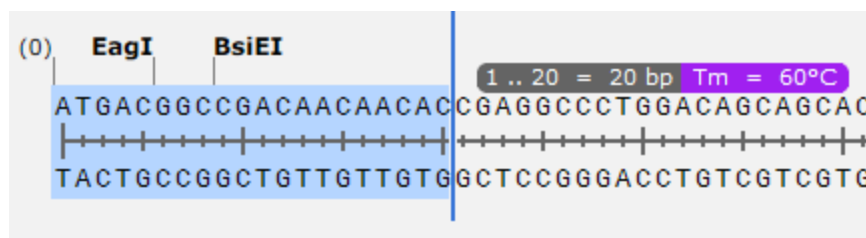
Or like this:



Imported sequence:

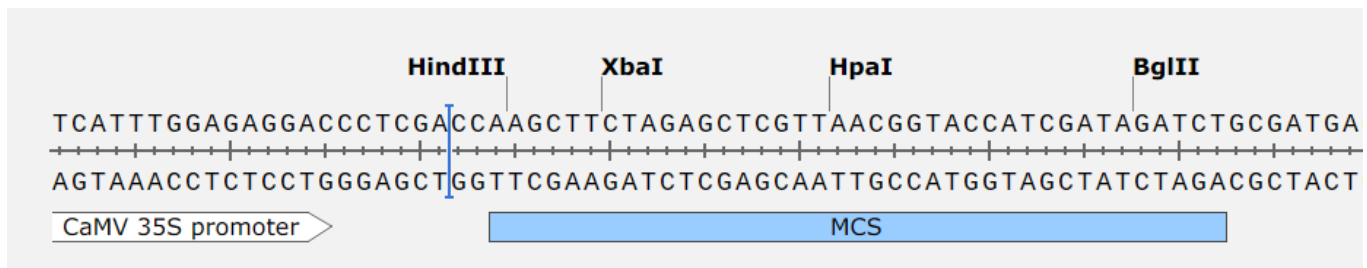


2. Select the first few nucleotides (until melting temperature 60 °C is reached) and click “Add primer”.



***Click the tab “Insertions” and select HindIII sequence to be attached to the 5’ end to the forward primer and since restriction will be more efficient if recognition site is not at the beginning of the DNA, add additional restriction site of any other restriction enzyme.**

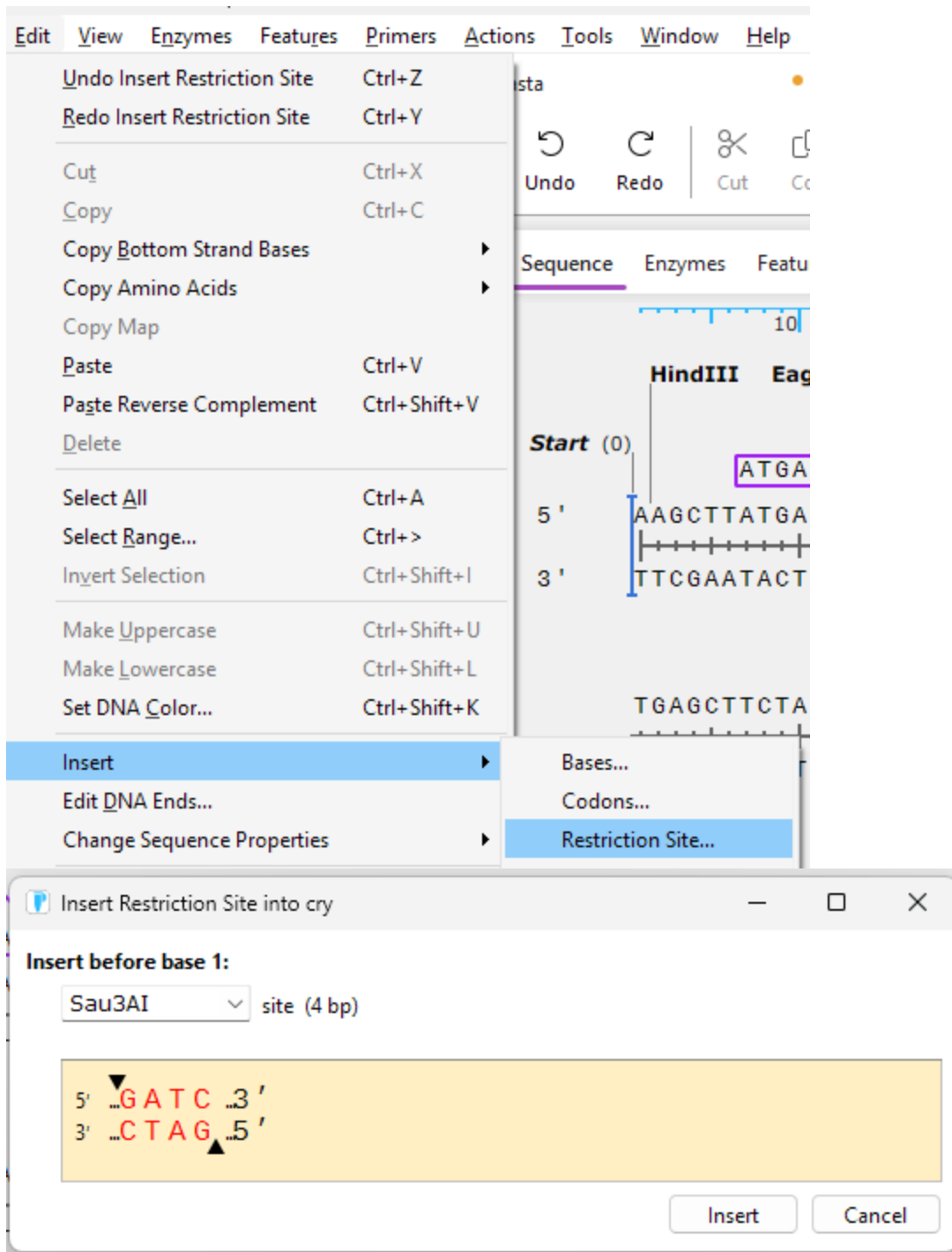
Primer zakaj izbiramo te encime:



Izločila se bo MCS regija (multiple cloning site)...

Klikneš na začetek zaporedja, najdeš HindIII in potem še za Sau3AI (ker je kratek in naredi sticky ends idk, to sem dal na začetek).

PAZI, DA DAŠ ZA PRIMERJEM, KER MI HOČEMO FRAGMENT NAMNOŽITI S TEMI REGIJAMI (na sliki je napačno).



Repeat the process for the reverse primer, except that the restriction site at the other site of DNA should be BglII (Bg"l"2).

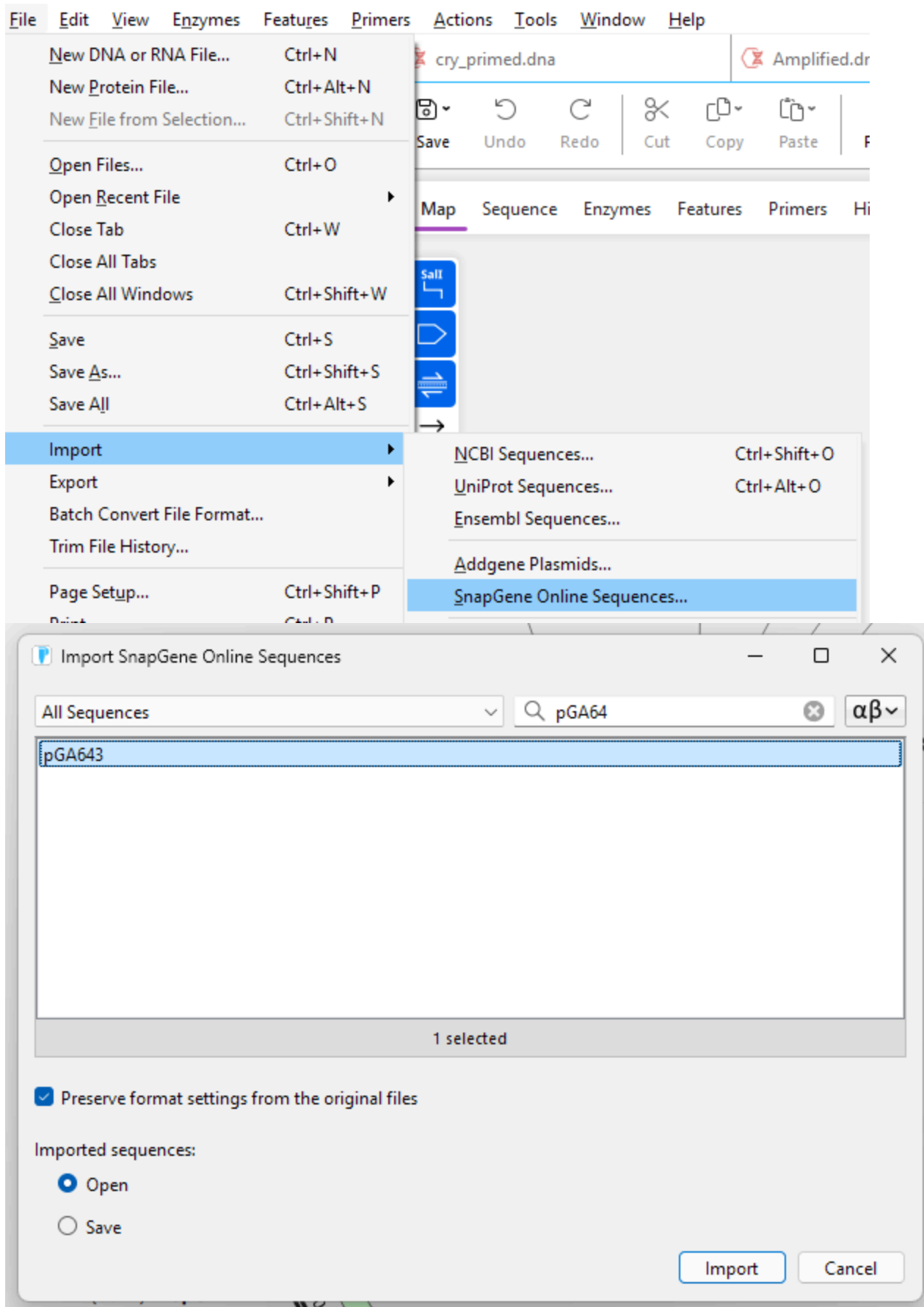
Enako, tudi tu sem dal `Sau3AI` na konec.

3. Simulate PCR to get DNA with elongated primers.

Actions → PCR in izbereš primerja, datoteka `Amplified.dna`.

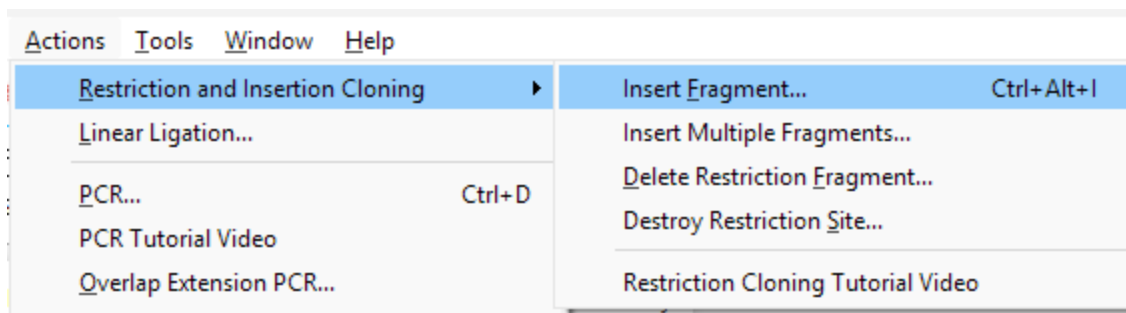
4. Import pGA643 plasmid.

Odpreš prenešeno datoteko ali pa...



5. Click “Actions”, then “Restriction and Ligation cloning” and “Insert fragment”. Select vector in the first tab, amplified DNA in the second

tab and press “Clone”



6. Select window with pGA643 and inserted DNA. Select inserted sequence from ATG to the TAG codon

Click “Features”, “Add feature”, rename the inserted sequence and select CDS for the “type”

Add Feature for Cloned

Feature: coding sequence

αβ

Type: CDS

☒ Translate this feature in Sequence view

Feature Translation Options...

1794 bp / 1 segment

2830

4623

Split Feature...

Merge Segments

Delete Segment

Segment	Location	Size (bp)	Color
1	coding sequence	2830 .. 4623	1794

/product

=

+

-

☐ Prioritize display of this feature in maps

OK

Cancel

