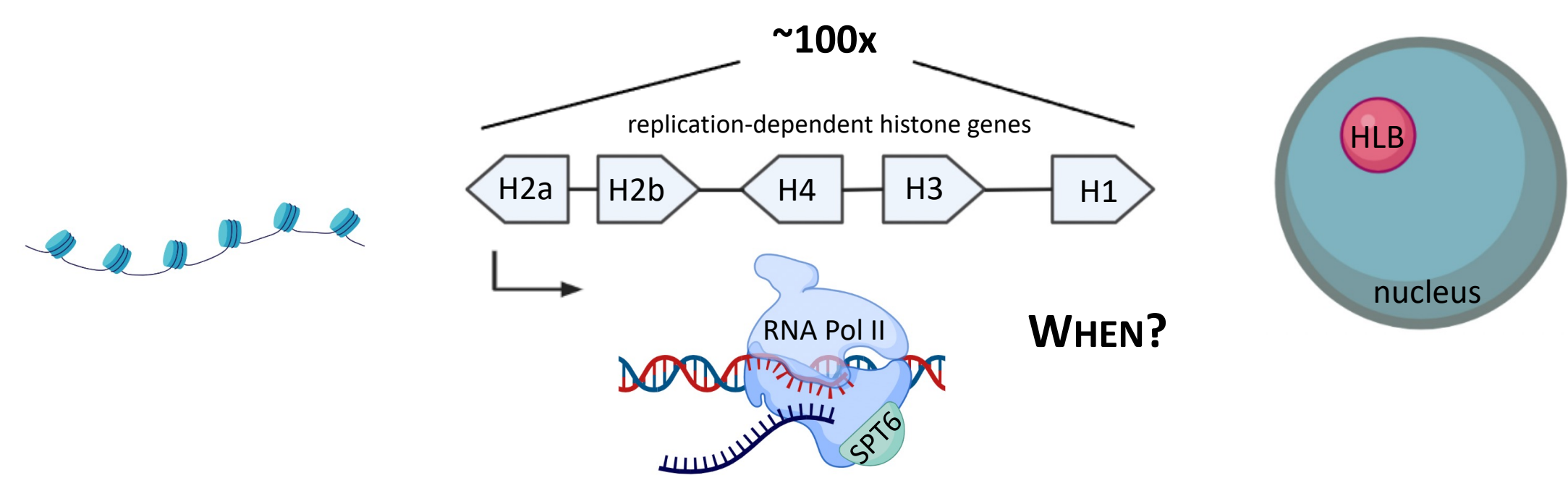


# Dynamics of recruitment of transcription elongation factor SPT6 to the Histone Locus Body during the activation of replication-dependent histone genes in early embryogenesis

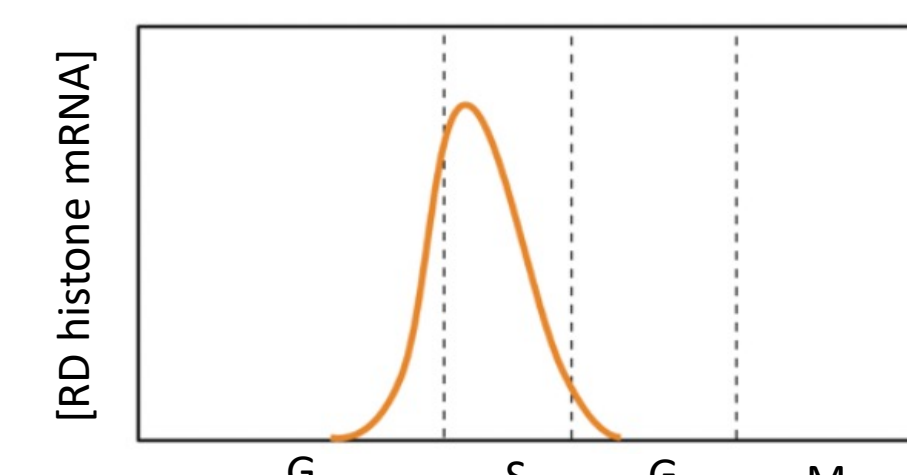
Mia C. Hoover<sup>1</sup>, James P. Kemp, Jr.<sup>2</sup>, Robert J. Duronio<sup>1-4</sup>

Department of Biology<sup>1</sup>, Integrative Program for Biological and Genome Sciences<sup>2</sup>, Lineberger Comprehensive Cancer Center<sup>3</sup>, Department of Genetics<sup>4</sup>, University of North Carolina, Chapel Hill, 27516

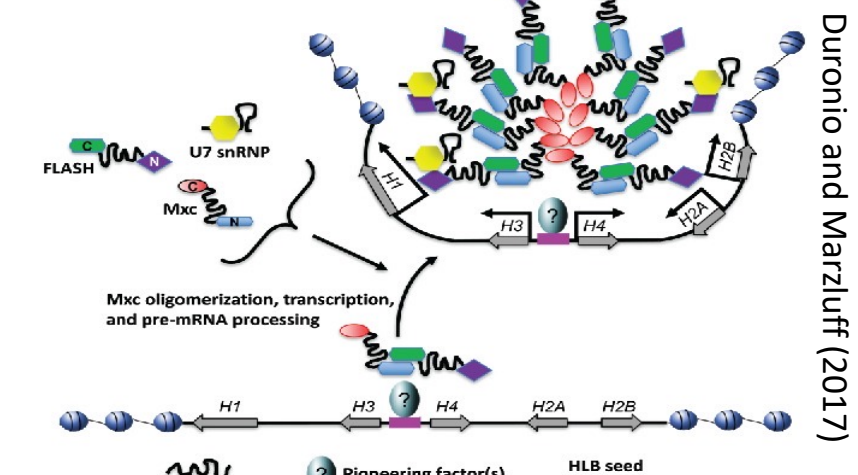
## BACKGROUND



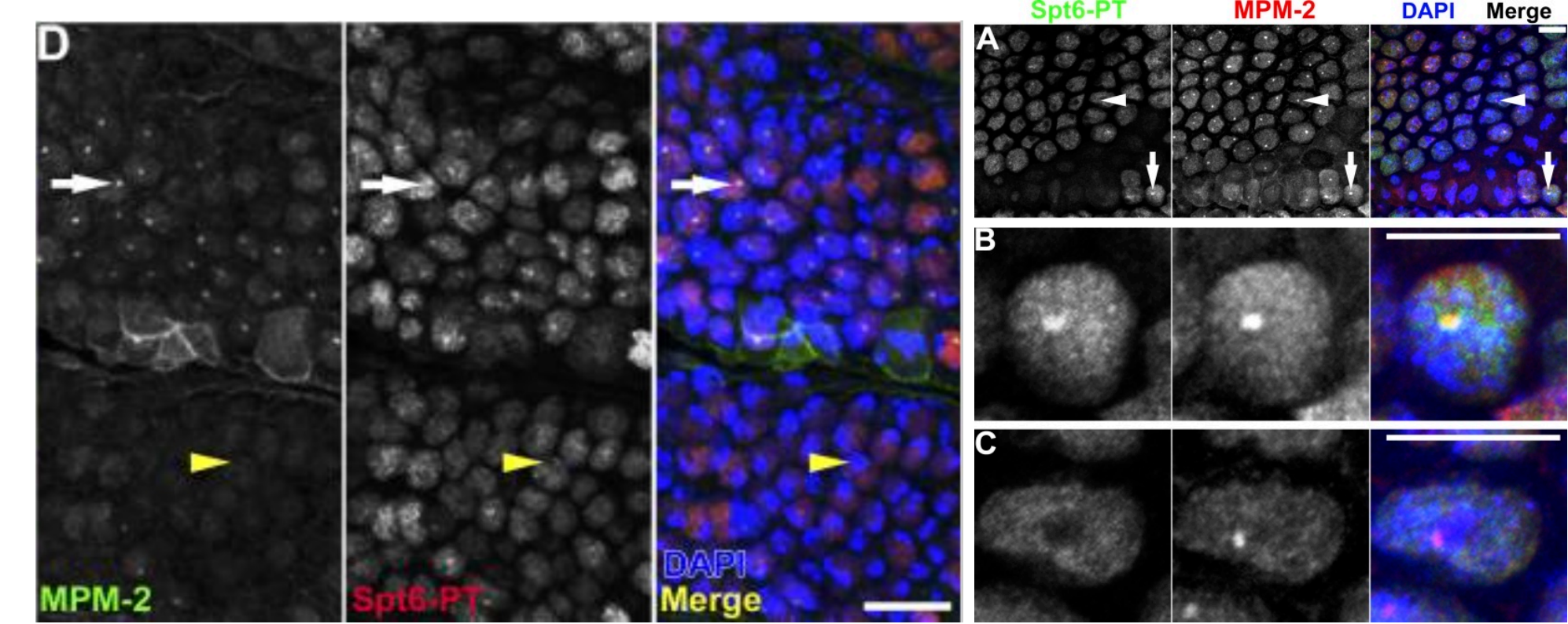
Histones are essential for packaging DNA, and their expression is tightly controlled and coupled to DNA replication. In *Drosophila*, replication-dependent (RD) histone mRNAs are transcribed from a single locus at which an evolutionarily conserved, phase-separated nuclear body forms — the histone locus body (HLB).



Current model of RD histone expression



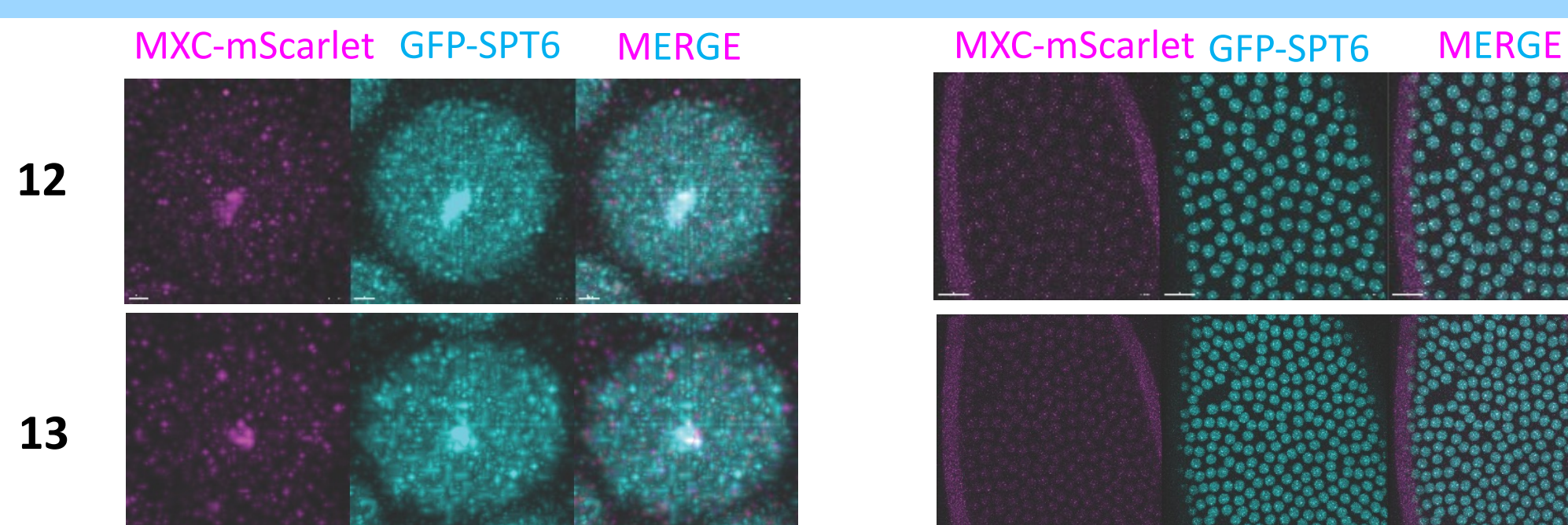
Hierarchical recruitment of components



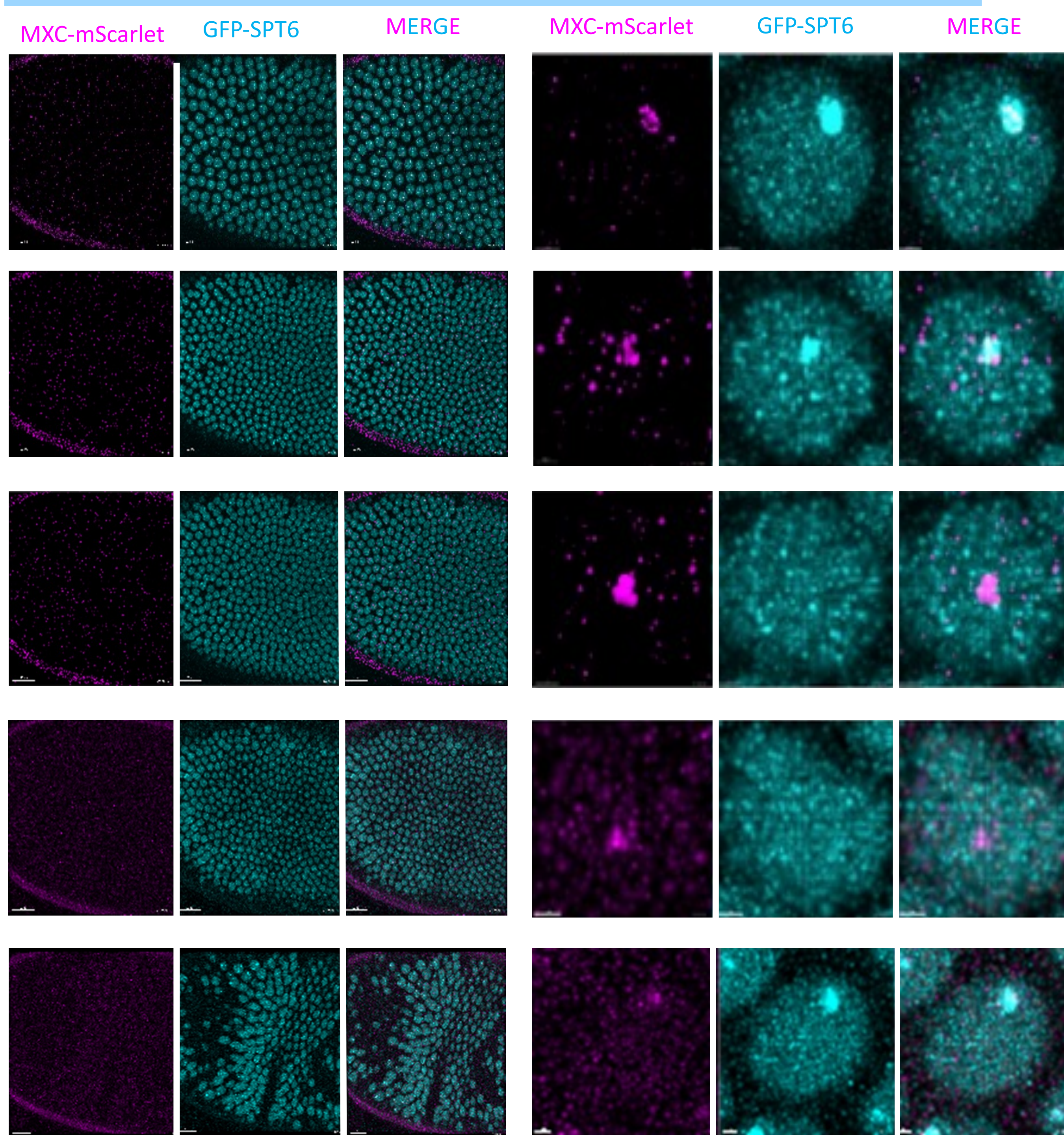
Previous understanding: SPT6 (visualized via  $\alpha$ -SPT6) is a cell-cycle dependent component of the HLB, present when histone genes are being transcribed

Here, using antibodies and fluorescently tagged proteins we investigate the spatiotemporal localization of the transcription elongation factor SPT6 and RNA Pol II at the HLB in developing *Drosophila* embryos. By combining this approach with a sensitive FISH probe that can detect nascent RD histone transcripts, we examined the relationship between the localization of these factors and actively transcribing RD histone genes.

## SPT6 LOCALIZATION IN SYNCYTIAL NUCLEAR CYCLES



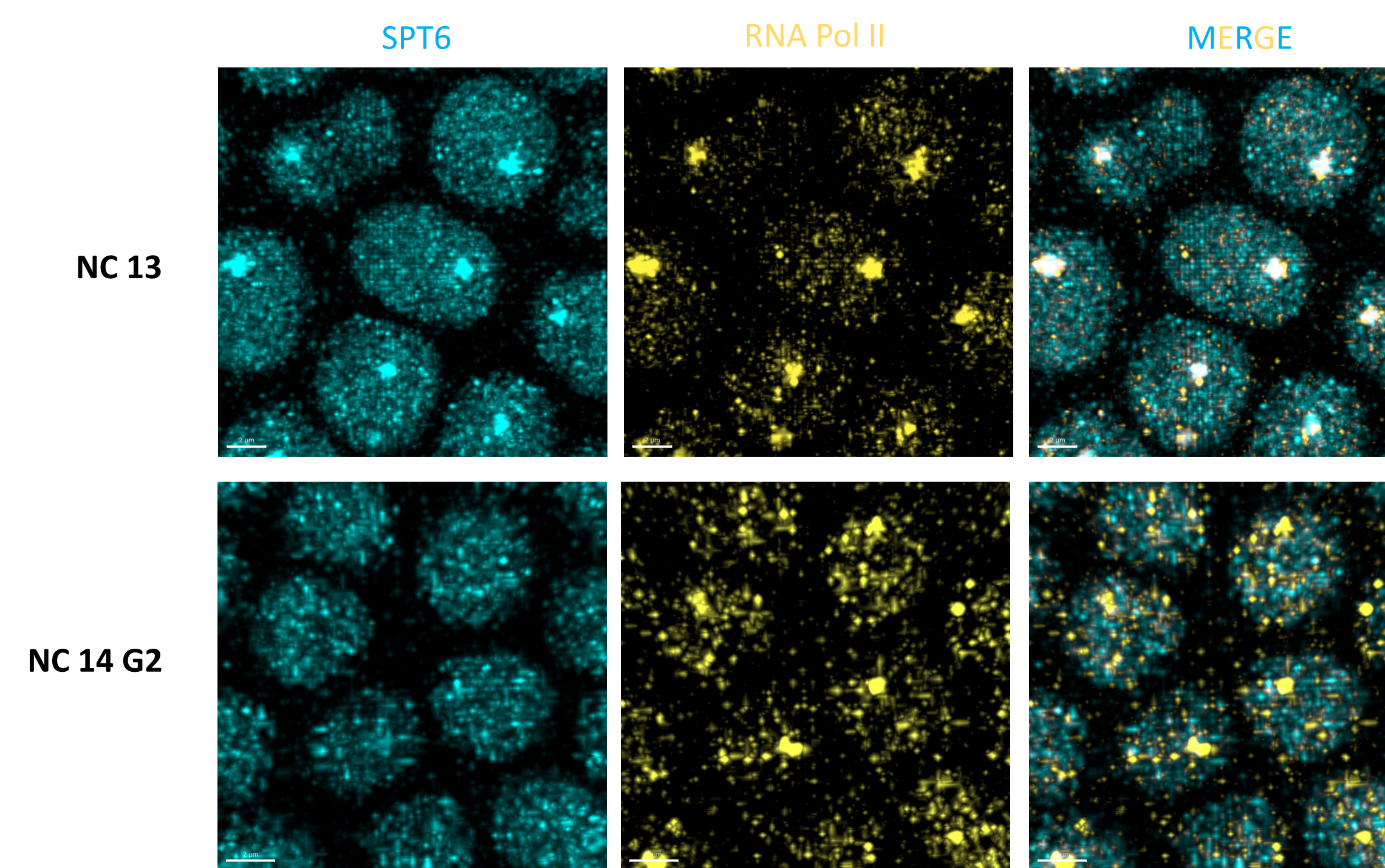
## SPT6 IS PRESENT ONLY TRANSIENTLY AT THE BEGINNING OF NC 14



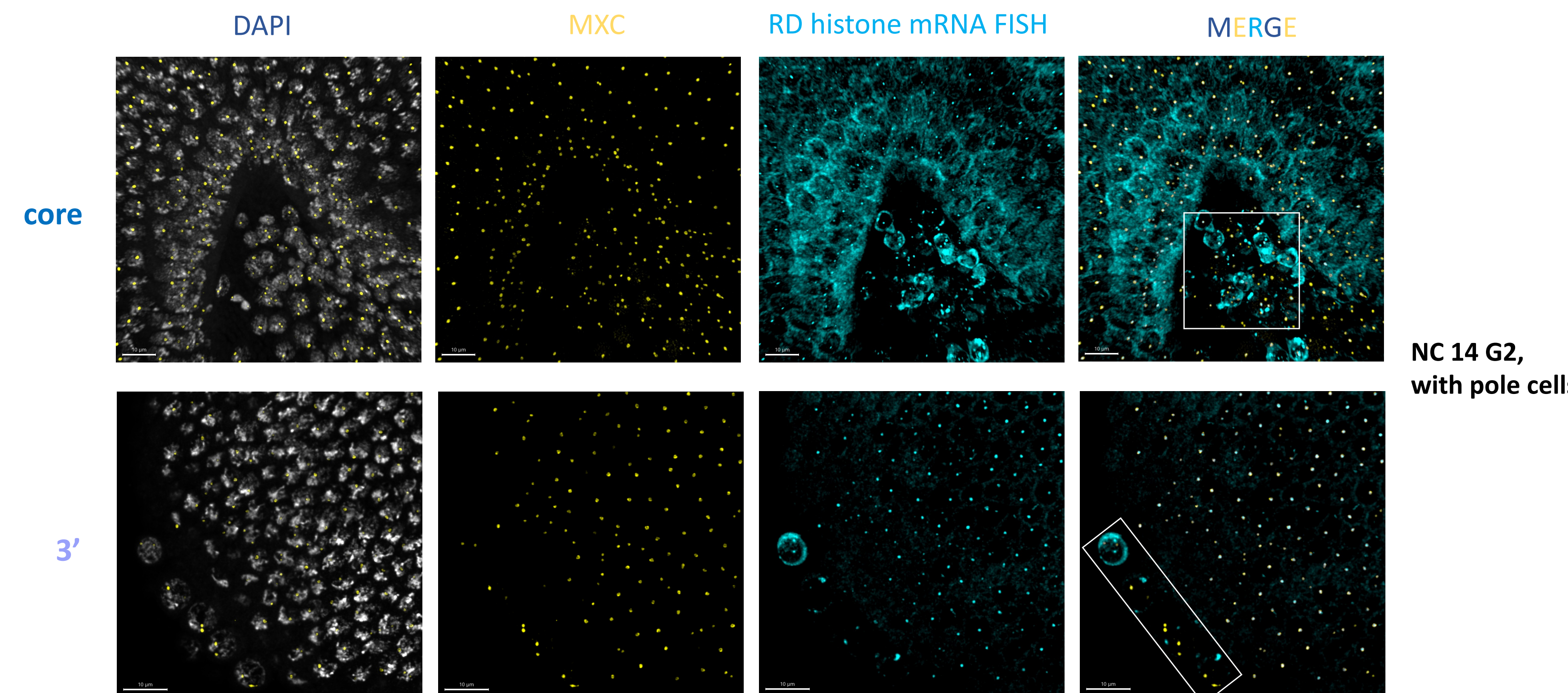
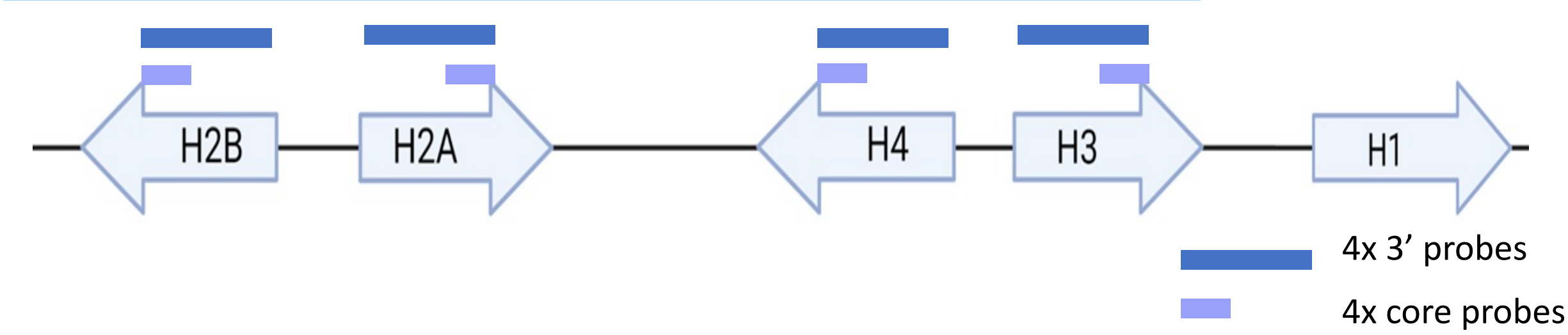
Representative stills from live imaging of developing embryo

## NASCENT RD HISTONE MRNA ACCUMULATES IN THE HLB INDEPENDENT OF DETECTABLE SPT6 ENRICHMENT DURING G2 OF NC 14

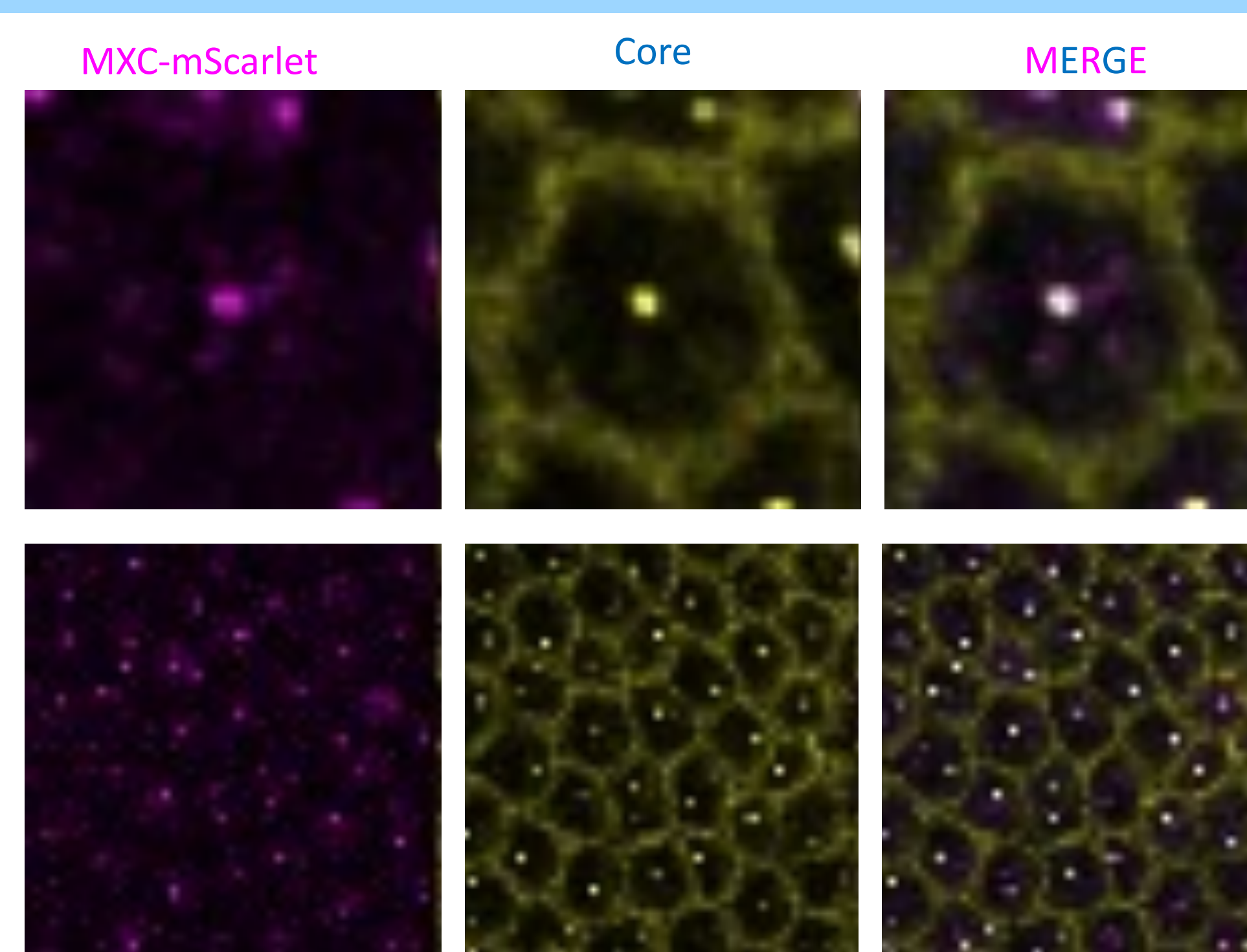
### Decoupling of SPT6 and RNA Pol II in NC 14 G2



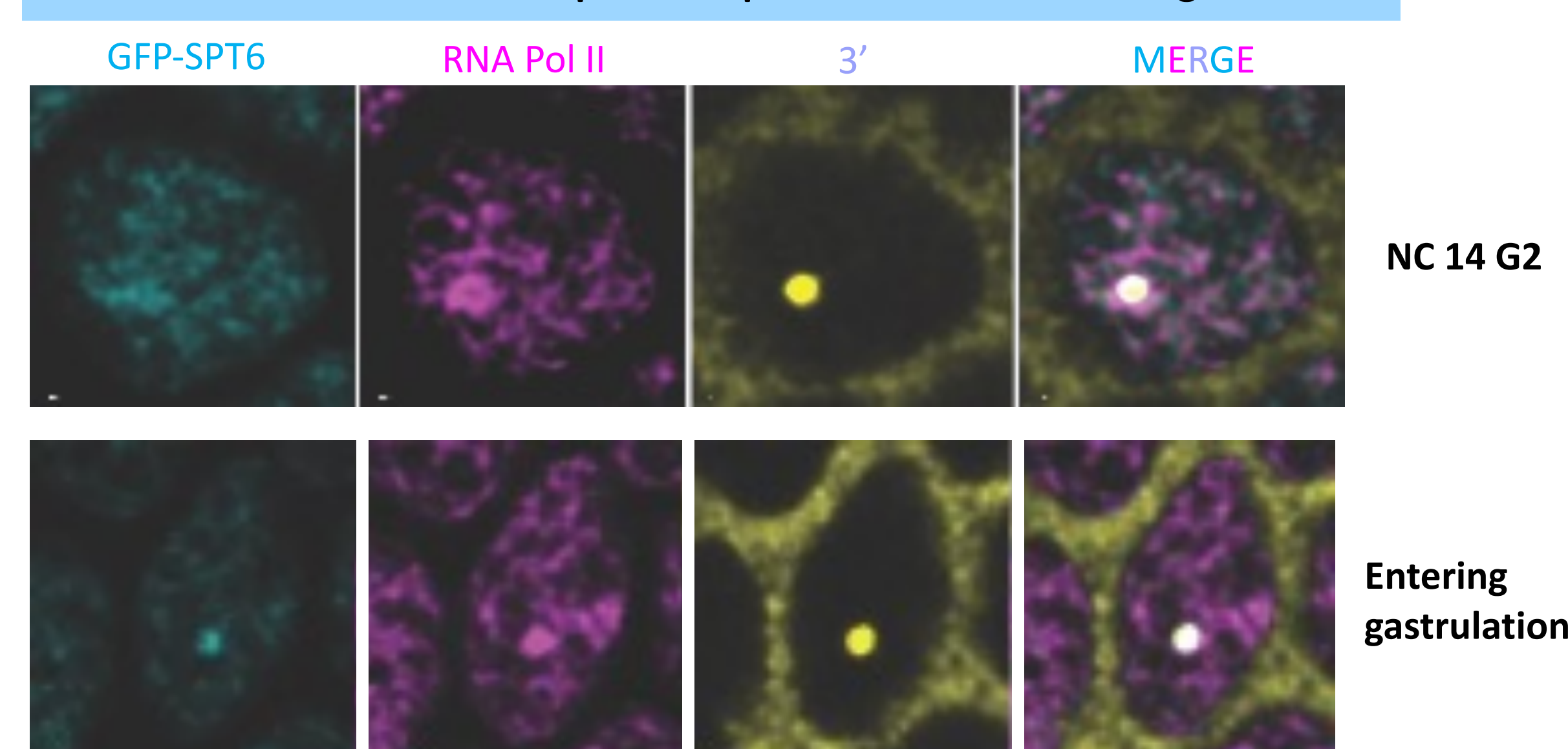
### Different probes reveal paused polymerase at RD histone genes



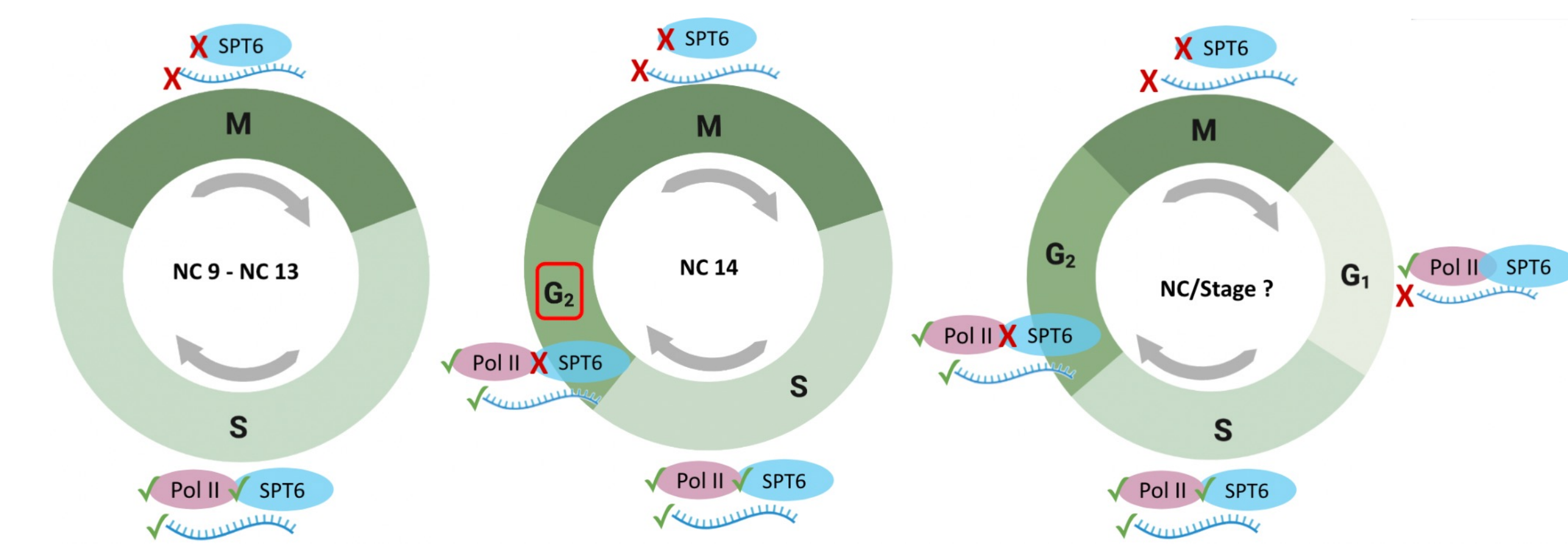
### Core nascent RD histone transcript is present at the HLB during NC 14 G2



### 3' nascent RD histone transcript is also present at the HLB during NC 14 G2



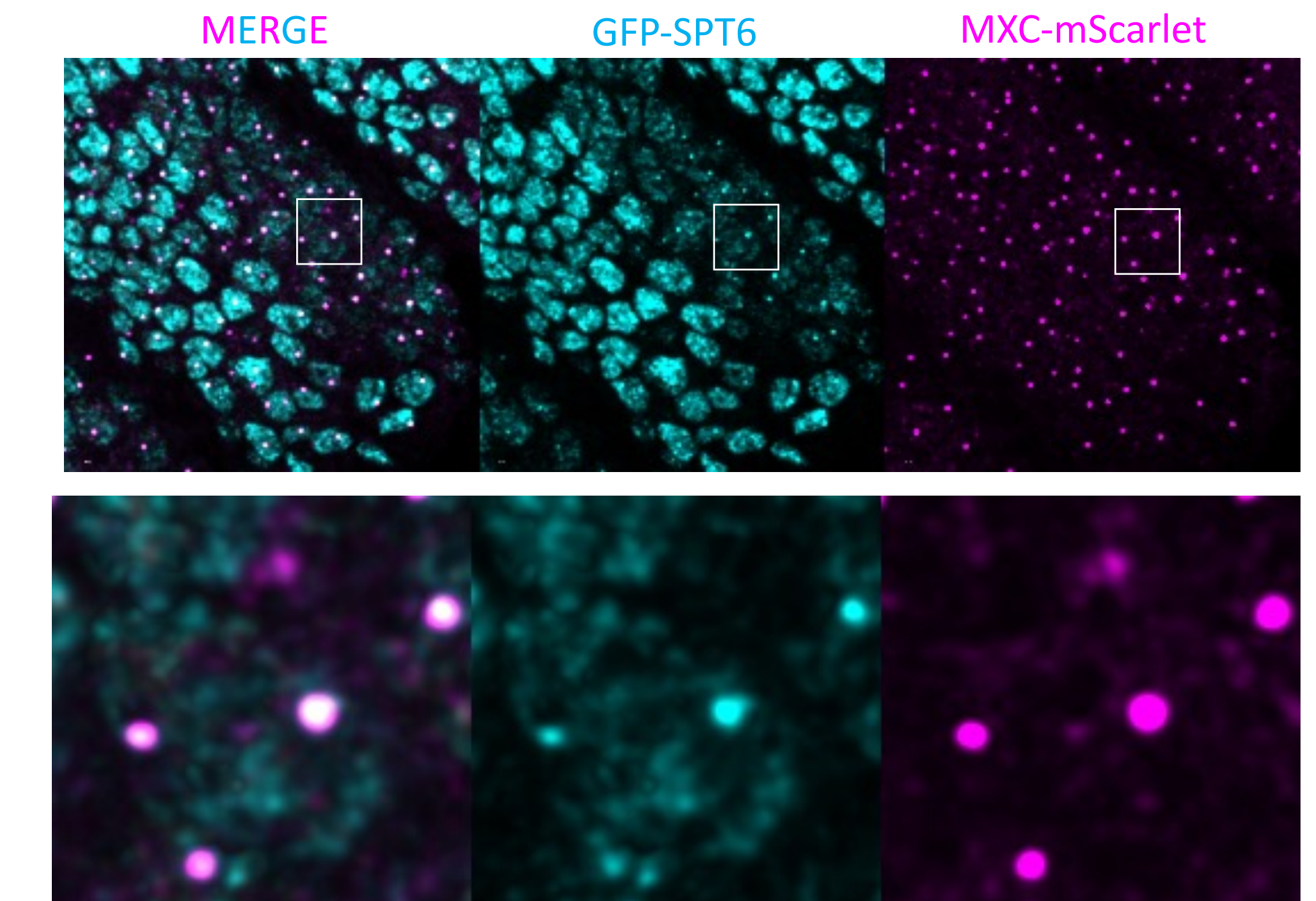
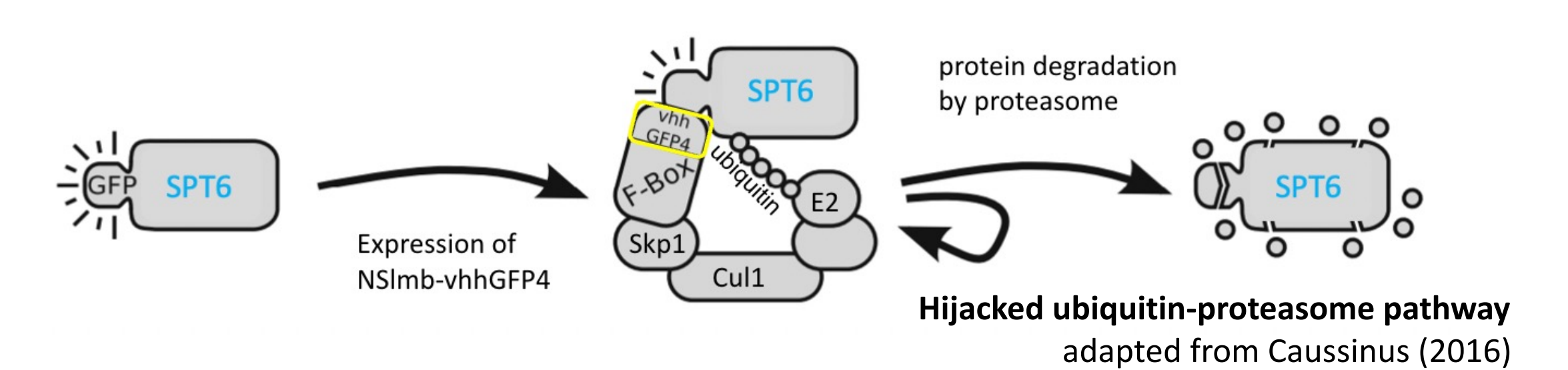
## RD HISTONE TRANSCRIPTION & THE CELL CYCLE



**Result:** Uncoupling of SPT6 association with RNA Pol II and active transcription of RD histone mRNA. RNA Pol II is not paused during transcription of RD histone mRNA in NC 14 G2.

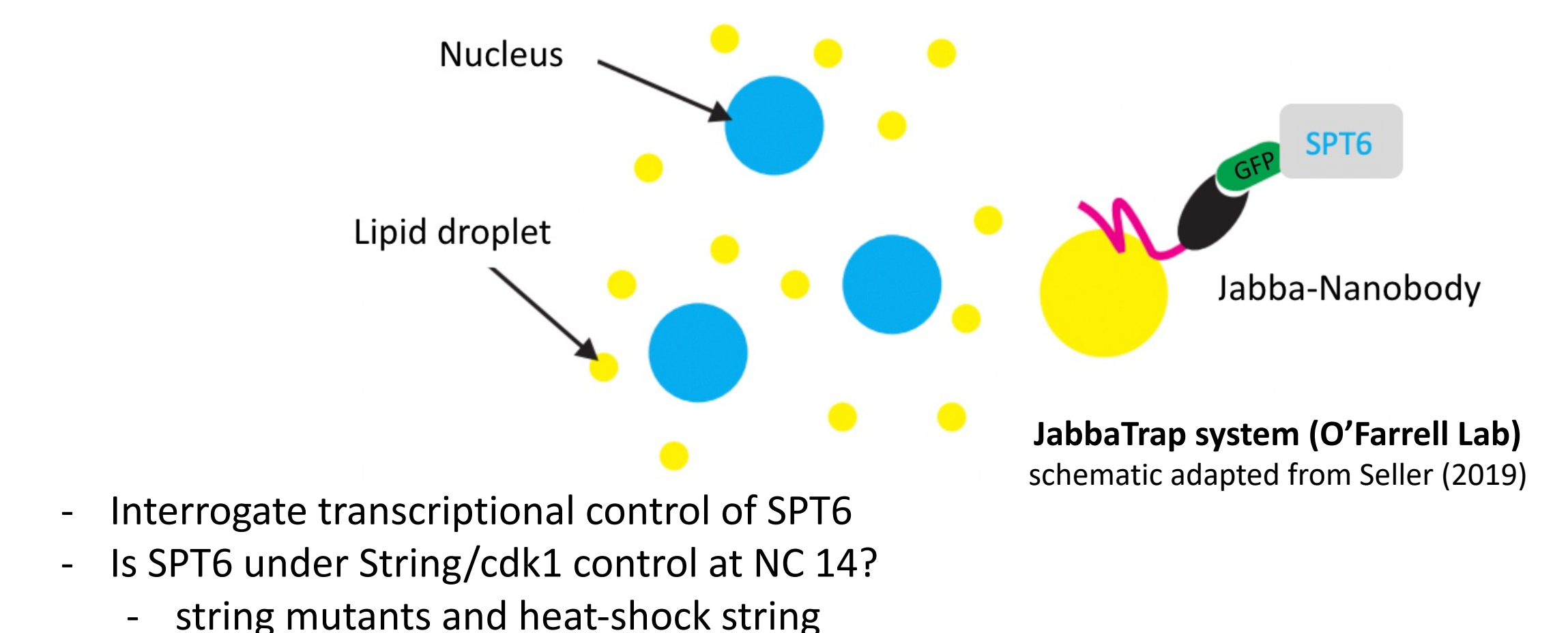
## IS SPT6 NECESSARY FOR HLB FORMATION OR ACTIVATION?

Attempted depletion of SPT6 from nucleus via DeGradFP system



Stage 14 post-gastrulation embryo

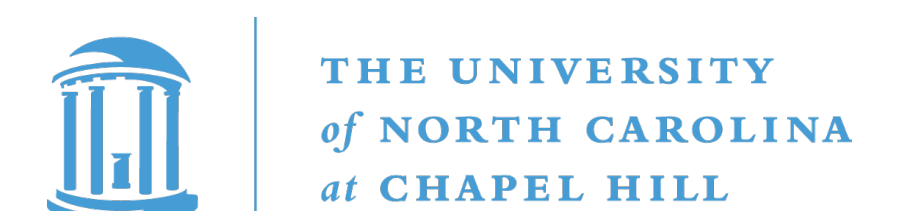
## FUTURE DIRECTIONS



- Interrogate transcriptional control of SPT6
- Is SPT6 under String/cdk1 control at NC 14?
- string mutants and heat-shock string

## ACKNOWLEDGEMENTS

Many thanks to the Duronio Lab for their years of mentorship, teaching me how to do science, and friendship — especially Bob Duronio, for the opportunities to learn in a wonderful lab environment, and to my mentor Jim Kemp, for his patience and teaching me confocal microscopy.



This work is funded by R35 GM145258, and in part by a SURF grant from the UNC Office of Undergraduate Research.

## REFERENCES

White, A. E., Burch, B. D., Yang, X.-C., Gasdaska, P. Y., Dominski, Z., Marzluff, W. F., & Duronio, R. J. (2011). *Drosophila* histone locus bodies form by hierarchical recruitment of components. *The Journal of Cell Biology*, 193(4), 677–694. <https://doi.org/10.1083/jcb.201012077>

White, A. E., Leslie, M. E., Calvi, B. R., Marzluff, W. F., & Duronio, R. J. (2007). Developmental and cell cycle regulation of the *Drosophila* histone locus body. *Molecular Biology of the Cell*, 18(7), 2491–2502. <https://doi.org/10.1091/mbc.E06-11-1033>

Hur, W., Kemp, J. P., Tarzia, M., Deneke, V. E., Marzluff, W. F., Duronio, R. J., & Di Talia, S. (2020). CDK-Regulated Phase Separation Seeded by Histone Genes Ensures Precise Growth and Function of Histone Locus Bodies. *Developmental Cell*, 54(3), 379–394.e6. <https://doi.org/10.1016/j.devcel.2020.06.003>

Causinus, E., & Affolter, M. (2016). deGradFP: A System to Knockdown GFP-Tagged Proteins. *Methods in Molecular Biology*, 1478, 177–187. [https://doi.org/10.1007/978-1-4939-6371-3\\_9](https://doi.org/10.1007/978-1-4939-6371-3_9)