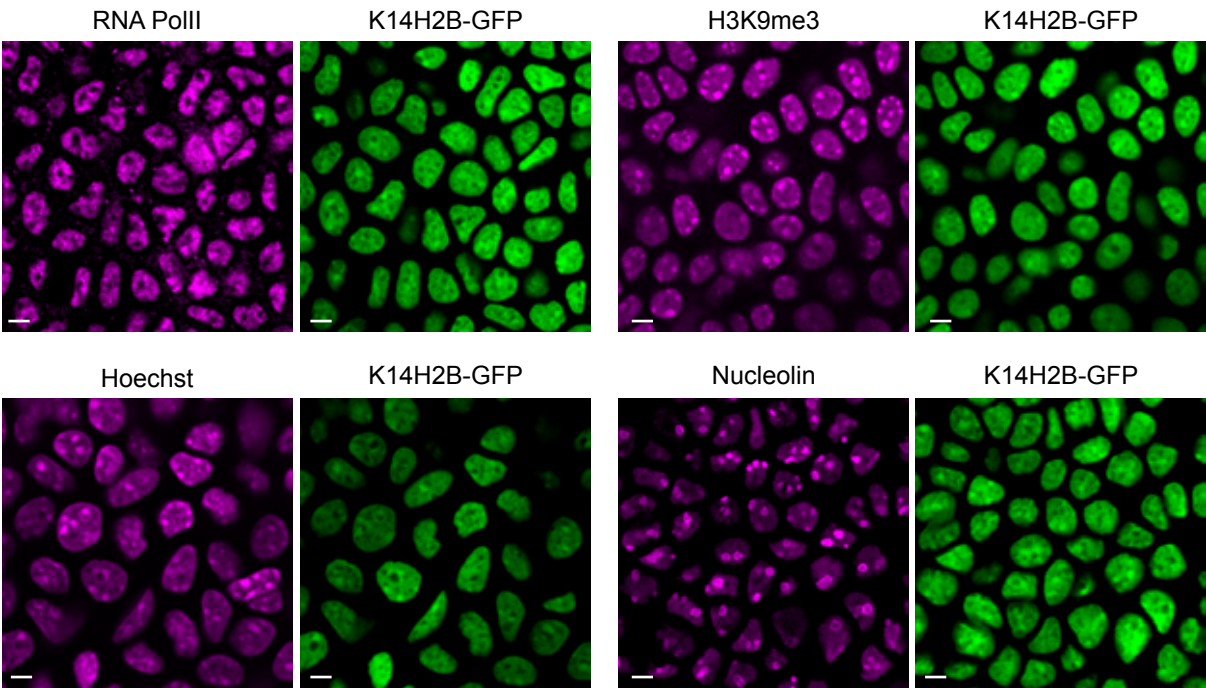
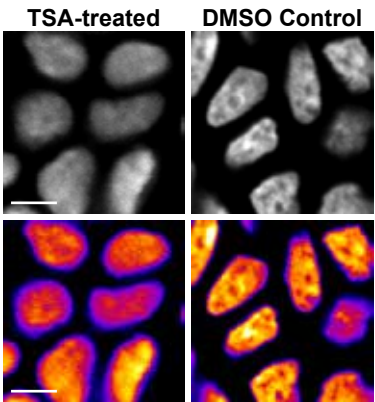


**A**



**B**



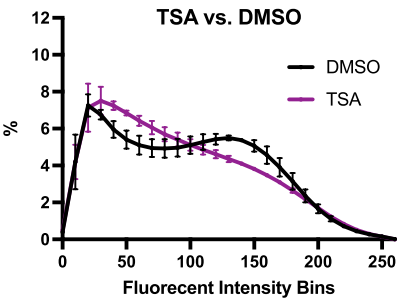
**C**

Example comparison of data organization between RNA-seq and chromatin compaction

	Gene 1	Gene 2	Gene 3	...
Cell 1	0 reads	3 reads	12 reads	
Cell 2	4 reads	1 read	3 reads	
Cell 2	0 reads	0 reads	8 reads	
...				

	Bin 1	Bin 2	Bin 3	...
Nucleus 1	0.4 %	2.0 %	1.8 %	
Nucleus 2	1.1 %	3.4 %	4.1 %	
Nucleus 3	1.6 %	0.9 %	6.2 %	
...				

**D**



**D'**

