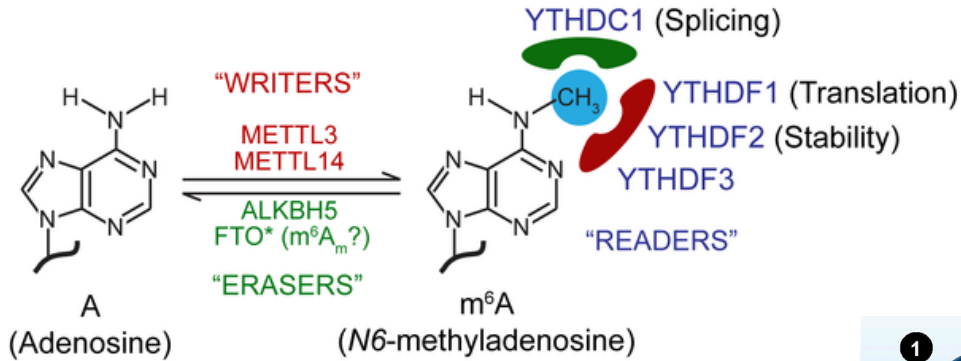


# RNA modifications: methodology and examples in disease

Evgenia Ntini  
evgenia.ntini@imbb.forth.gr  
RNA & Gene Regulation  
IMBB-FoRTH

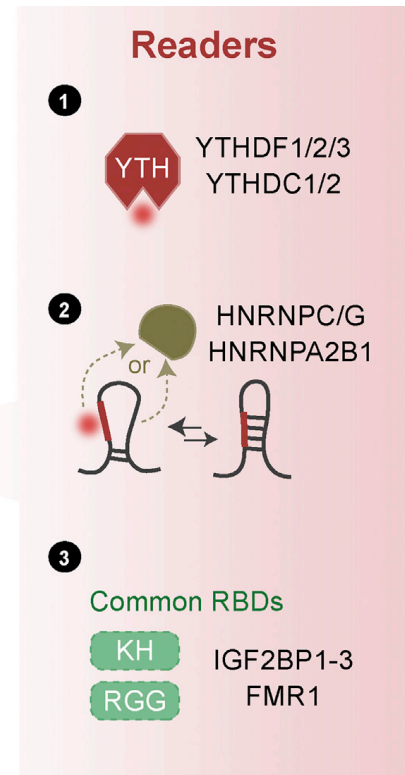
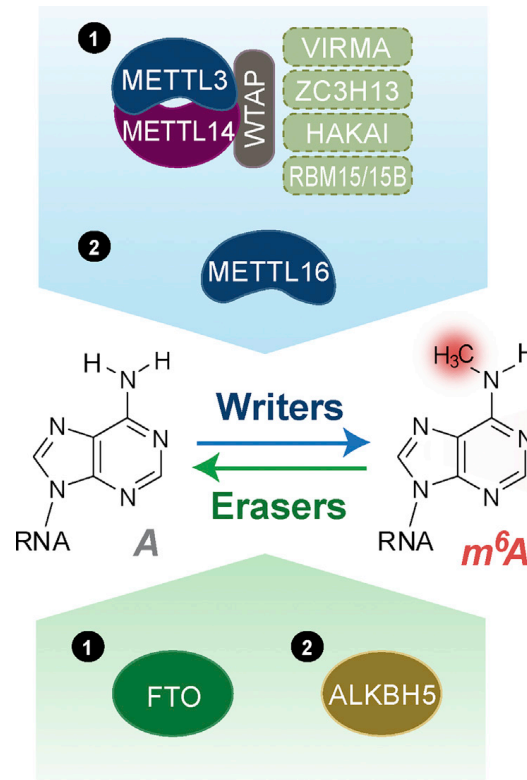
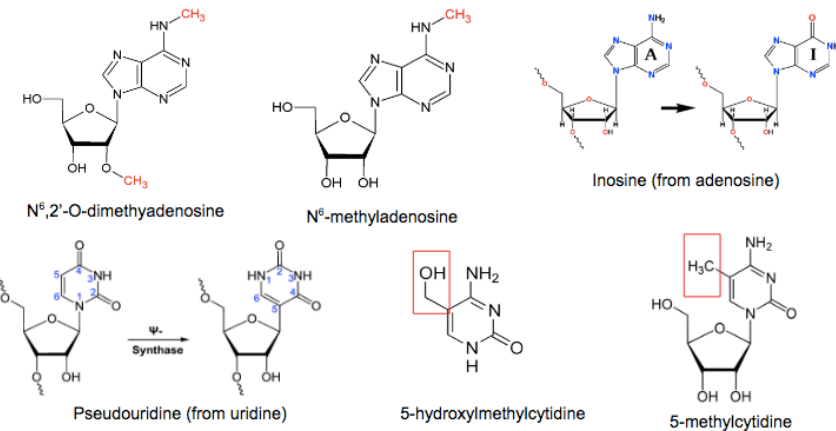


# RNA modifications: dynamic and reversible

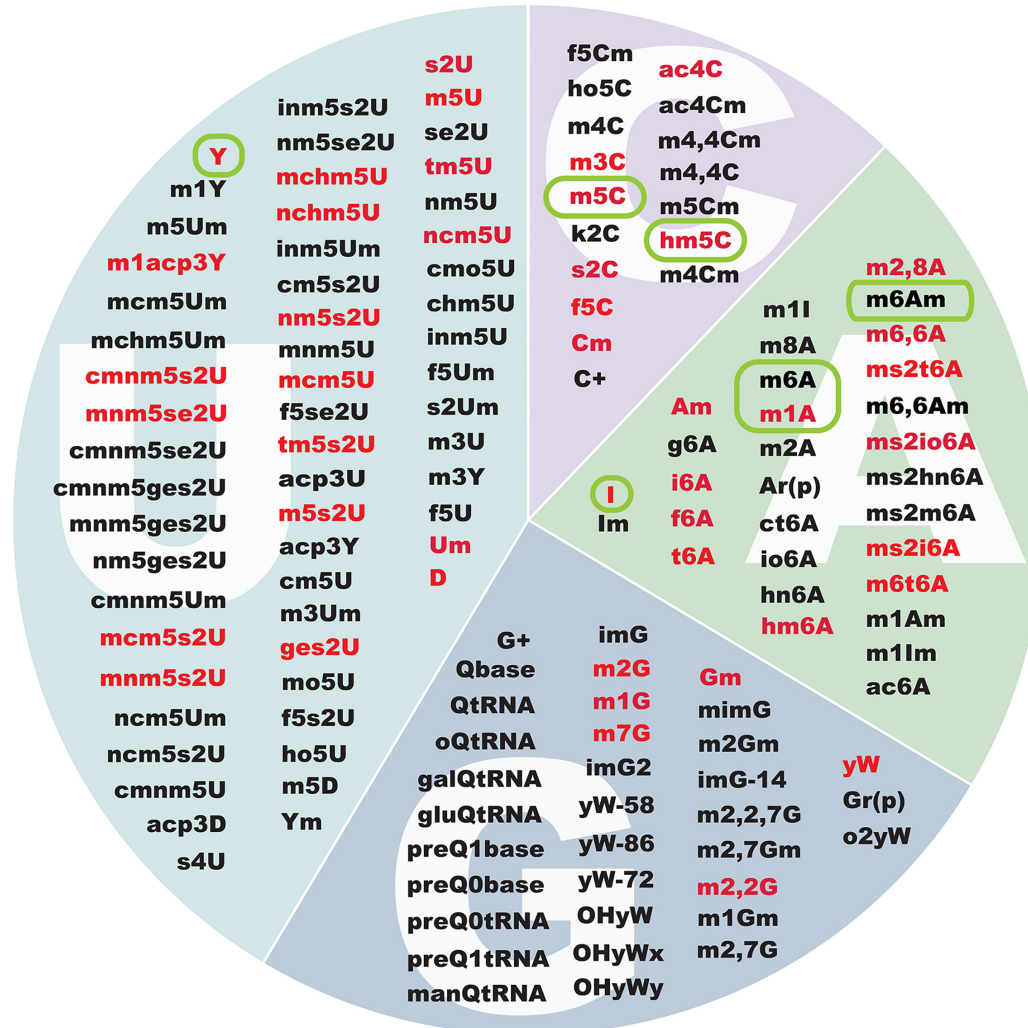
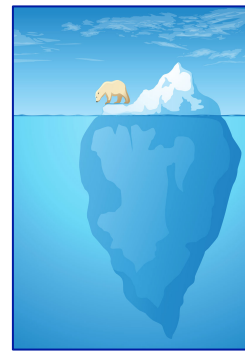


Gokhale and Horner SM, PLOS Pathogens 2017

Shi, Wei and Chuan He, Mol Cell 2019



# RNA modifications: dynamic and reversible



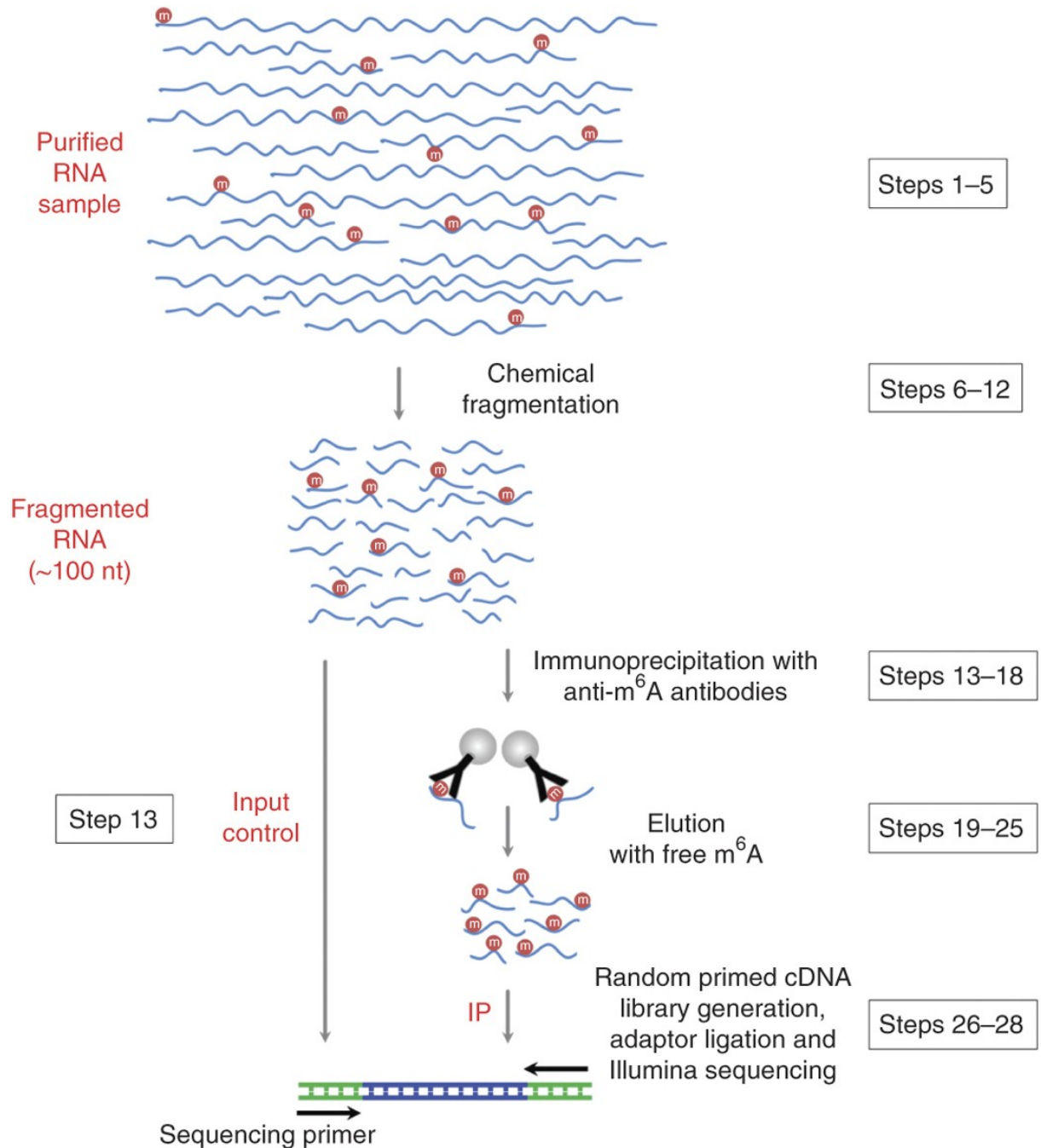
Disease related

Detection technique available

## Me-RIP/m6A-seq

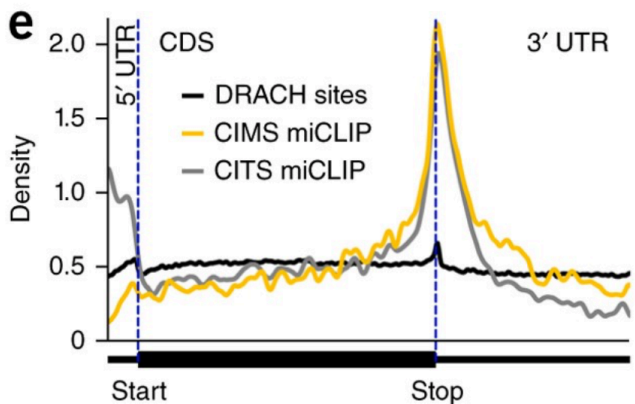
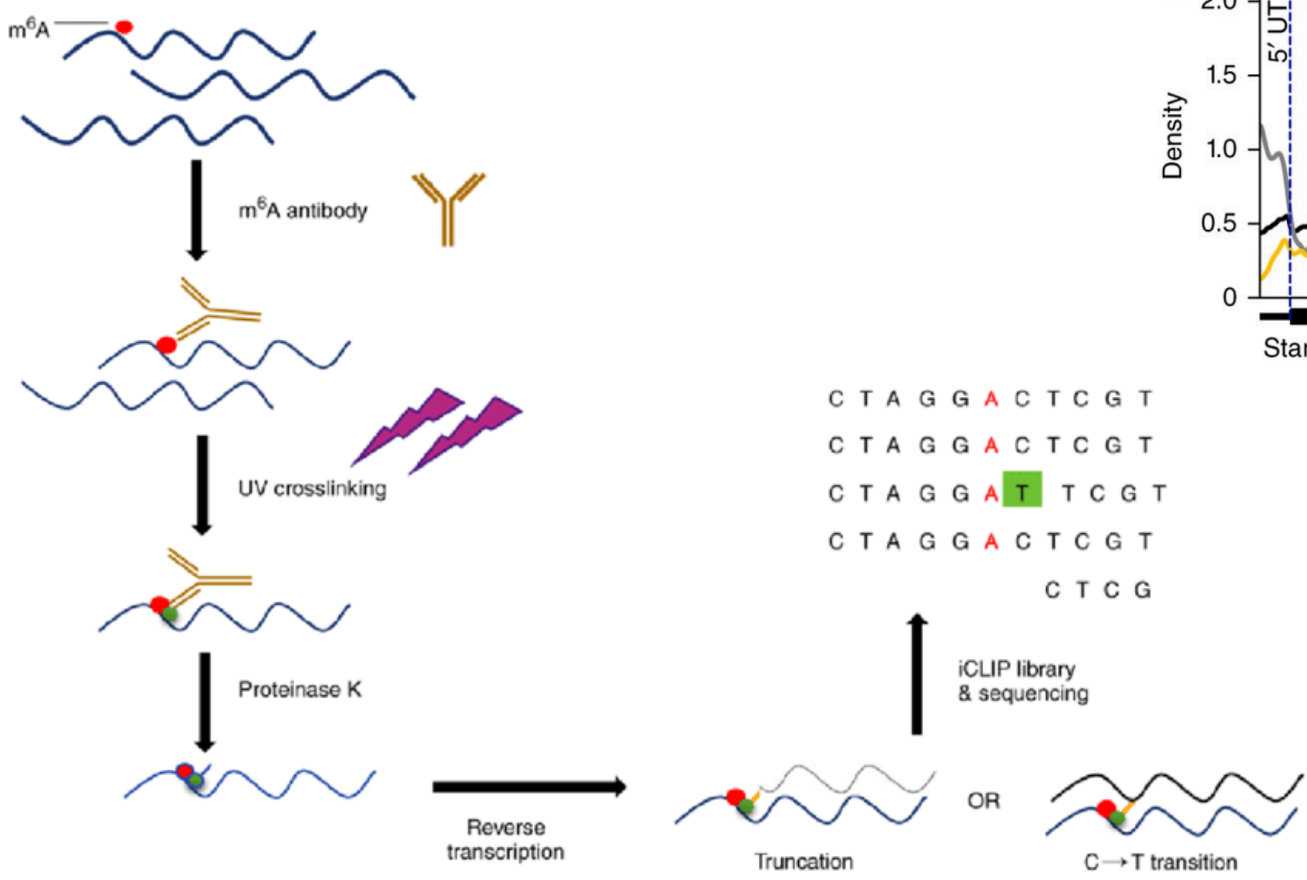
- MeRIP-Seq: Meyer K. D., Saletore Y., Zumbo P., Elemento O., Mason C. E., et al. (2012) Comprehensive analysis of mRNA methylation reveals enrichment in 3' UTRs and near stop codons. *Cell* 149: 1635-1646
- m6A-Seq: Dominissini D. et al. (2012) Topology of the human and mouse m6A RNA methylomes revealed by m6A-seq. *Nature* 485: 201-206
- Dominissini D. et al. (2013) Transcriptome-wide mapping of N6-methyladenosine by m6A-seq based on immunocapturing and massively parallel sequencing

# Me-RIP/m6A-seq



# miCLIP –m6A

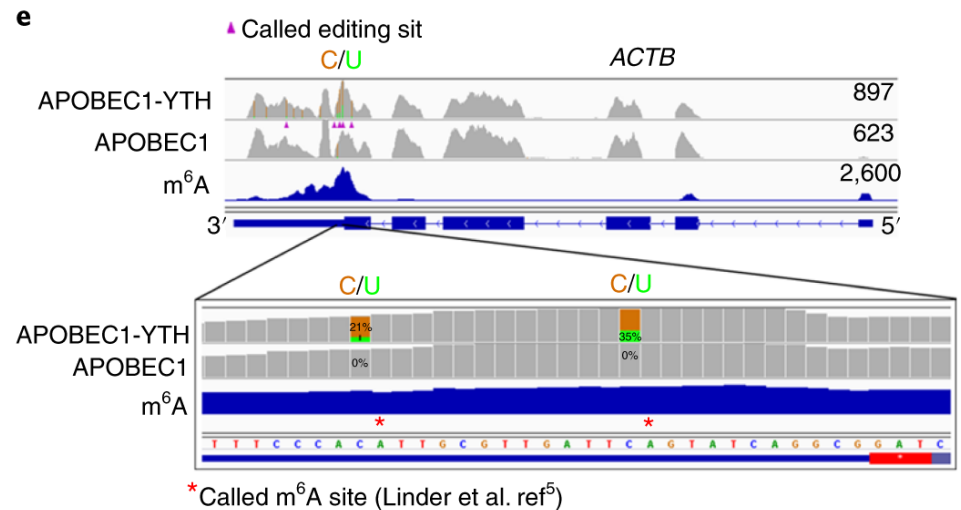
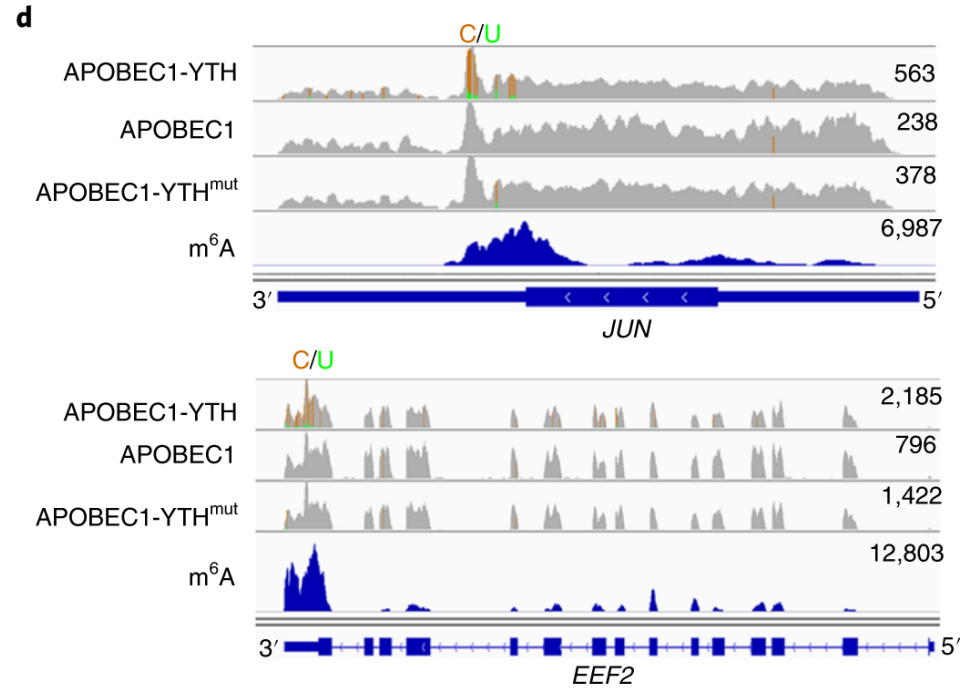
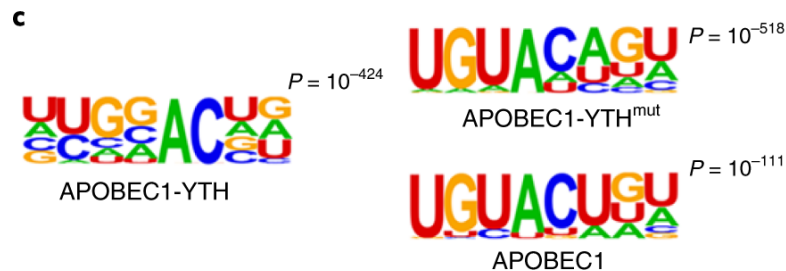
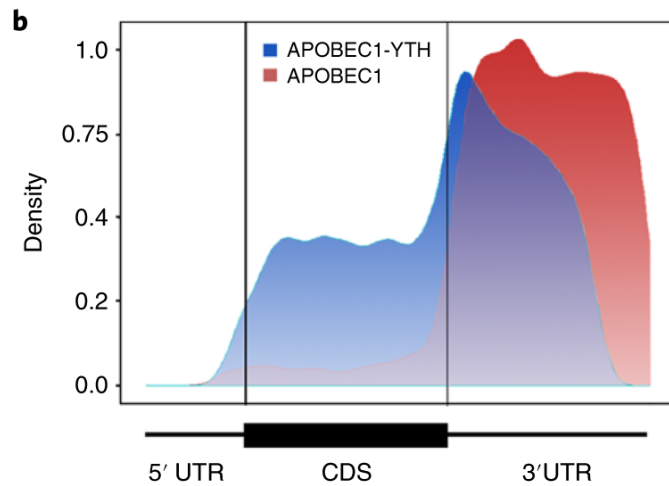
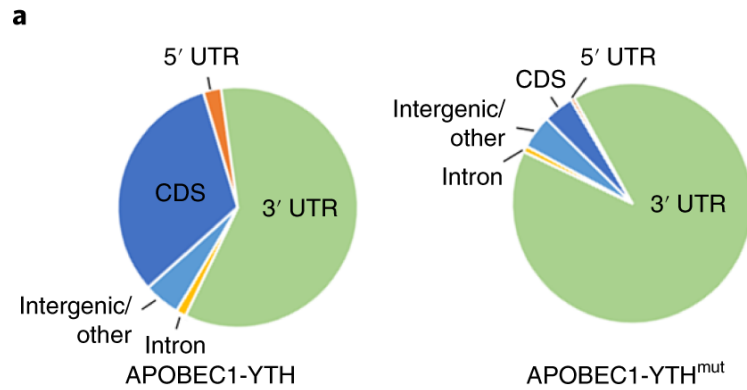
- miCLIP-m6A: Linder, et al. (2015) Single-nucleotide-resolution mapping of m6A and m6Am throughout the transcriptome. Nat Methods 12, 767–772 (2015)



# Antibody-free methods:

## Meyer KD. DART-seq: an antibody-free method for global m6A detection

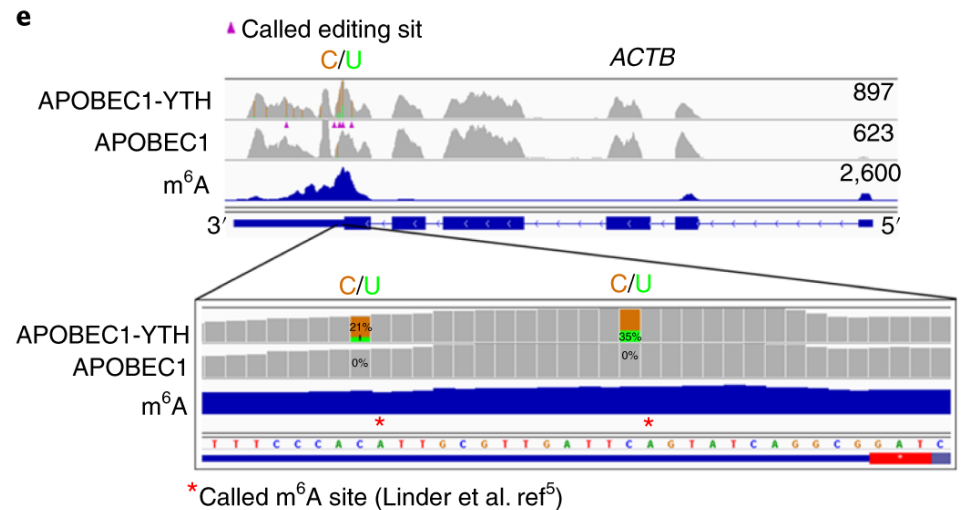
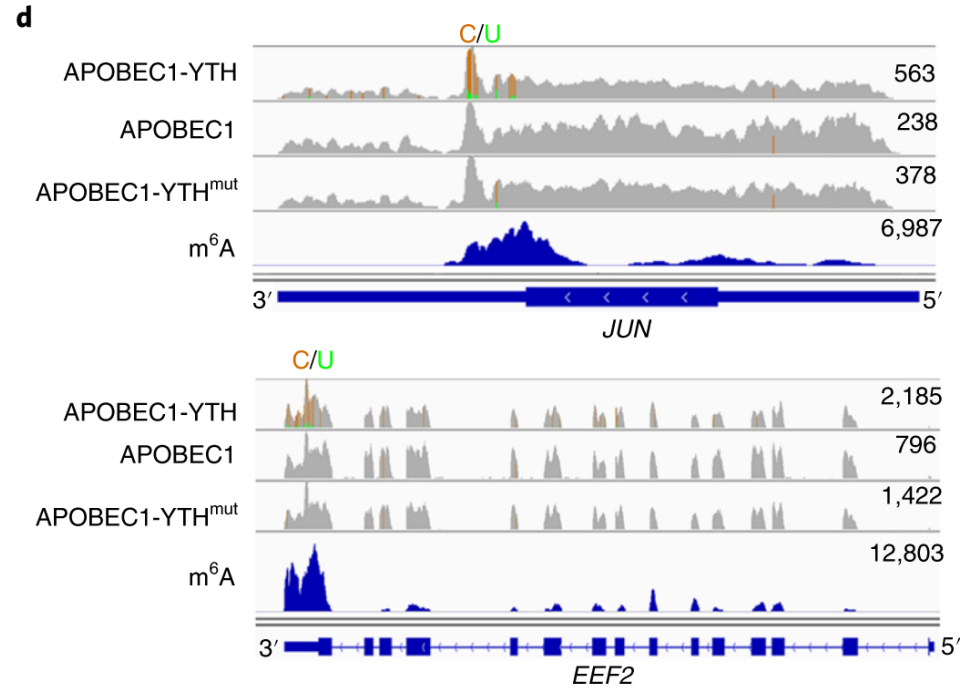
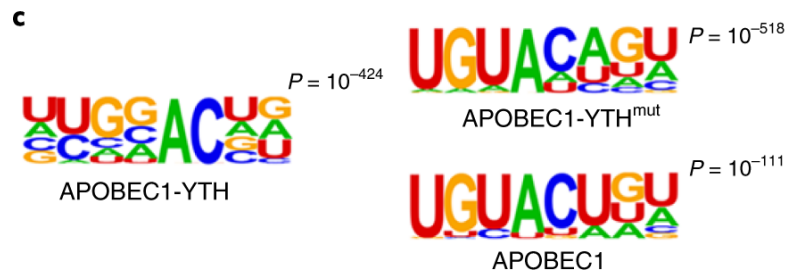
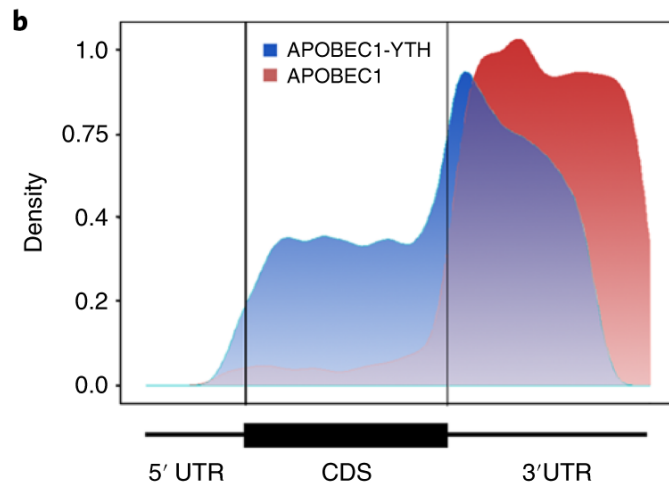
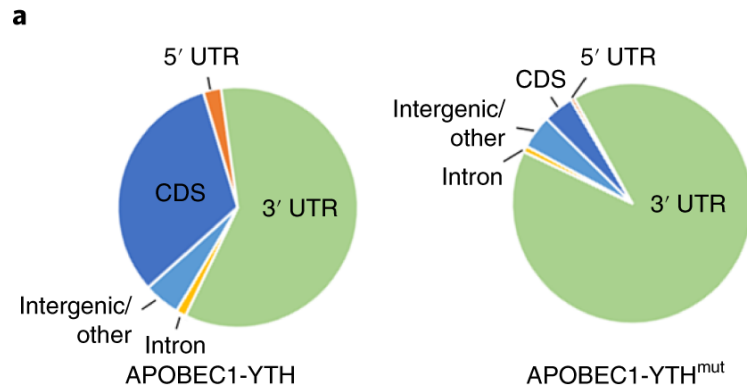
Nature Methods v 16, p 1275–1280 (2019)



# Antibody-free methods:

## Meyer KD. DART-seq: an antibody-free method for global m6A detection

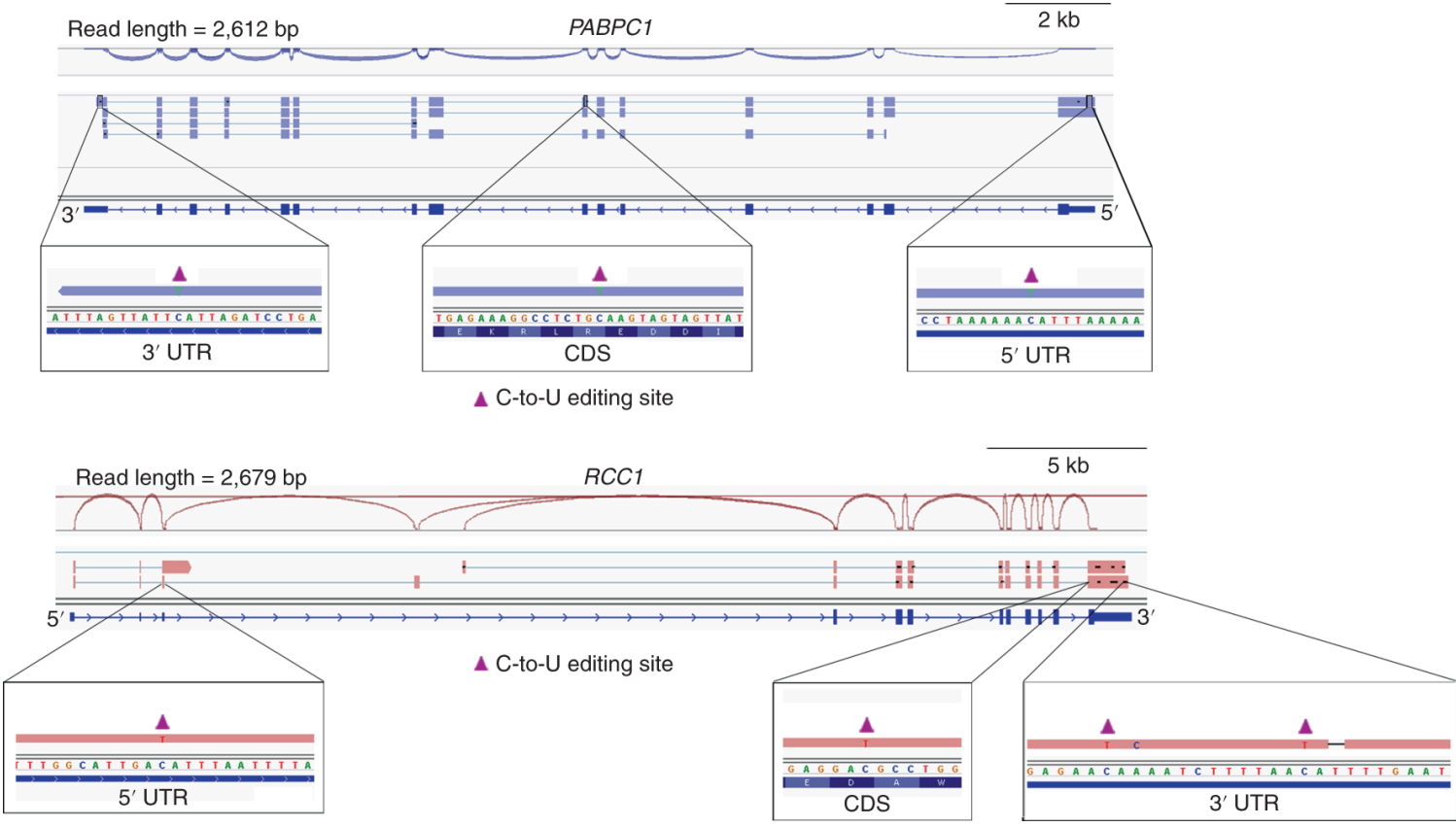
Nature Methods v 16, p 1275–1280 (2019)



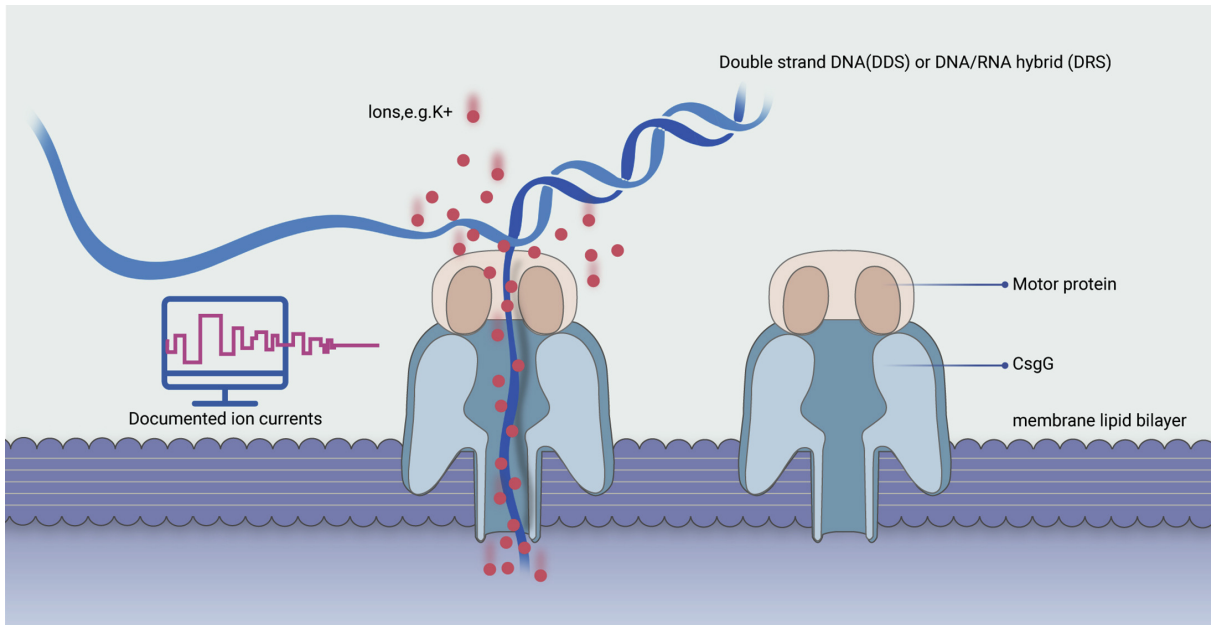


Meyer KD. DART-seq: an antibody-free method for global m6A detection  
Nature Methods v 16, p 1275–1280 (2019)

Long-read DART-seq (with PacBio) reveals m6A distribution within individual RNA molecules.



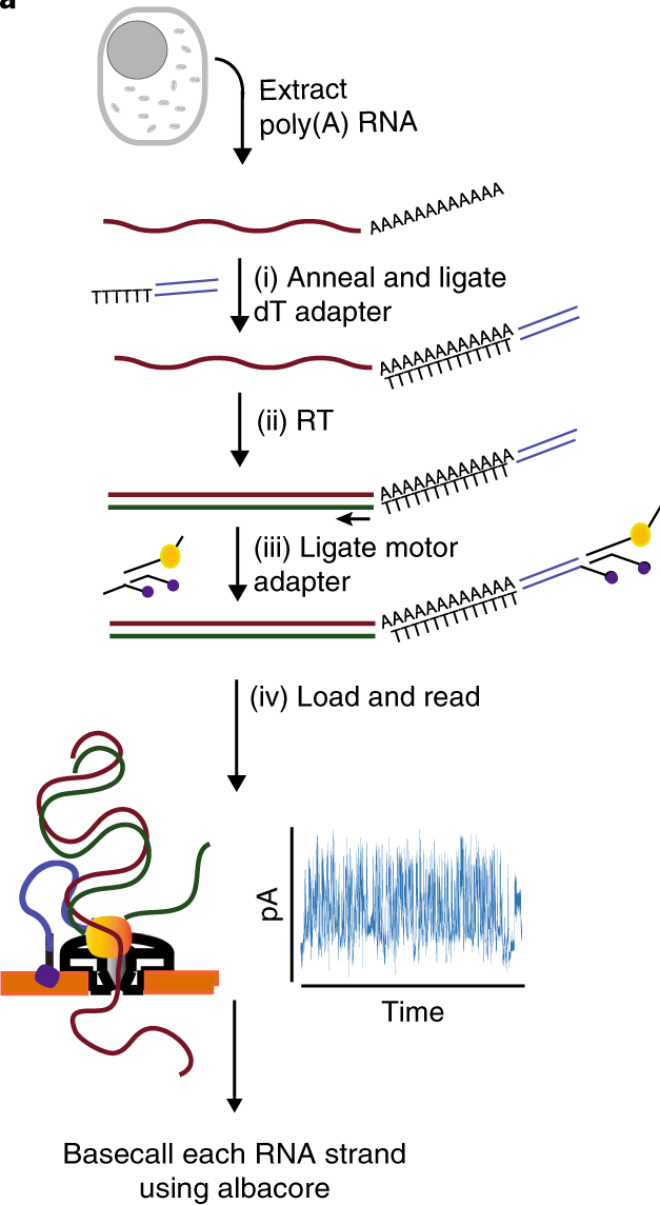
# Direct RNA sequencing with Oxford Nanopore Technology



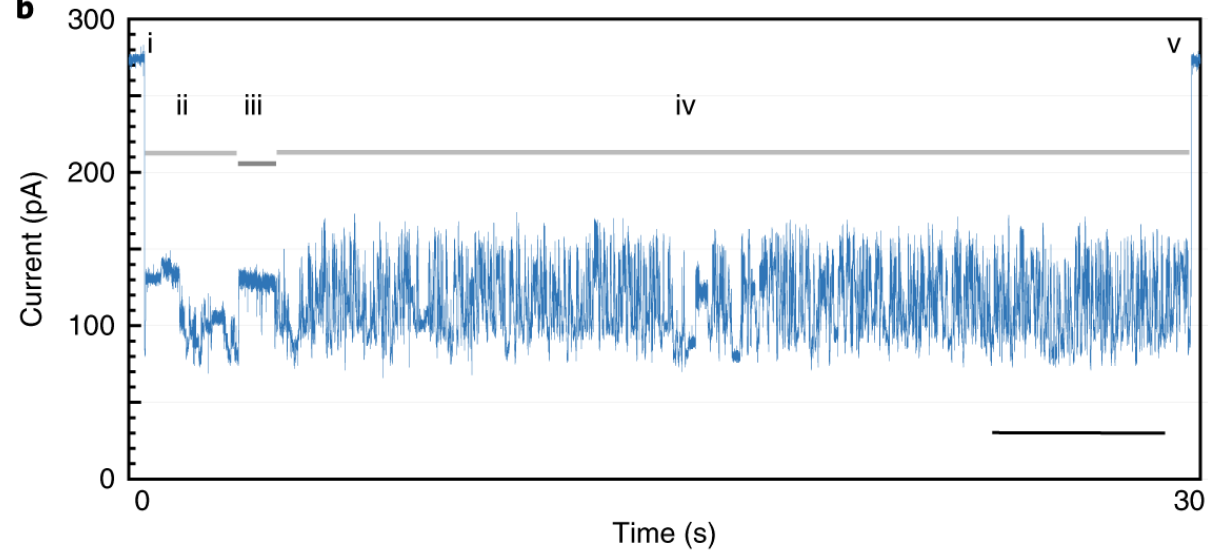
<https://doi.org/10.1016/j.xinn.2021.100153>

# Oxford Nanopore Direct RNA sequencing

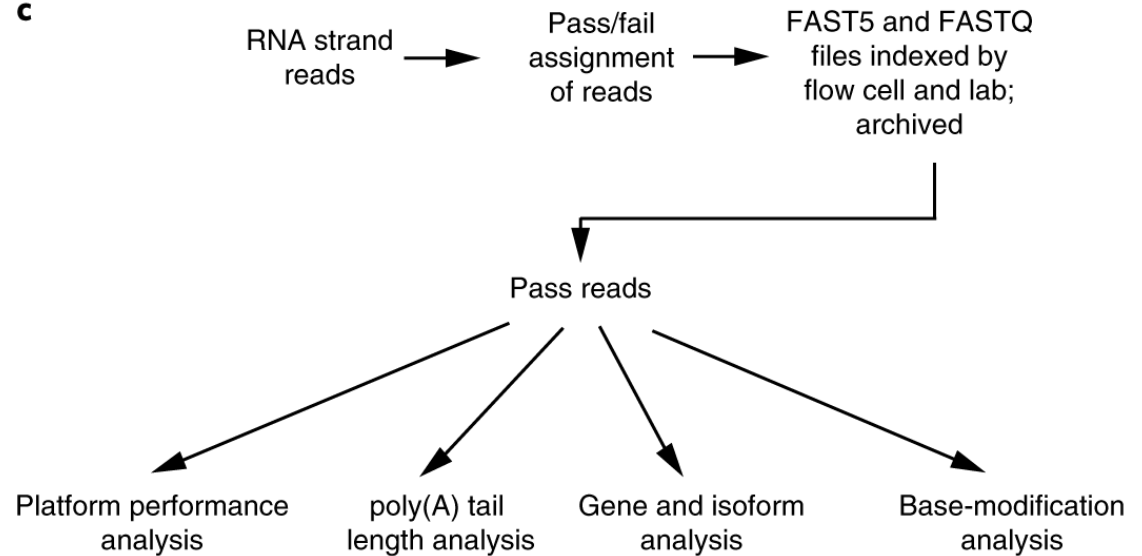
**a**



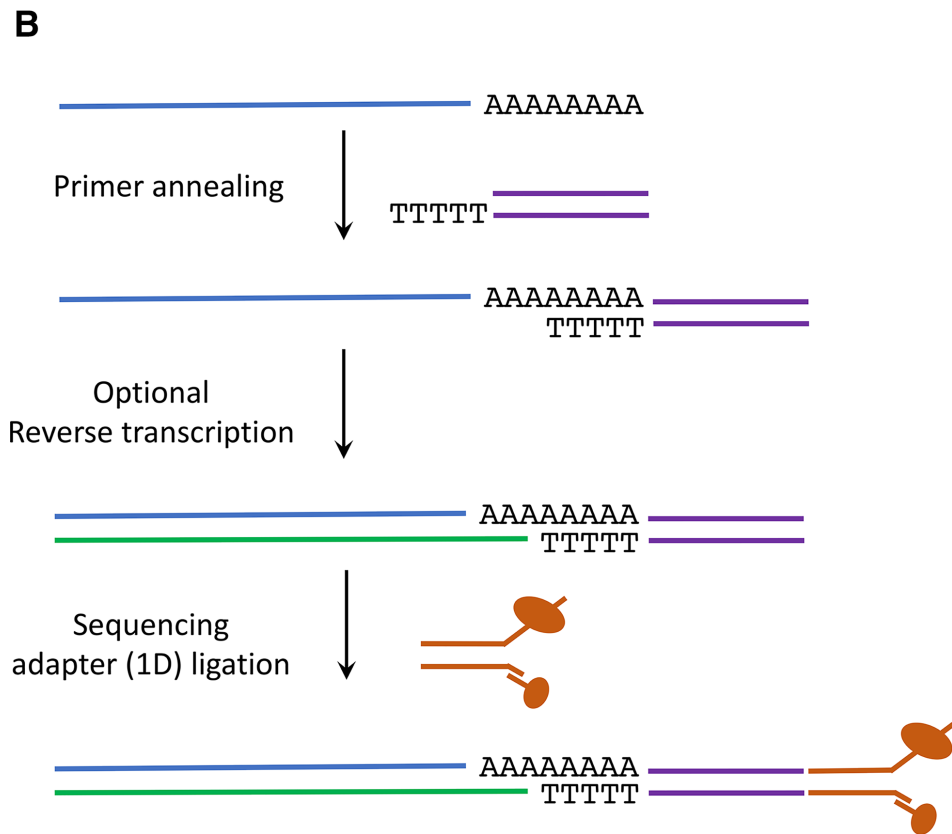
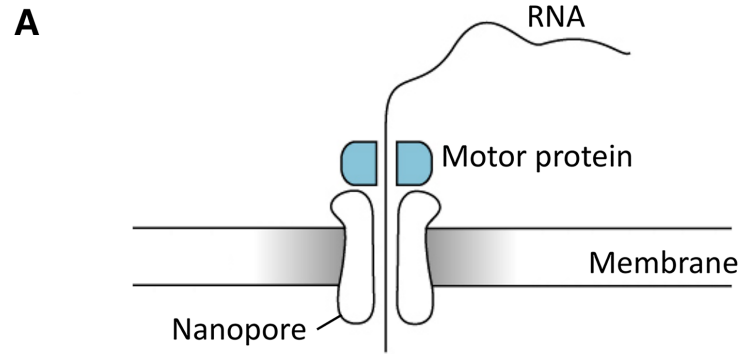
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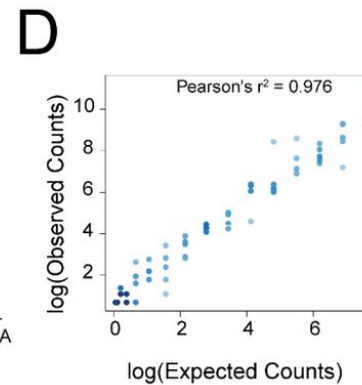
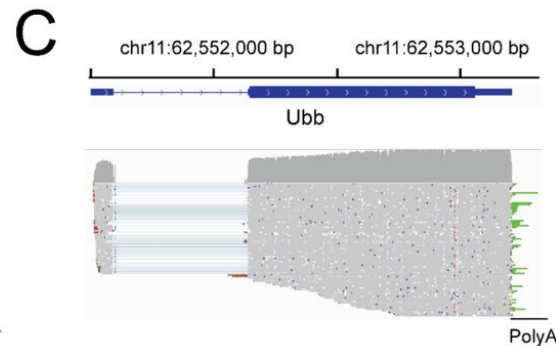
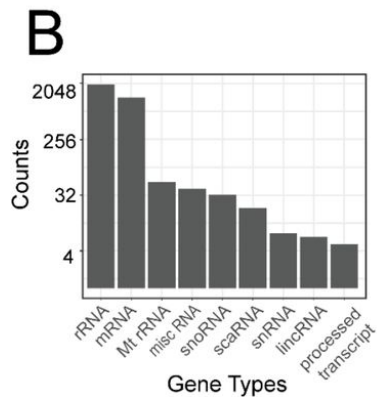
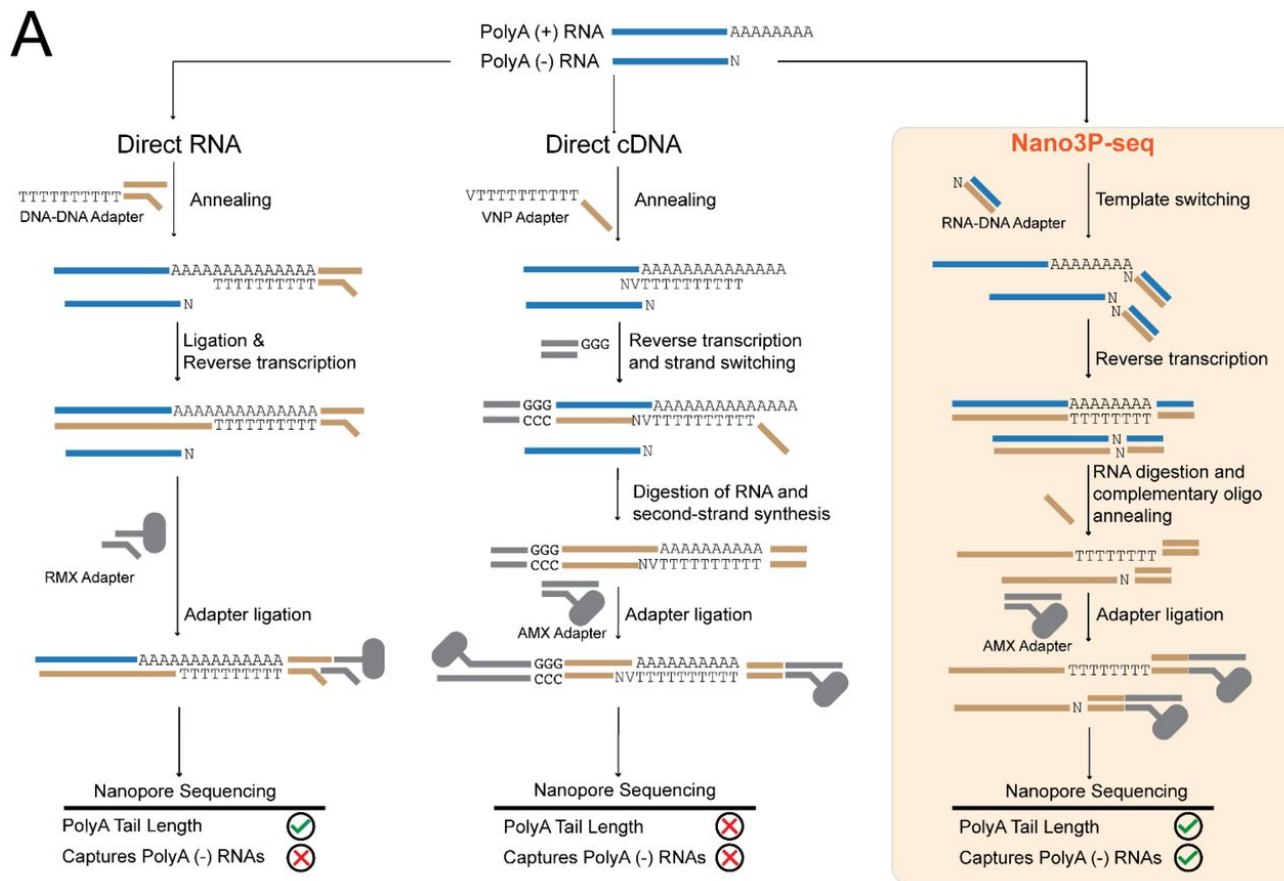


**c**

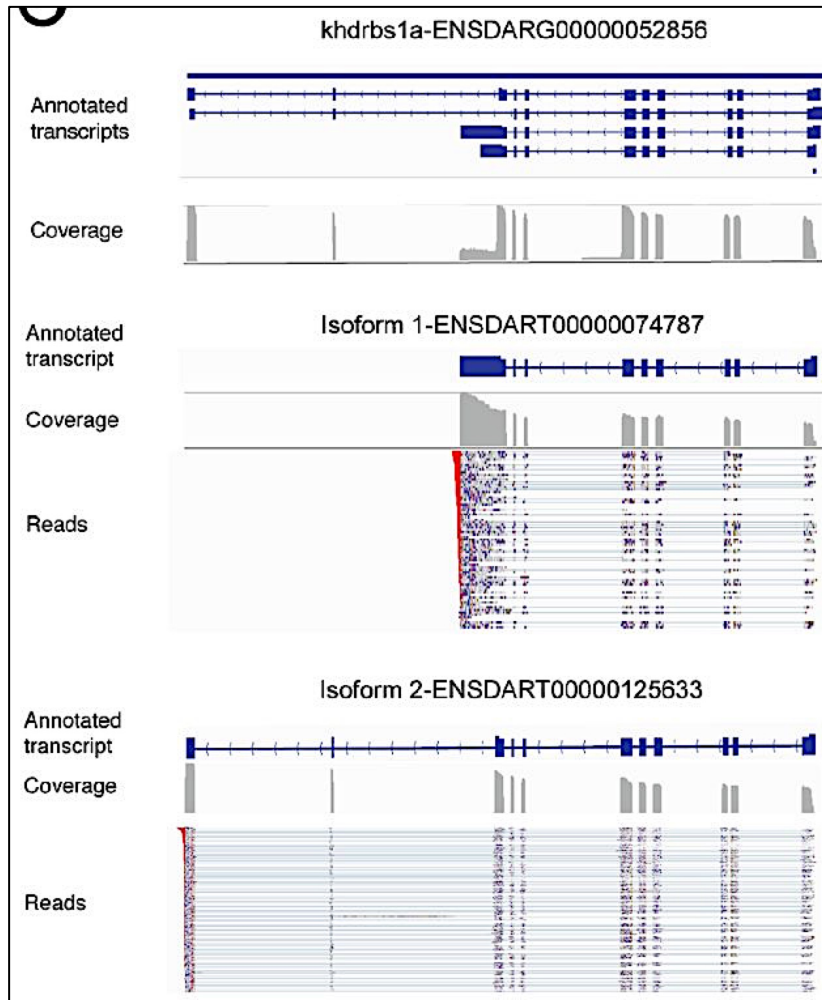


# Oxford Nanopore Direct RNA sequencing

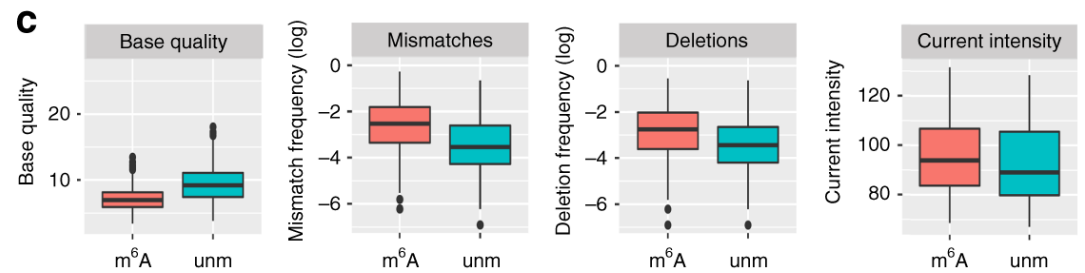
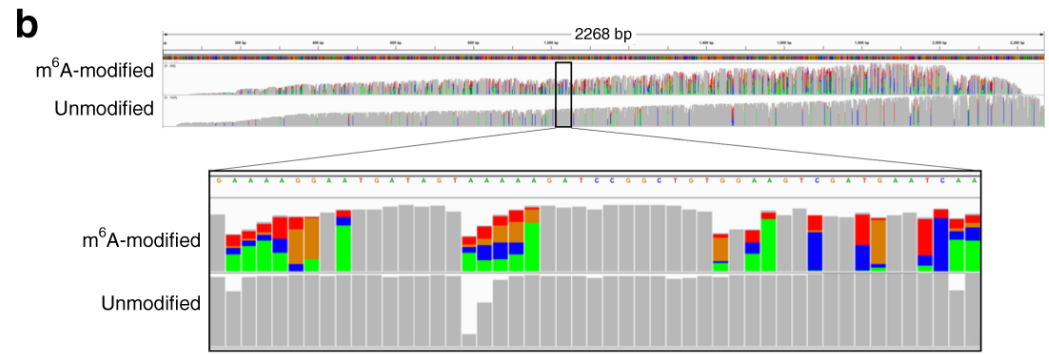
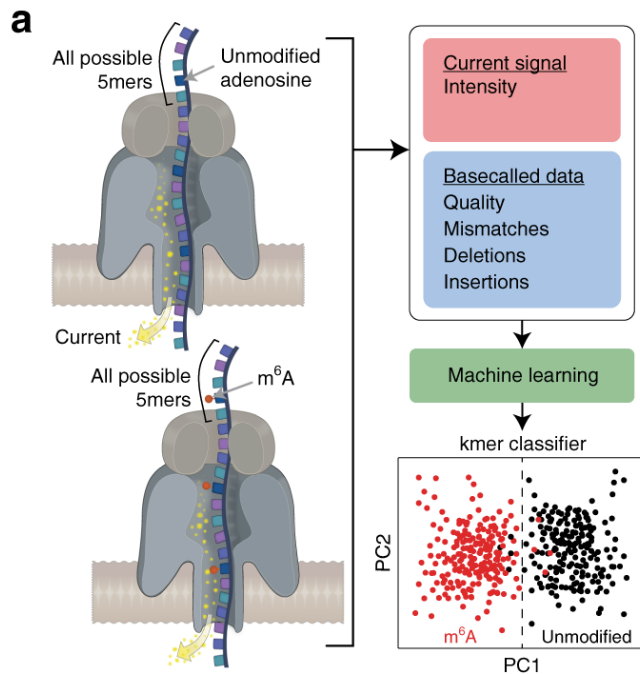




# ✓ Study isoform-specific polyA-tail dynamics

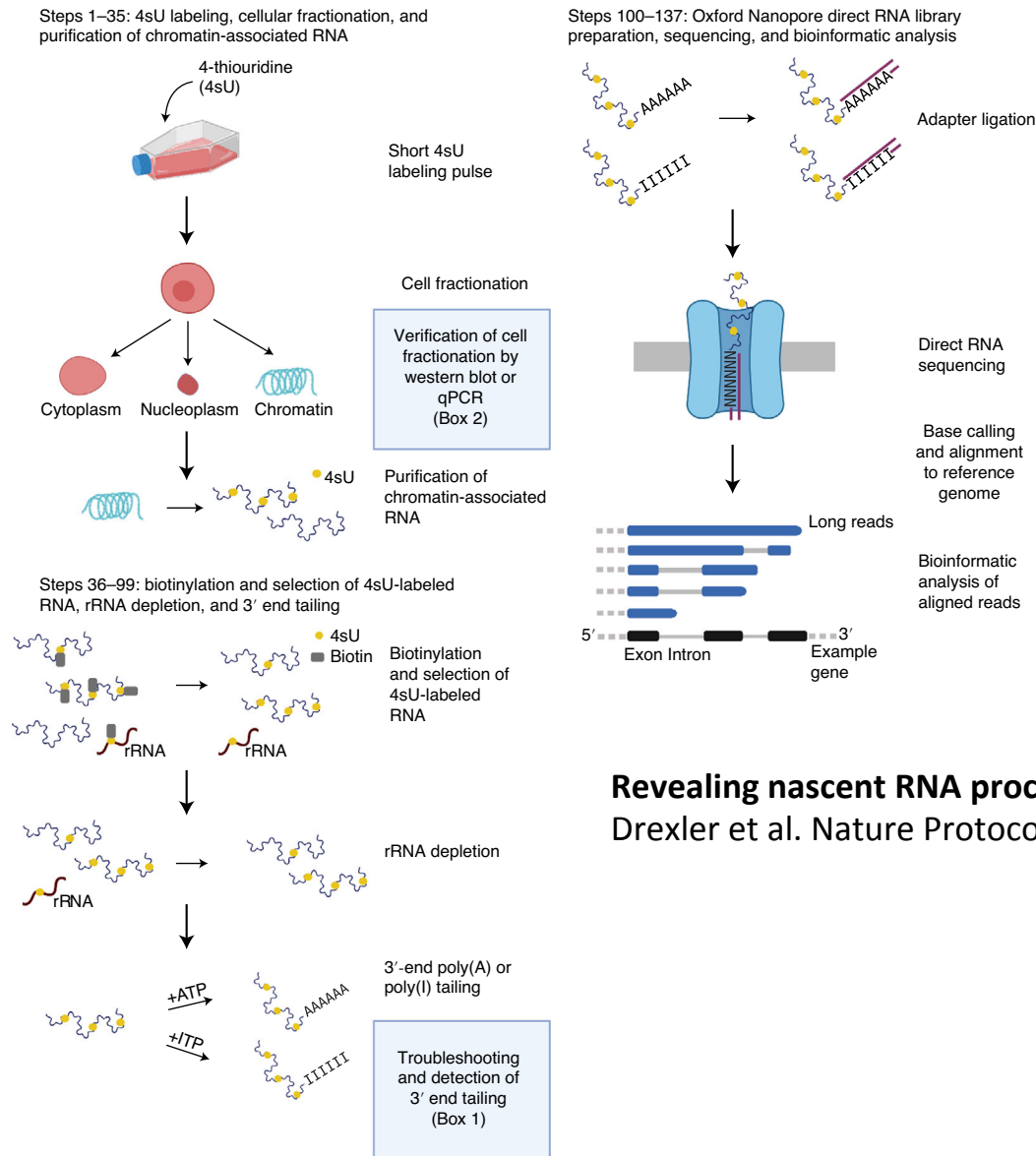


# ✓ Predict RNA modifications



**Accurate detection of m<sup>6</sup>A RNA modifications in native RNA sequences**  
Liu et al. Nature Communications 10: 4079 (2019)

# Long-read sequencing of nascent RNA (combine long read-sequencing with ~chromatin fractionation and ~metabolic labeling)

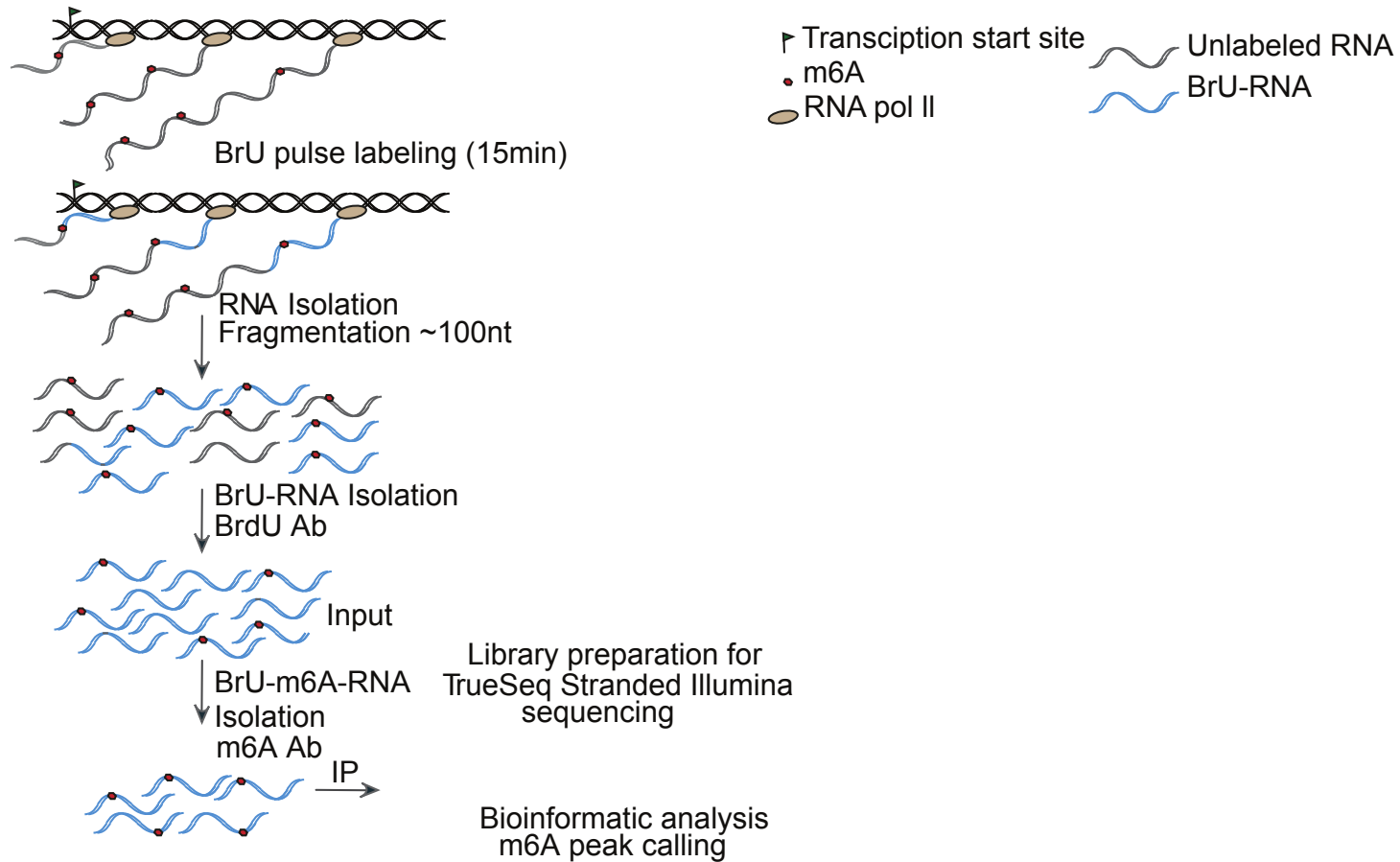


**Revealing nascent RNA processing dynamics with nano-COP**  
 Drexler et al. Nature Protocols 16, 1343–1375 (2021)

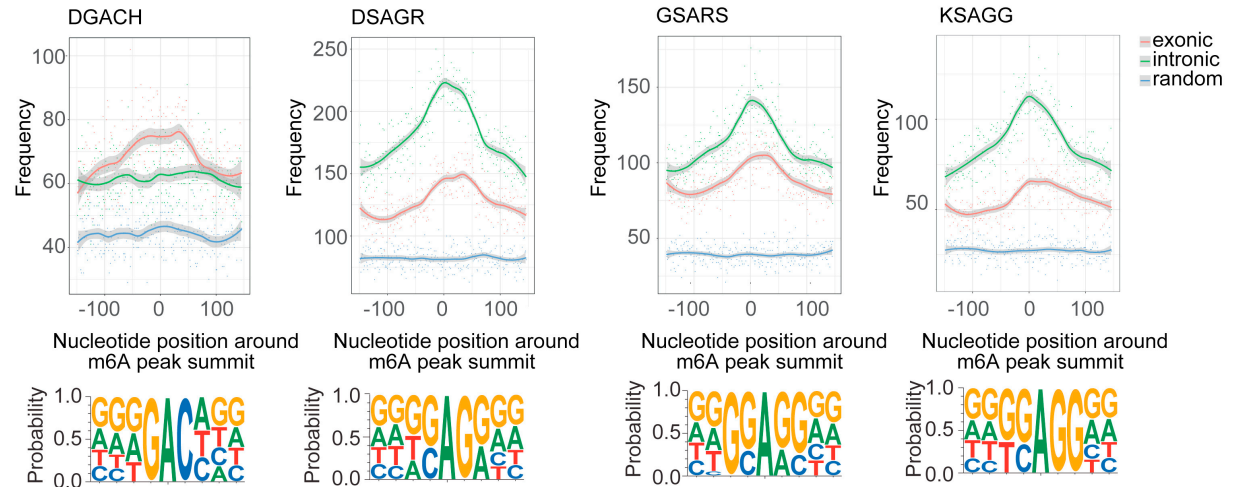
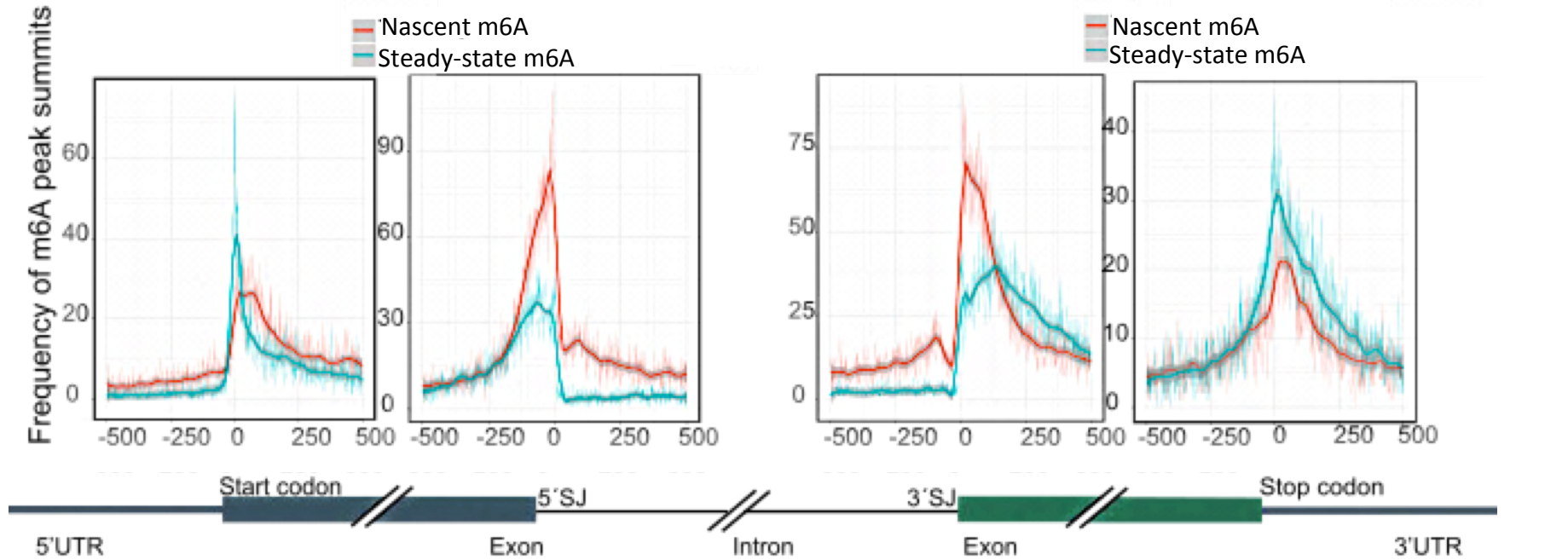


# Functions of co-transcriptional RNA modifications

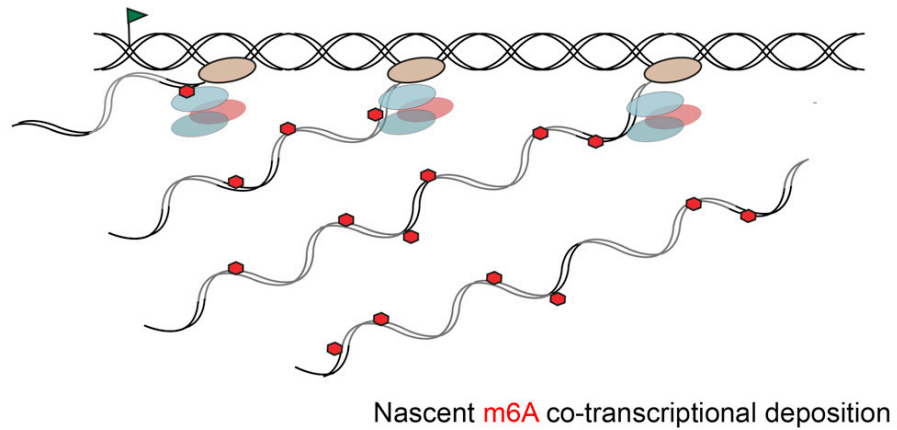
Capturing **nascent** N6-methyladenosine (m6A) RNA modifications:



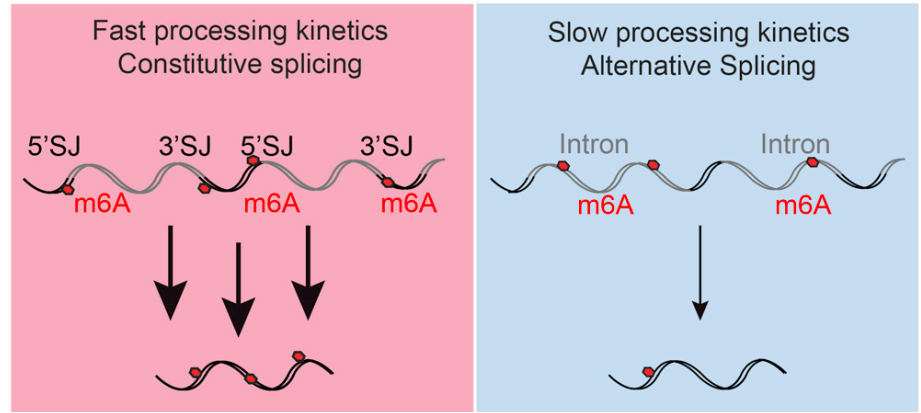
# Nascent m6A affects co-transcriptional RNA processing



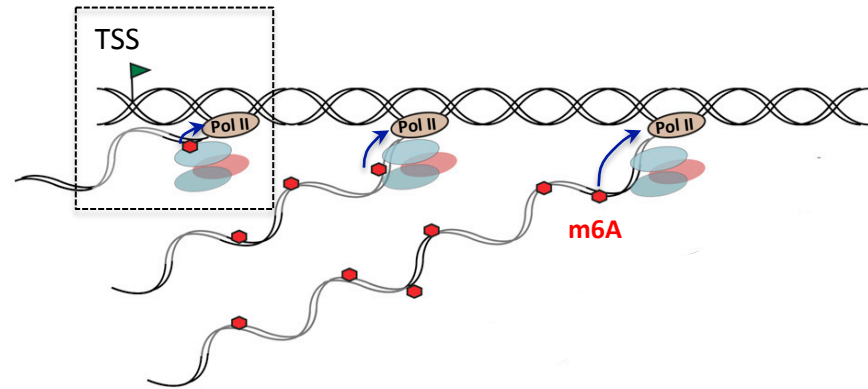
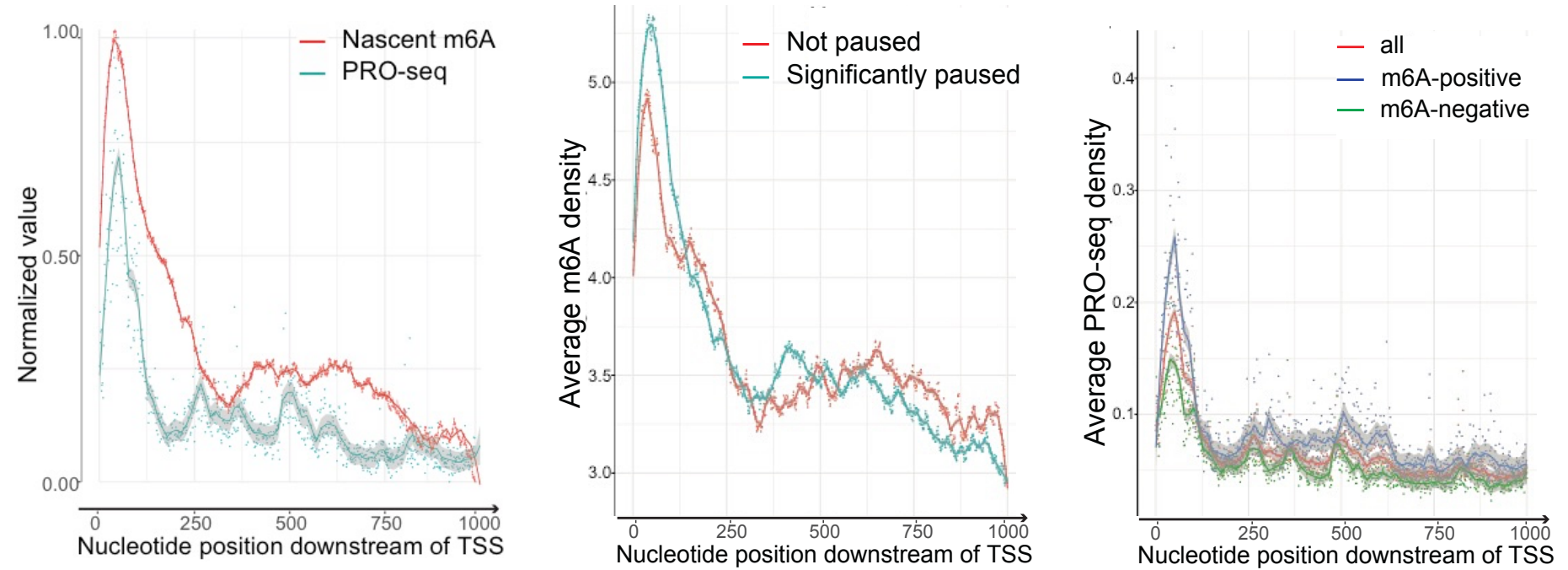
# Nascent RNA m6A modification affects co-transcriptional RNA processing



## Model:

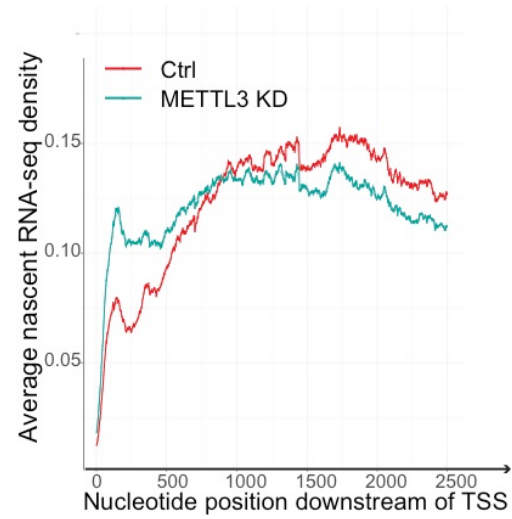
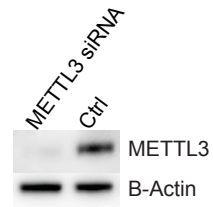
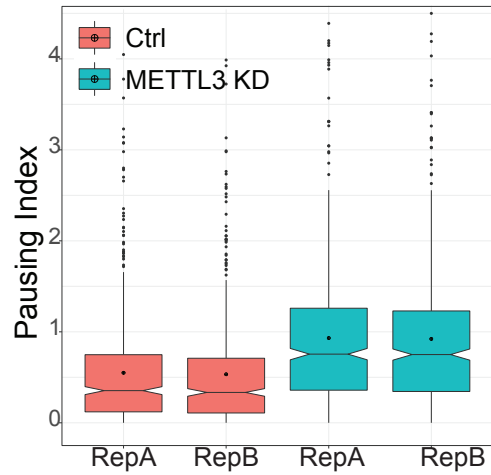


# Nascent RNA modifications in crosstalk with transcription



# Nascent m6A in crosstalk with transcription

Pausing Index: Read density [0\_+300]/[+300\_+3000]



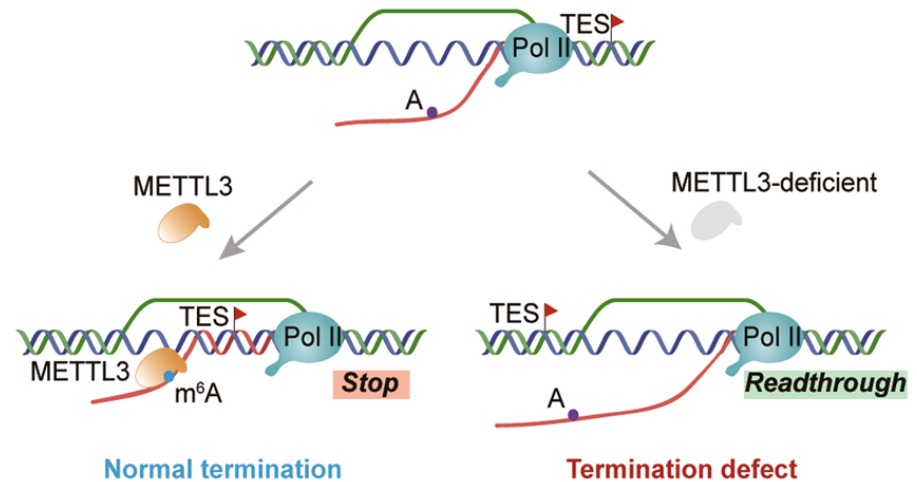
# Nascent m6A in crosstalk with transcription

## LETTER TO THE EDITOR

## m<sup>6</sup>A promotes R-loop formation to facilitate transcription termination

*Cell Research* (2019) 0:1–4; <https://doi.org/10.1038/s41422-019-0235-7>

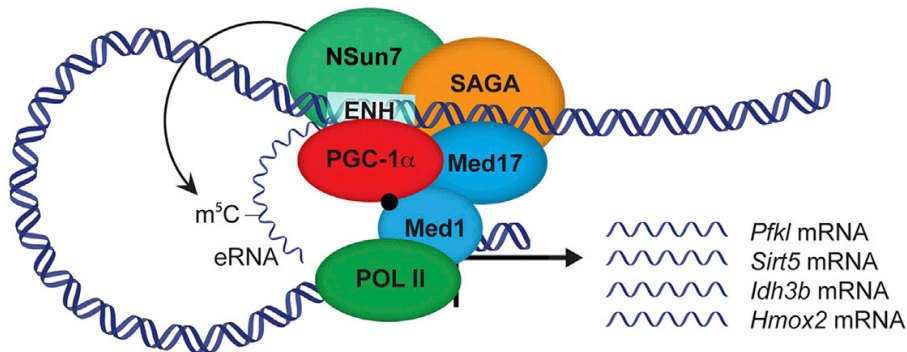
Yang et al. *Cell Res* 2019



## Cell Reports

### Deposition of 5-Methylcytosine on Enhancer RNAs Enables the Coactivator Function of PGC-1 $\alpha$

Aguilo et al., Cell Rep 2016



- METTL3-dependent RNA m6A dysregulation contributes to neurodegeneration in Alzheimer's disease through aberrant cell cycle events  
Zhao et al., *Molecular Neurodegeneration* 16: 70 (2021)
- m6A mRNA methylation-directed myeloid cell activation controls progression of NAFLD and obesity (in mice)  
Kin et al., *Cell Rep* 2021, 37:6 doi.org/10.1016/j.celrep.2021.109968

Review: Epitranscriptomics in liver disease: Basic concepts and therapeutic potential  
Zhao et al., *Journal of Hepatology* 2020





Yankova et al., Nature 2021 Small-molecule inhibition of METTL3 as a strategy against myeloid leukaemia, Nature 2021

*<https://www.stormtherapeutics.com>*