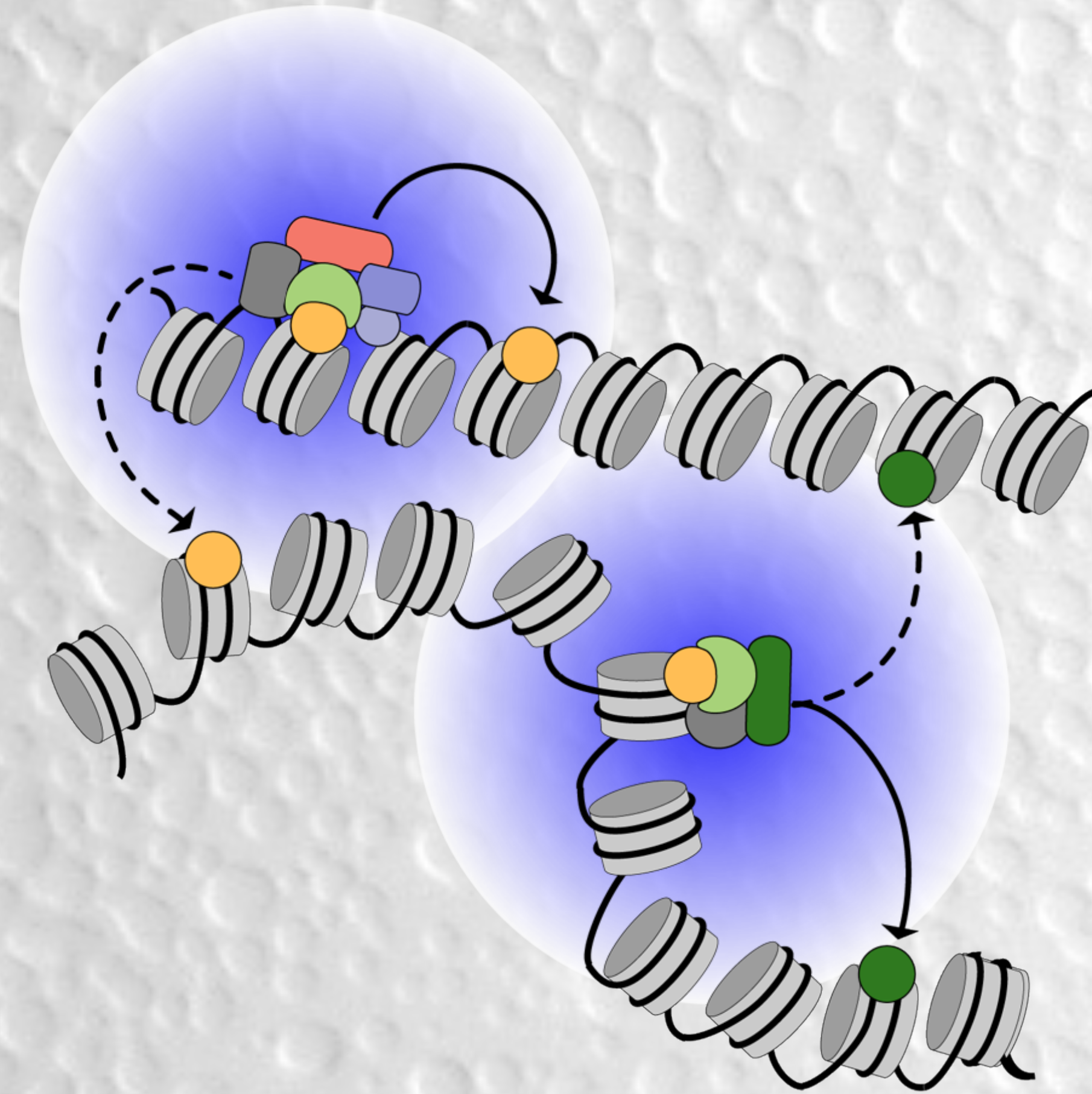
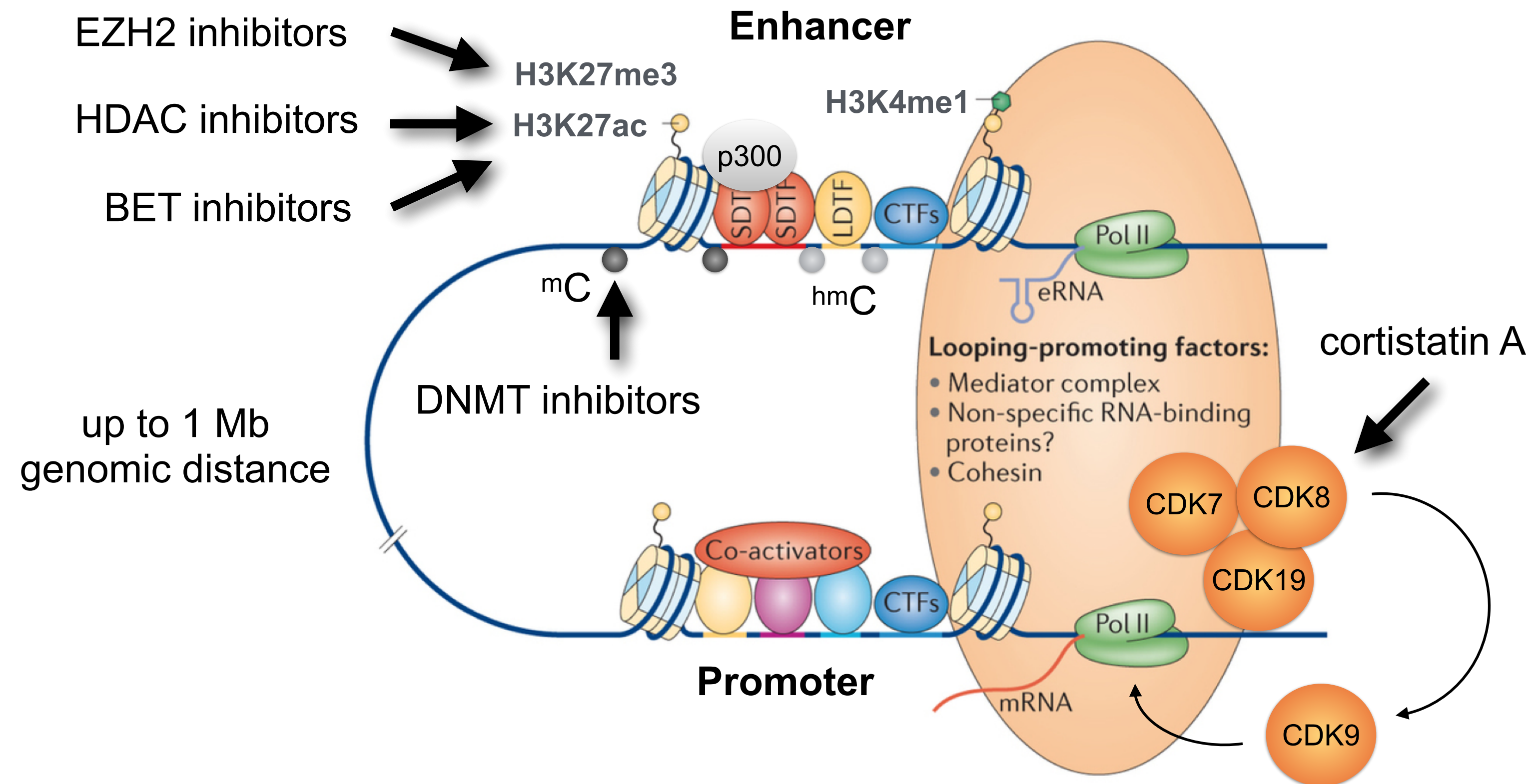


Interaction of DNA-bound proteins at a distance



More than **300 000** putative enhancers have been annotated in the human genome



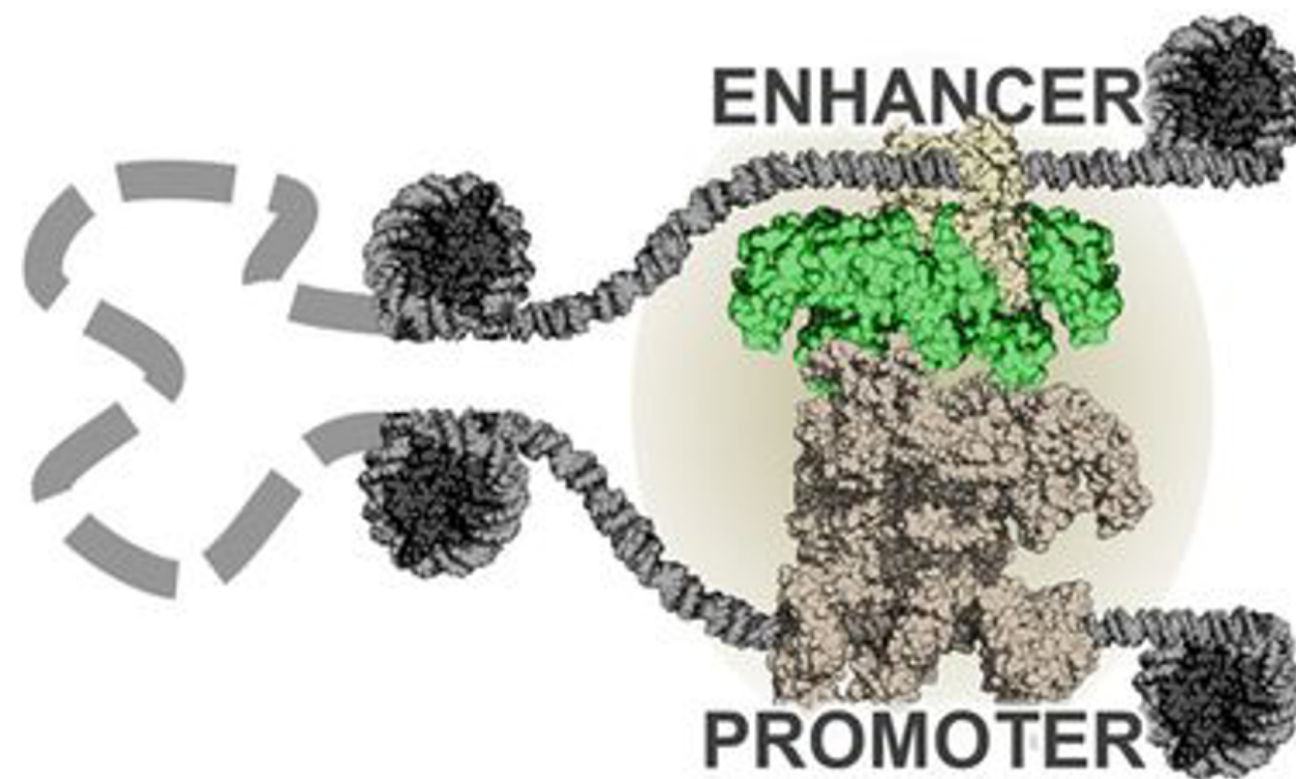
Cell type specific activity estimates

- ~20 000 coding genes (~54 000 including lncRNAs etc)
- 80 000 - 240 000 active enhancers
- typical: 1-2 target promoters per enhancer (~10 for some enhancers)
- multiple enhancers for single promoter
- 300-500 “super-enhancers” > 10 kb

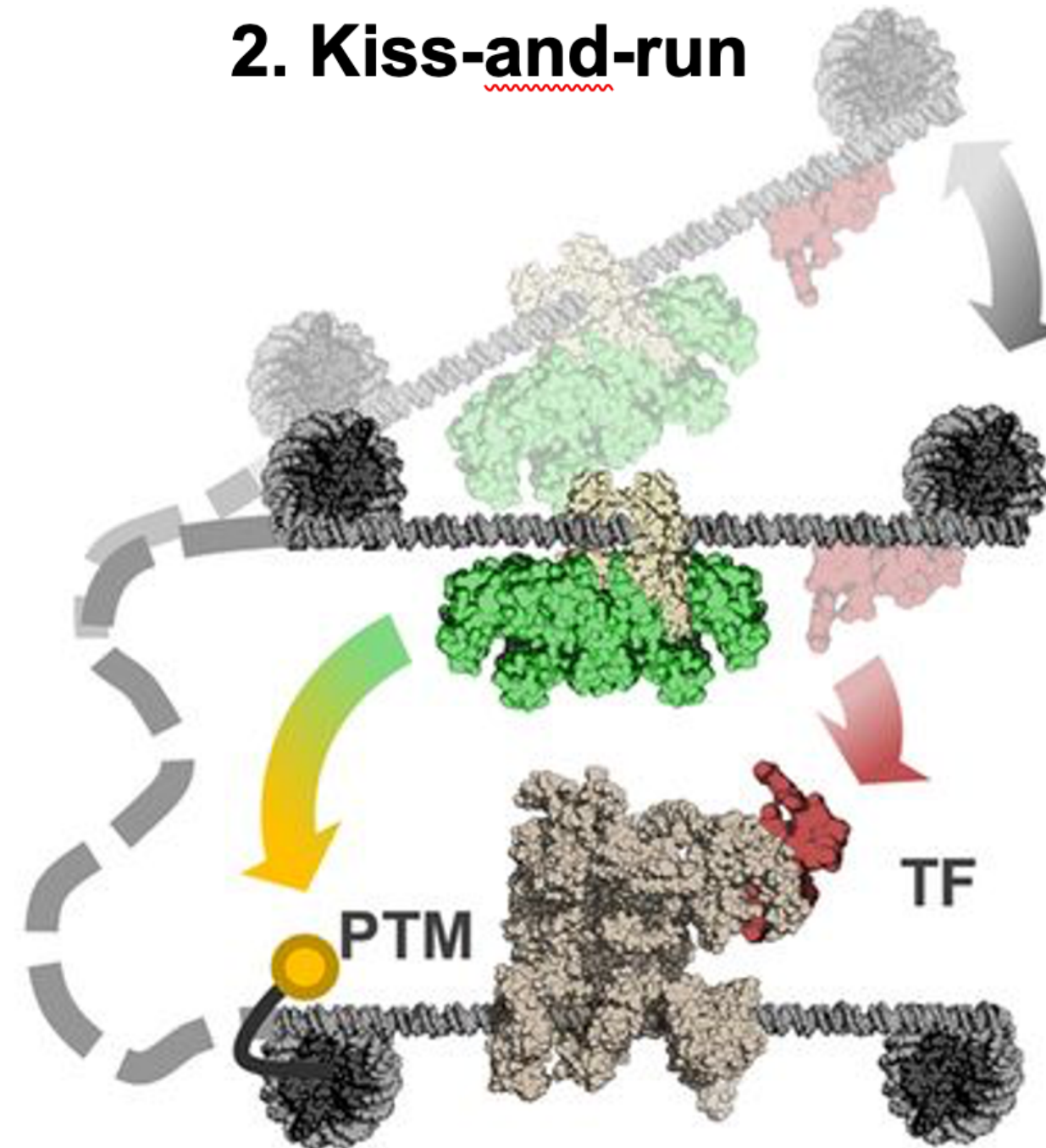
Heinz 2015, Nat Rev Mol Cell Biol
Roadmap Epigenomics Consortium 2015, Nature
FANTOM Consortium 2015, Nature

How do enhancers activate transcription?

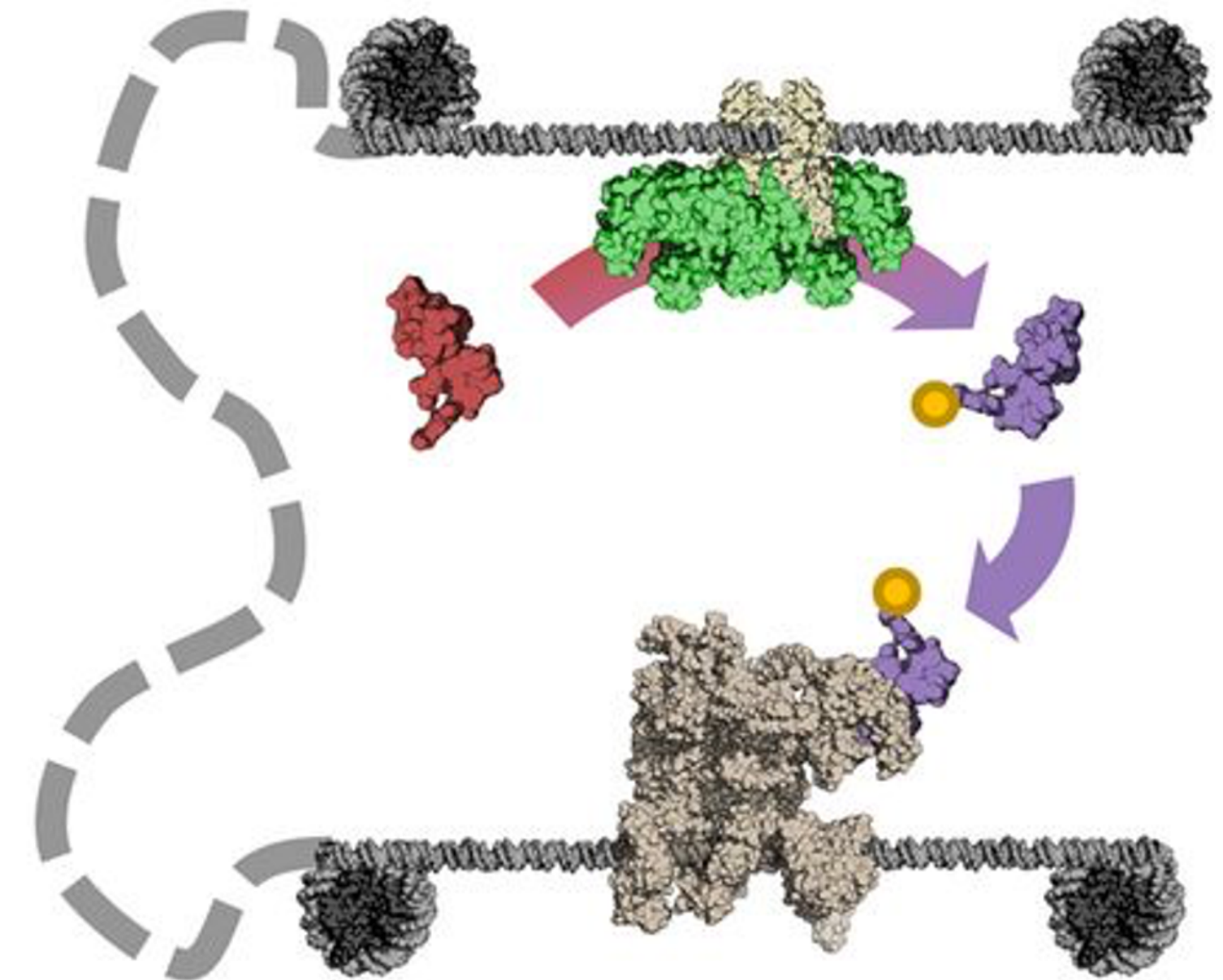
1. Stable contact



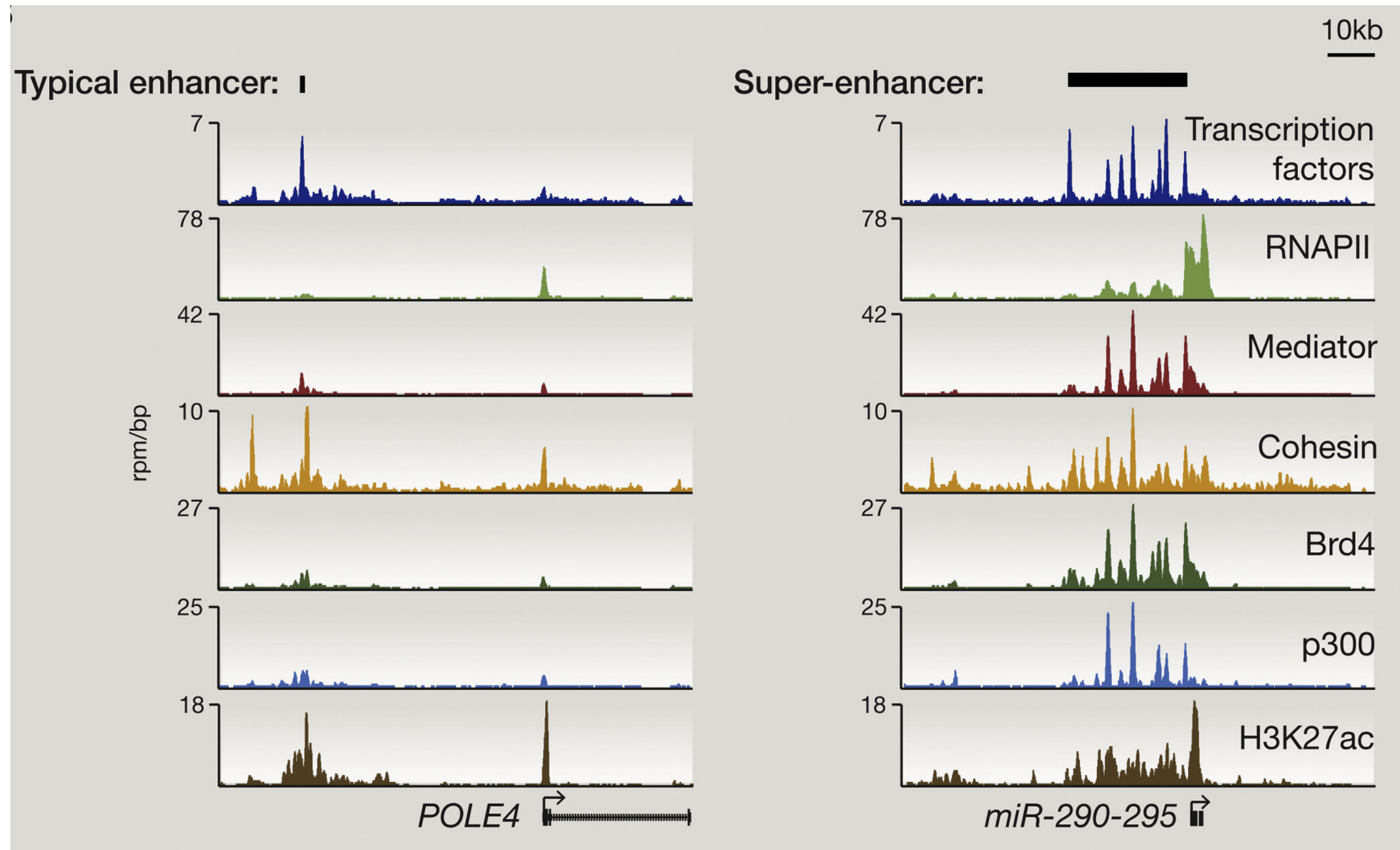
2. Kiss-and-run



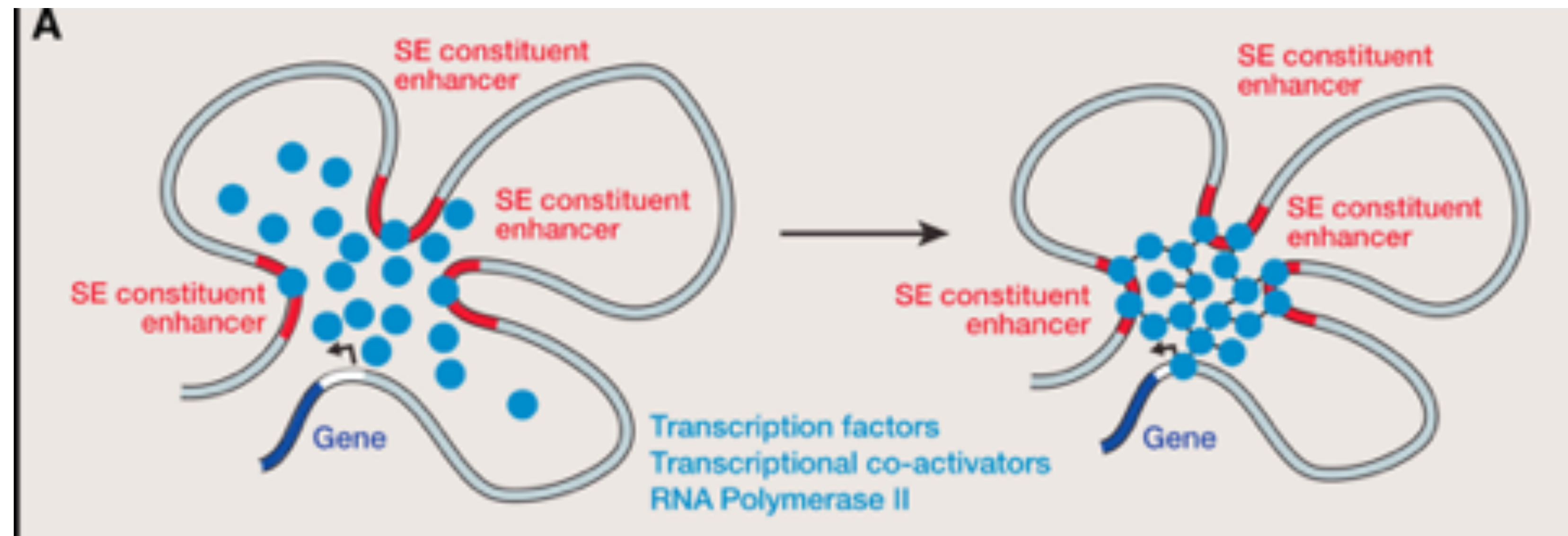
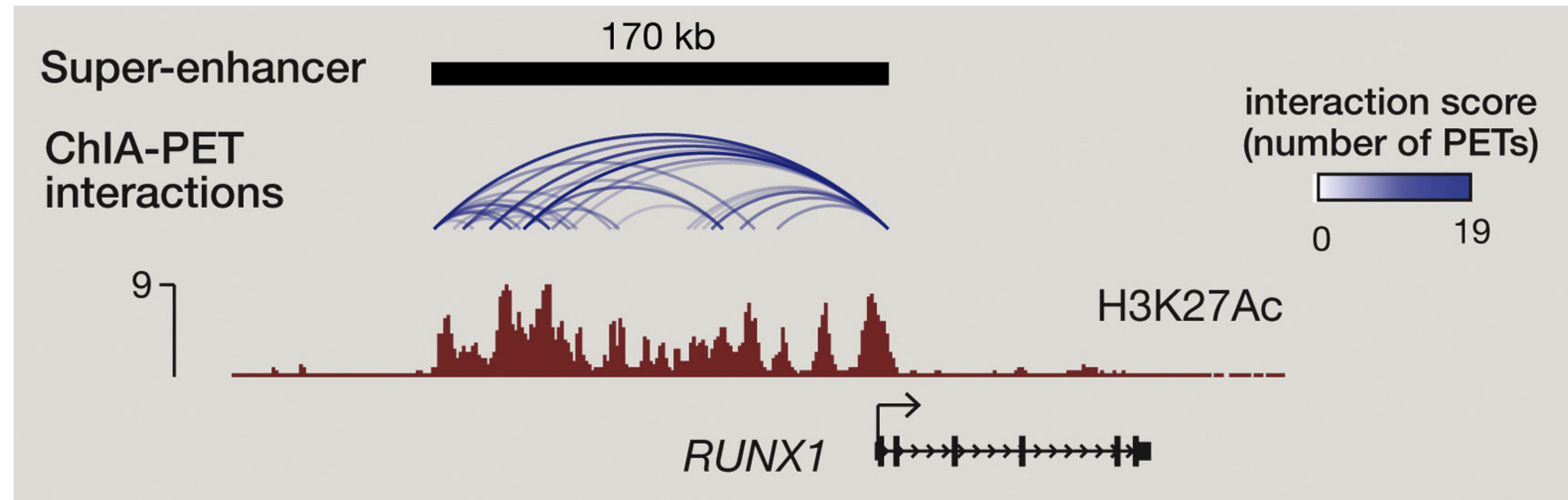
3. Transcription factor activity gradient



Enhancers vs “super-enhancers” (≥ 10 kb)



Do super-enhancers activate via a phase separation mechanism?

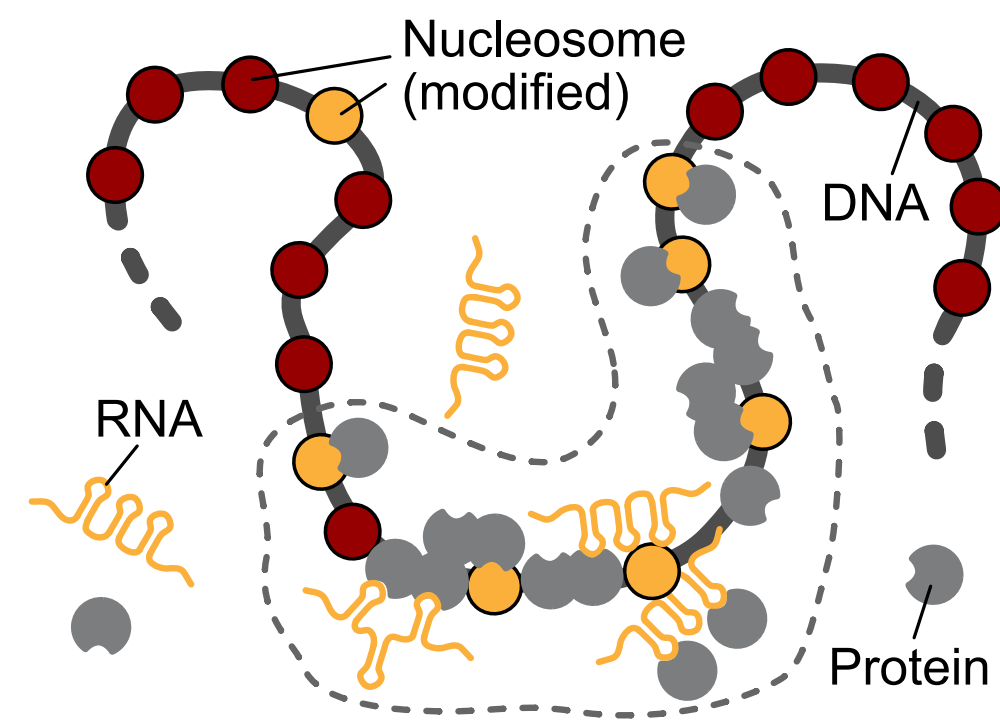


Hnisz D, Shrinivas K, Young RA, Chakraborty AK, Sharp PA (2017) A Phase Separation Model for Transcriptional Control. *Cell* **169**: 13-23

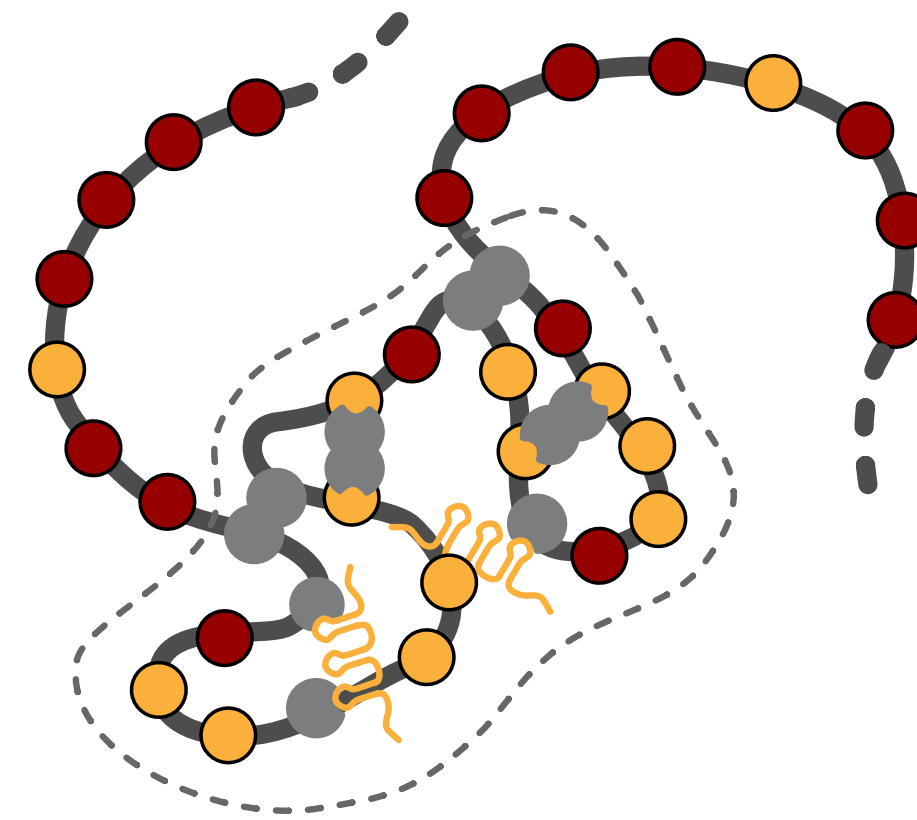
Mechanisms to form chromatin subcompartments

The “null hypothesis”

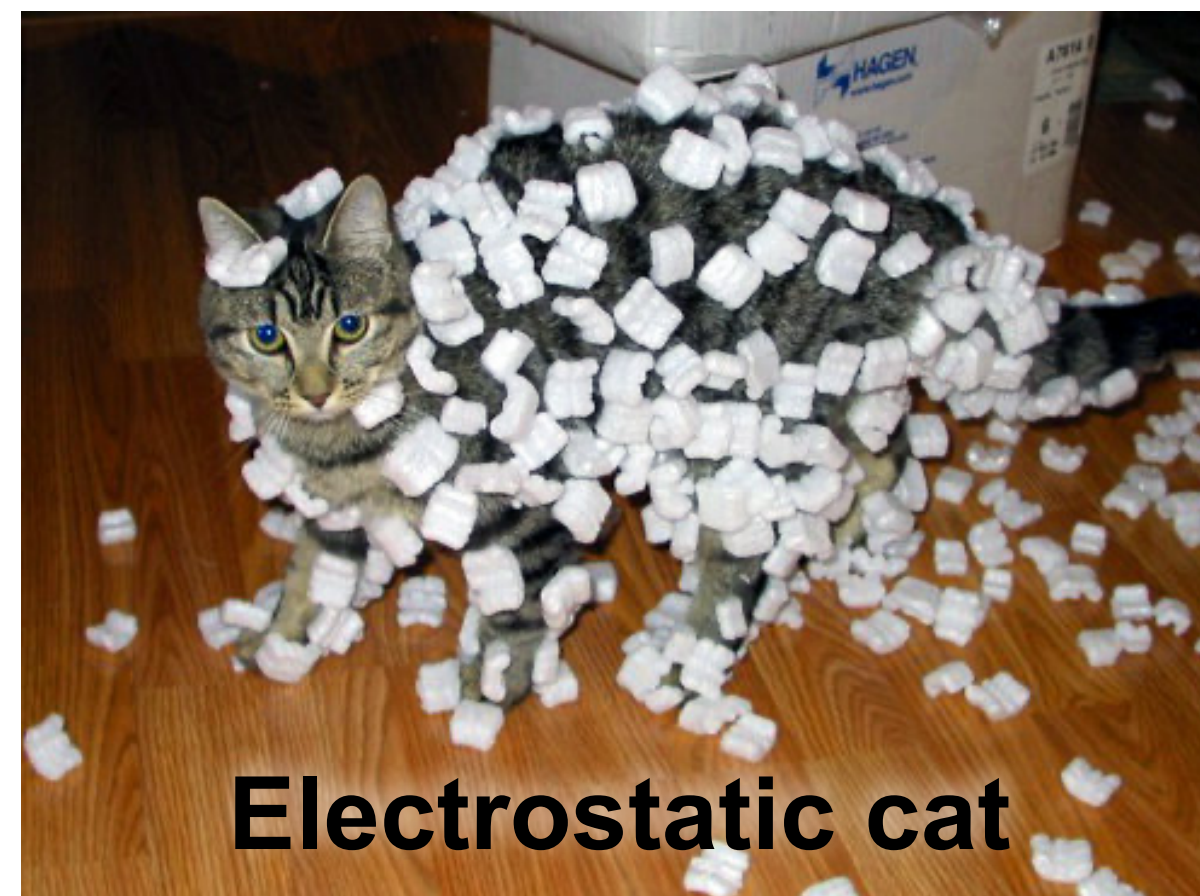
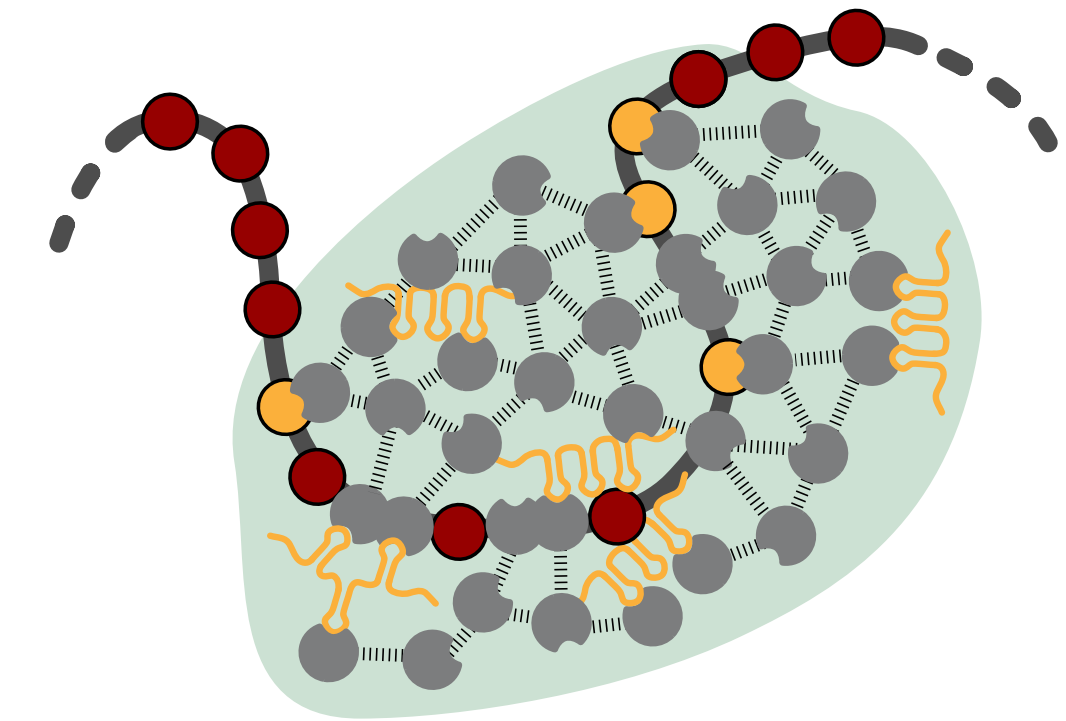
Binding site cluster



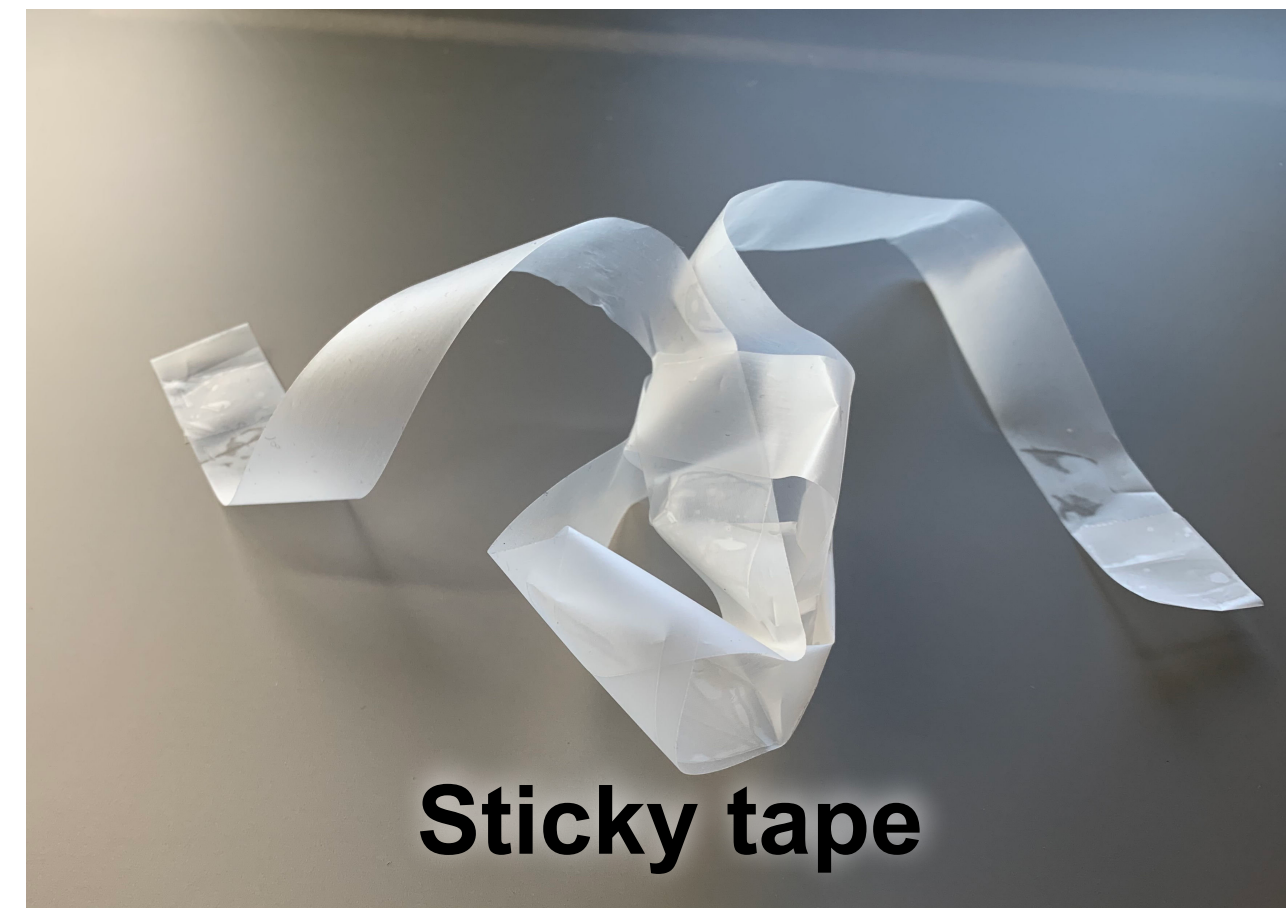
Chromatin bridging interactions



Liquid-liquid phase separation



Electrostatic cat



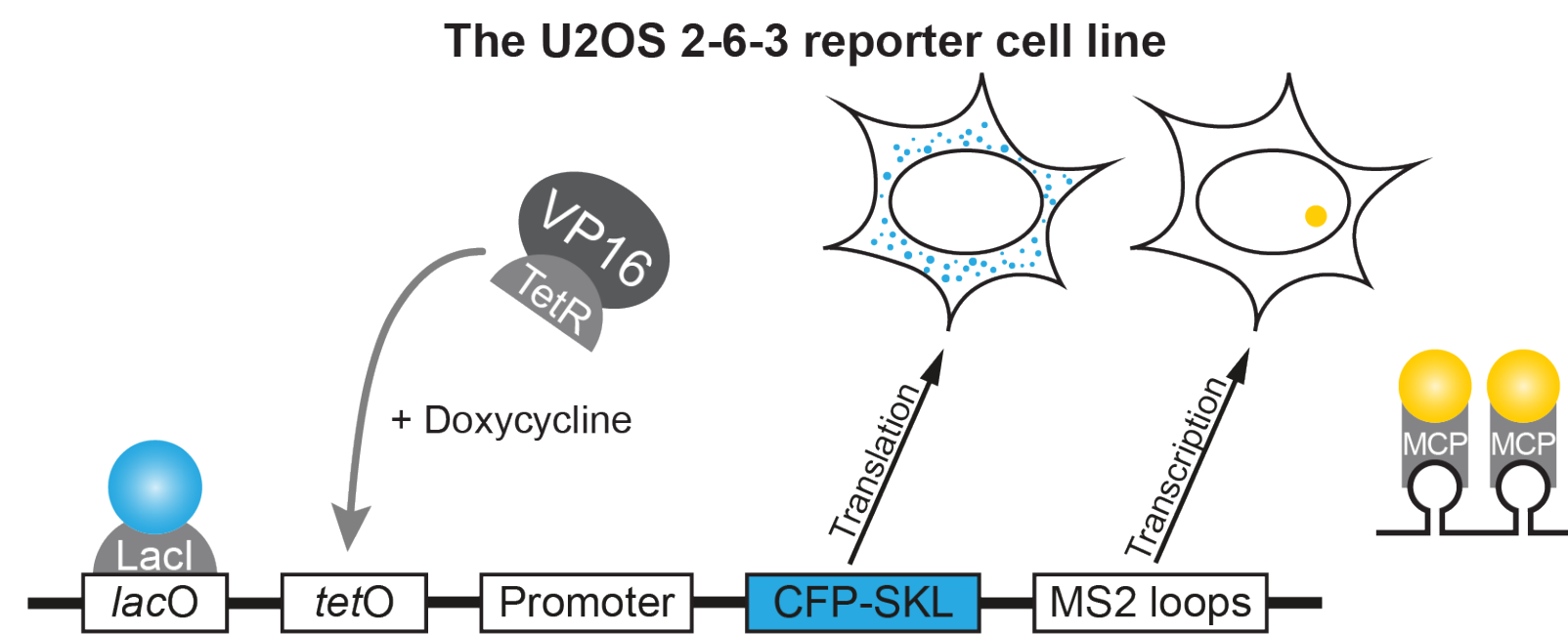
Sticky tape



Drops on a spider net

Image by Mathew Spolin

Studying transcription factor droplets in living cell at a reporter gene array



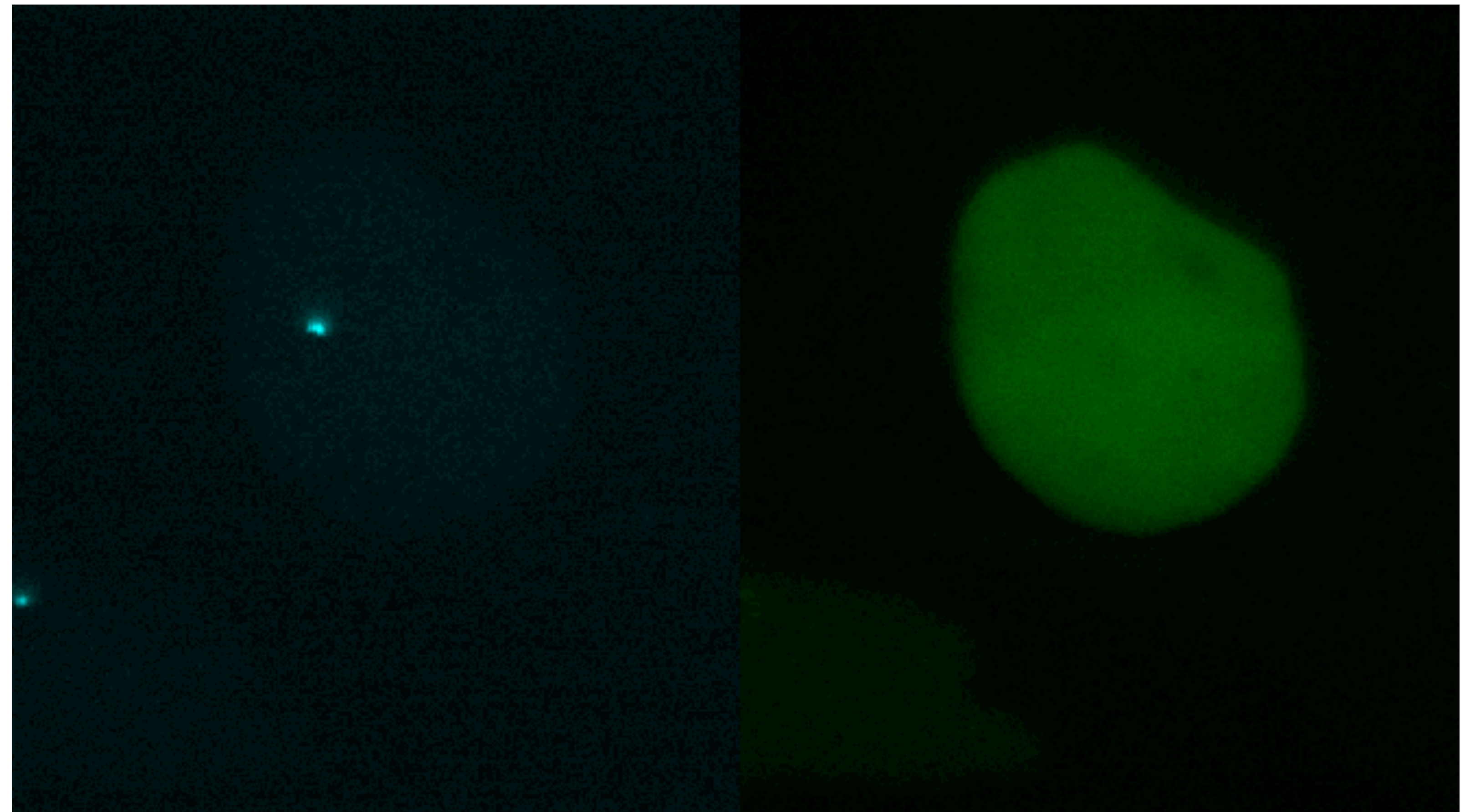
Heterochromatic locus:

- array with ~200 repeats
- compact chromatin state
- H3K9me3
- HP1 enriched
- Activated by VP16/VPR

Janicki *et al.* (2004) *Cell* 116:683-698.

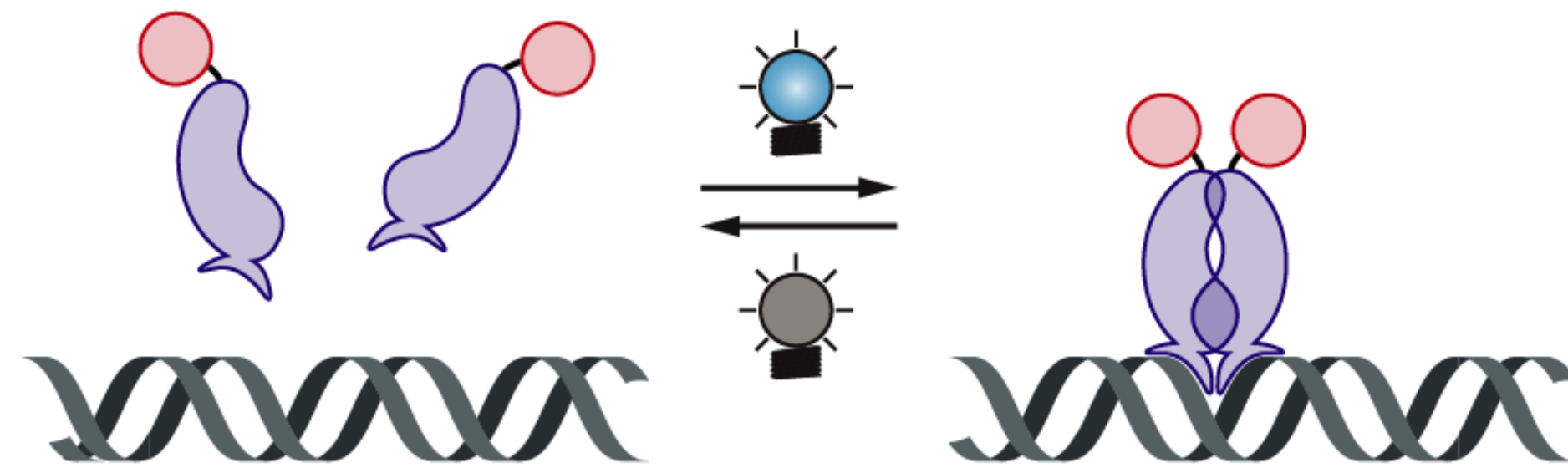
CFP-LacI (chromatin)
CFP (translated protein)

MS2-YFP (RNA)

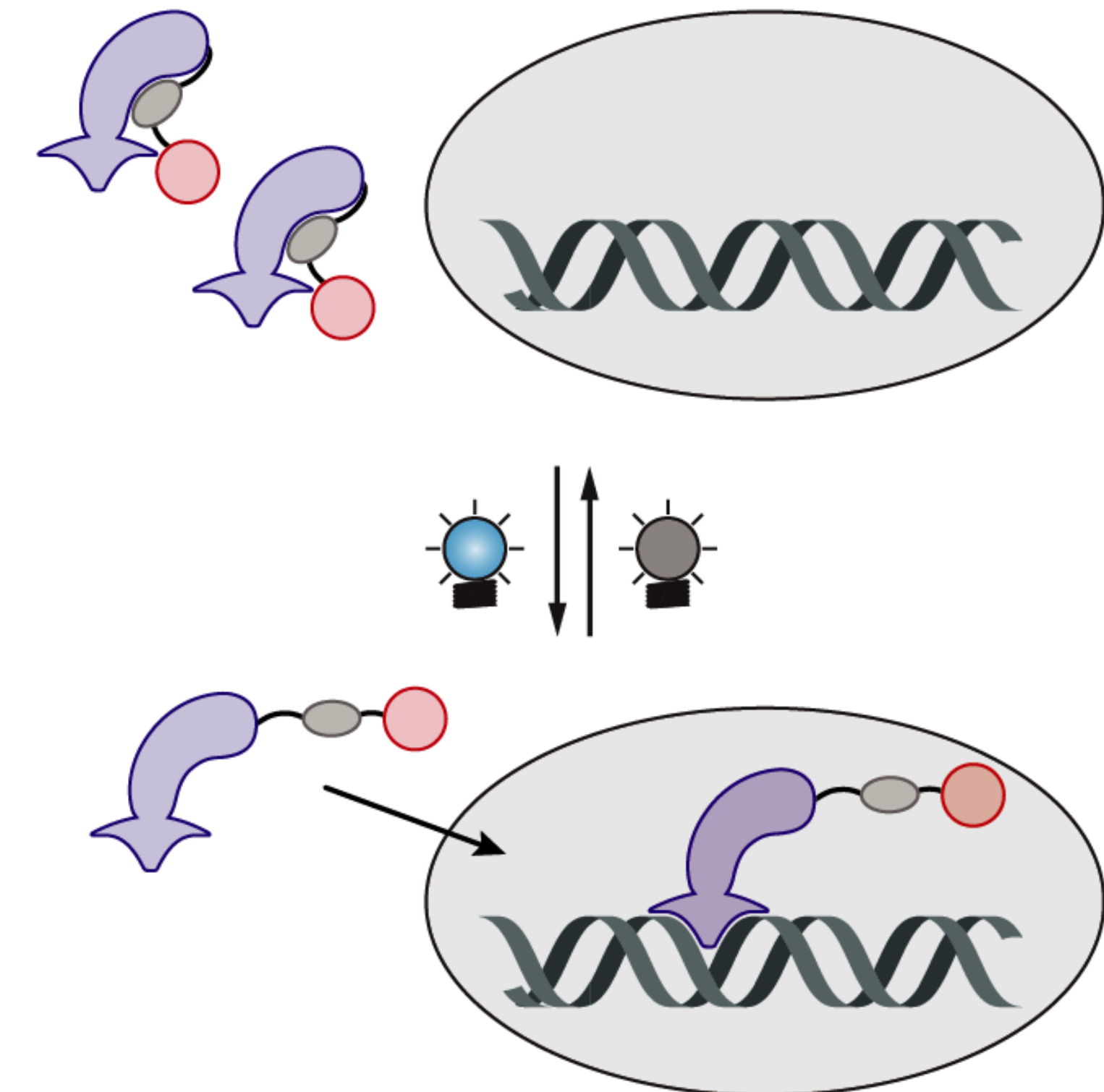


Optogenetics to control proteins in living cells with light

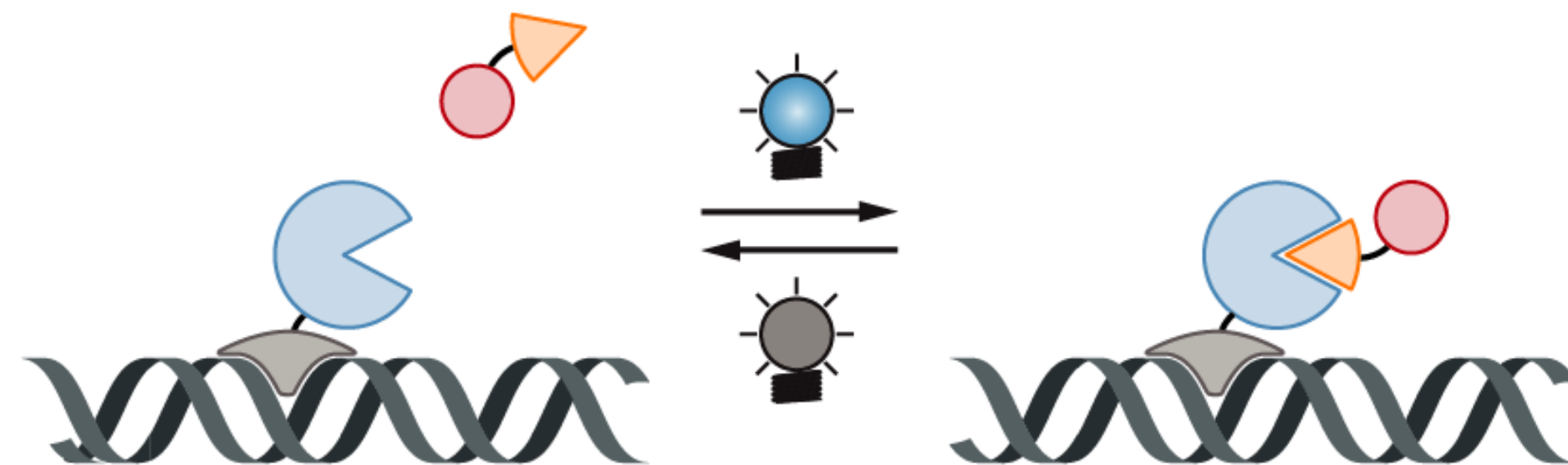
Homodimerization



Translocation to the nucleus



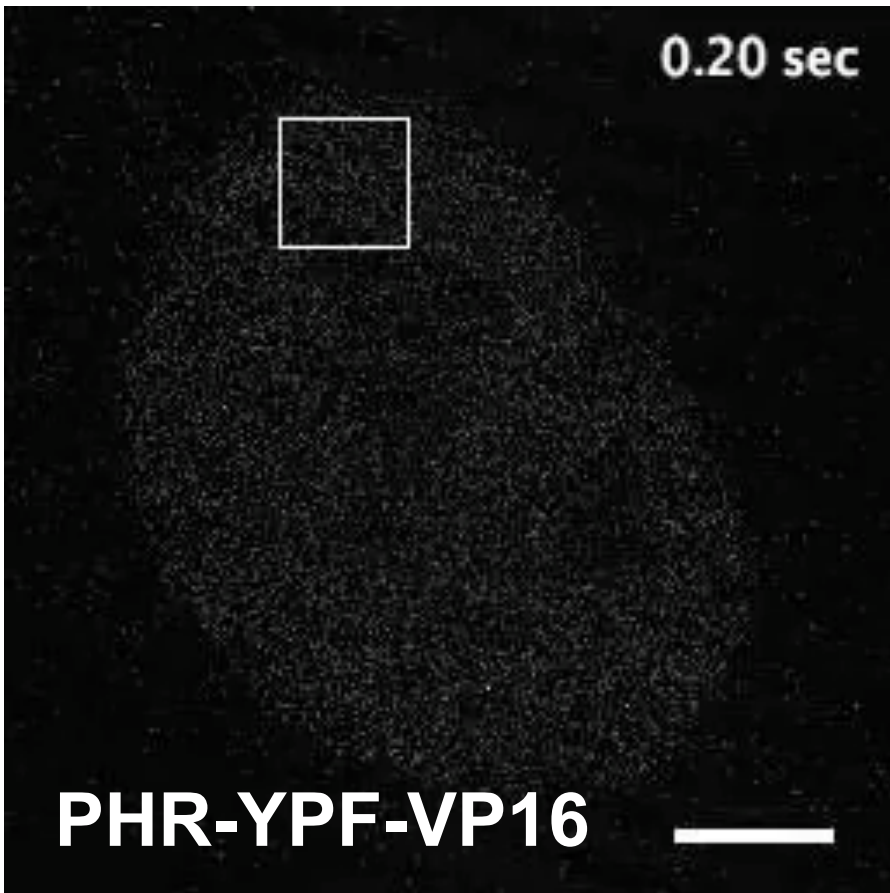
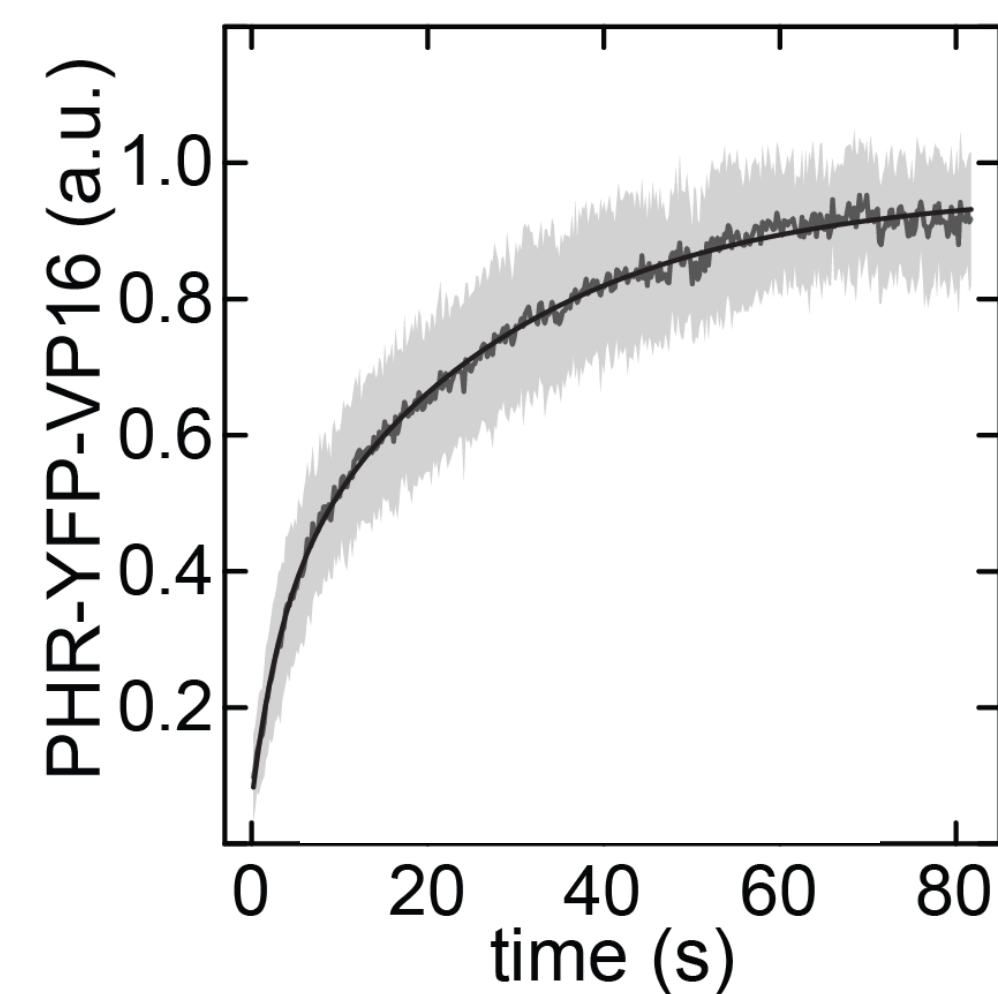
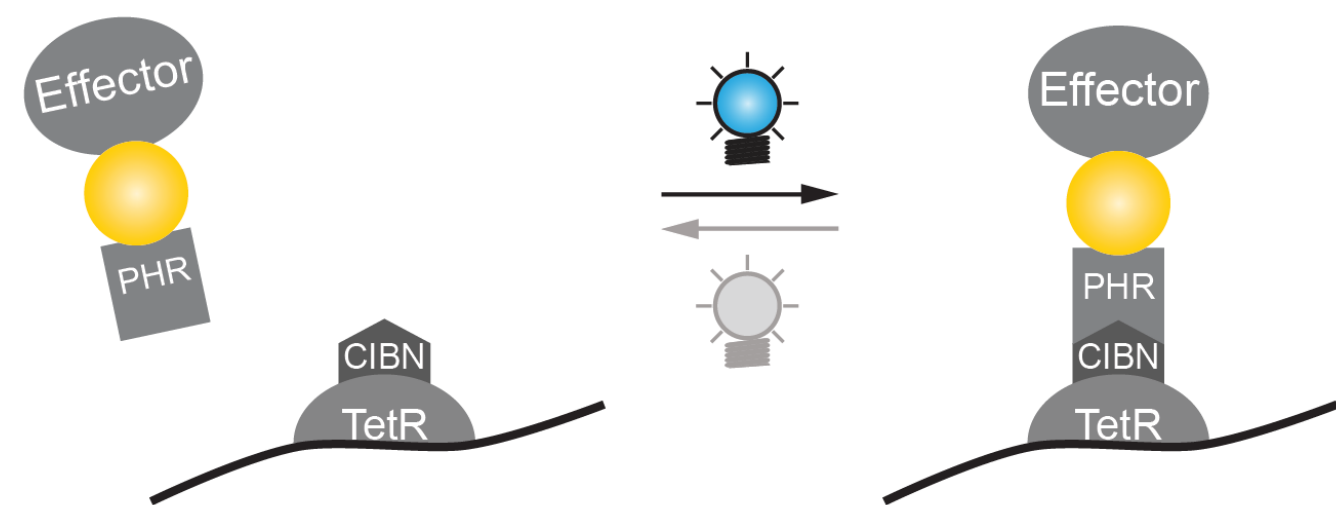
Heterodimerization



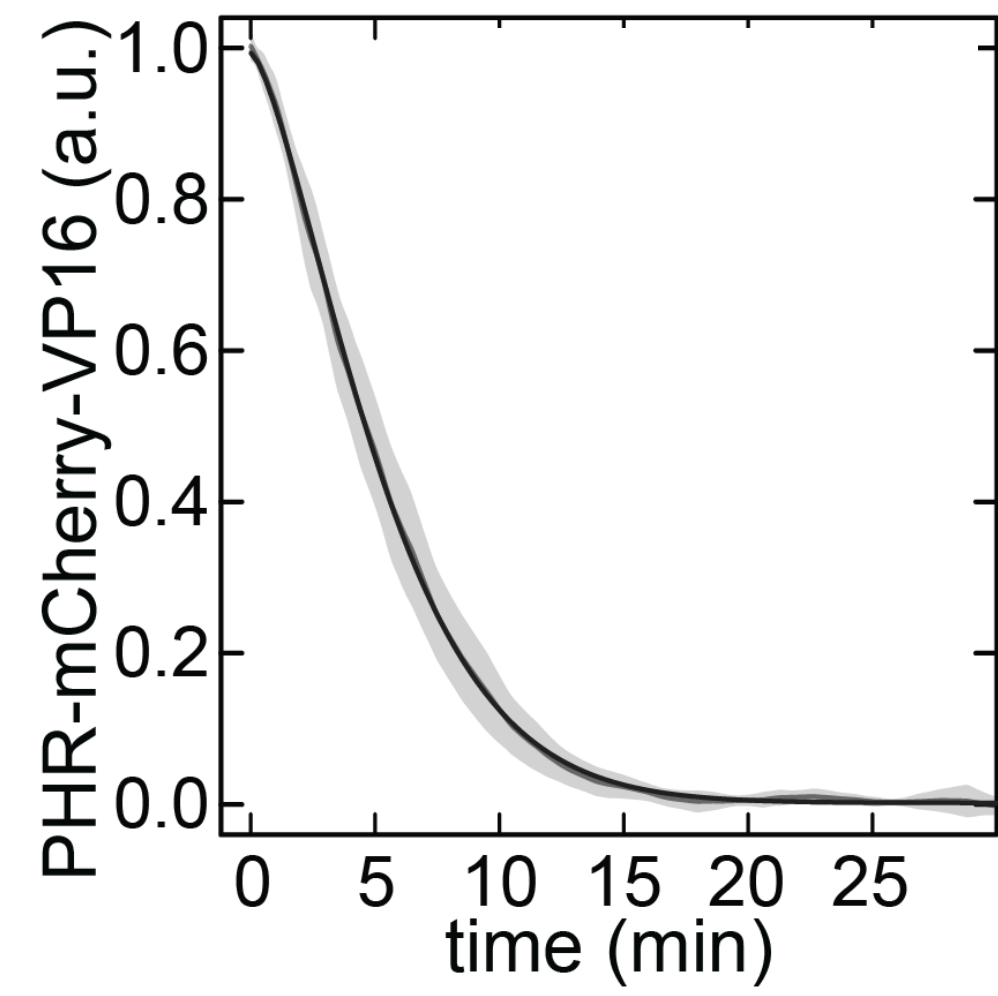
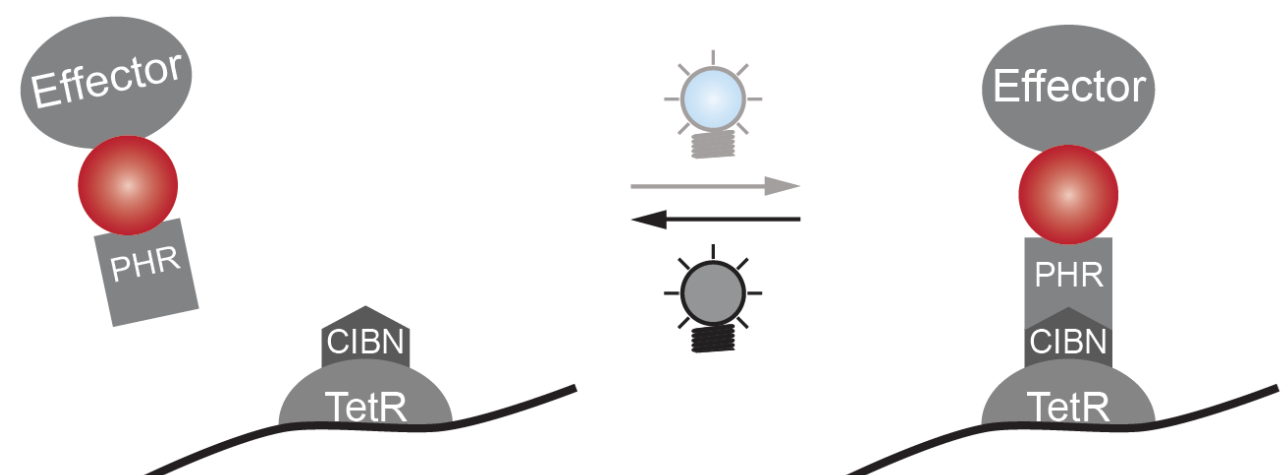
Tischer & Weiner (2014) *Nature Rev Mol Cell Biol* 15, 551-558.

Blue light induced recruitment of transcription activators within seconds

Binding within seconds



Dissociation within minutes

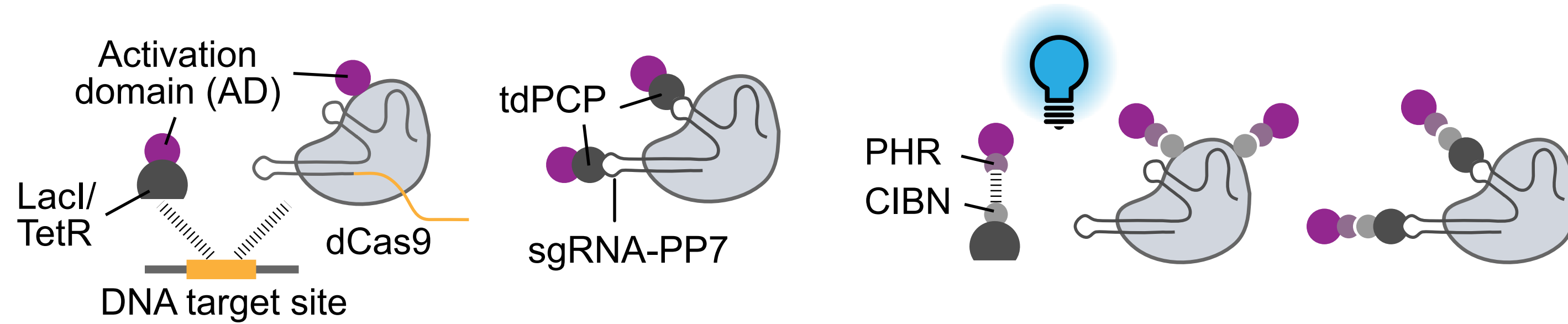


CIBN TetR iRFP713 localizer + PHR-mCherry effector

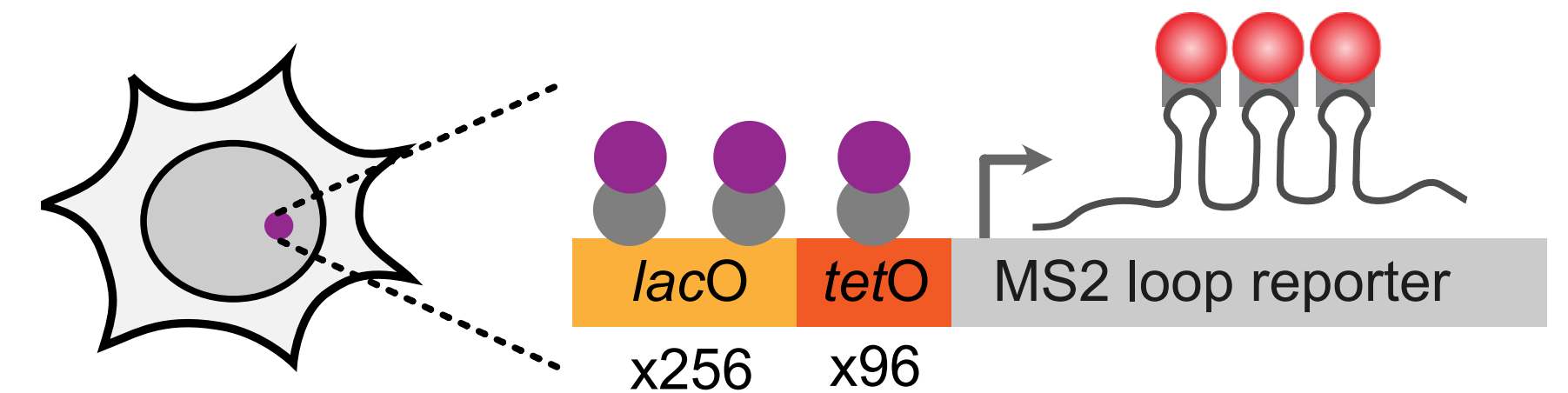
	Effector	Time $t_{on/off}$
Binding	VP16	8.0 ± 5.3 s
	NLS	9.2 ± 3.9 s
Disso- ciation	VP16	4.4 ± 0.8 min
	NLS	5.1 ± 0.5 min

Controlling the recruitment of transcription activators to a reporter array

TF constructs with different ADs und DNA binding modules



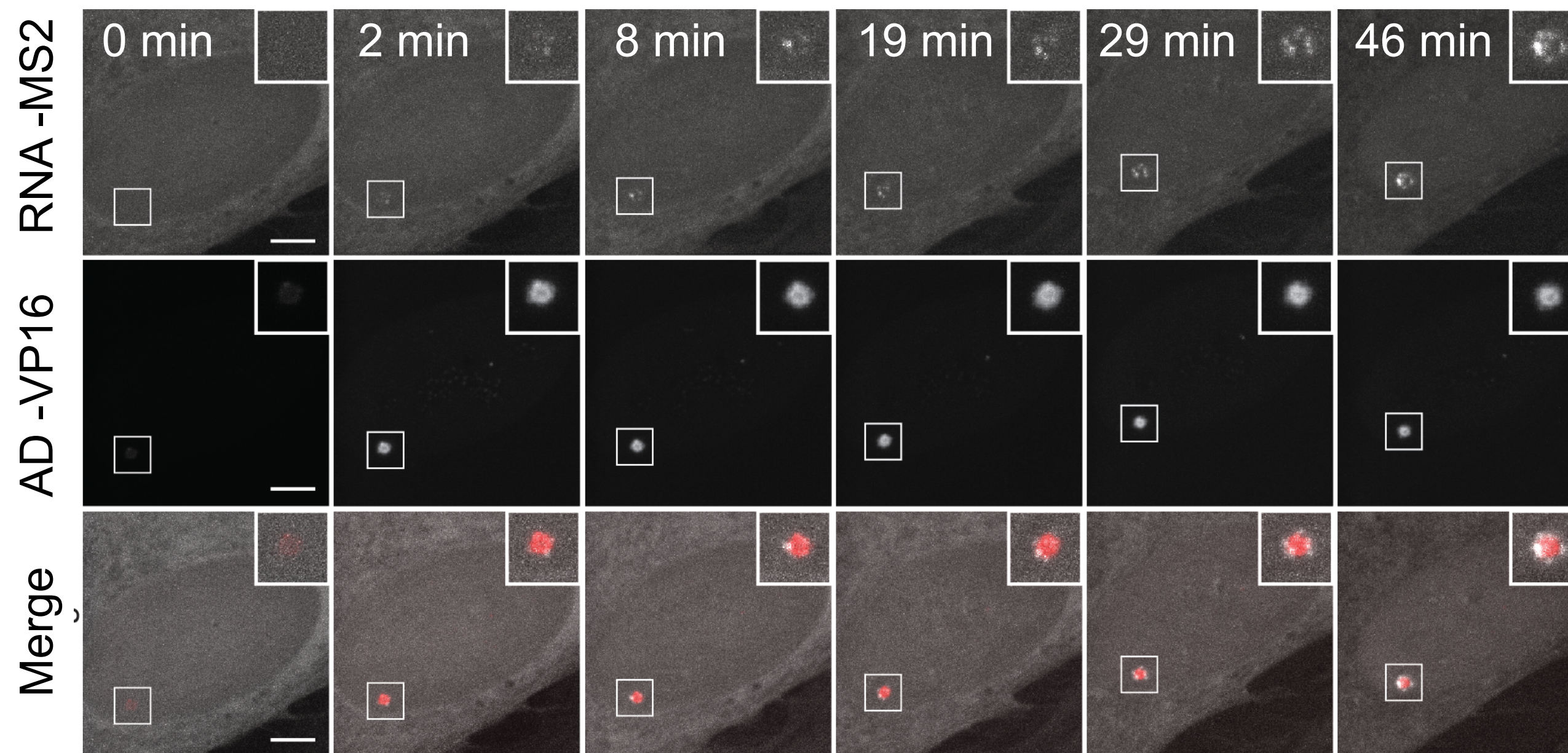
U2OS cell line with reporter array (~200 copies)



Janicki 2004 *Cell*



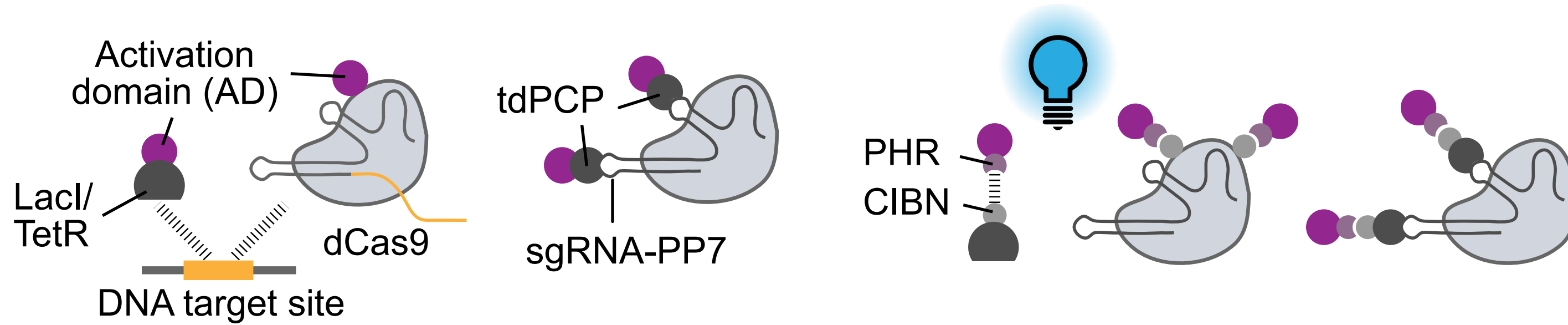
Transcription induction by light induced binding of the VP16 AD



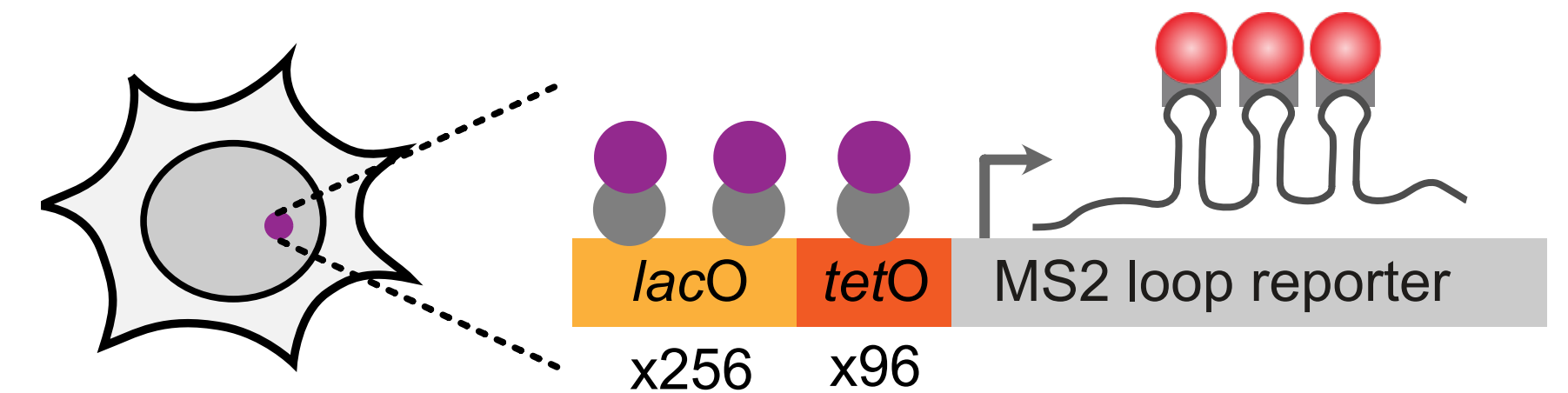
Rademacher 2017 *J Cell Sci*, Trojanowski 2019 *Meth Mol Biol*

Controlling the recruitment of transcription activators to a reporter array

TF constructs with different ADs und DNA binding modules



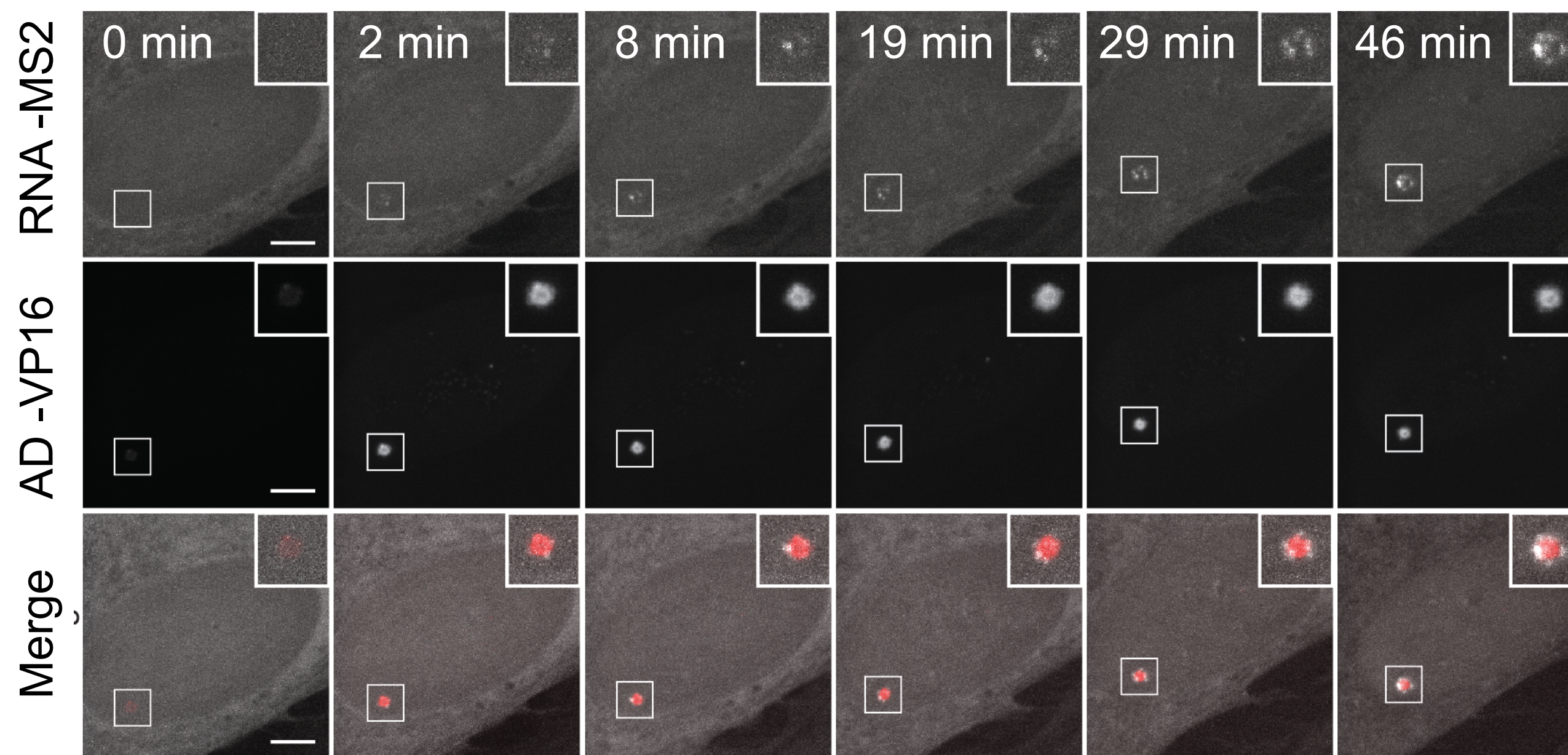
U2OS cell line with reporter array (~200 copies)



Janicki 2004 *Cell*

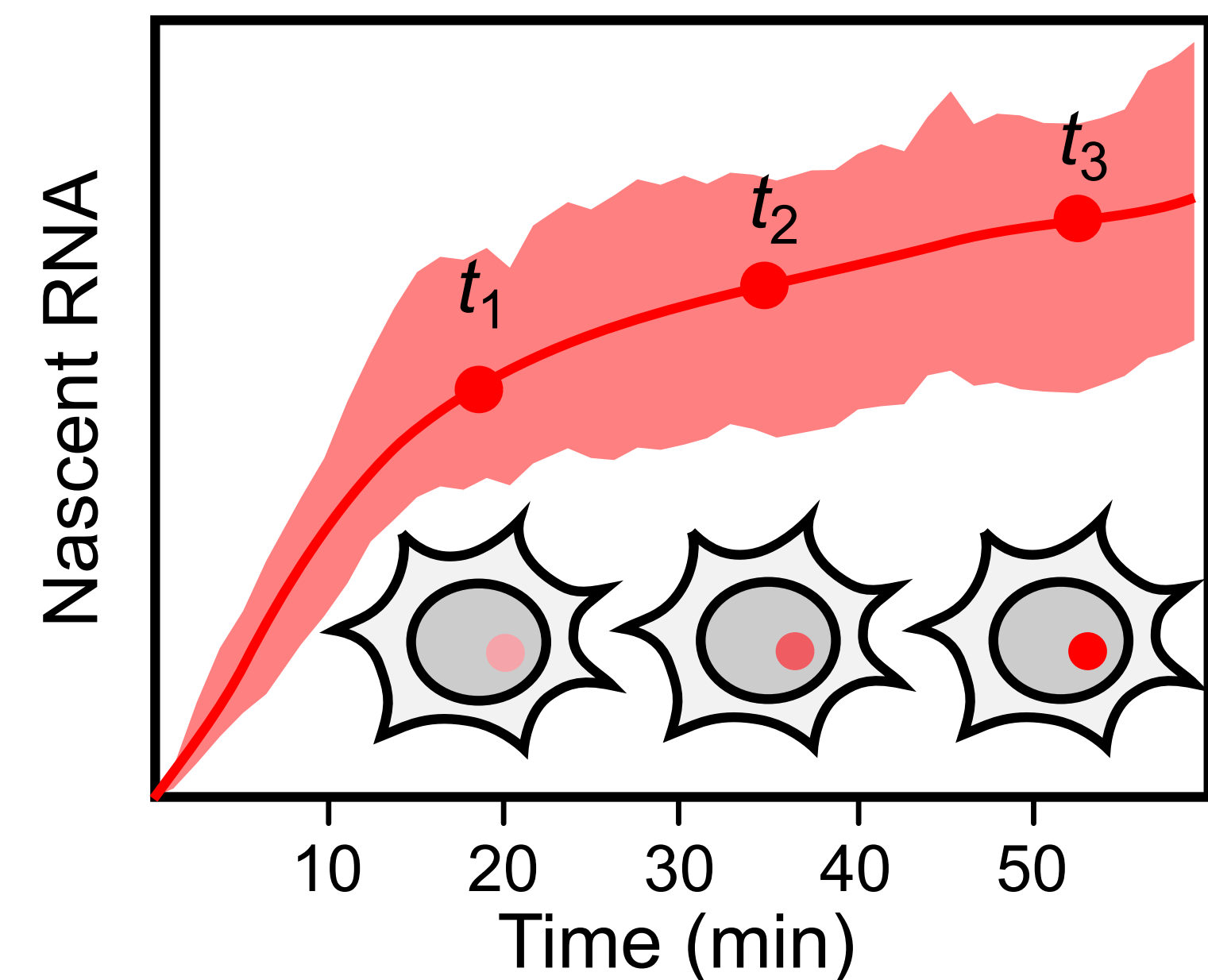


Transcription induction by light induced binding of the VP16 AD



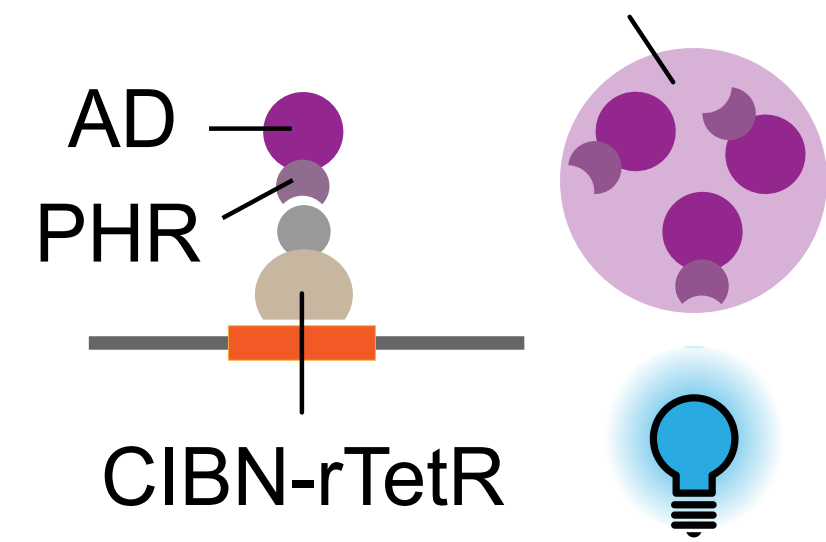
Rademacher 2017 *J Cell Sci*, Trojanowski 2019 *Meth Mol Biol*

Transcription activation kinetics

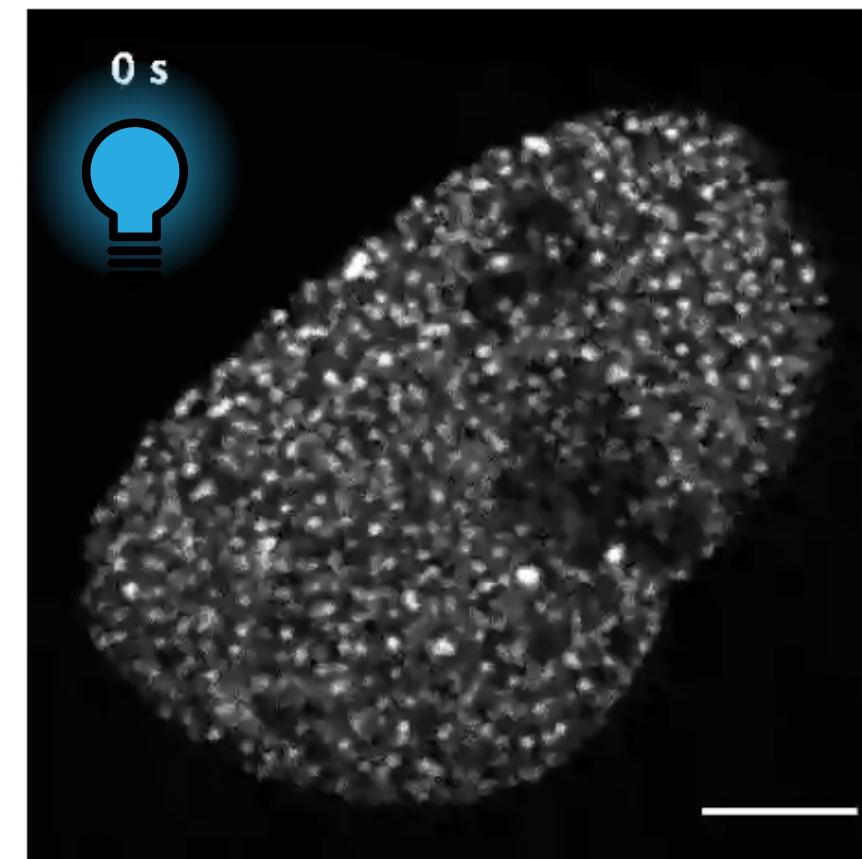


The activation domains (ADs) have different propensities to form optodroplets

PHR-AD optodroplets

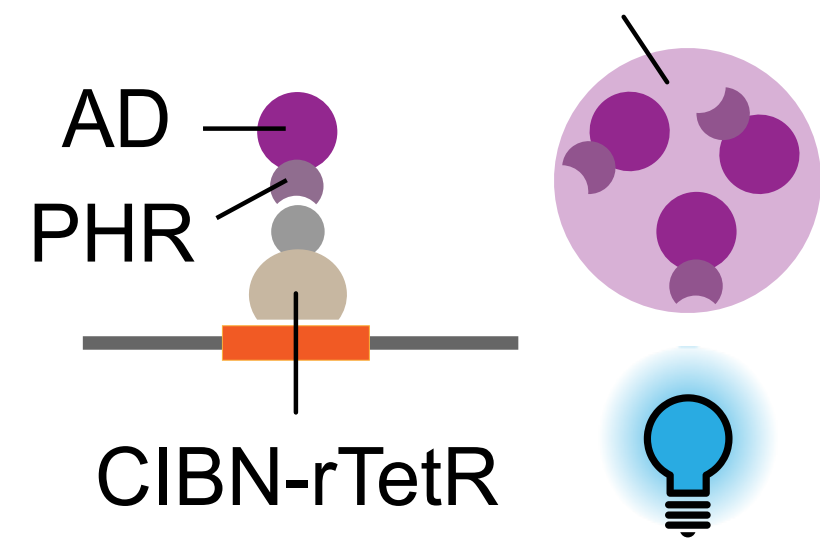


FUSN droplets after
blue light pulse

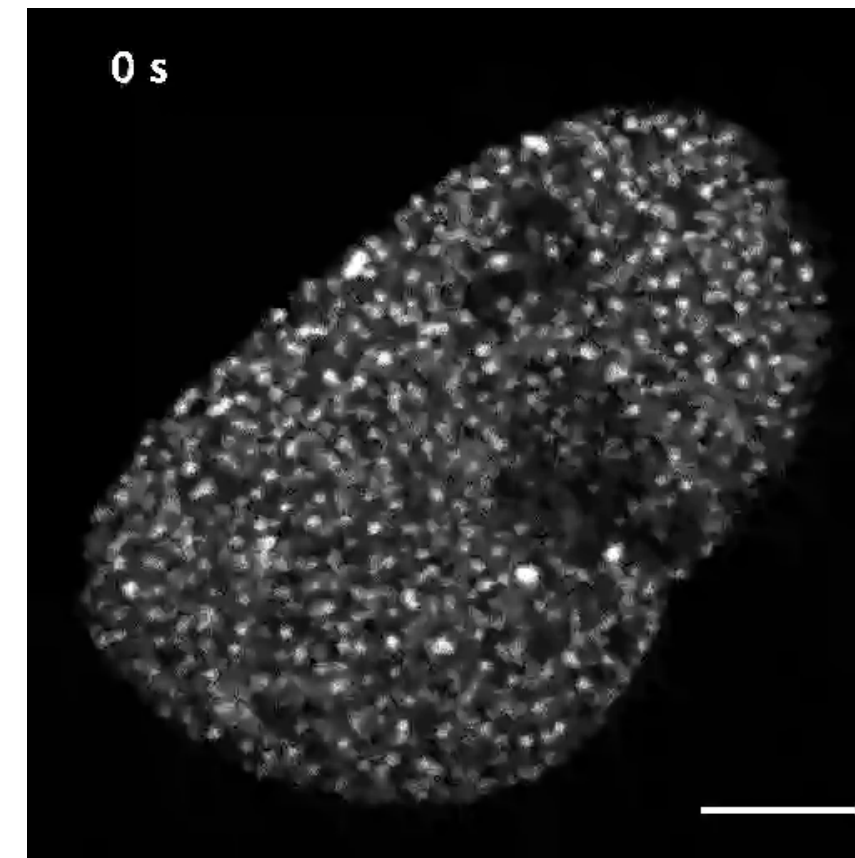


The activation domains (ADs) have different propensities to form optodroplets

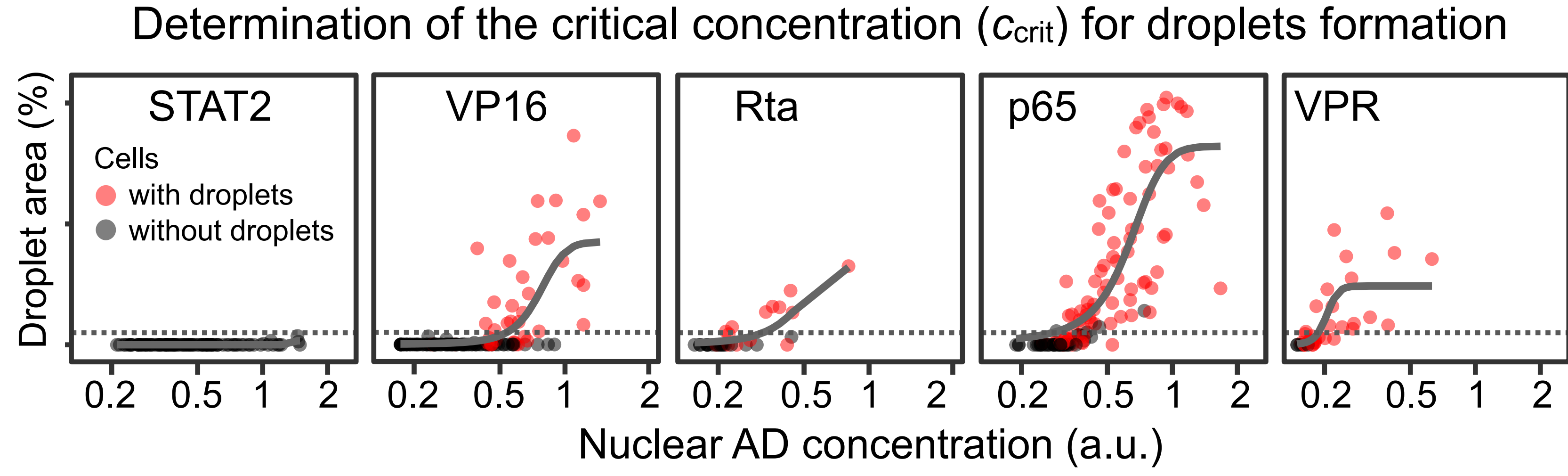
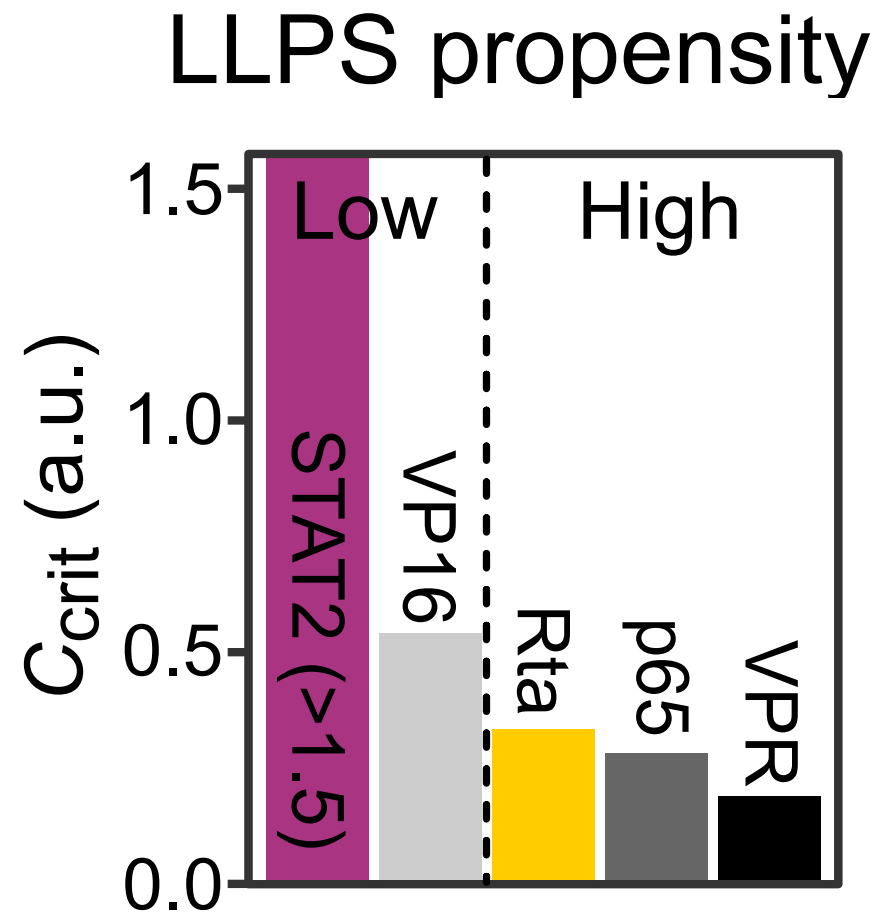
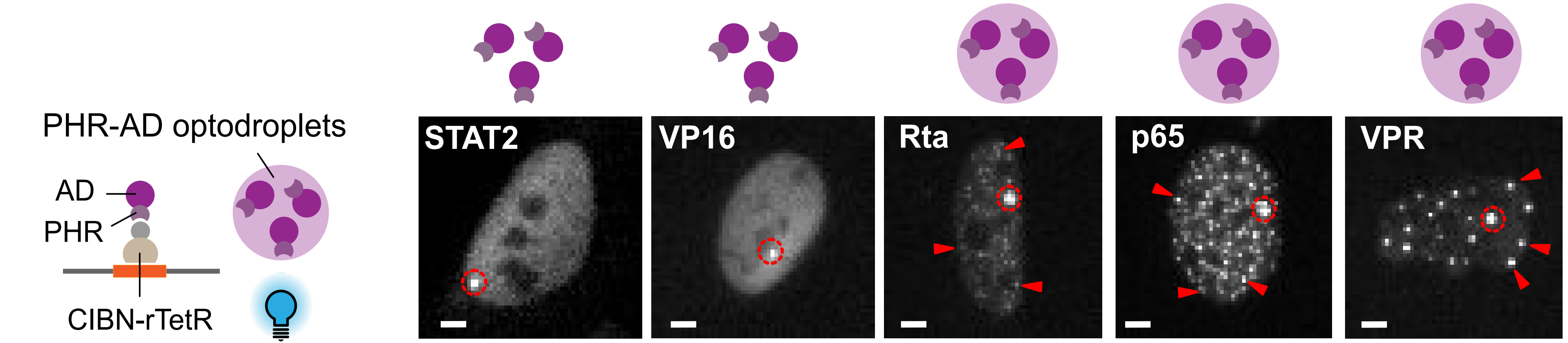
PHR-AD optodroplets



PHR-FUSN droplets
after blue light pulse

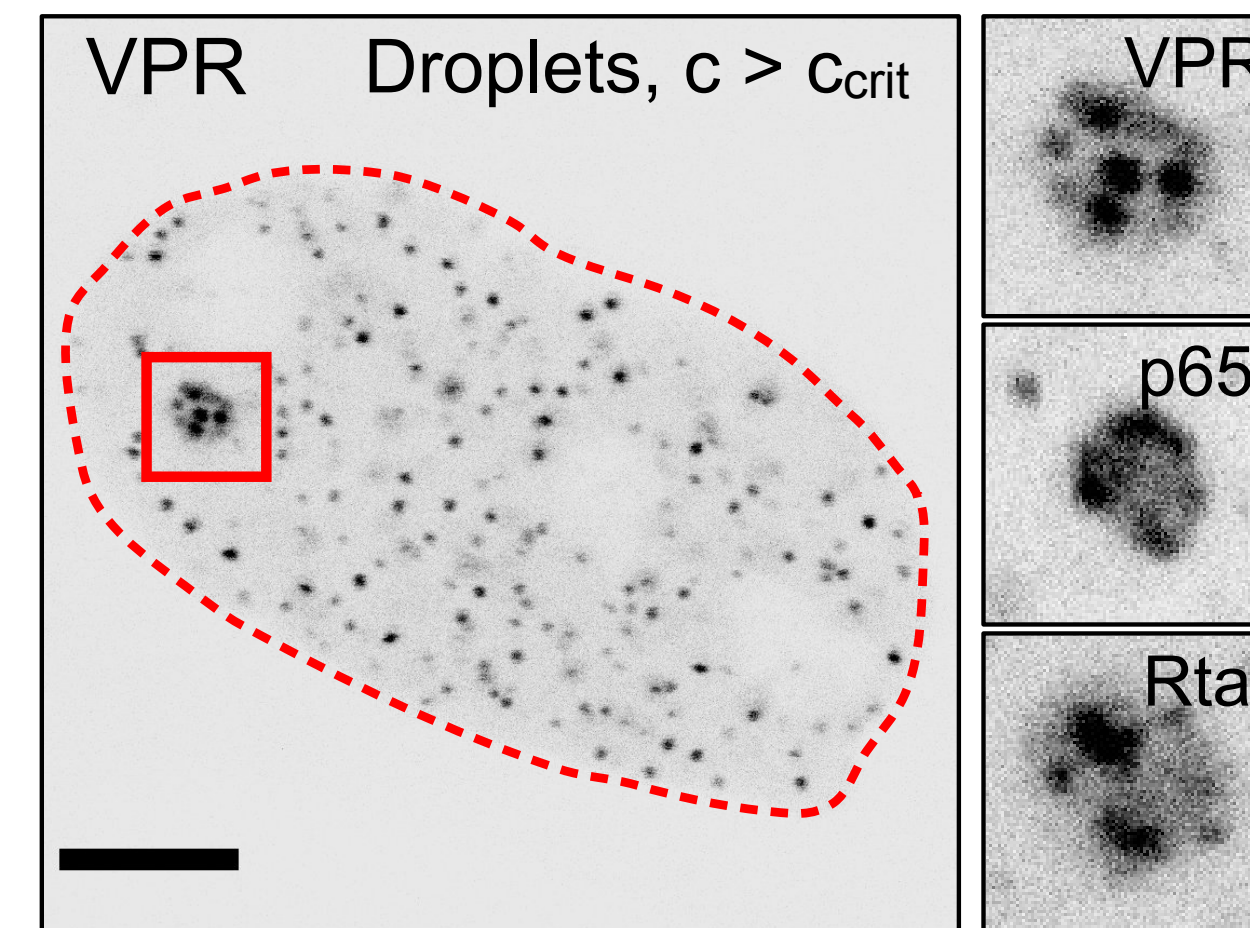
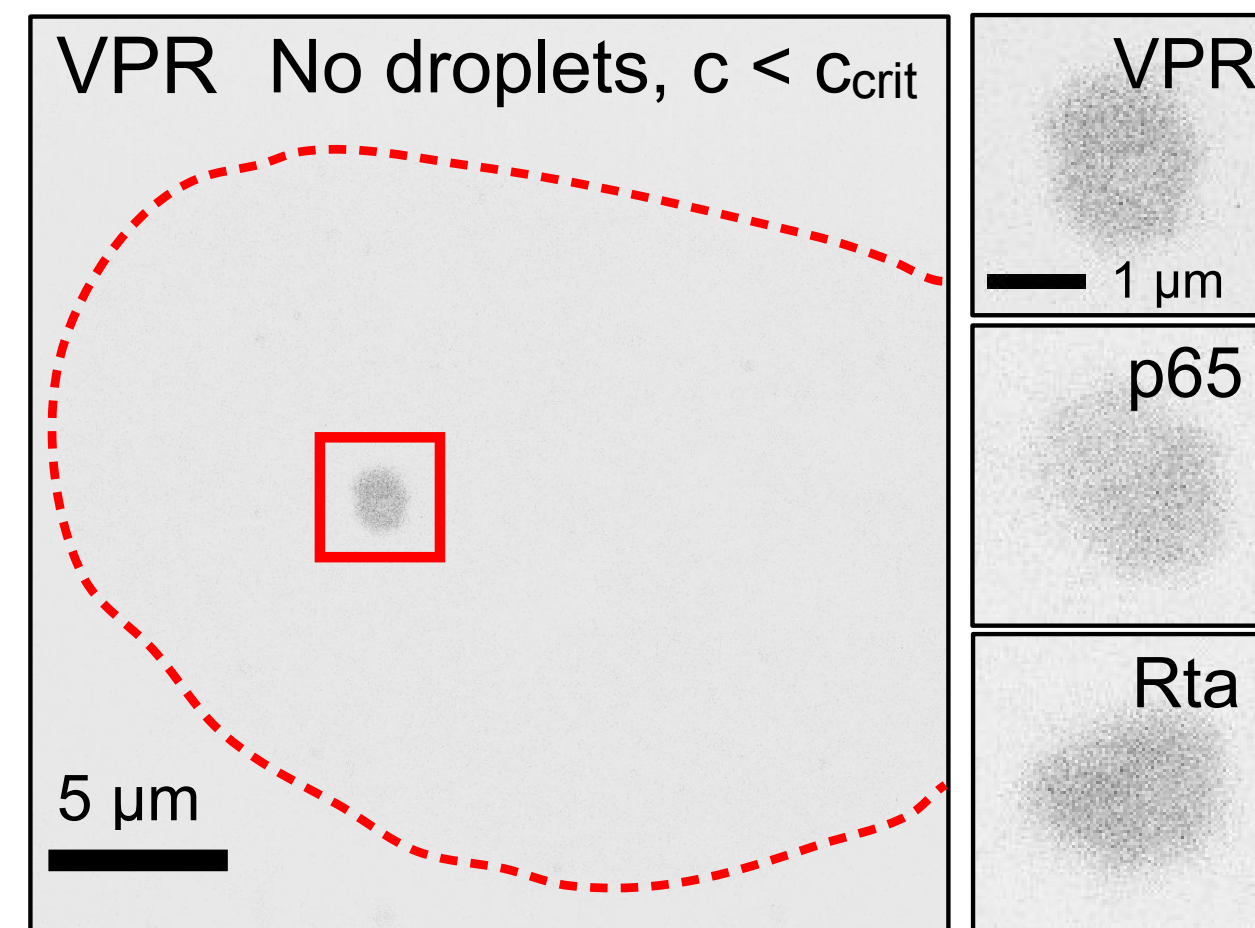


The activation domains (ADs) have different propensities to form optodroplets

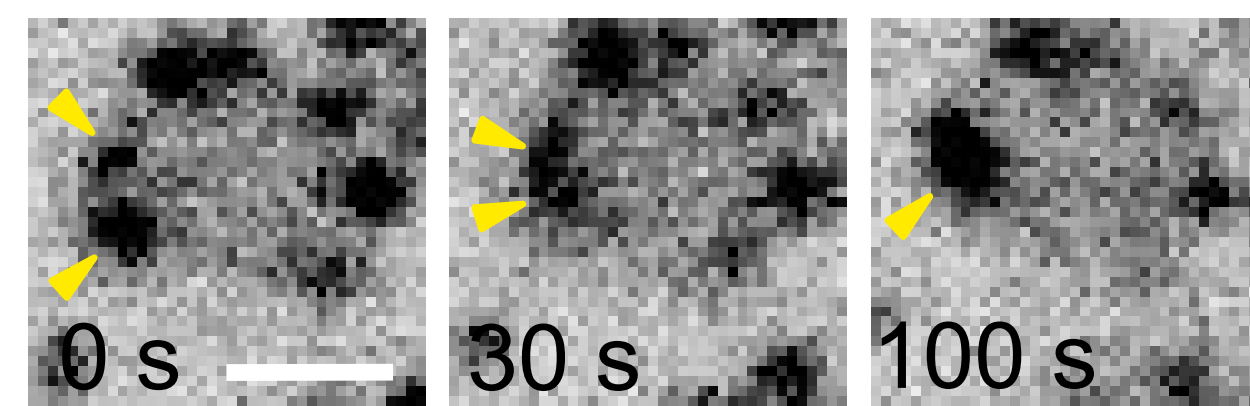


ADs accumulate into liquid like subdomains at the reporter above c_{crit}

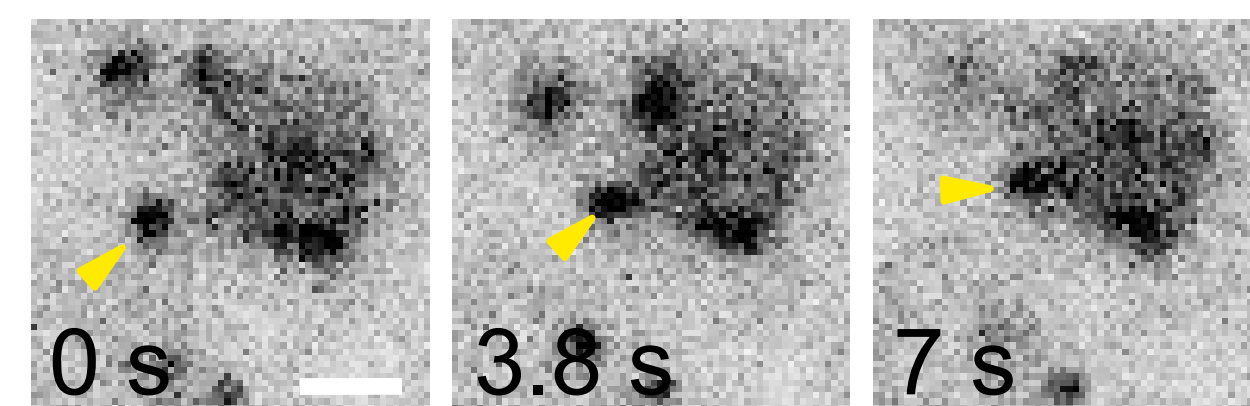
Living cell confocal microscopy



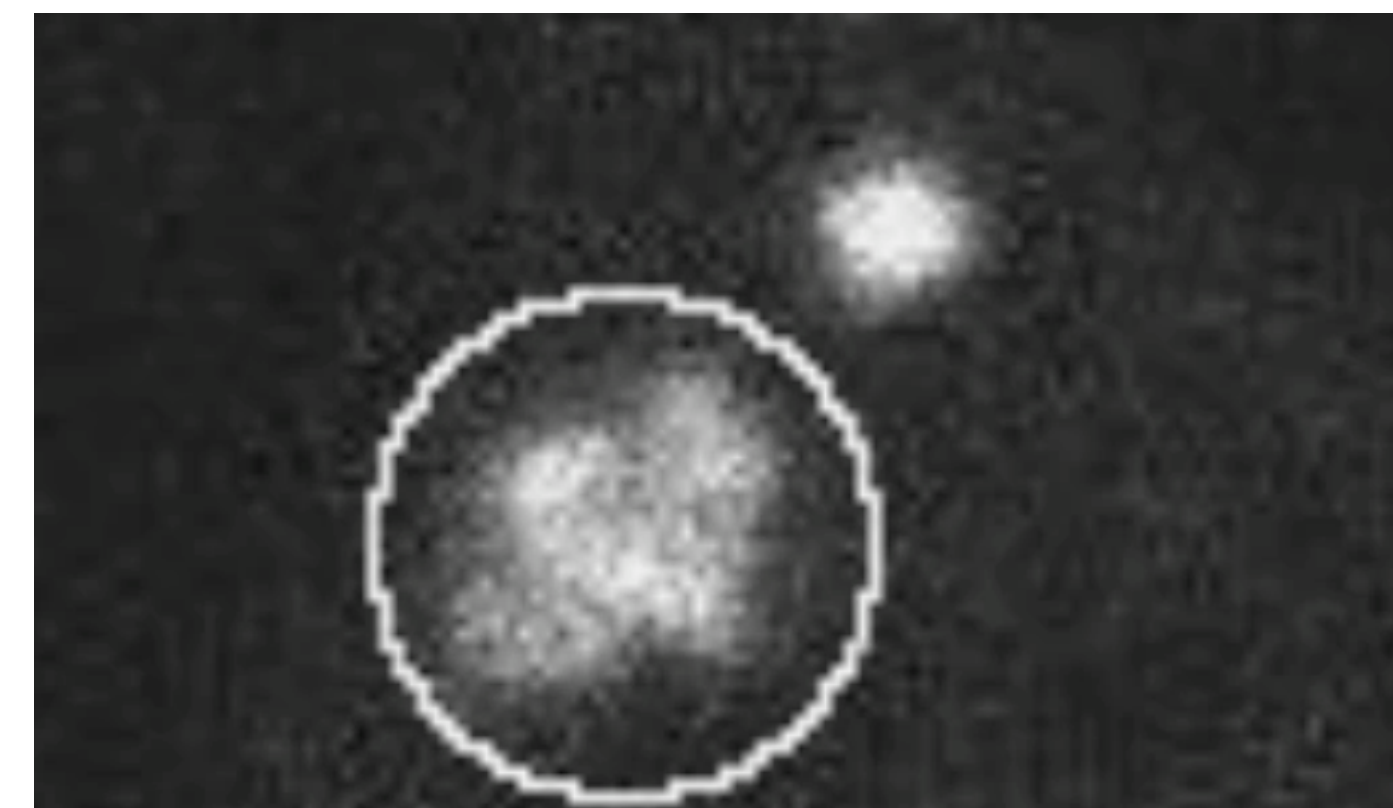
Rta: subdomain coalescence



Rta: fusion with reporter array

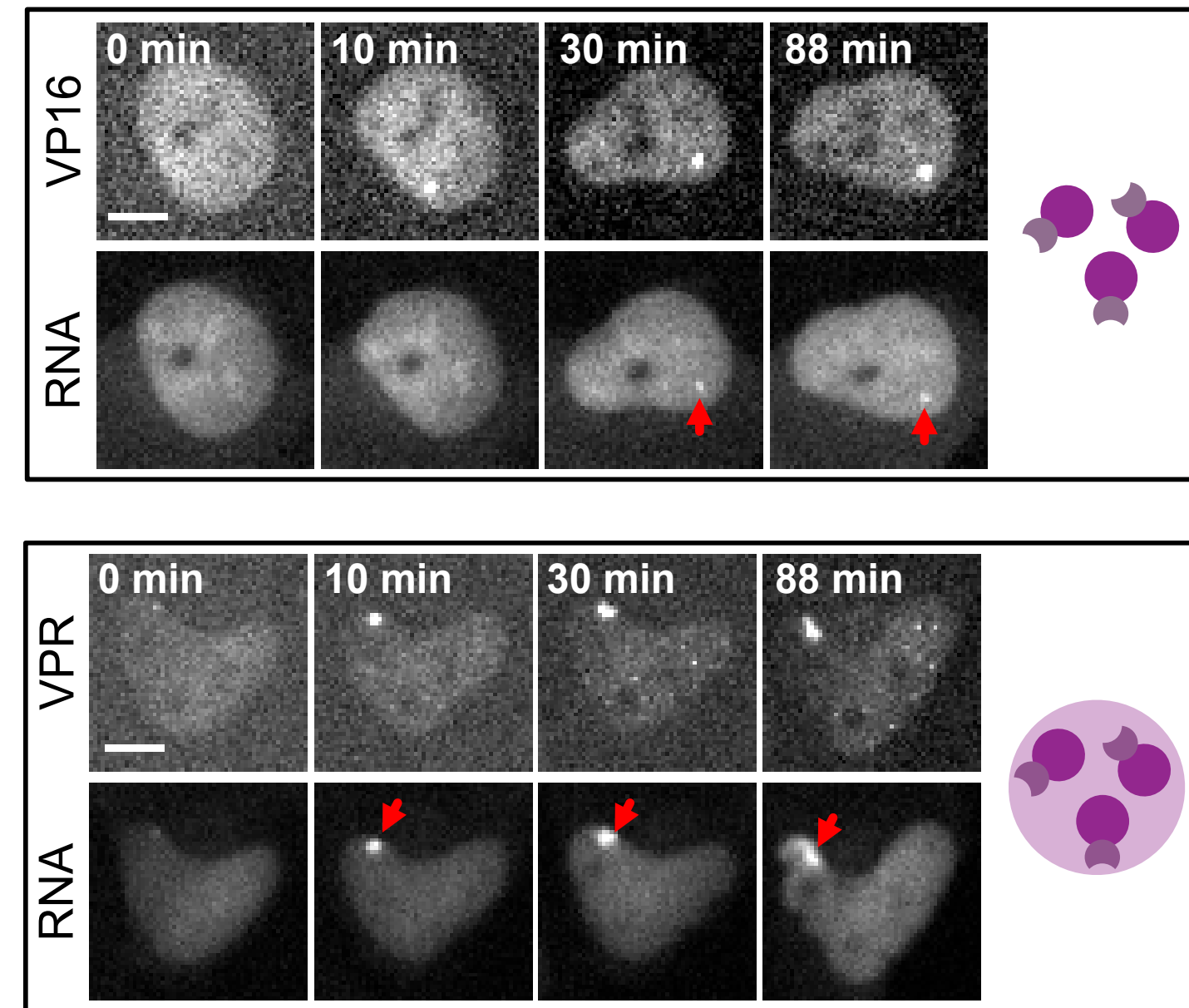
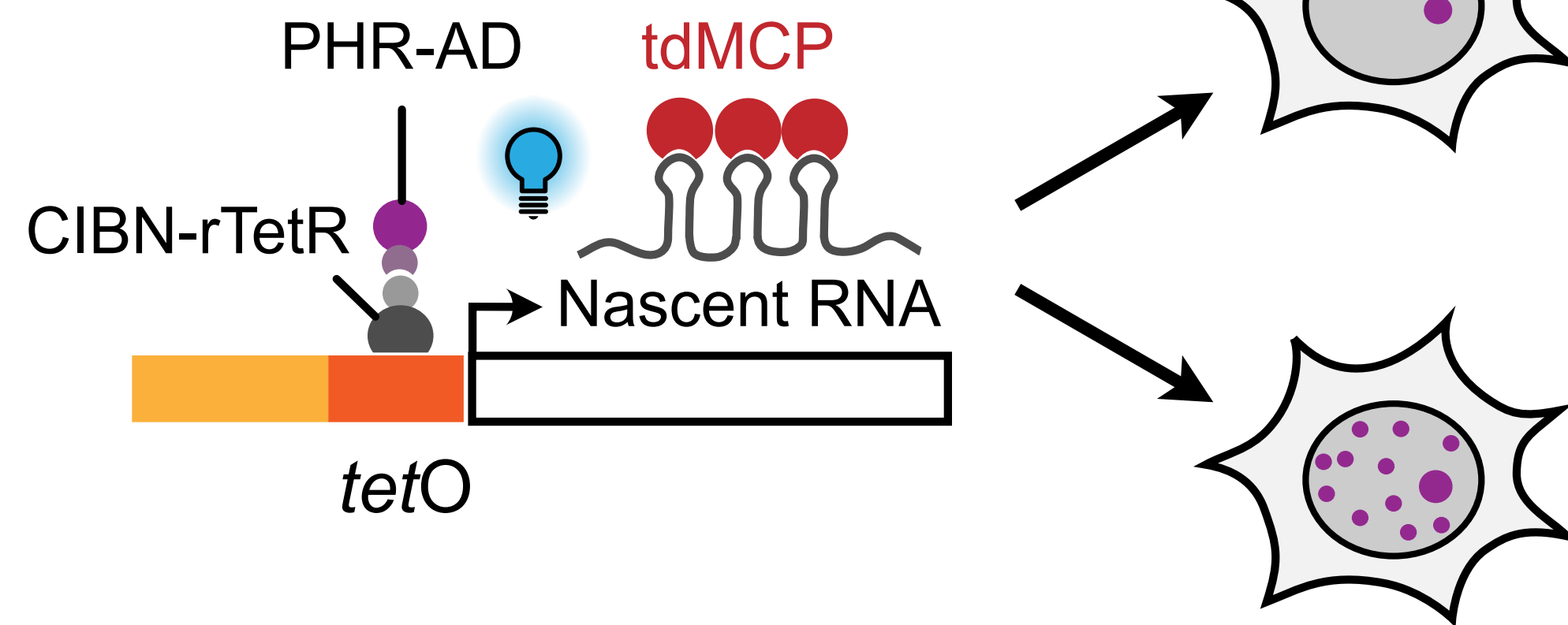


VPR: fusion with reporter array (21 s)

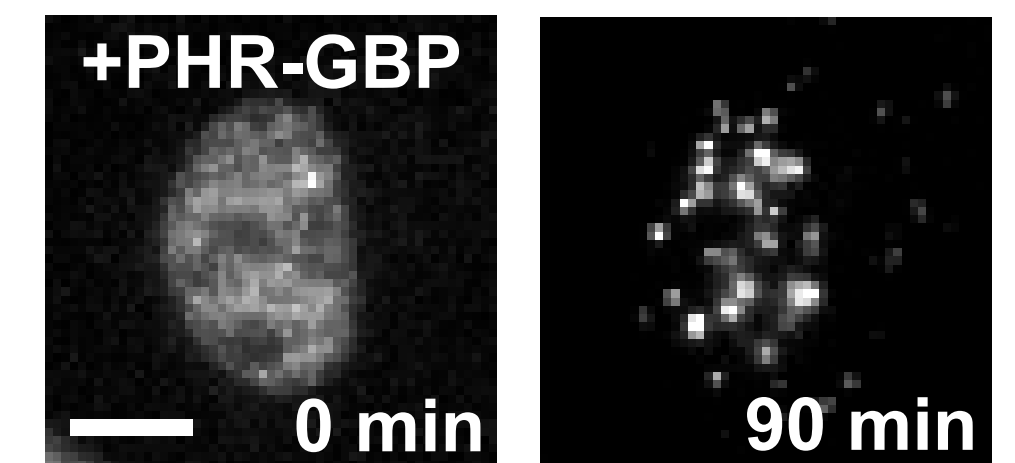
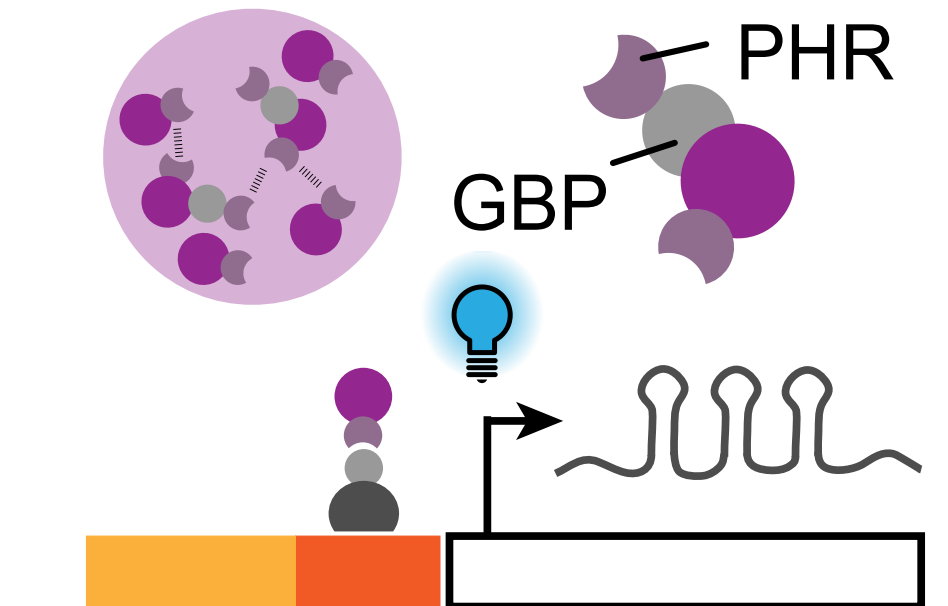


Droplet formation does not enhance transcription activation

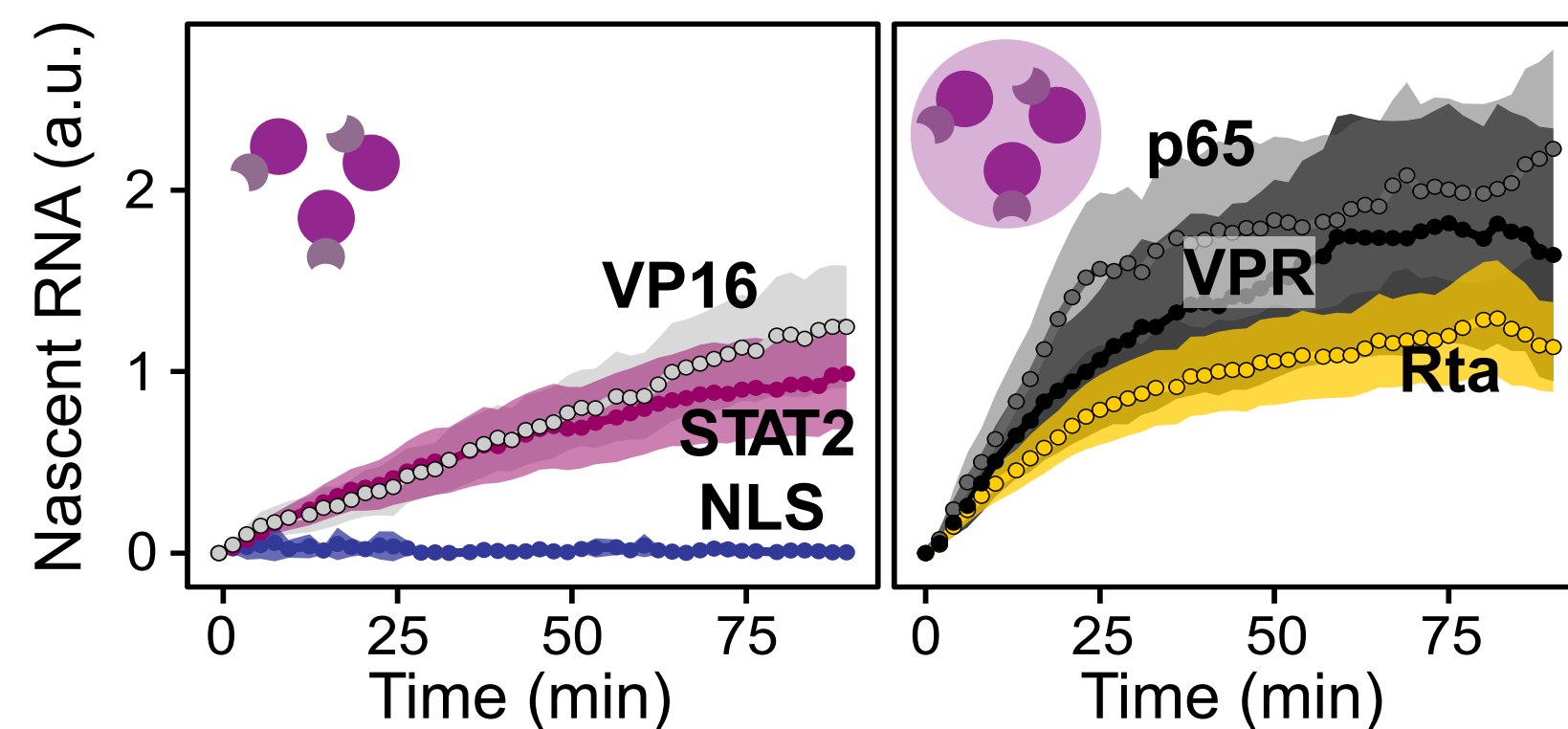
Comparing transcription in cells with/without droplets



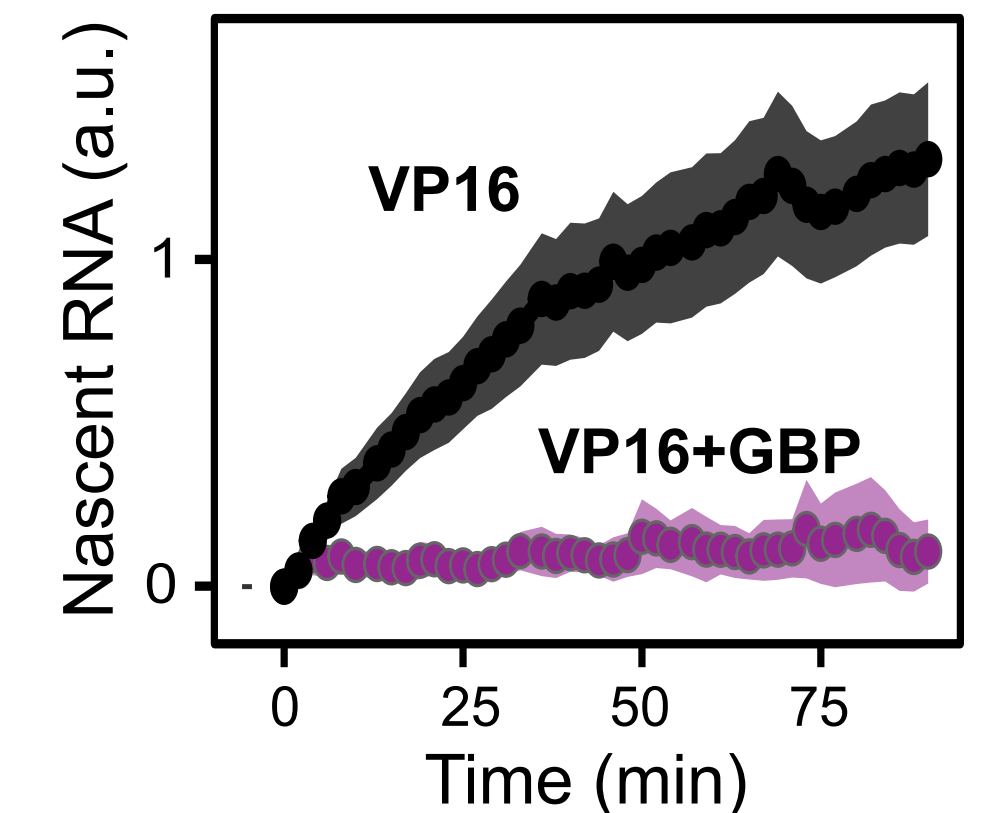
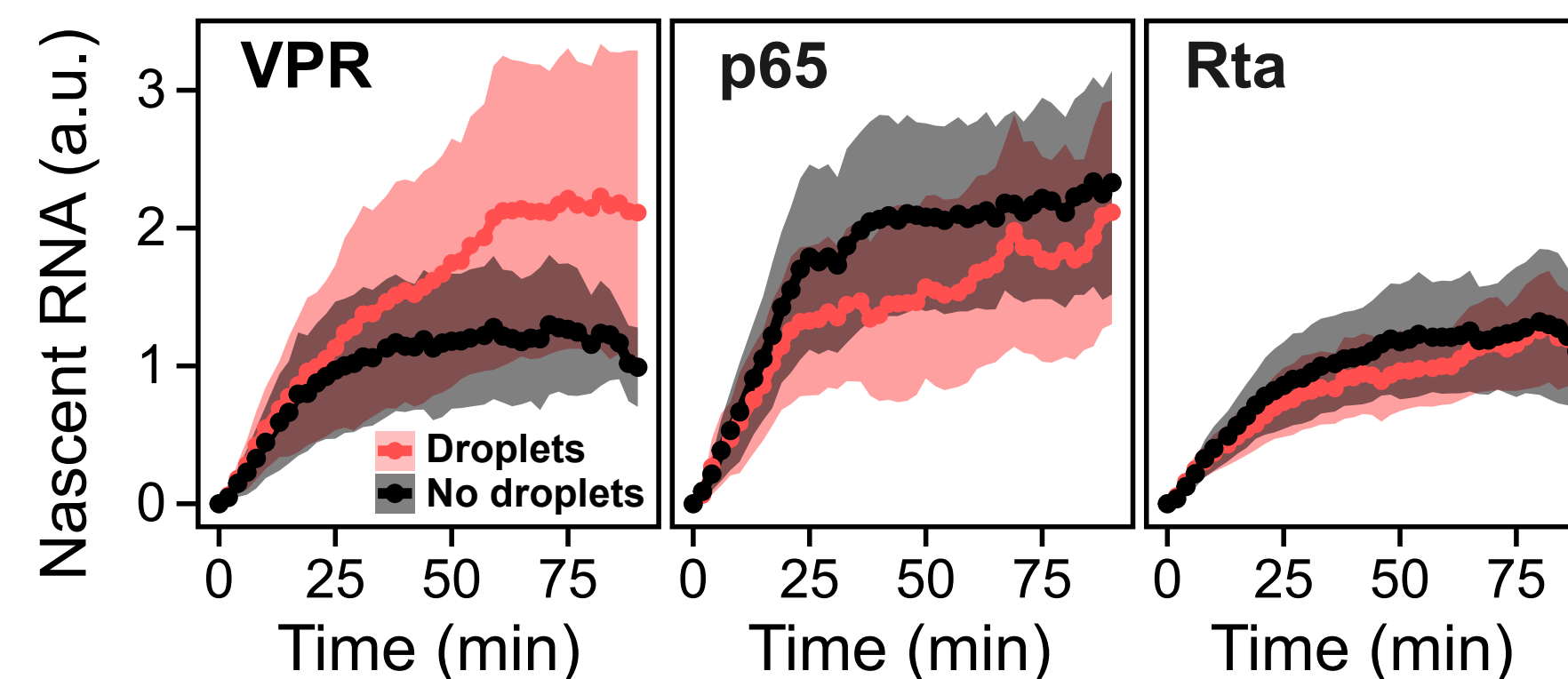
Repression of VP16 by enhanced droplet formation



Nascent RNA kinetics

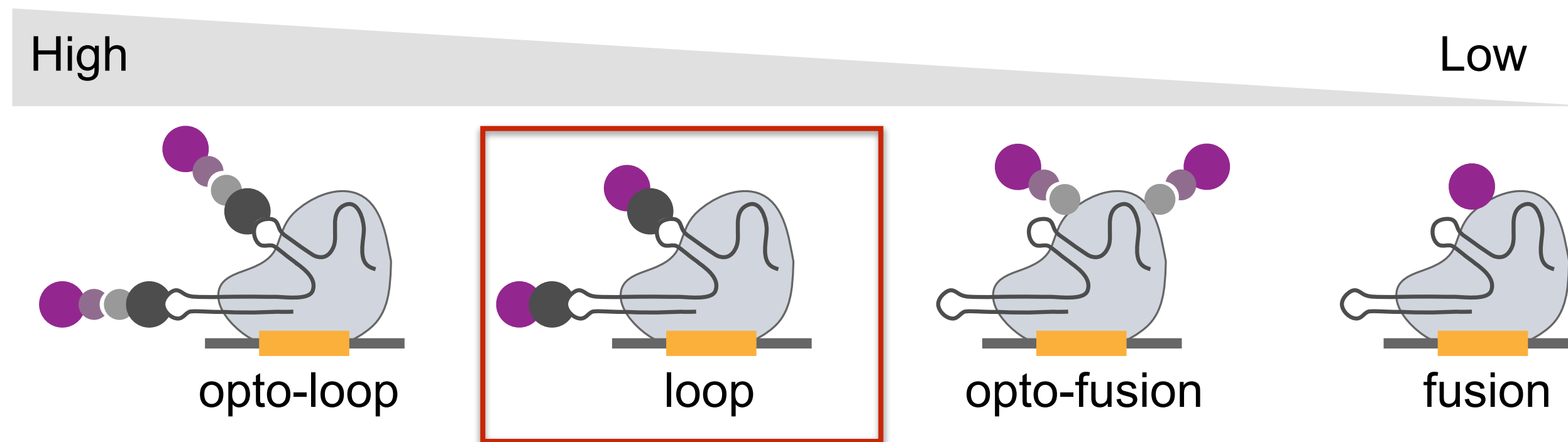


Nascent RNA kinetics above/below c_{crit}

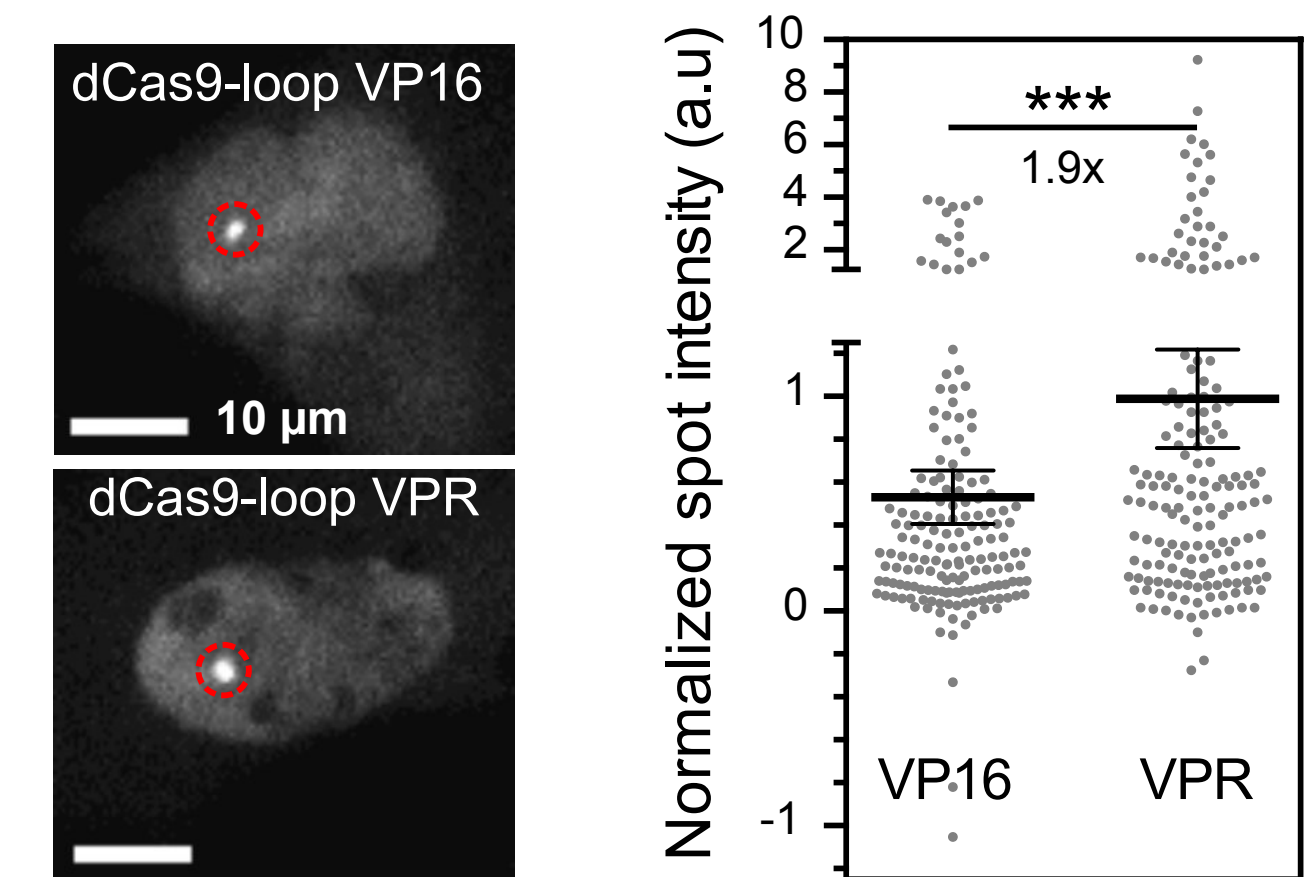


TF binding is stabilized by higher AD multivalency/LLPS propensity

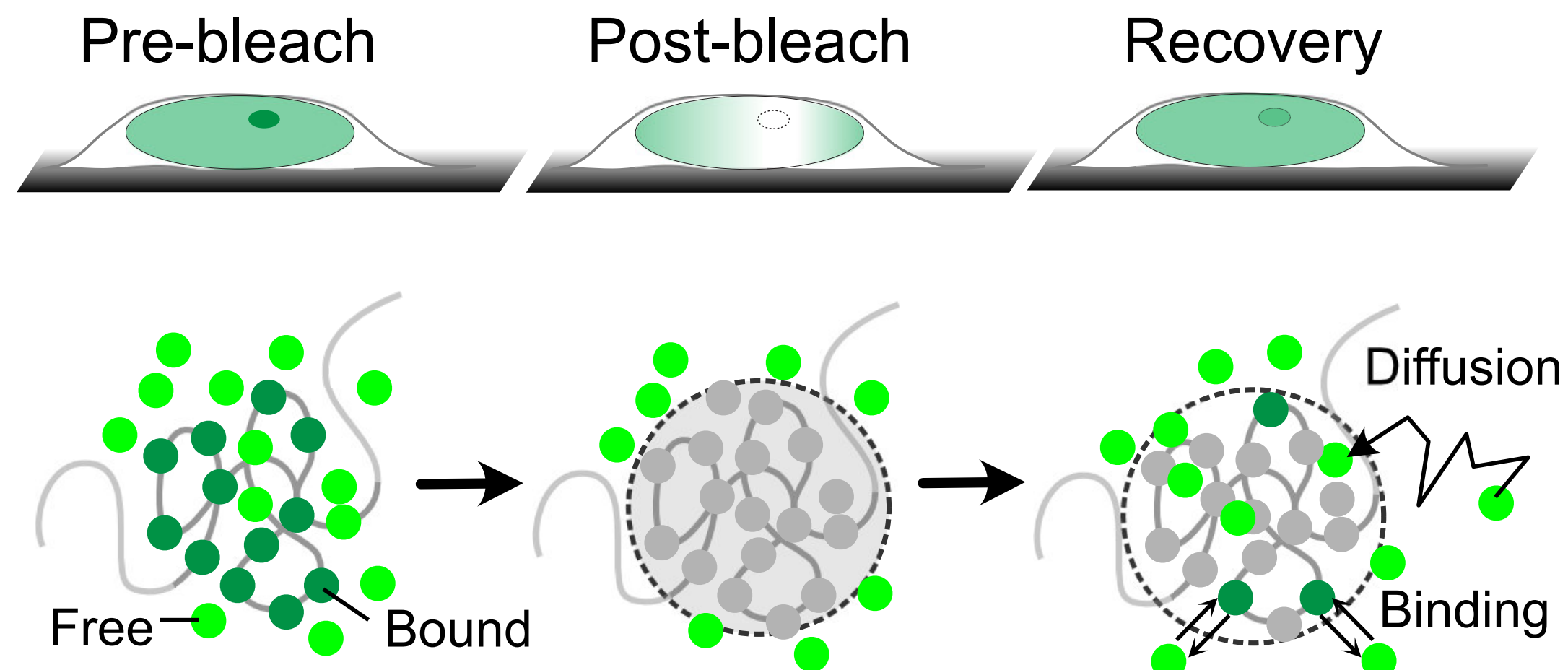
Activation domain turnover



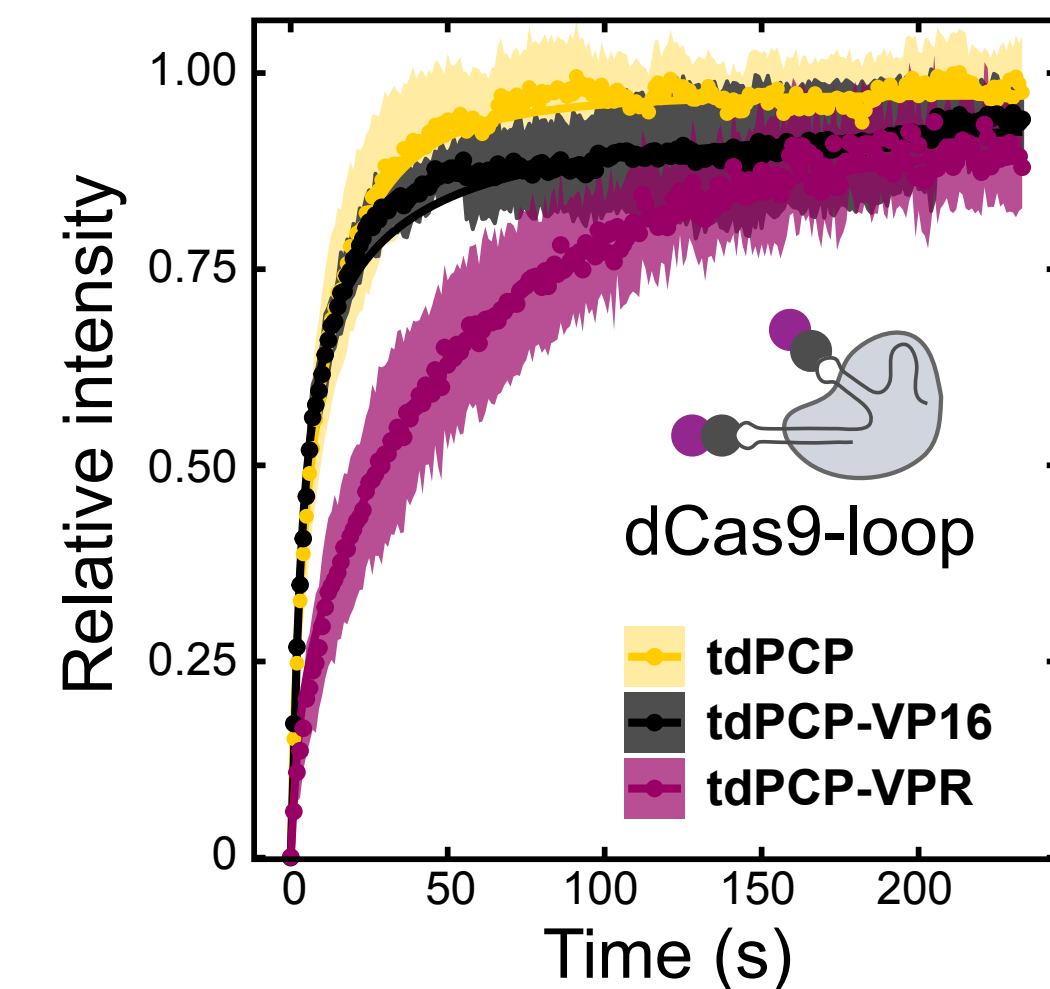
2-fold more VPR than VP16 at the array



Fluorescence recovery after photobleaching FRAP



Longer residence time of VPR than VP16



Conclusions: transcription activation is enhanced by multivalent interactions independent of phase separation

