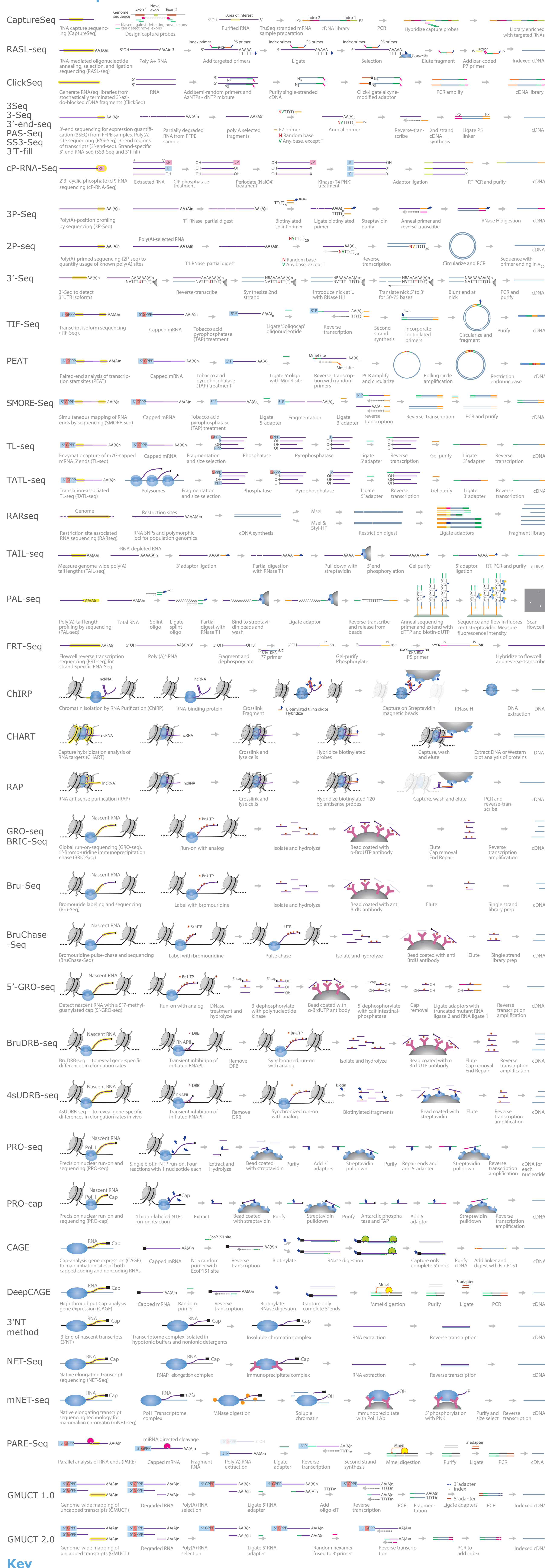


# For all you seq...

# RNA

## RNA Transcription

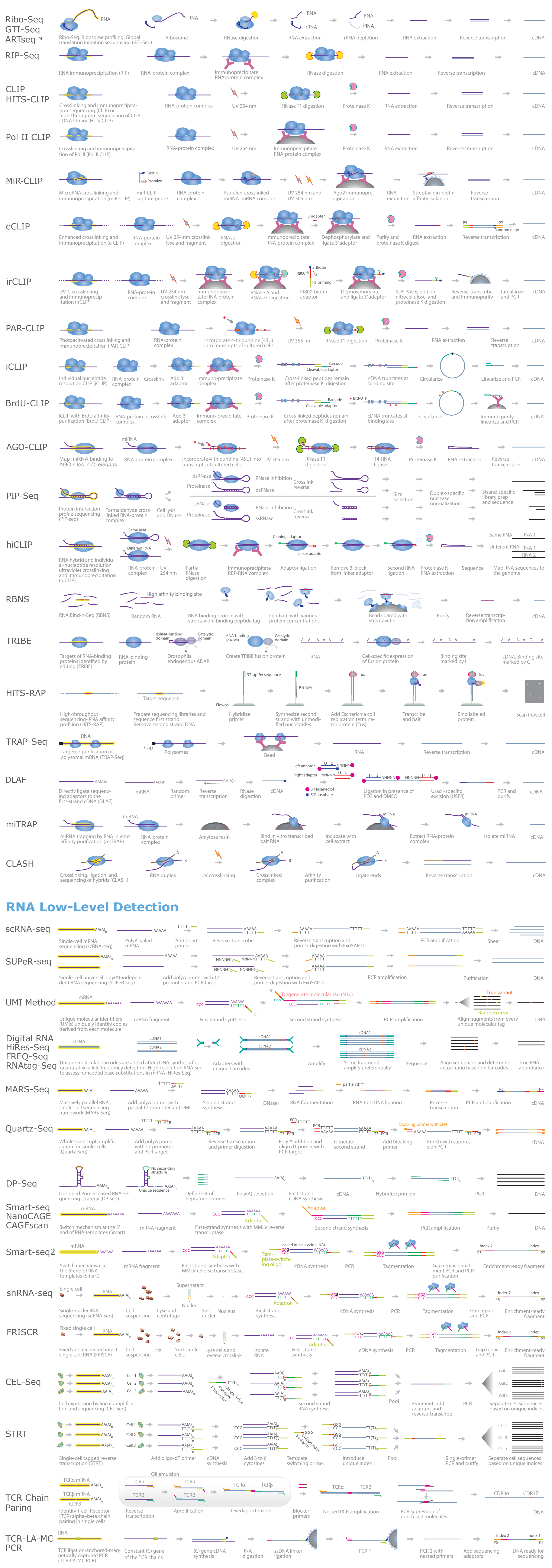


**Key**  
Yellow highlights indicate the target of the protocol

## References

35-Seq: Saito N. et al. (2012) Genome Res 22: 2076-2086  
 3-Seq: Hwang J. et al. (2012) Nature Methods 9: 639-643  
 3'-end-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 PAS-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 S3-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 3'-IT-1: Saito N. et al. (2012) Nature Methods 9: 639-643  
 CP-RNA-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 3P-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 2P-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 3-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 TIF-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 PEAT: Saito N. et al. (2012) Nature Methods 9: 639-643  
 SMORE-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 TI-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 TATL-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 RARseq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 TAIL-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 PAL-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 FRT-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 ChIRP: Saito N. et al. (2012) Nature Methods 9: 639-643  
 CHART: Saito N. et al. (2012) Nature Methods 9: 639-643  
 RAP: Saito N. et al. (2012) Nature Methods 9: 639-643  
 GRO-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 BRIC-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 Bru-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 BruChase-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 5'-GRO-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 BruDRB-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 4sUDBR-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 PRO-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 PRO-cap: Saito N. et al. (2012) Nature Methods 9: 639-643  
 CAGE: Saito N. et al. (2012) Nature Methods 9: 639-643  
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 3'NT method: Saito N. et al. (2012) Nature Methods 9: 639-643  
 NET-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 mNET-seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 PARE-Seq: Saito N. et al. (2012) Nature Methods 9: 639-643  
 GMUCT 1.0: Saito N. et al. (2012) Nature Methods 9: 639-643  
 GMUCT 2.0: Saito N. et al. (2012) Nature Methods 9: 639-643

## RNA-Protein Interactions



**Key**  
Yellow highlights indicate the target of the protocol

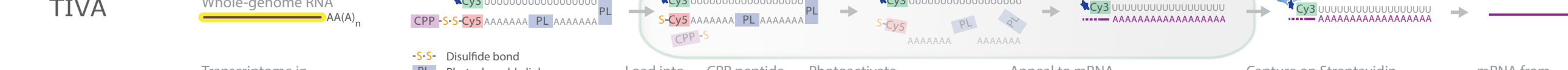
## References

Ribo-Seq: Ingolia N. et al. (2012) Science 326: 686-691  
 GTI-Seq: Ingolia N. et al. (2012) Science 326: 686-691  
 ART-seq: Ingolia N. et al. (2012) Science 326: 686-691  
 CLIP: He Y. et al. (2012) Nature Methods 9: 639-643  
 HITS-CLIP: He Y. et al. (2012) Nature Methods 9: 639-643  
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 PAR-CLIP: He Y. et al. (2012) Nature Methods 9: 639-643  
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 AGO-CLIP: He Y. et al. (2012) Nature Methods 9: 639-643  
 PIP-Seq: He Y. et al. (2012) Nature Methods 9: 639-643  
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 scRNA-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 SUPER-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
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 FREQ-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 RNAtag-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 MARS-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 Quartz-Seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 DP-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 Smart-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 NABC-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 CAGEscan: He Y. et al. (2012) Nature Methods 9: 639-643  
 Smart-seq2: He Y. et al. (2012) Nature Methods 9: 639-643  
 snRNA-seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 FRISCR: He Y. et al. (2012) Nature Methods 9: 639-643  
 CEL-Seq: He Y. et al. (2012) Nature Methods 9: 639-643  
 STRT: He Y. et al. (2012) Nature Methods 9: 639-643  
 PCR Chain: He Y. et al. (2012) Nature Methods 9: 639-643  
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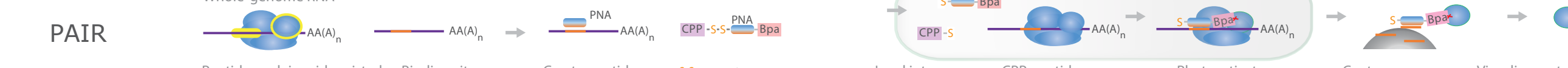
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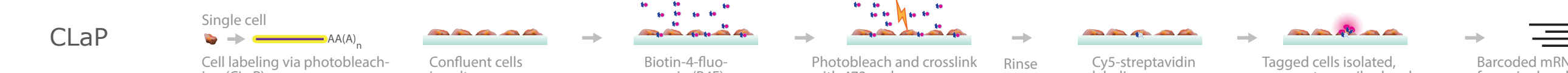
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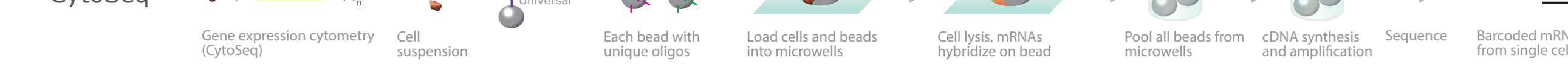
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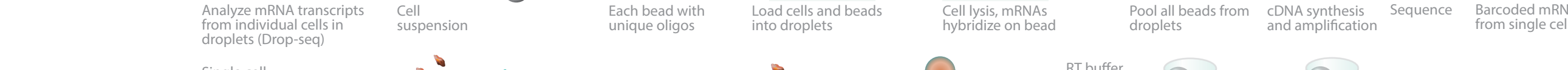
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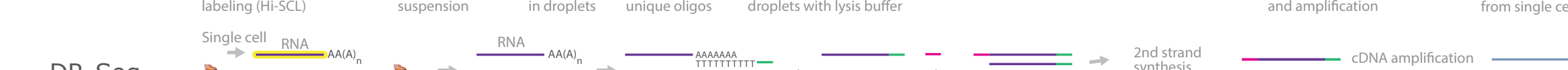
## CytoSeq



## Drop-Seq



## Hi-SCL



## DR-Seq



## inDrop



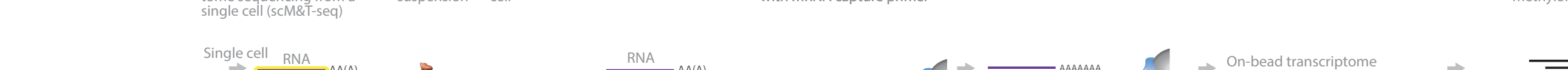
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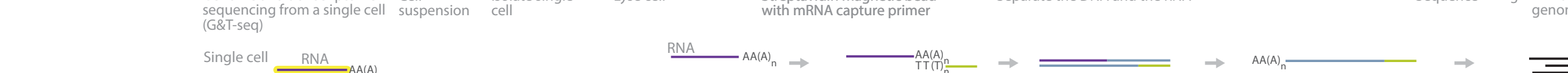
## Div-Seq



## SCRB-Seq



## scMT-Seq



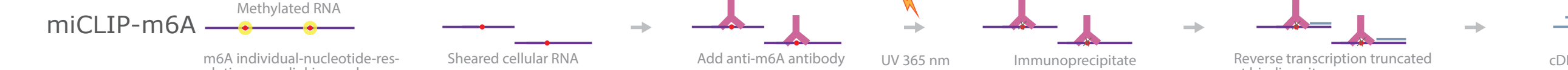
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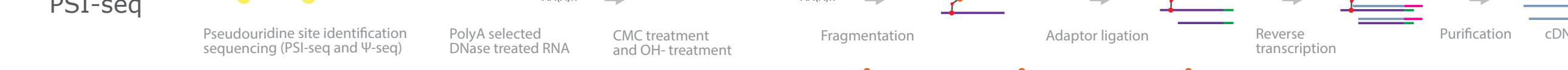
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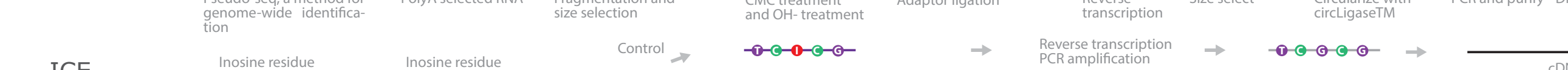
## RNA Modifications



## U-seq



## PSI-seq



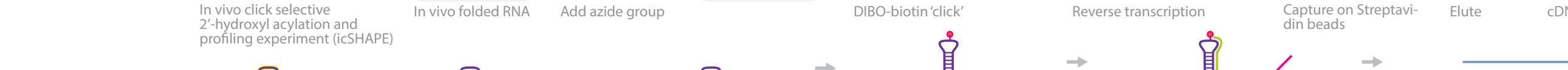
## Pseudo-seq



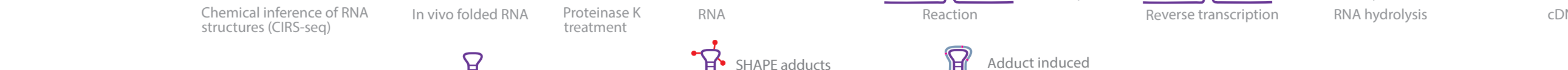
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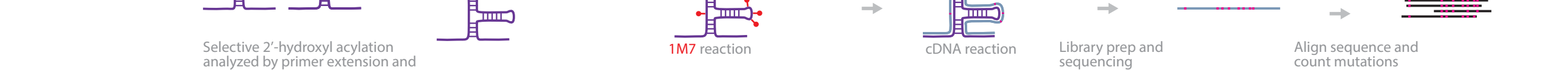
## RNA Structure



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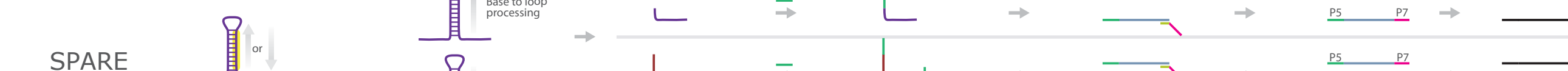
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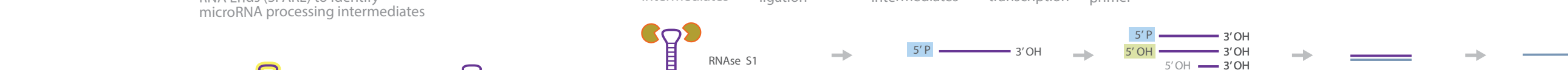
## CIRS-seq



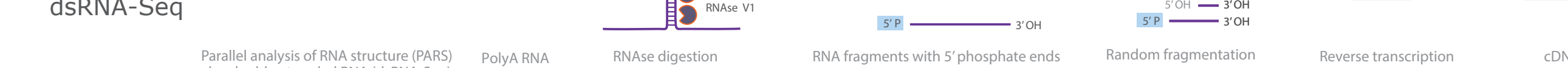
## SHAPE-MaP



## Struc-Seq



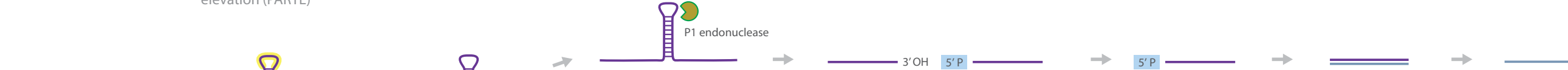
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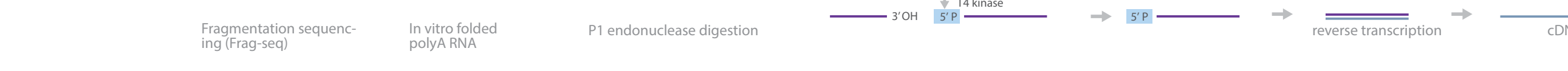
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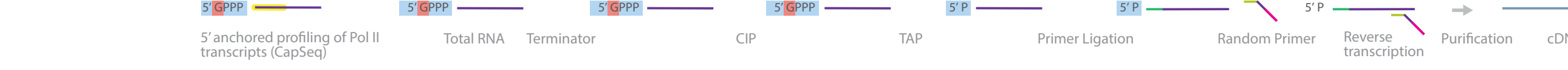
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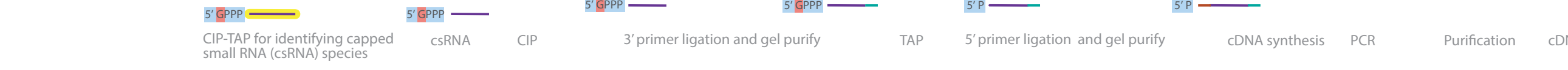
## dsRNA-Seq



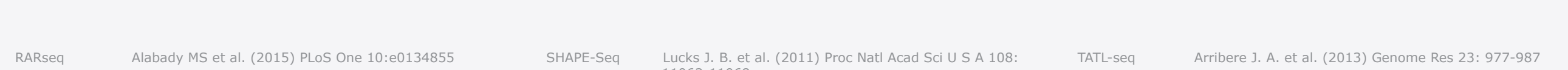
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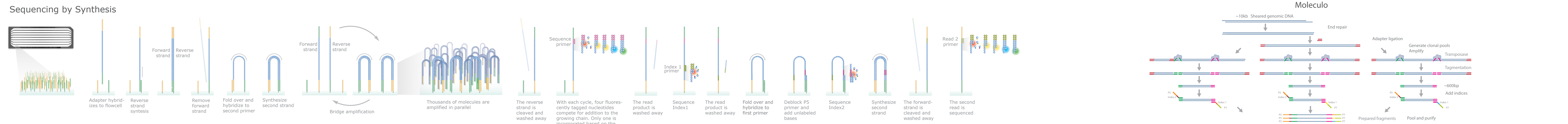
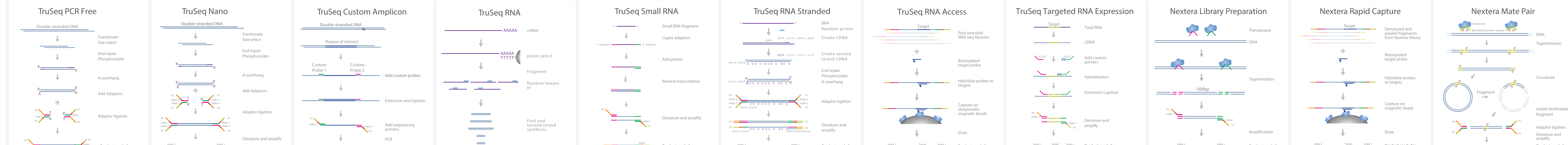
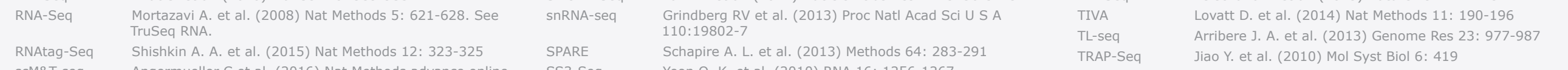
## Frag-seq



## CapSeq



## CIP-TAP



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