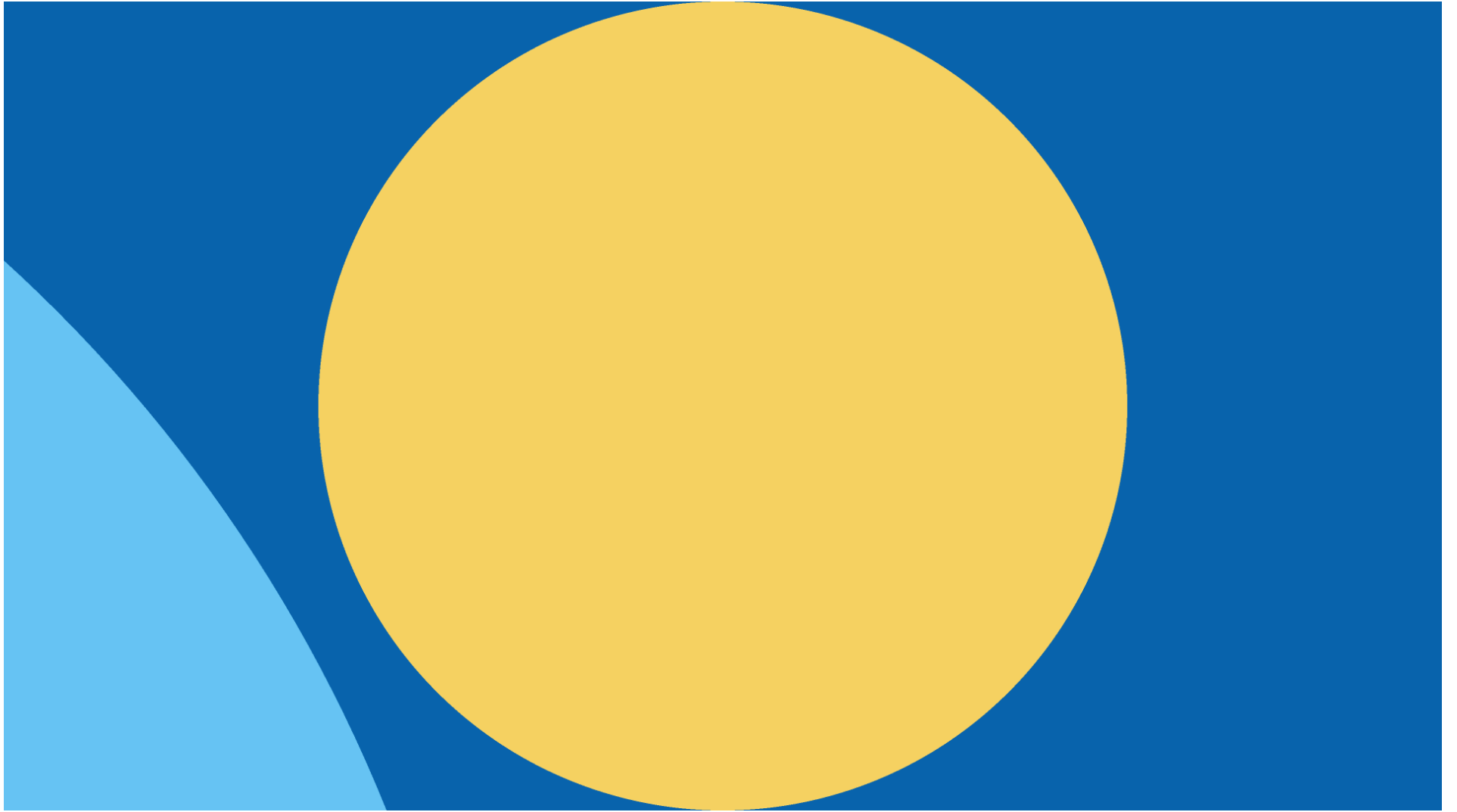


Having high quality RNA is the MOST important thing you can do to insure the success of your experiment!

RNA Handling

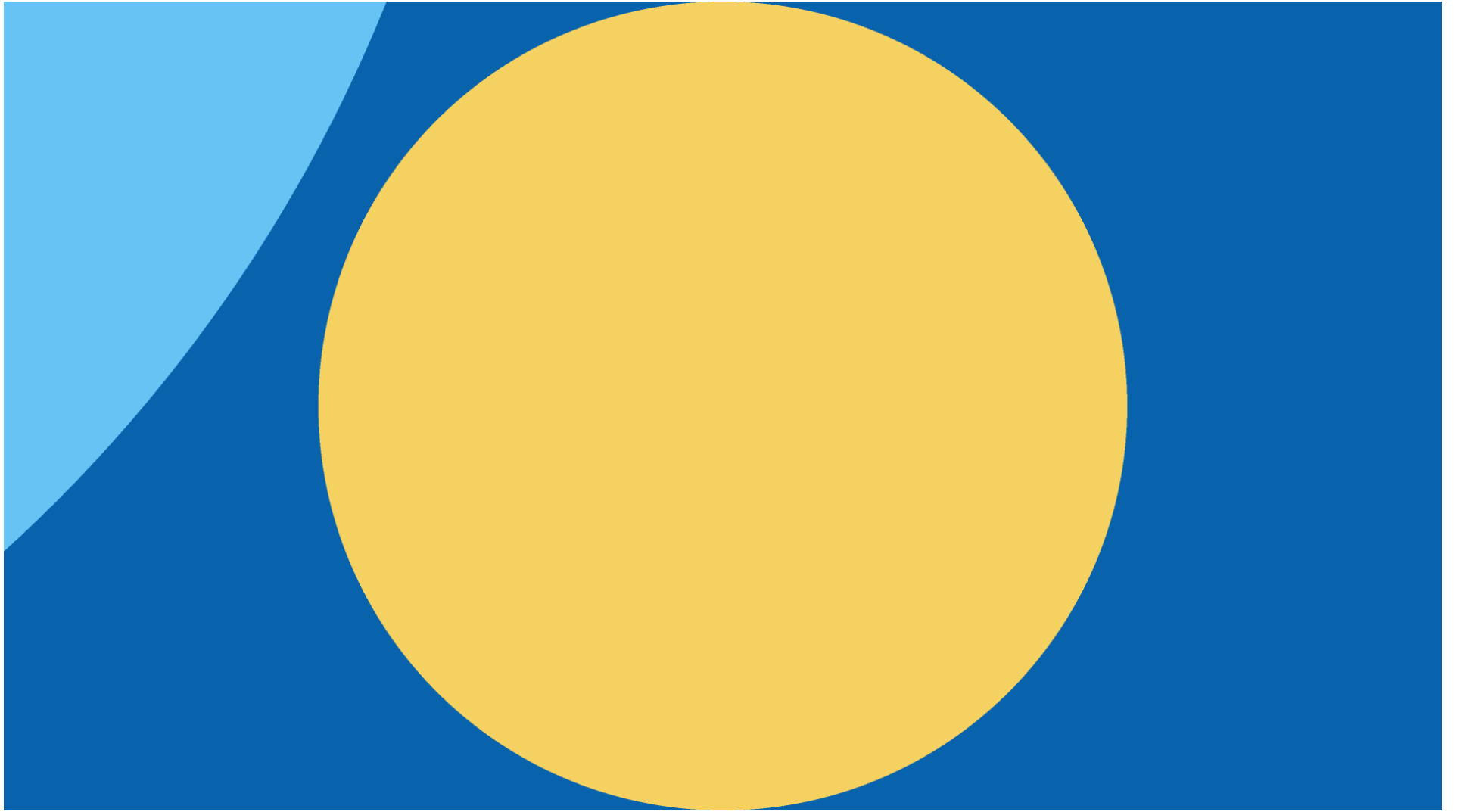
RNA Extraction

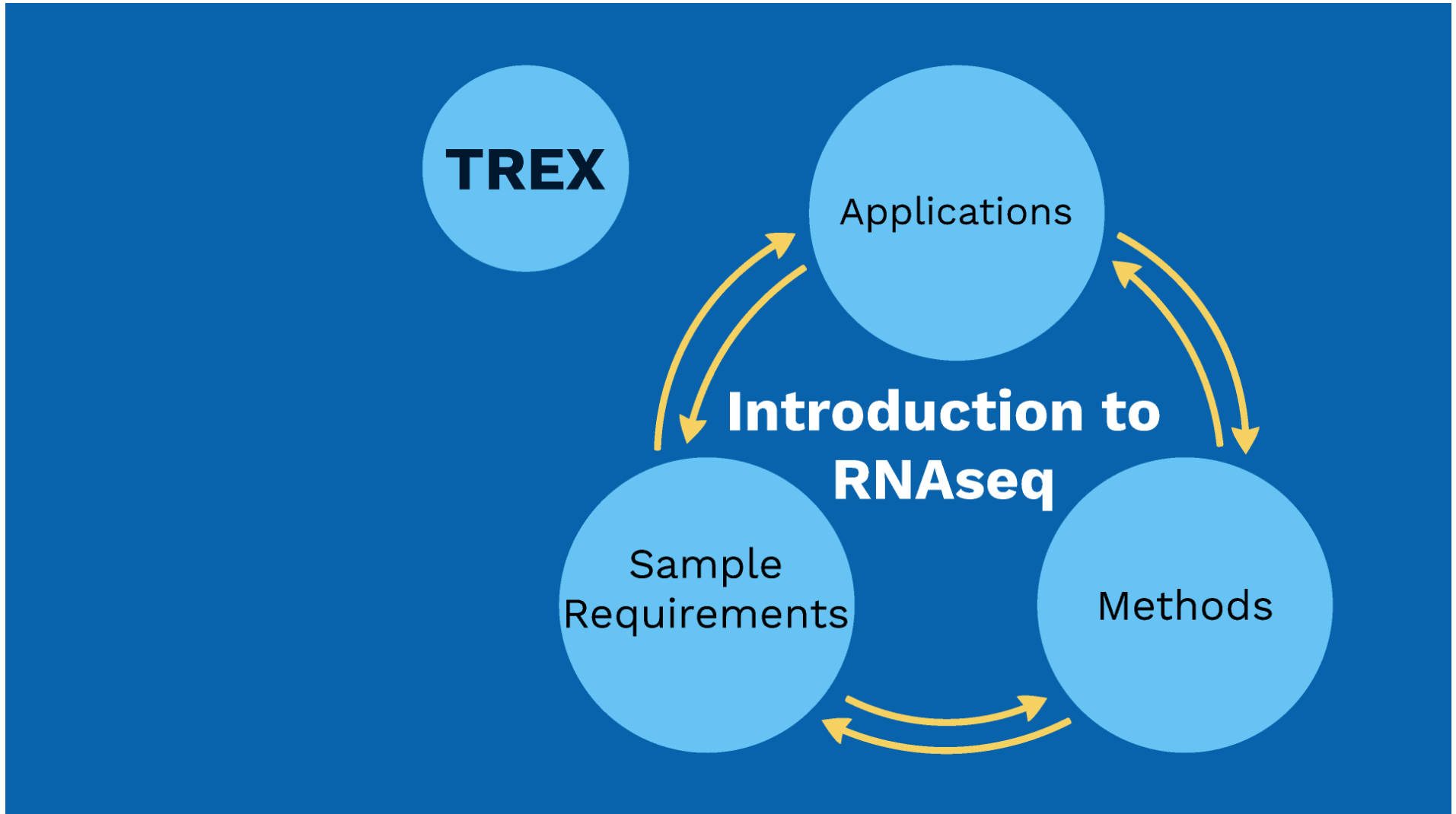
RNA QC





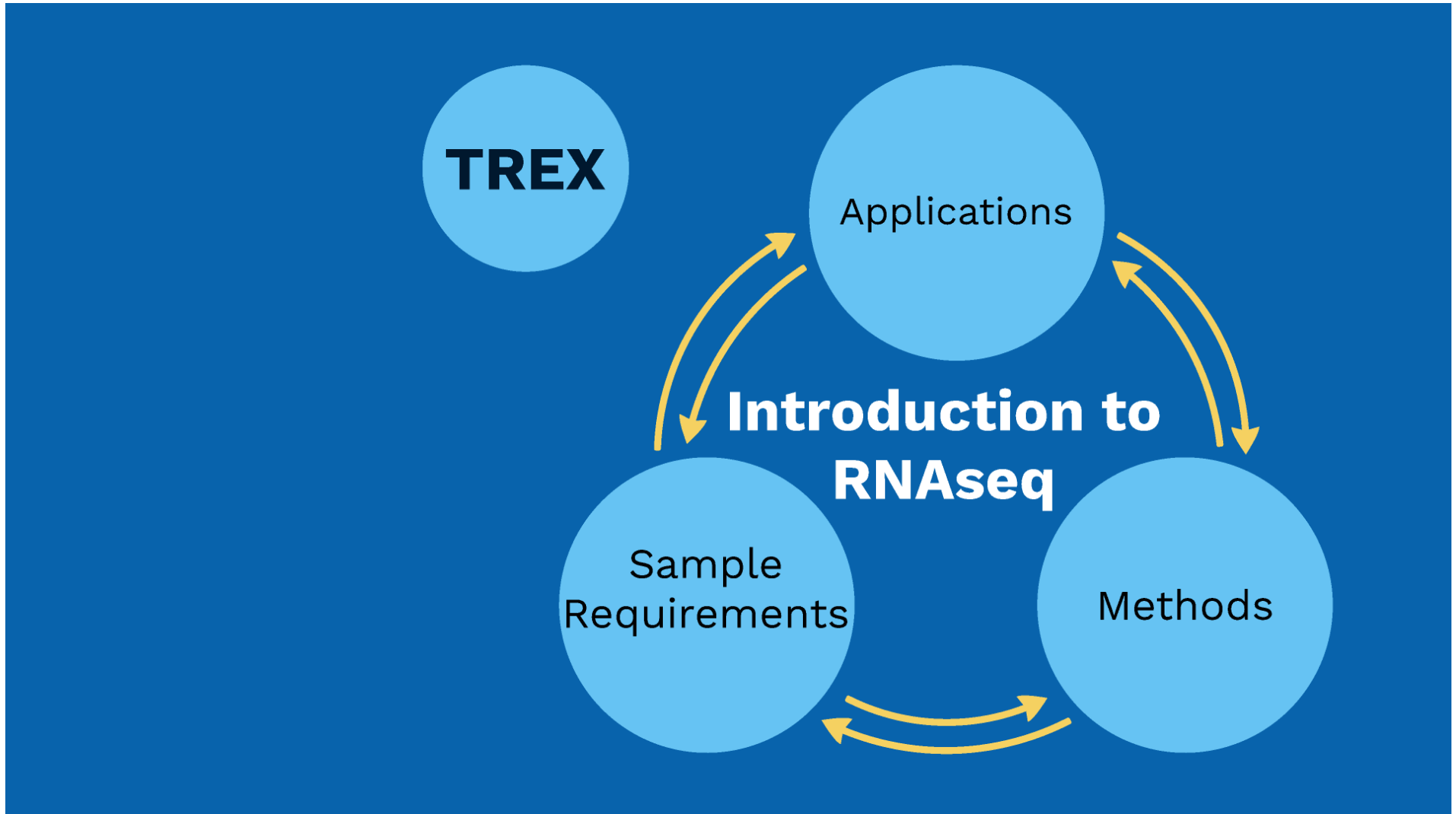
Subtopic





# Why do RNASeq ?

- Gene Expression Profiling
  - Reference (Annotated)
  - De Novo (discovery)
- Variant Analysis or Discovery
- Pathogen ID





## Methods

- What are my research goals?
- What is my RNA quality?
- How many samples do I have?
- How much RNA do I have?

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Lexogen  
3'  
RNASeq

Illumina  
Truseq  
RNA

NEBNext  
Ultra II

NEB  
Small  
RNA

## Methods

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Lexogen  
3'  
RNASeq

**BRC**

Illumina  
Truseq  
RNA

NEBNext  
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NEB  
Small  
RNA

**TREX**

## Why Choose Lexogen 3'RNA Seq?

High Throughput: BRC Service requires >32 samples

Experimental design tolerant of dropouts

The information you are interested in is at the 3' end of the RNA strand

Tolerant of:

- Input concentration diversity
- RNA quality diversity

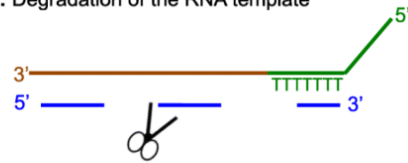
Prep  
Chemistry

### 3' method (LEXO)

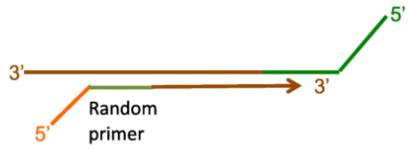
**Step 1:** 1<sup>st</sup> strand synthesis of polyA tailed RNA from total RNA using oligo dT primers



**Step 2:** Degradation of the RNA template



**Step 3:** 2<sup>nd</sup> strand synthesis with random primers containing 5' Illumina-compatible linker sequences



**Step 4:** Amplification using random primers that add barcodes and cluster generation sequences



**Step 5:** Sequencing

# Why Choose Truseq RNA

## Flexible:

- Input type:
  - Total RNA
  - Ribosomal Depleted RNA
  - Poly A Selected RNA
- Input Concentration: 1000-100ng
- RNA Integrity: Intact OR Degraded
- Sample Number: <384

Directional Or Non-Directional

Highly Supported



Chemistry

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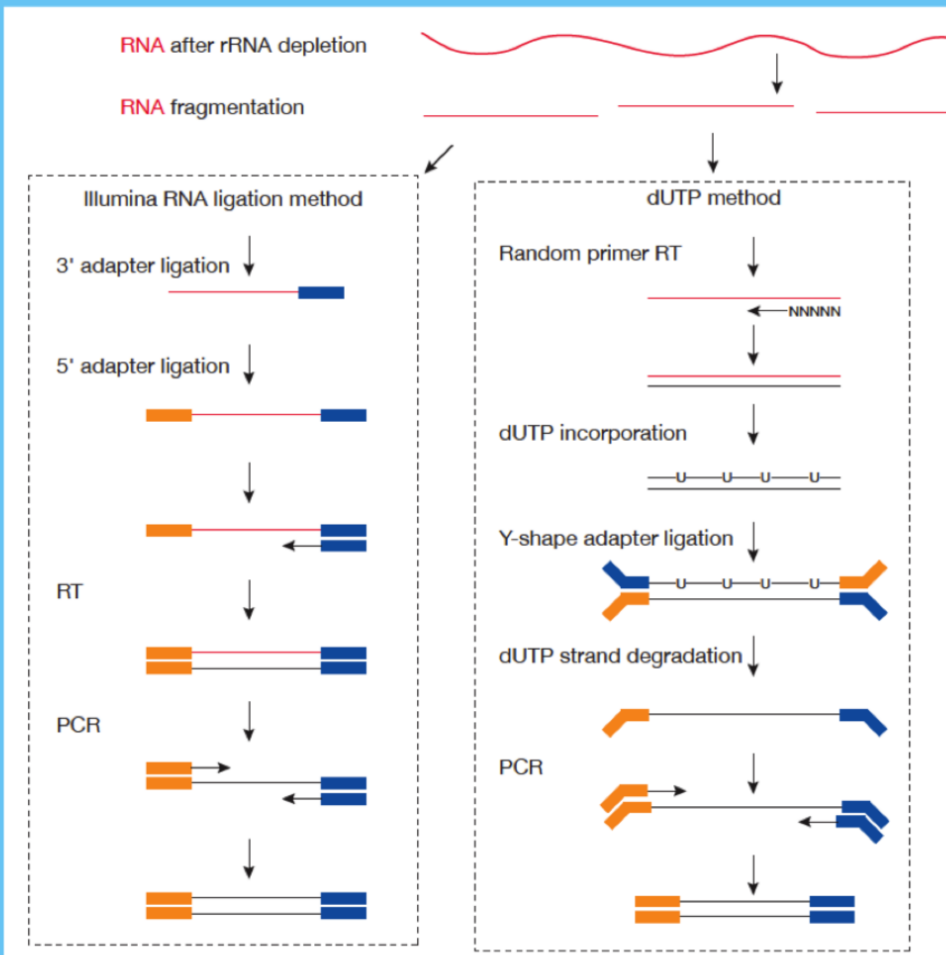
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Chemistry





## Why choose NEB Next Ultra II

### Flexible:

- Input type:
  - Total RNA
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  - Poly A Selected RNA
- Input Concentration: 1000-100ng
- RNA Integrity: Intact OR Degraded
- Sample Number: <384

Fully Supported

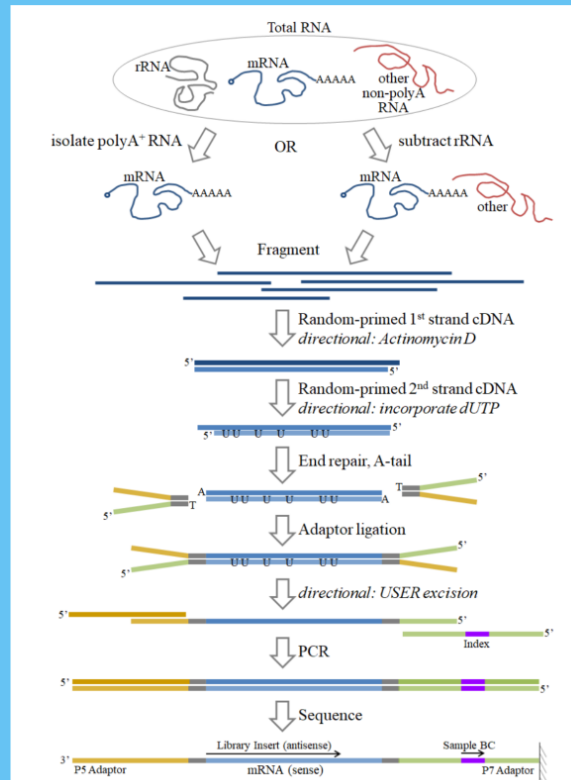
Directional Or Non-Directional

Modular

Prep  
Chemistry

Decision Tree

# NEB Next Ultra II



Poly A

Ribosomal  
Depletion

Directional  
vs.  
Non-  
Directional

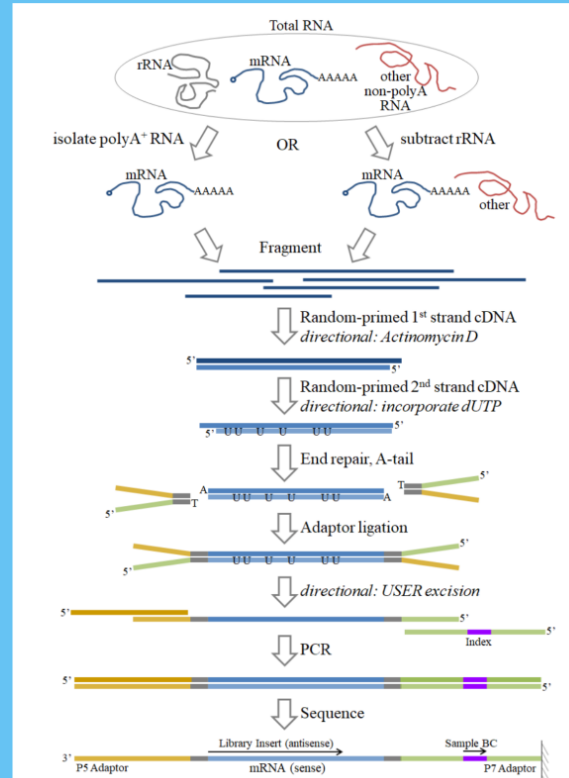
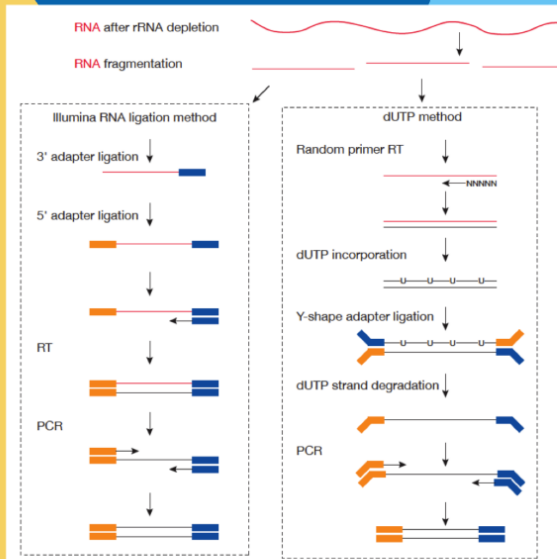
# NEB Next Ultra II

Poly A

Ribosomal Depletion

Directional vs. Non-Directional

## Illumina Truseq



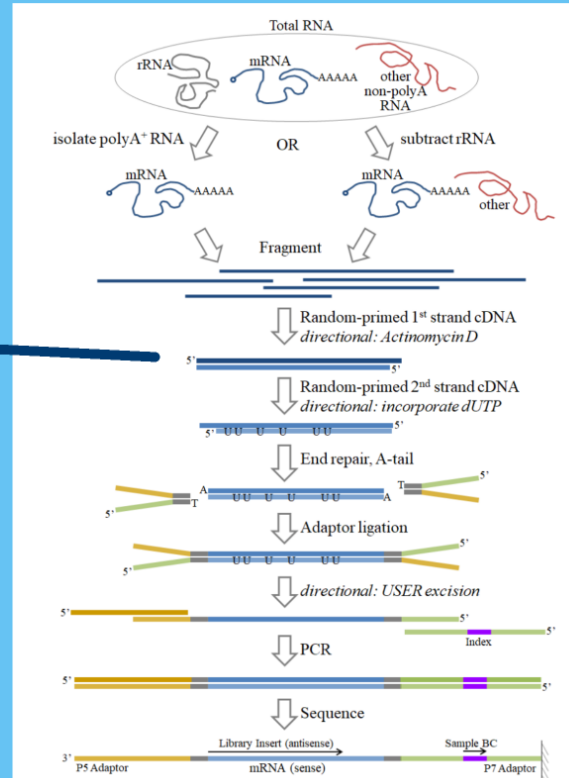
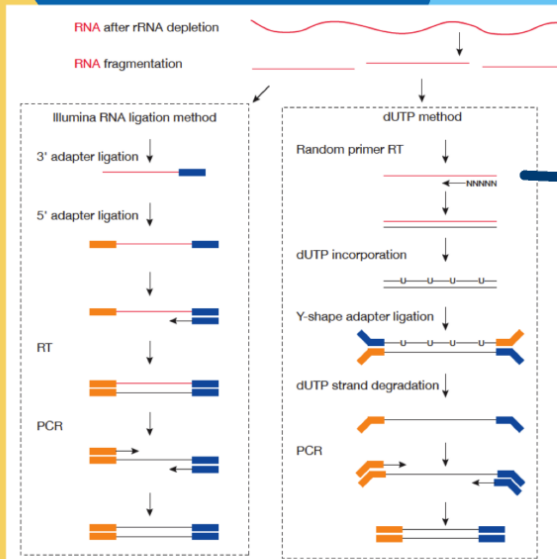
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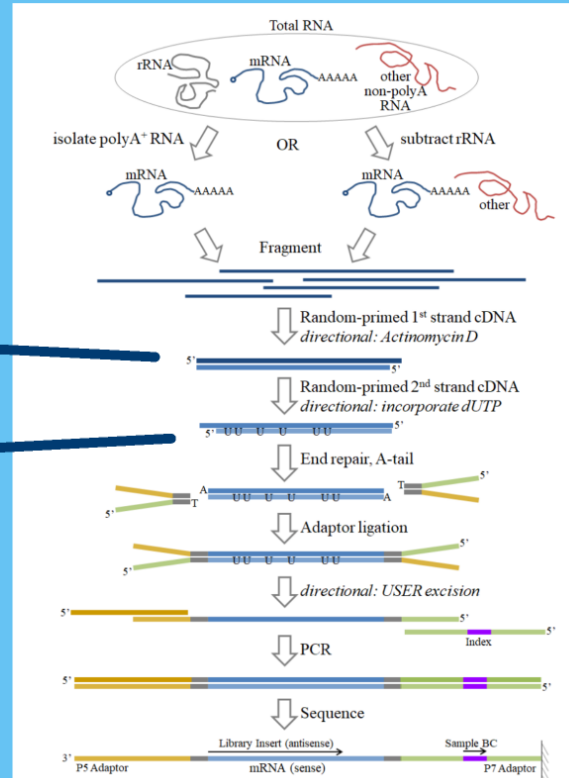
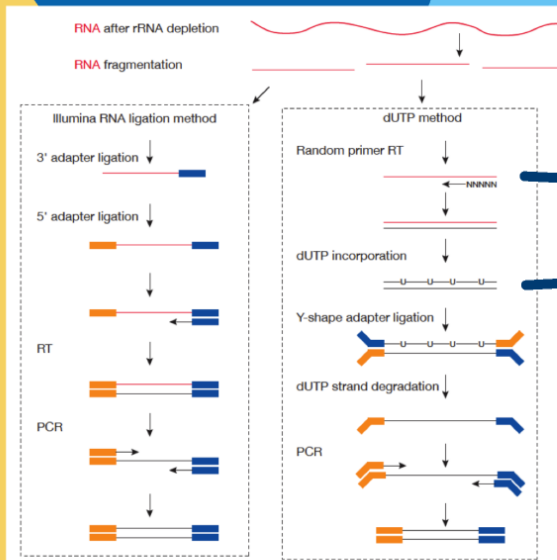
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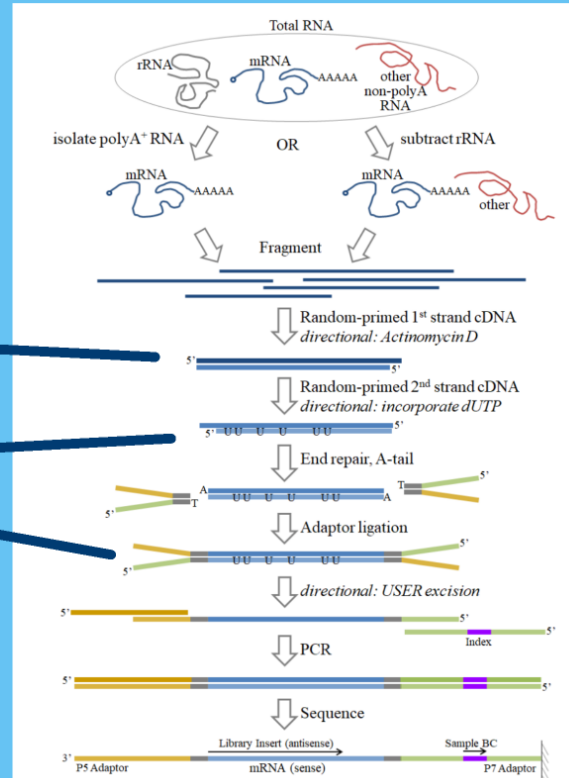
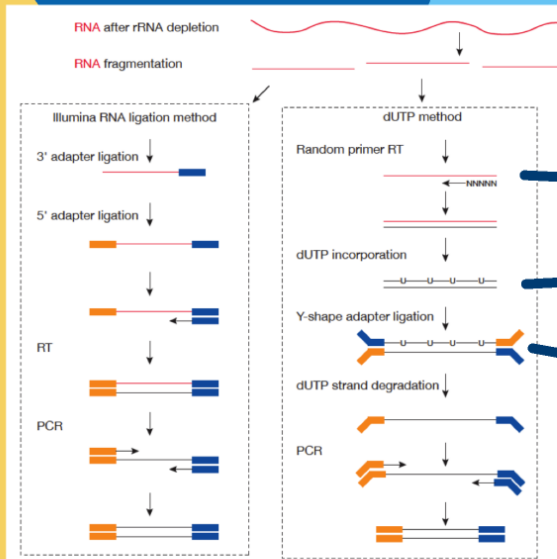
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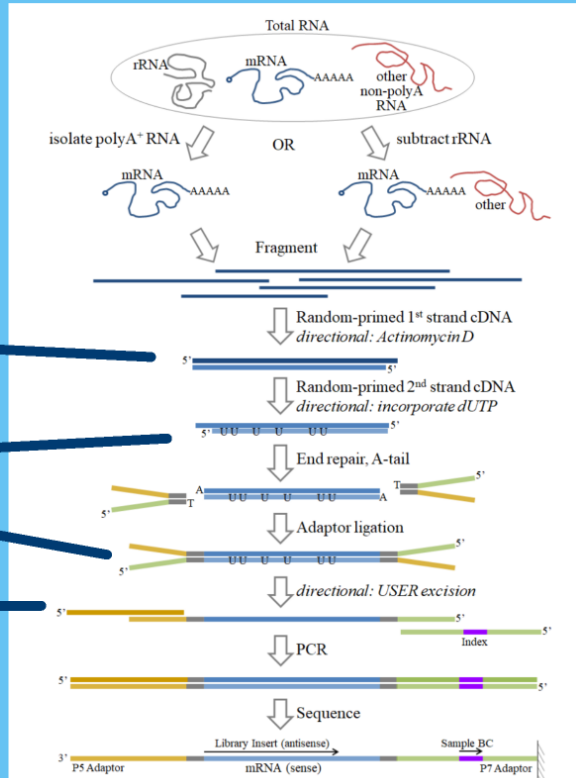
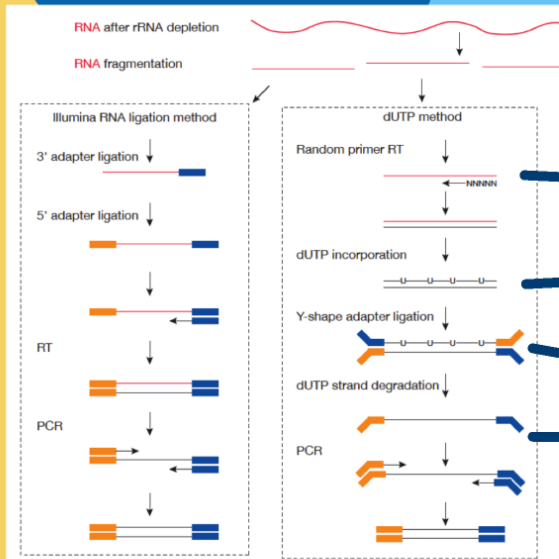
Directional vs. Non-Directional

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# NEB Next Ultra II

## Illumina Truseq



Poly A

Ribosomal Depletion

Directional vs. Non-Directional



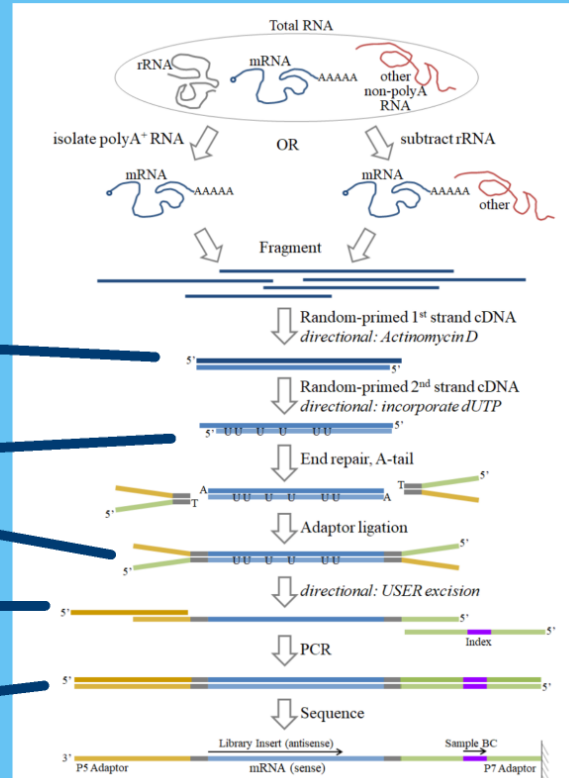
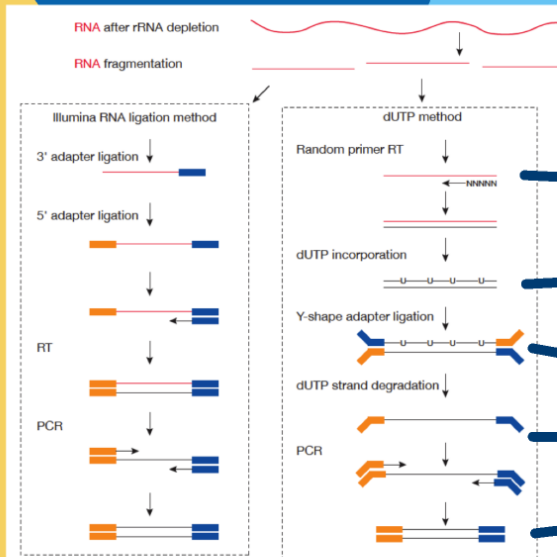
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Poly A

Ribosomal Depletion

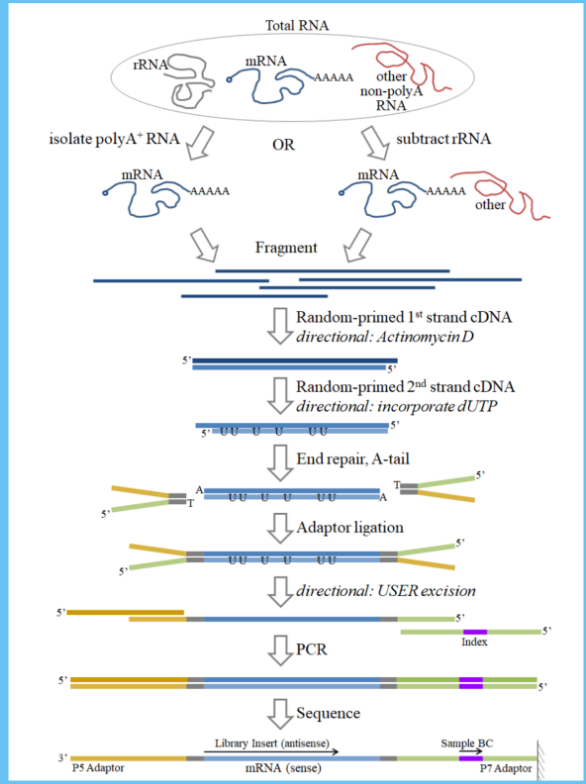
Directional vs. Non-Directional

## Illumina Truseq



# NEB Next Ultra II

Illumina Truseq



Poly A

Ribosomal Depletion

Directional vs. Non-Directional

# What is Poly A Selection?

Keep any RNA fragment with a Poly A stretch in it

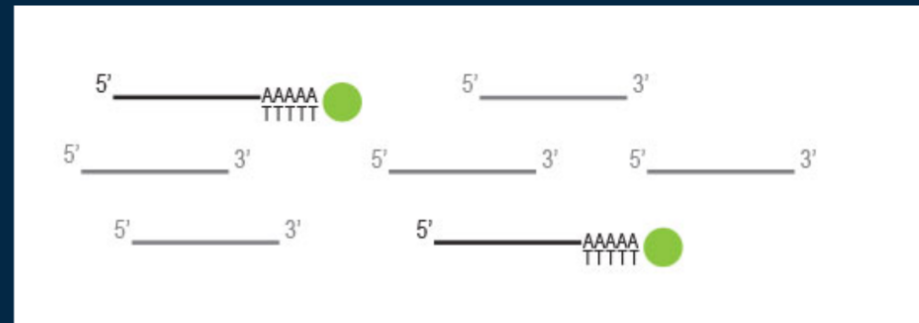
Discard everything else

- Ribosomal RNA

- Some lncRNA

- Other Housekeeping RNAs

- Degraded RNA - only keep the pieces with Poly A tails



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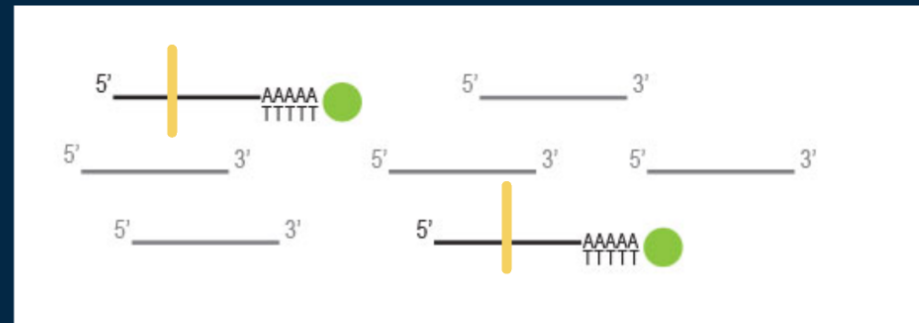
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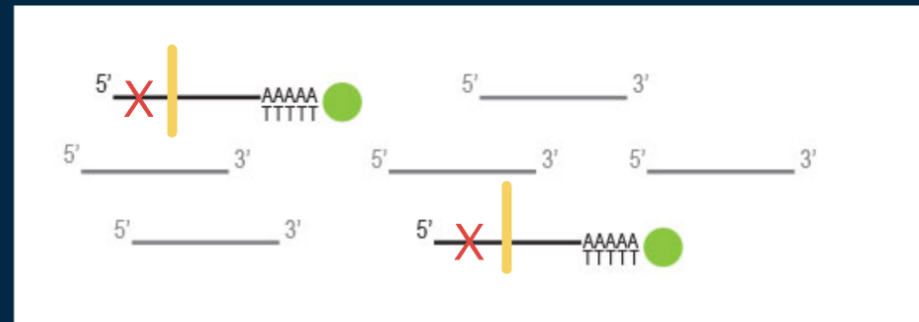
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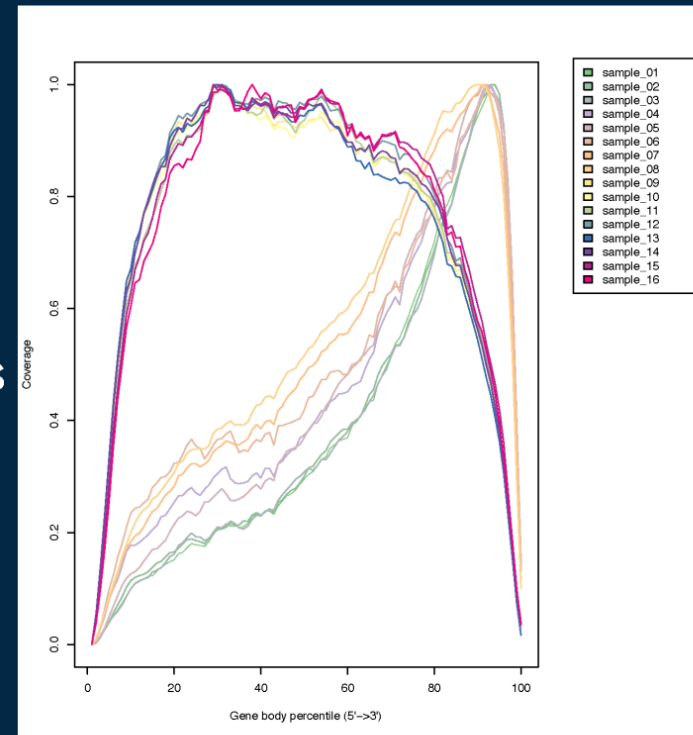
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3' Bias



# Why Choose Poly A?



# Why Choose Poly A?

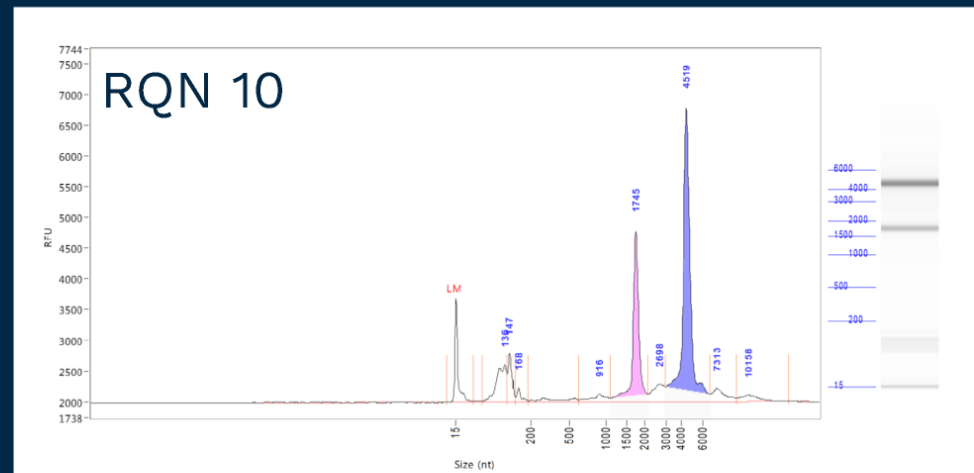
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  - If you are looking for RNA's that don't have a Poly A tail, ex: LNC RNA

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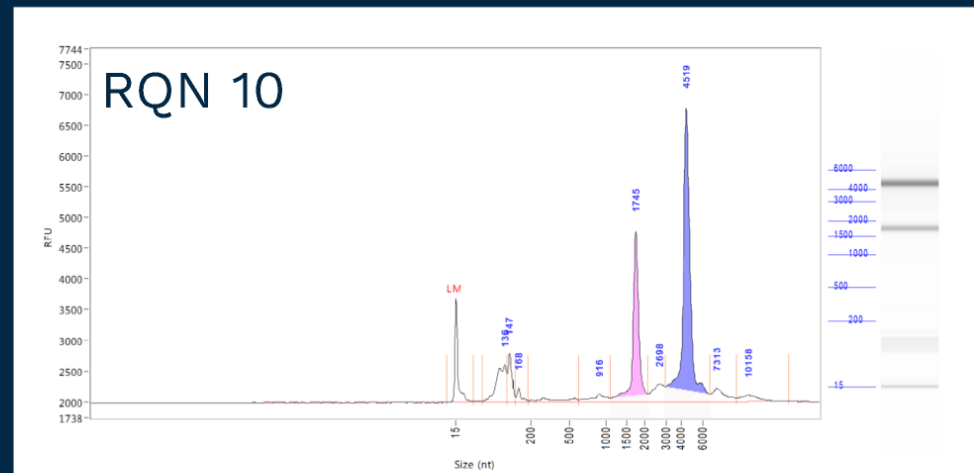
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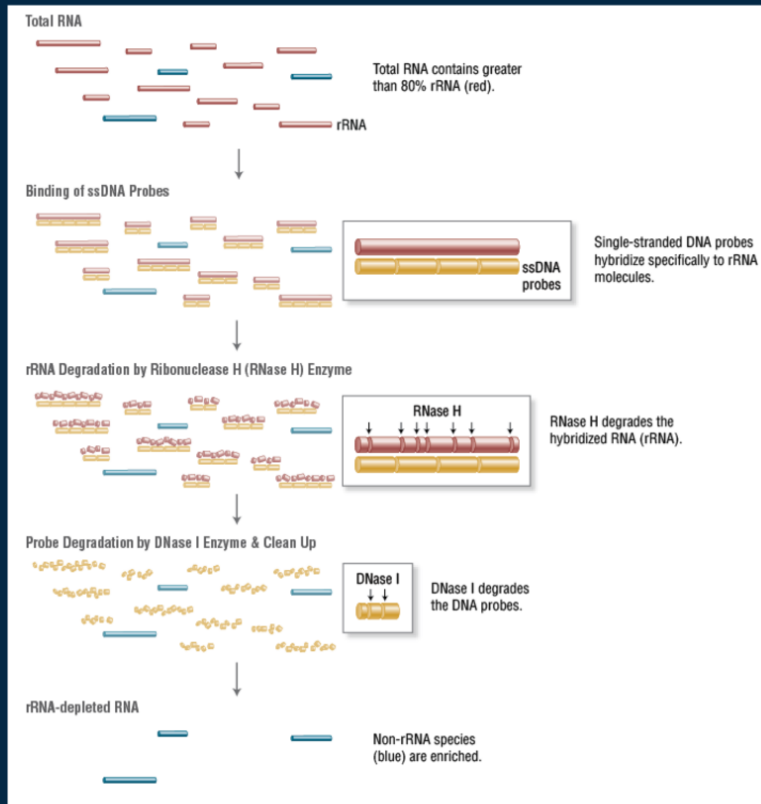


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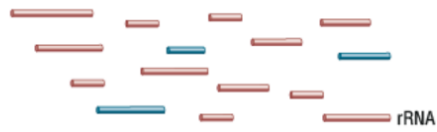
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- If your RNA is intact:
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- If you don't care about 3' bias



# What is Ribosomal Depletion



Total RNA



Total RNA contains greater than 80% rRNA (red).

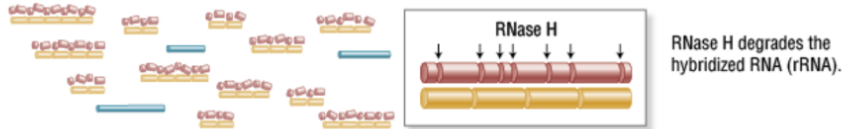
rRNA

Binding of ssDNA Probes



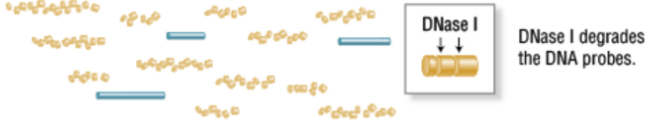
Single-stranded DNA probes hybridize specifically to rRNA molecules.

rRNA Degradation by Ribonuclease H (RNase H) Enzyme



RNase H degrades the hybridized RNA (rRNA).

Probe Degradation by DNase I Enzyme & Clean Up



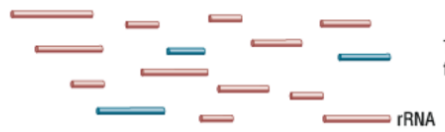
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rRNA-depleted RNA



Non-rRNA species (blue) are enriched.

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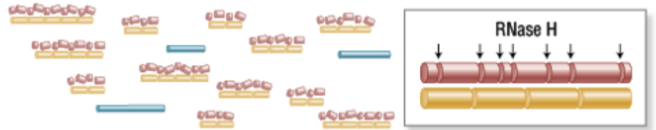
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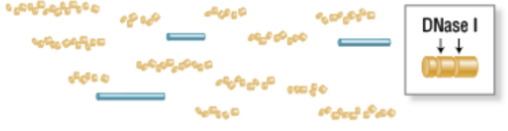
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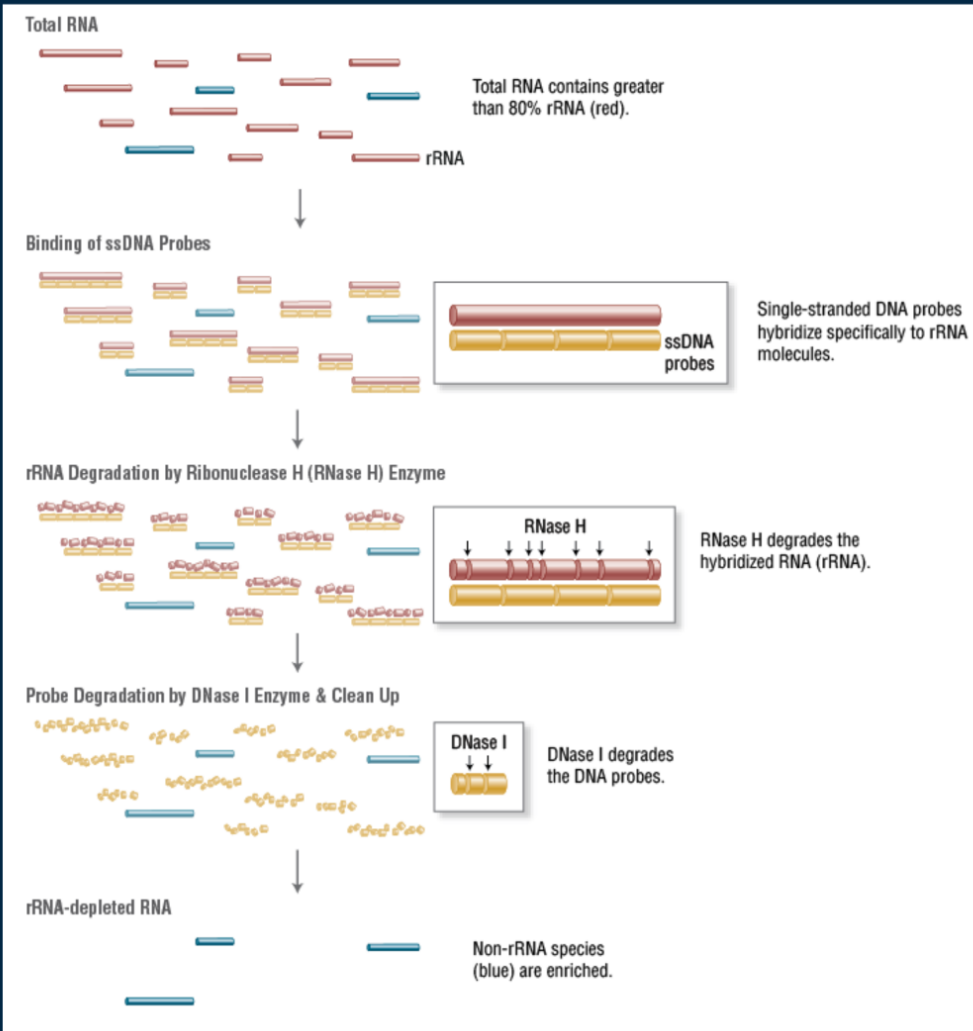
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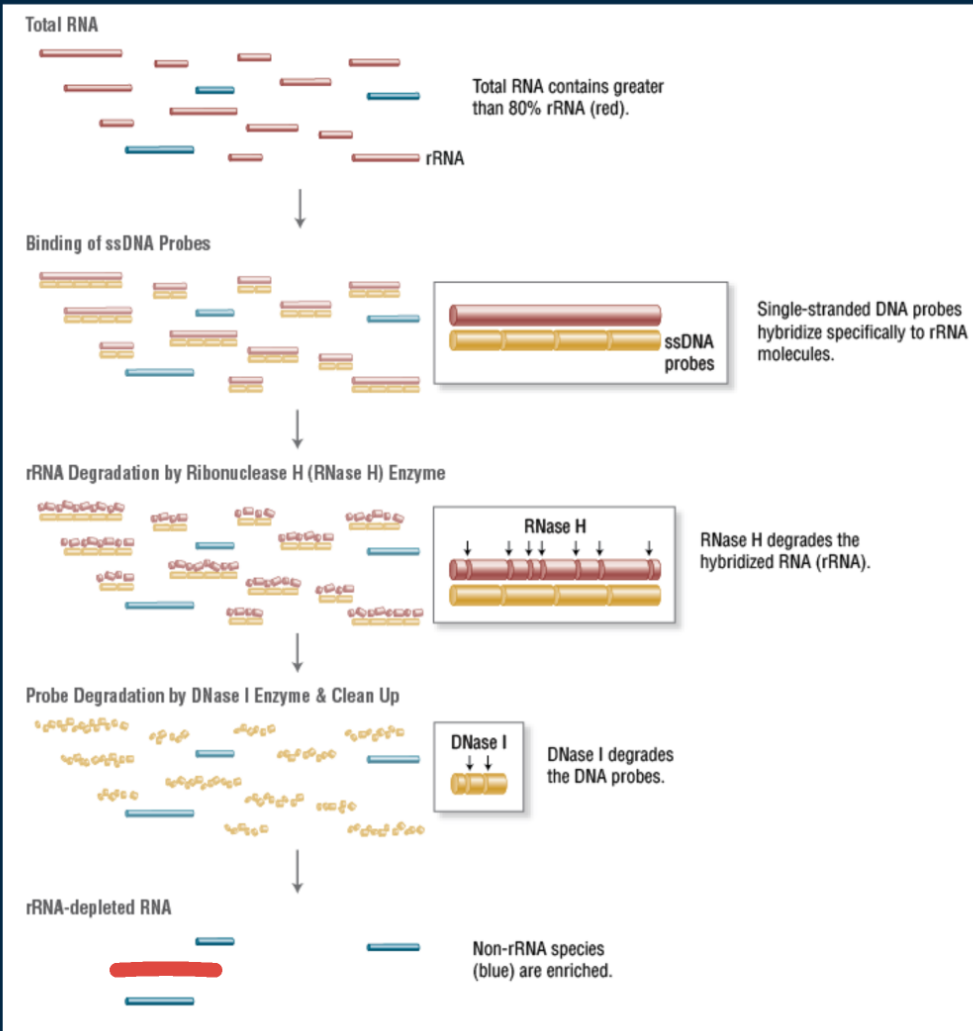
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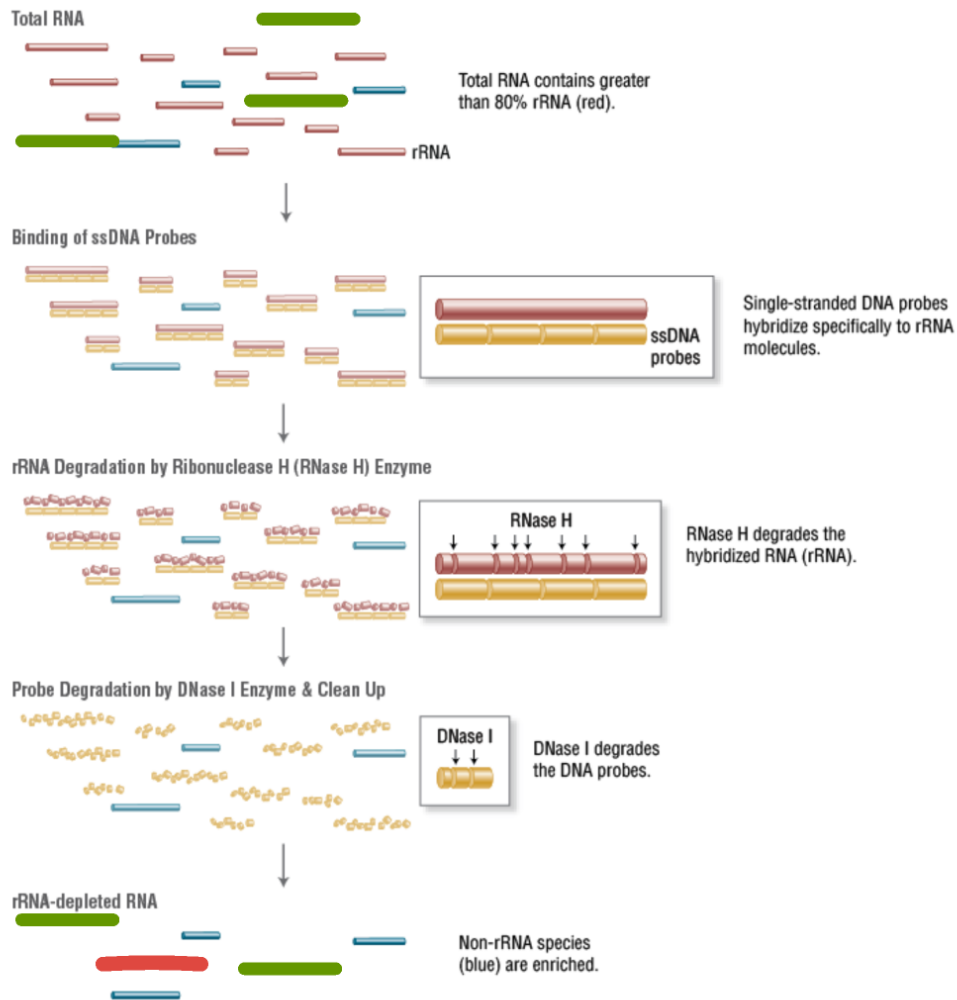
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Housekeeping RNAs

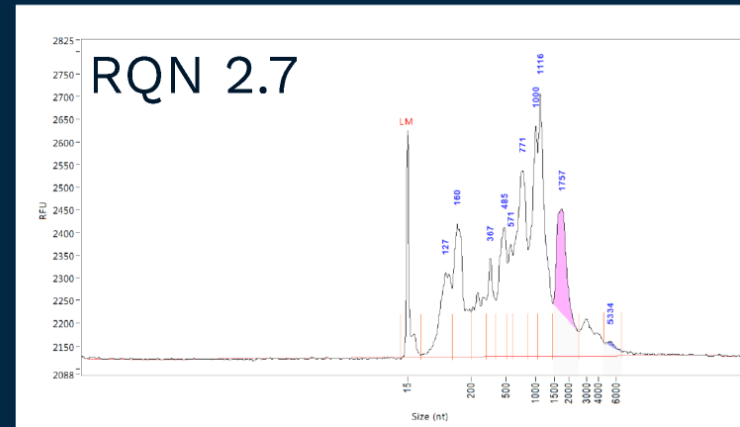
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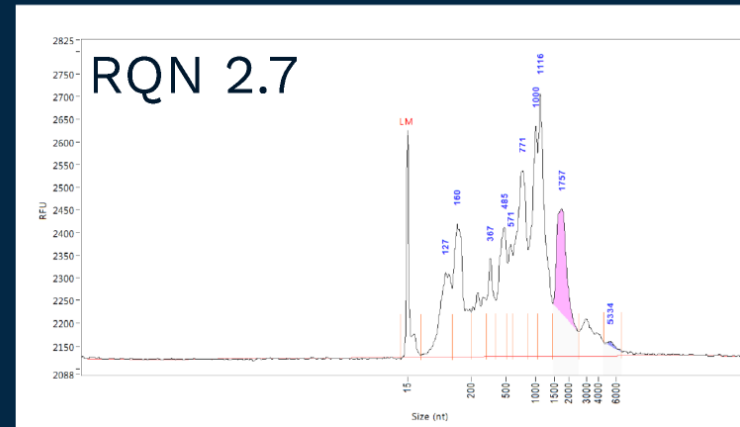
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If your organism is not compatible with Poly A

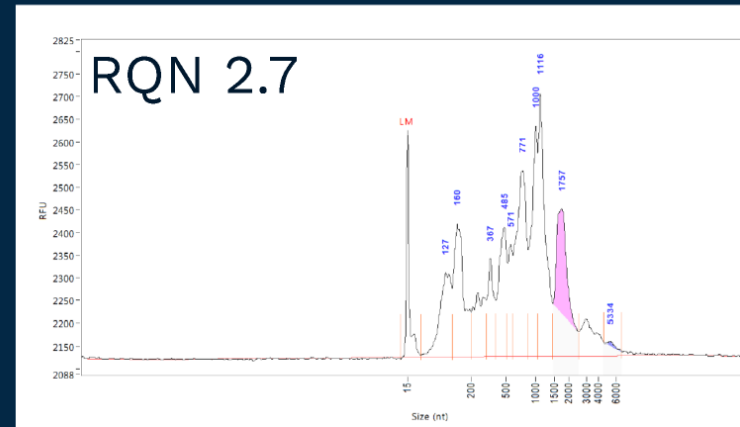


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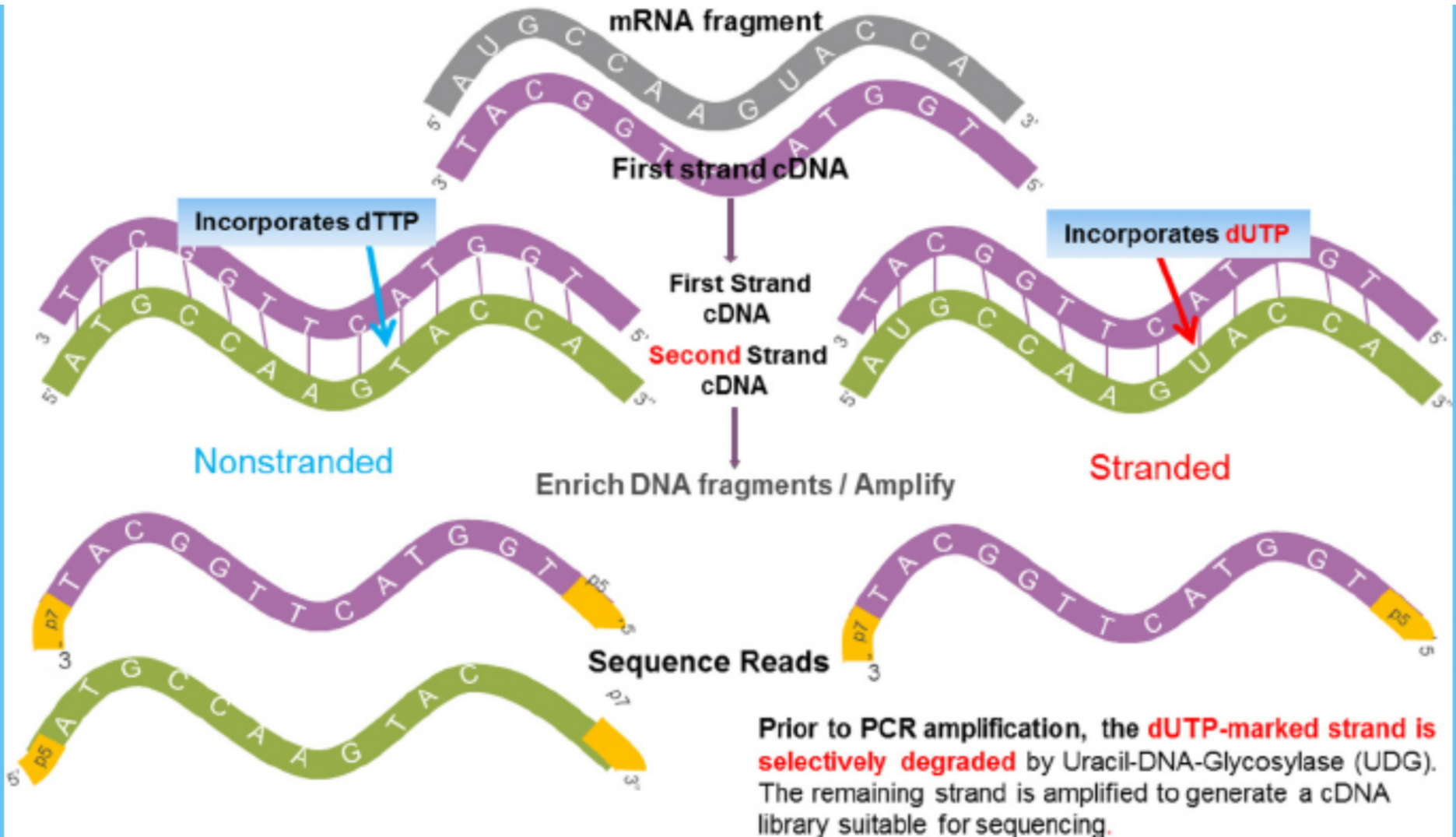
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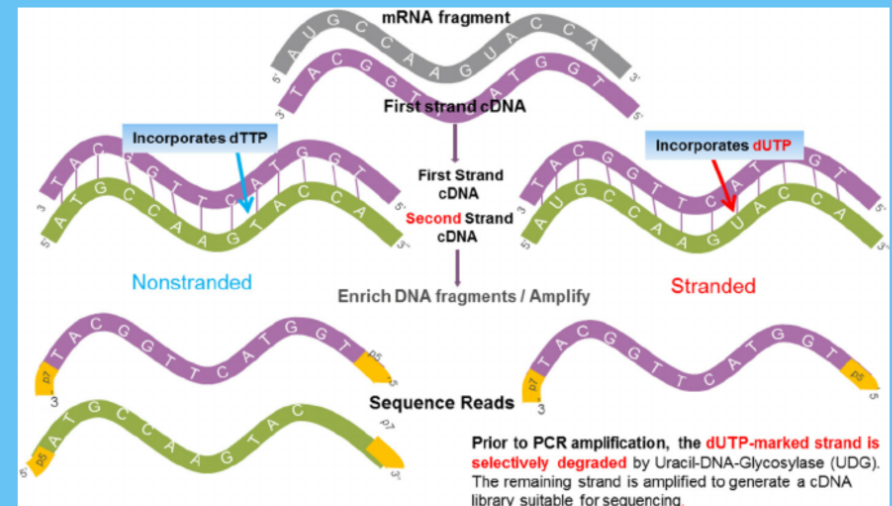
# What is the difference between Directional and Nondirectional?







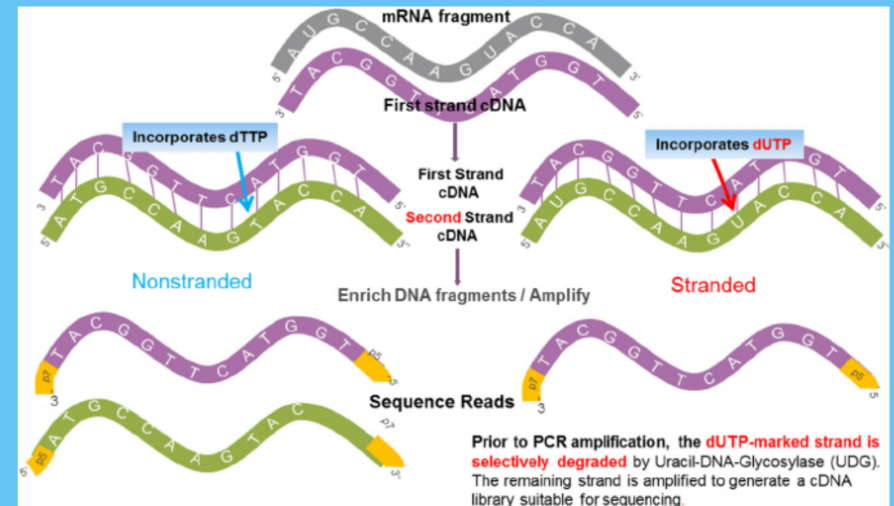
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  - Which strand your RNA is being transcribed from
- More accurate count of genes in differential expression analysis



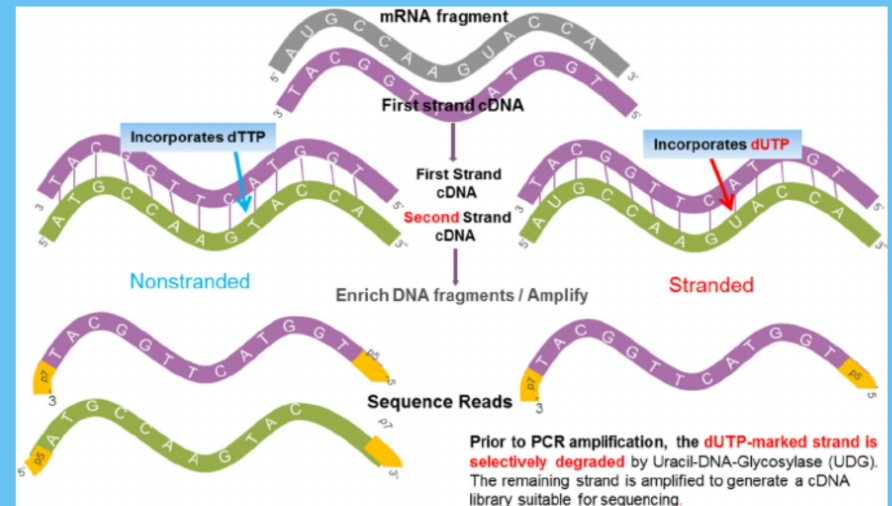
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- If you have <10ng of total RNA



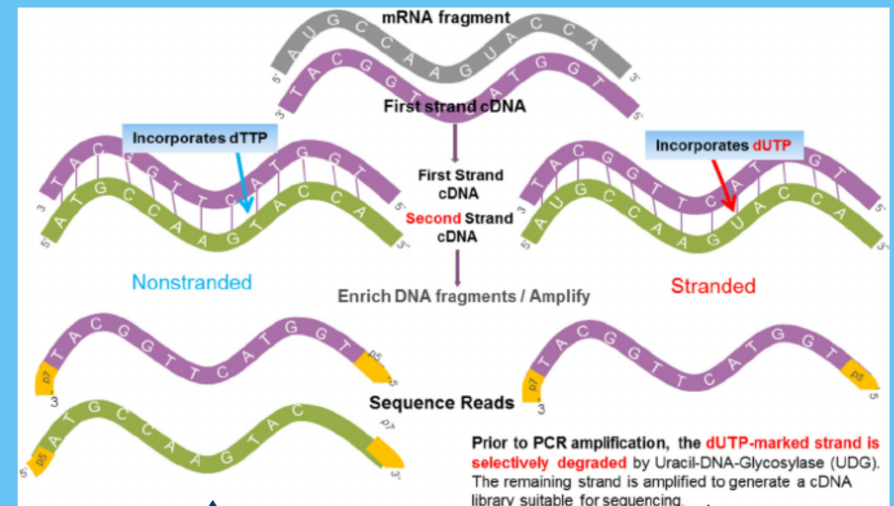
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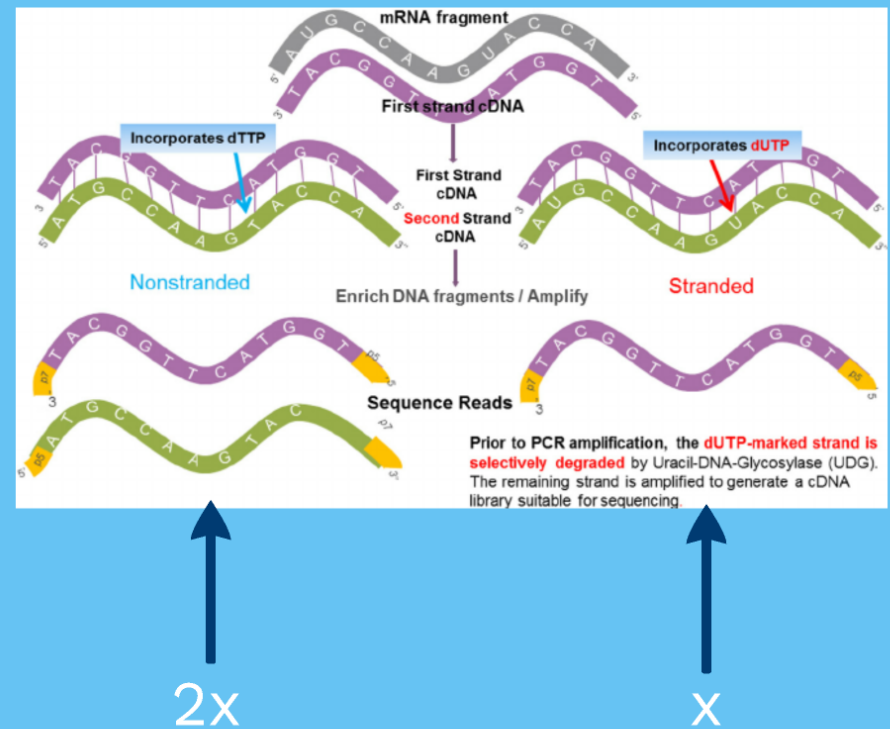
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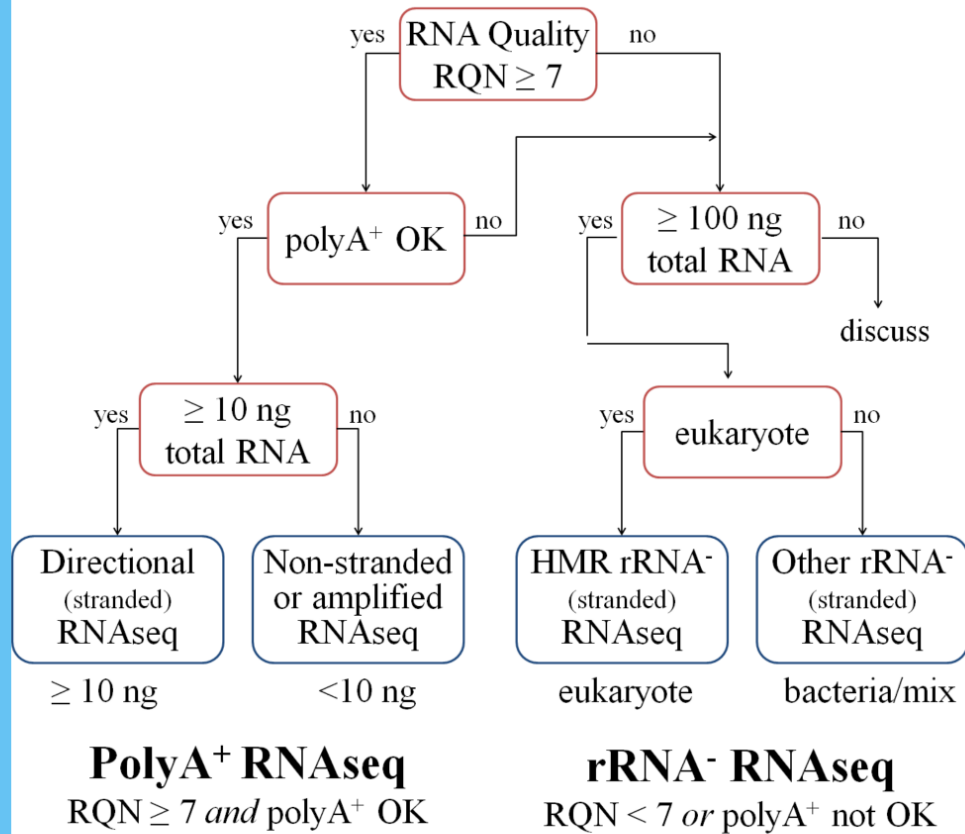
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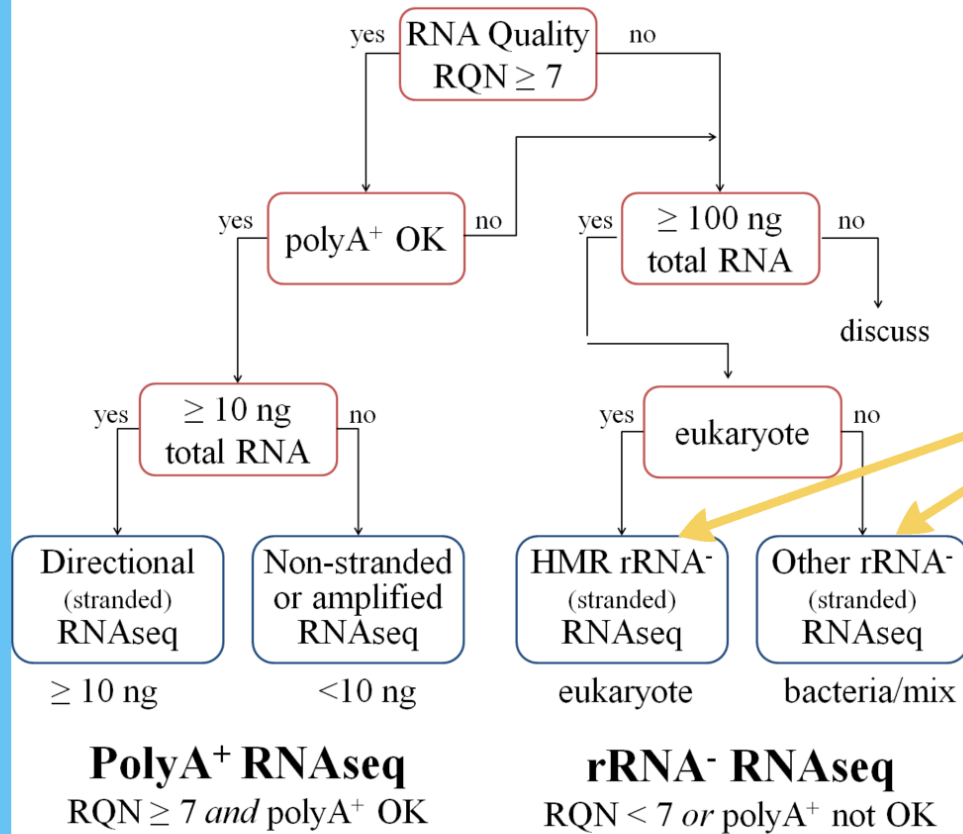
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## RNaseq Decision Tree



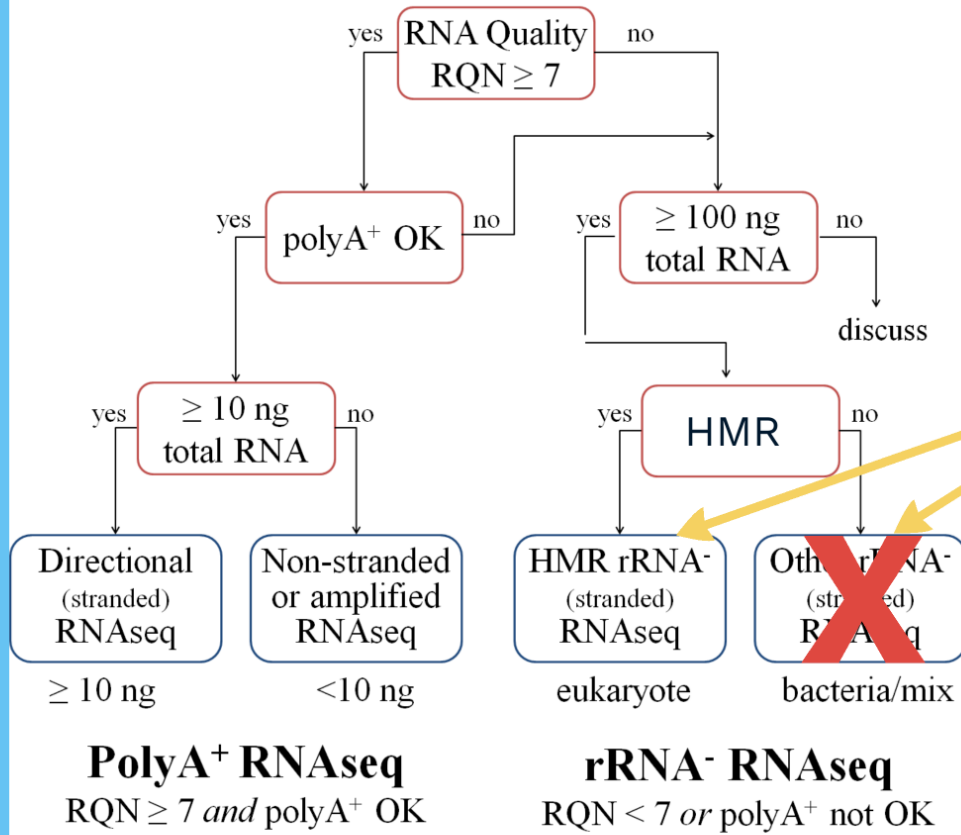
## RNaseq Decision Tree



Ribo Zero from illumina has been discontinued!



## RNaseq Decision Tree



Ribo Zero from  
illumina has been  
discontinued!

## Why choose Small RNA?

Analyzing microRNAs, siRNAs, piRNAs

- Selecting for 20-30nt small RNAs
- Minimum input of 100ng of cellular Total RNA

What is  
small RNA

Prep  
Chemistry

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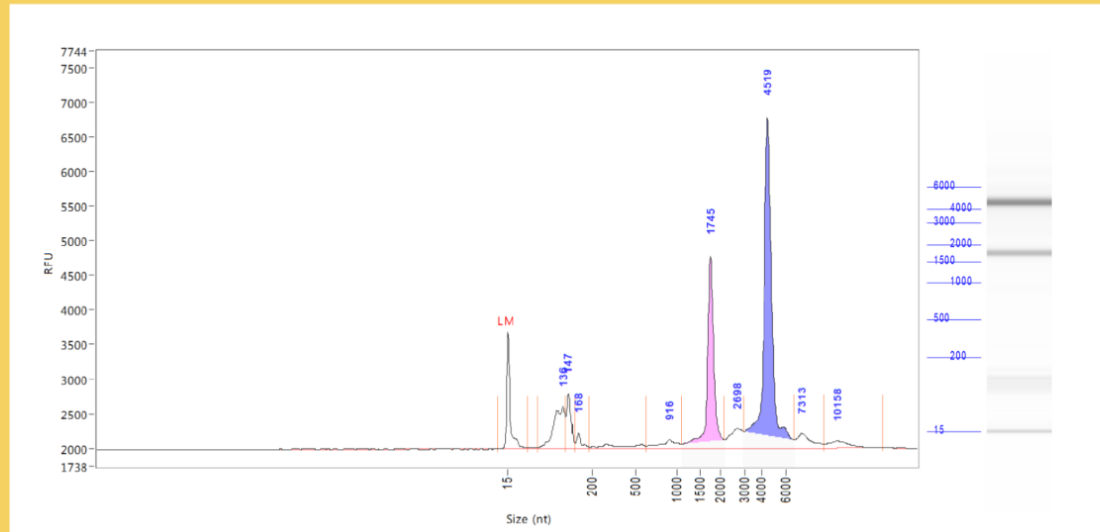
\* For RNA Extraction: make sure you use a method that keeps small RNA's \*

What is small RNA

Prep Chemistry

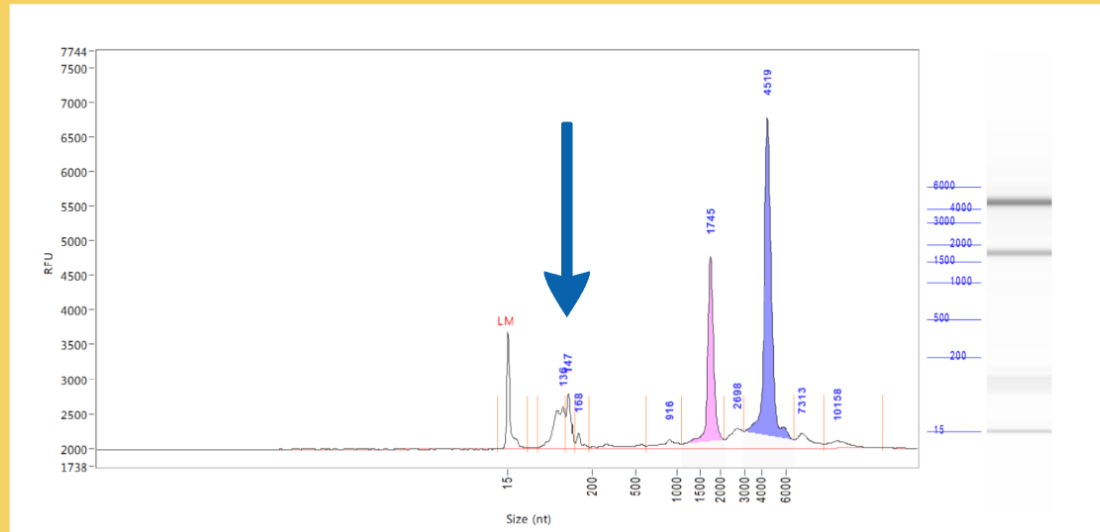
# What is small RNA

- Type of ncRNA
- Small, 25-250NT's
- Involved in regulating translation of target RNA's

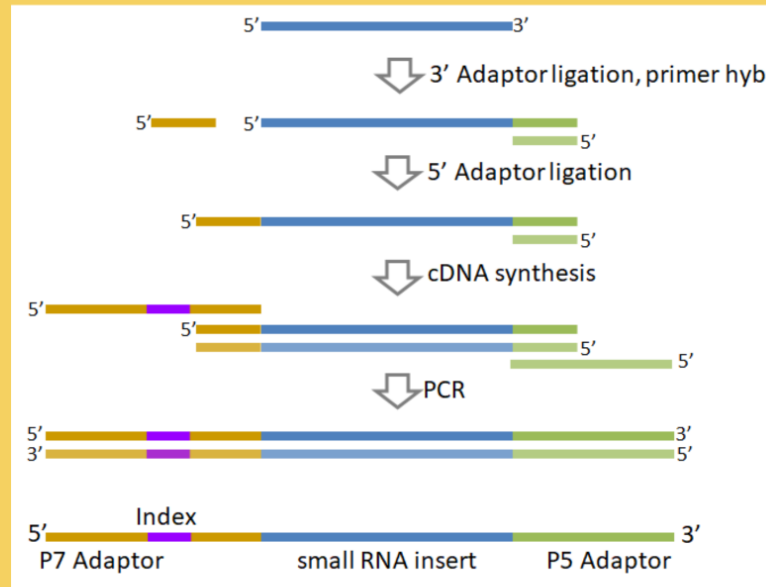


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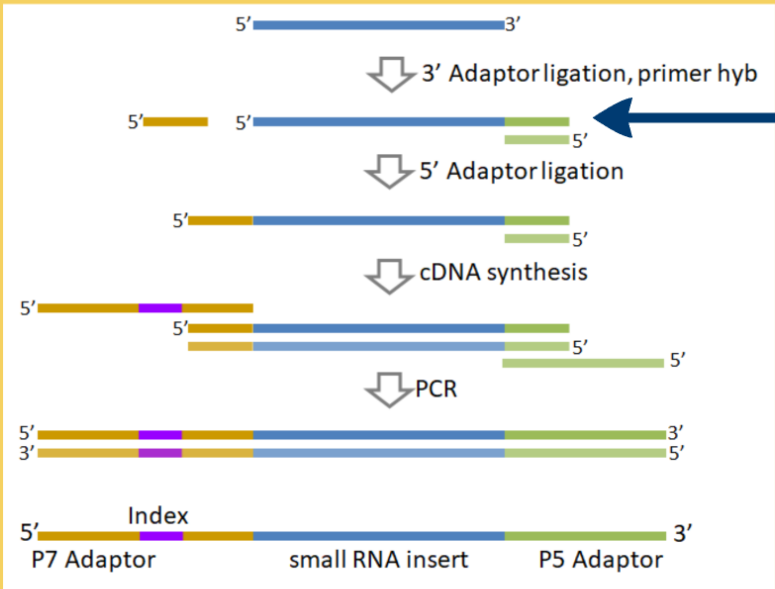
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# Small RNA Seq



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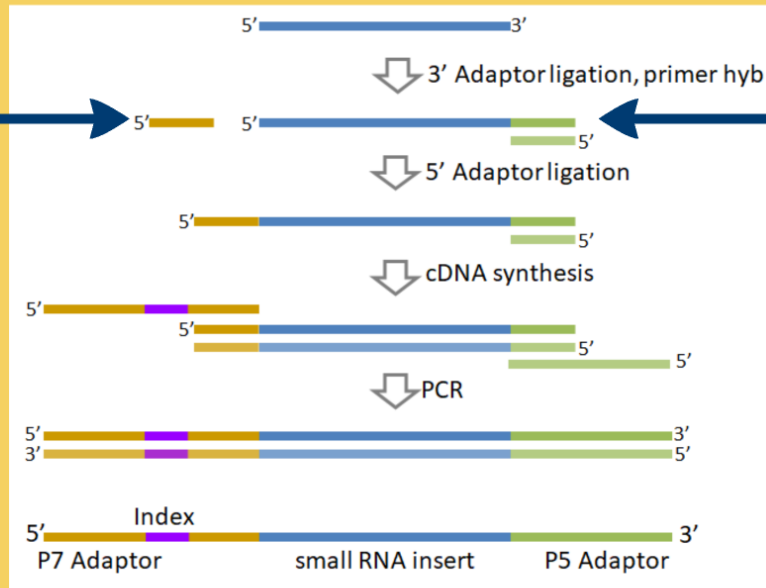


Hydroxyl Group

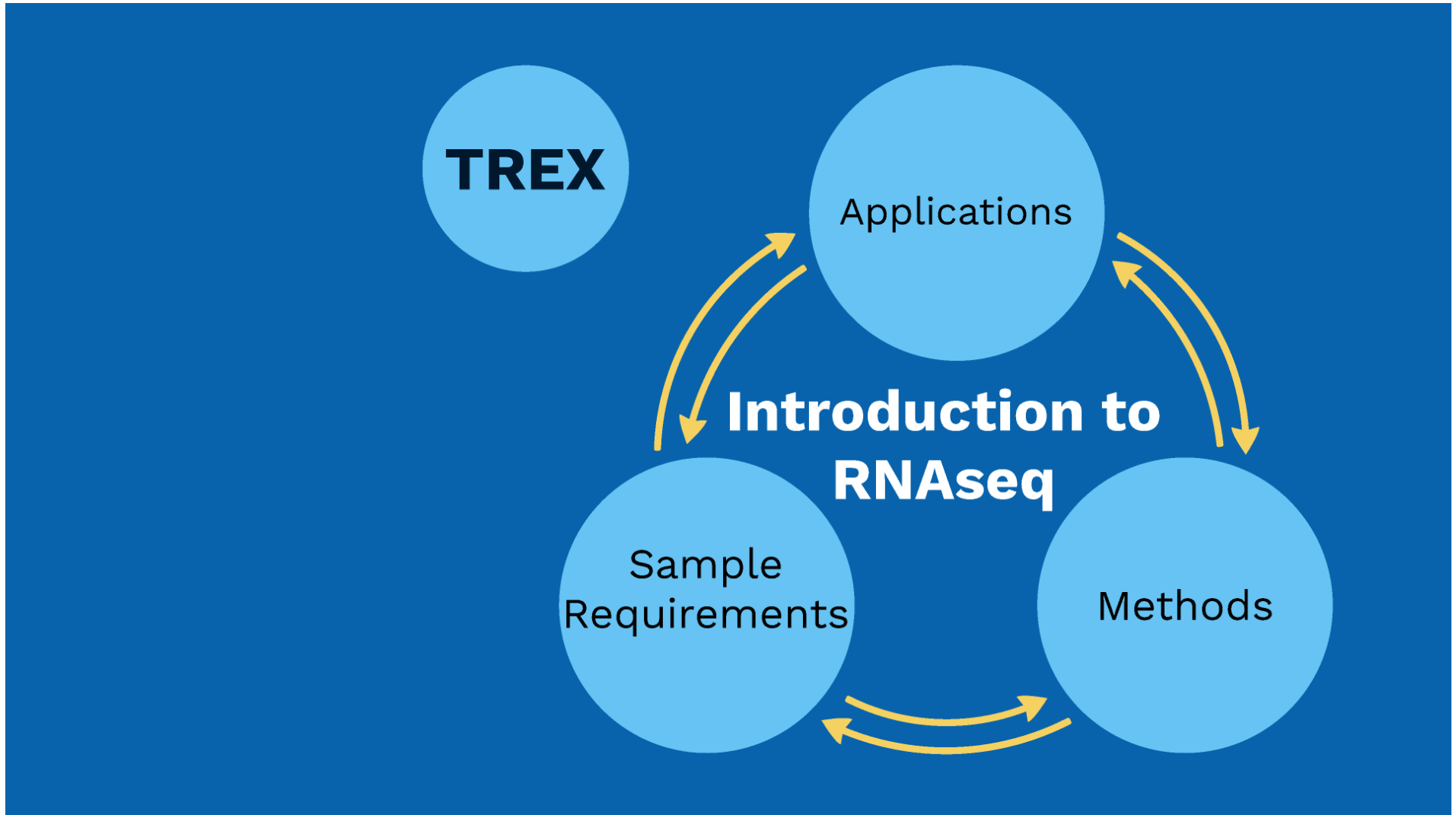
# Small RNA Seq

Mono-  
Phosphate

Hydroxyl  
Group







# Transcriptional Regulation and Gene Expression

**Mission:** Develop and provide high quality, project-oriented genomics services to the Cornell research community.

**Goal:** Enable successful research, from funding to publication

Jen Grenier

Chrissy Butler

Faraz Ahmed

Ann Tate

