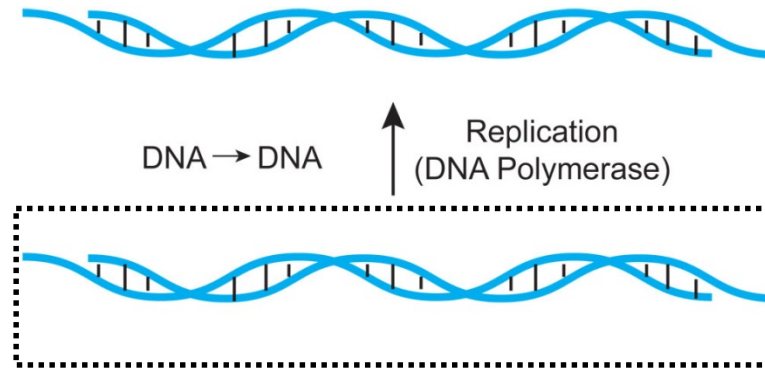


## Cloning-1

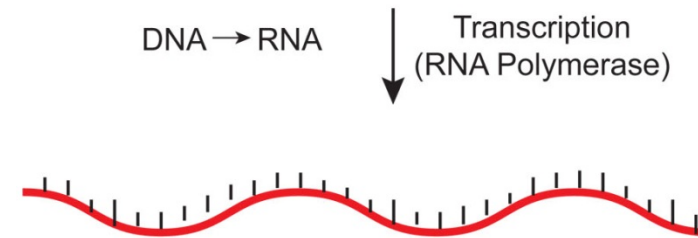
### Polymerase Chain Reaction and Endonucleases (Restriction Enzymes)

# Basics – the Central Dogma



Genetic Information is the blueprint for how to respond to any situation

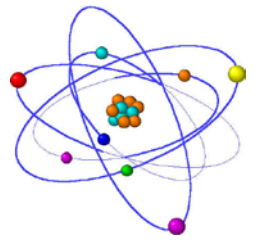
RNA is a short-lived intermediate that translates the genetic code



Proteins are the workhorse of the cell - carry out cellular functions.

# Molecular Cloning

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- What do we mean by “cloning”?

*Manipulating the genetic information present in an organism to give it new characteristics or function*

- As a society, why would we want to do this?

*Make farming easier/more efficient*

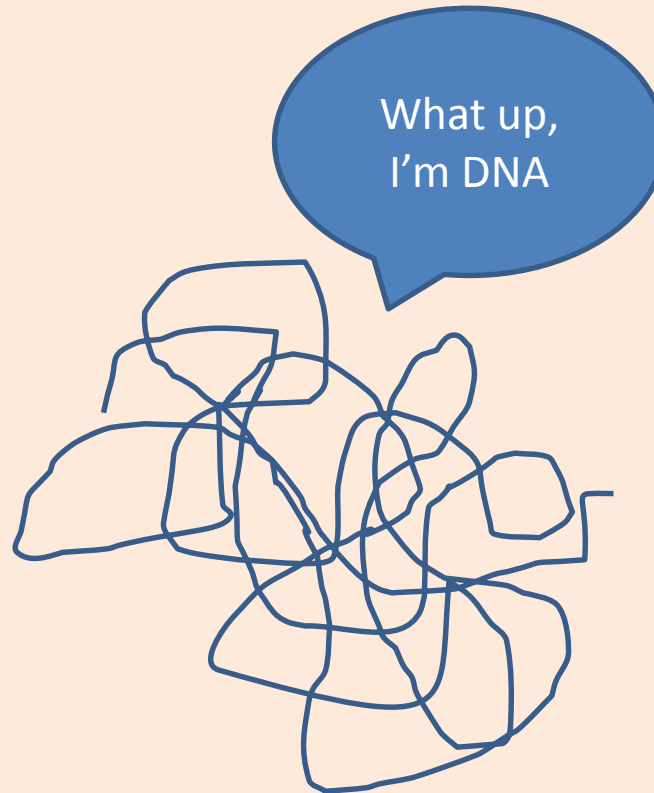
*Enhance nutrition value of food*

- As Biochemists, why would we want to do this?

*Take advantage of easily manipulated cells to increase protein yield.*

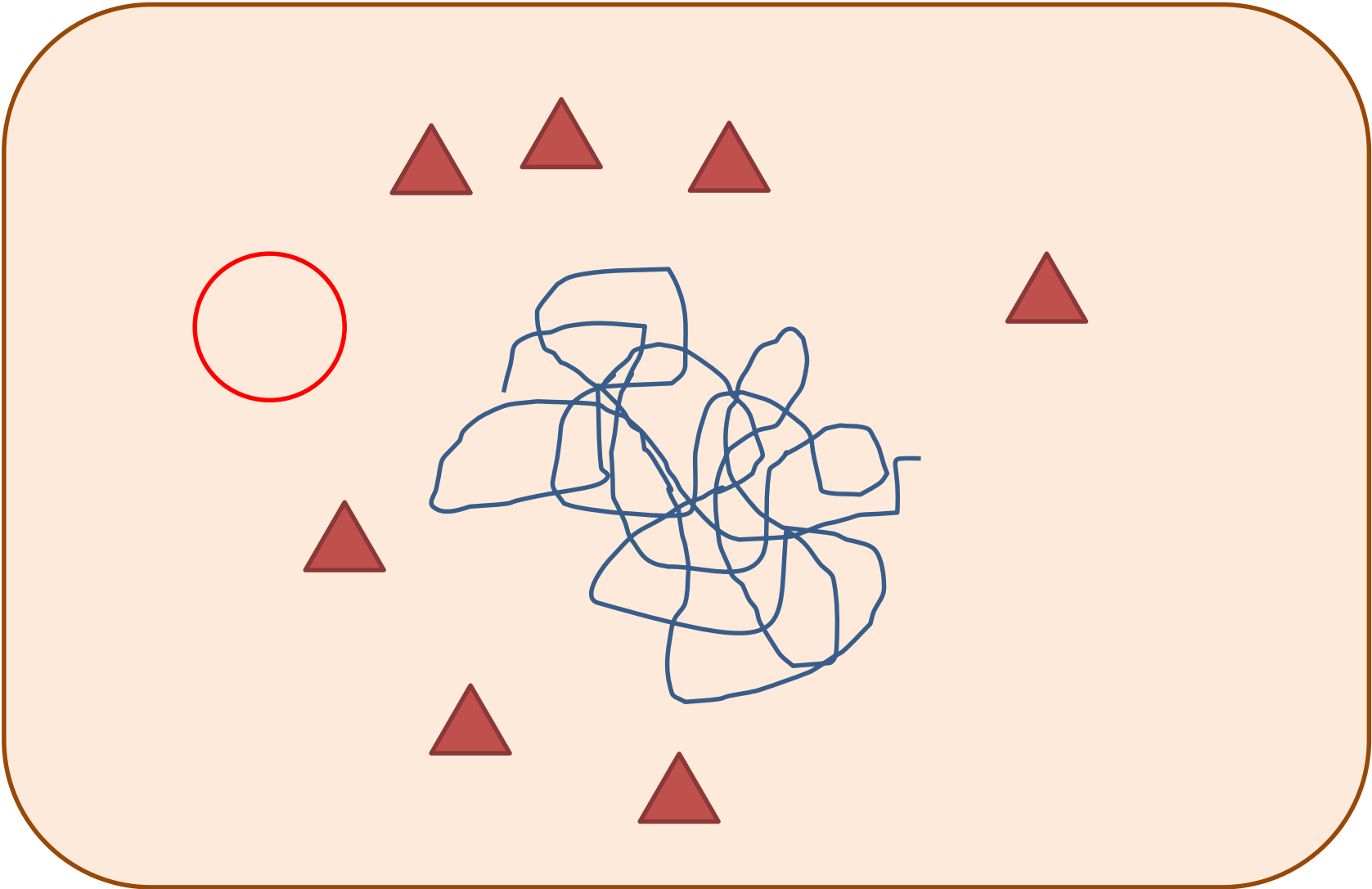
# Overall Goal

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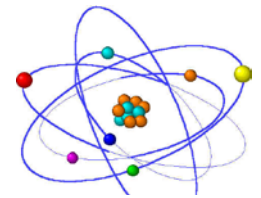


# Overall Goal

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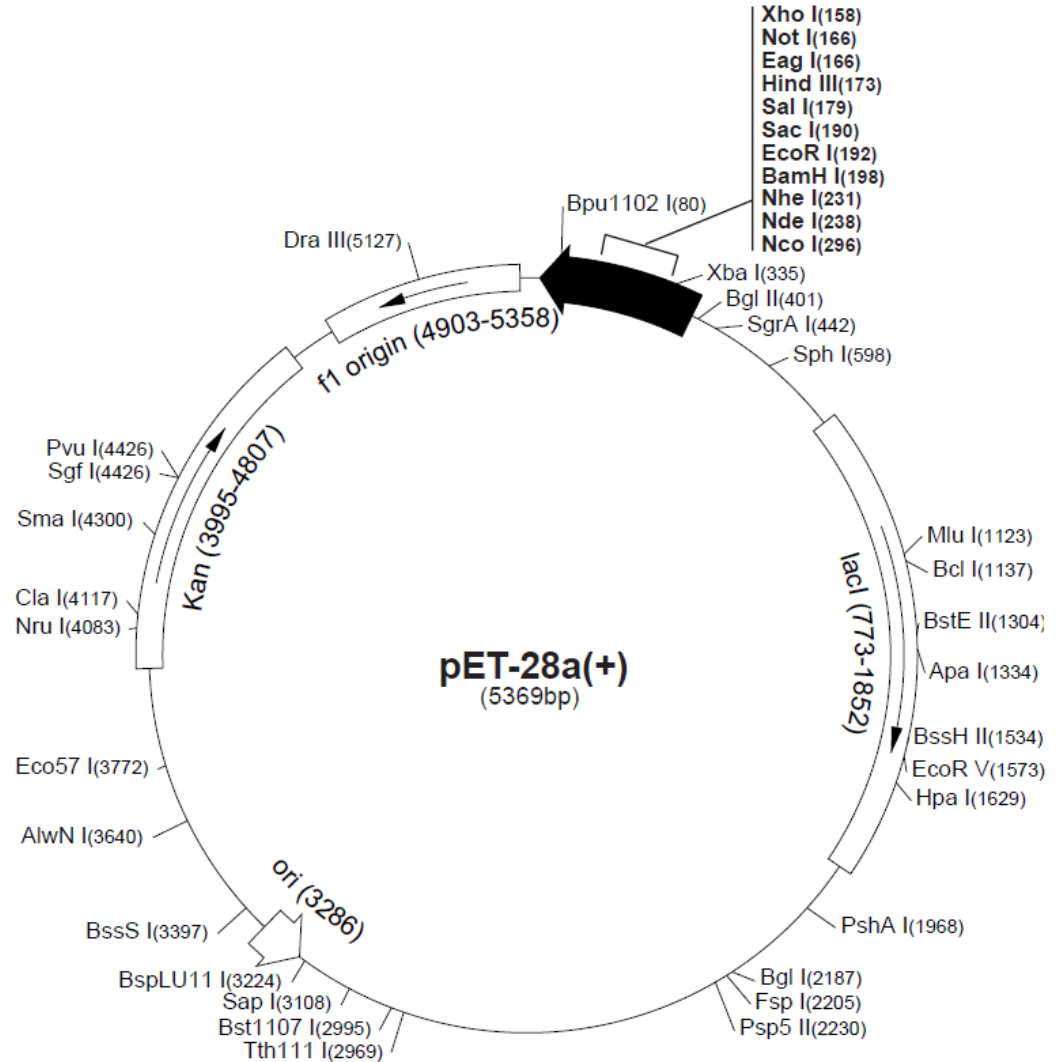
# Plasmid DNA



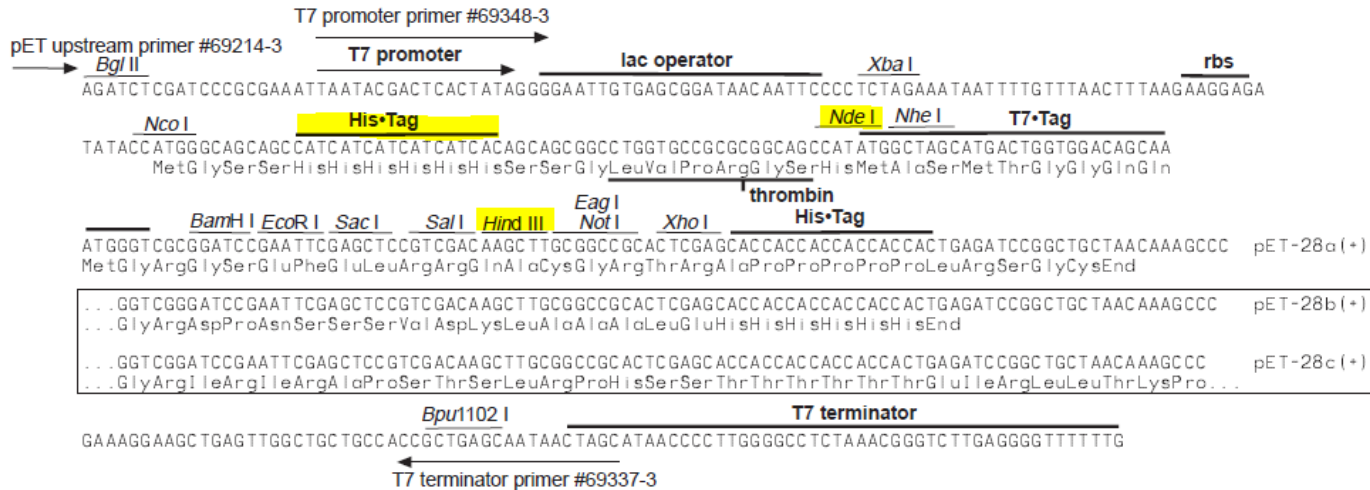
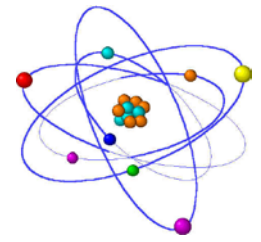
## pET-28a(+) sequence landmarks

T7 promoter	370-386
T7 transcription start	369
His•Tag coding sequence	270-287
T7•Tag coding sequence	207-239
Multiple cloning sites ( <i>Bam</i> H I - <i>Xho</i> I)	158-203
His•Tag coding sequence	140-157
T7 terminator	26-72
<i>lac</i> I coding sequence	773-1852
pBR322 origin	3286
Kan coding sequence	3995-4807
f1 origin	4903-5358

The maps for pET-28b(+) and pET-28c(+) are the same as pET-28a(+) (shown) with the following exceptions: pET-28b(+) is a 5368bp plasmid; subtract 1bp from each site beyond *Bam*H I at 198. pET-28c(+) is a 5367bp plasmid; subtract 2bp from each site beyond *Bam*H I at 198.

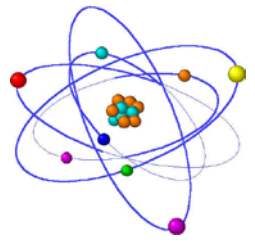


# Plasmid DNA



**pET-28a-c(+)** cloning/expression region

# Polymerase Chain Reaction



Role: Amplify a DNA sequence

Melting



Annealing



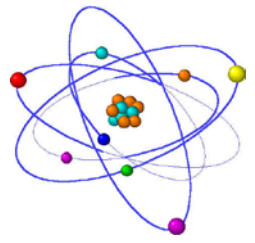
Elongation

*Total copies per unit of template =  $2^n$*



# Polymerase Chain Reaction

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So what do we need to carry out a PCR Reaction”?

1. Template DNA
2. Oligonucleotides - Also called “primers” – the sequence of these are critical to select the proper region of the template to amplify
3. dNTPs (dATP, dGTP, dCTP, dTTP) – monomers that build the polymer
4. DNA Polymerase – the enzyme that catalyzes the polymerization reaction
5. Proper solution conditions (buffer, salt, etc.)

# Digestion Overview

- HindIII

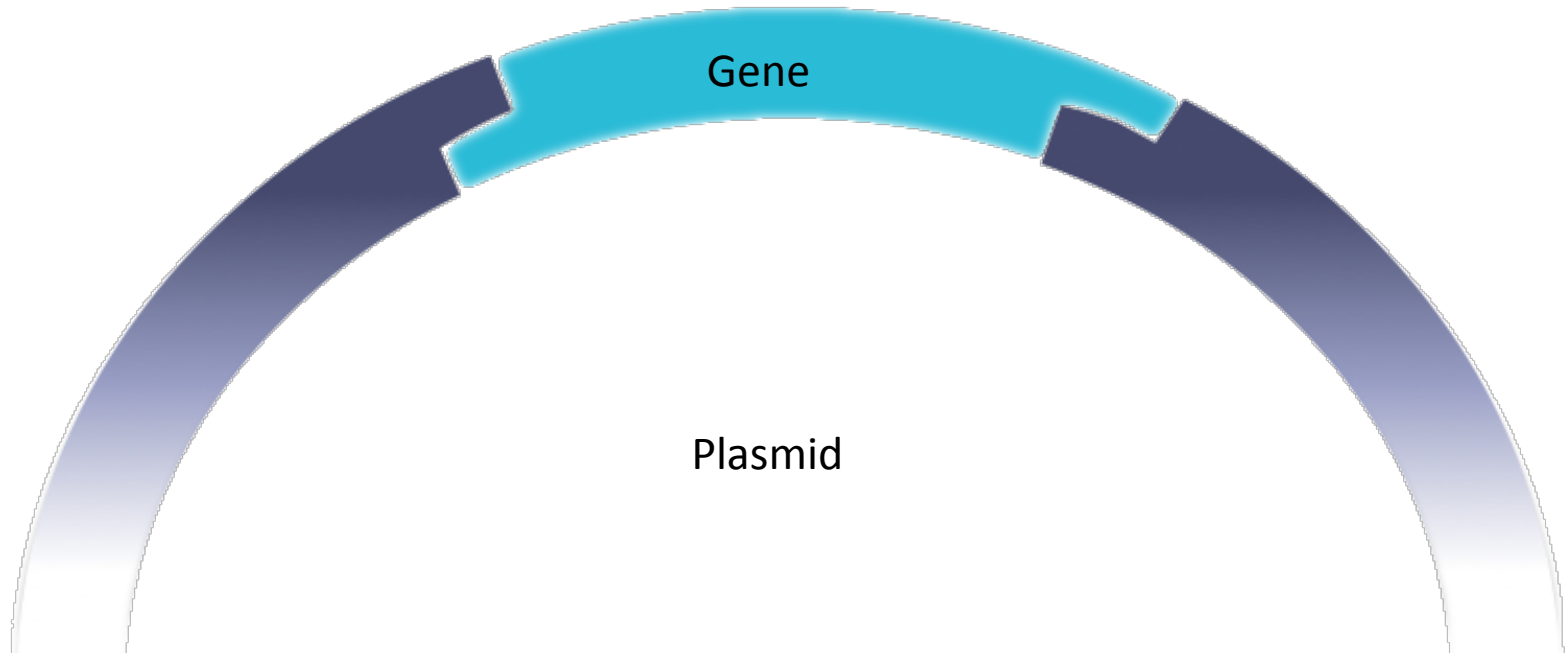


- NdeI



# Ligation

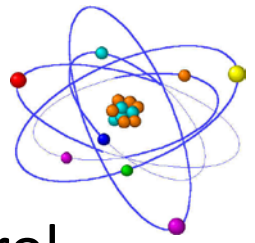
- Insert DNA into plasmid



- Transformation of DH5 $\alpha$  *E. coli* cells

# What's in a Primer?

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This is the part of the PCR reaction that you get to control – and the easiest way to make a PCR reaction fail!

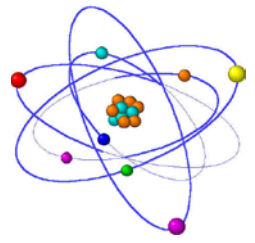
Primers need to “Bracket” the target DNA

Requirements:

- Specificity –
  - Correct Sequence!
  - Aiming for  $T_m \sim 60^\circ\text{C}$
- Directionality
- Include “add-ons”.
  - Probably need to include restriction enzyme sequences
  - Start or stop codon?
  - Protein “tag”

# In Lab

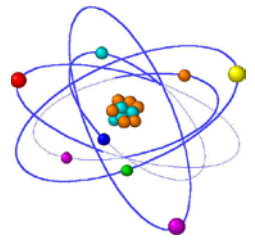
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- Set up a PCR reaction to amplify the LDH gene
  - Be prepared to do this quickly – the PCR reaction takes 1.5 hours
  - Always keep enzymes on ice
- Column purify (mini-prep) pET28 from *E.coli:DH5α::pET28a*
- Column purify your PCR sample
- Determine the concentration of your two samples using the NanoDrop
- Save 5μL of your purified PCR
- Set up two separate digestion reactions (using BamHI and HindIII):
  - pET28a
  - LDH

# Outside of Lab

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## Before Lab

- Complete the PreLab assignment – this will include exploring the sequence of the pET28 plasmid. I encourage you to download the [ApE program](#). You'll also be designing a set of primers to amplify a target DNA sequence. [You may want to use the tool here to help.](#)
- Know exactly what you'll be doing – if you are unprepared, you will not finish on time.
- Download the protocols in the resources link on the webpage – *these need to be copied or pasted in your lab notebook!*

## After Lab

- Think critically about what you did in lab. Understand the molecular underpinning of the reactions you conducted. This is all a very important part of your Introduction
- Begin writing your Introduction and Methods section – these are each upcoming assignments.