Imaging Nanoscale Chromatin compaction in vivo using FRET approach

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### Genome Organization



### Genome Organization



## In vivo single-cell approach

# FRET resolution in the range of the nucleosome array



Lleres et al. 2007. Current Protocols in Cytometry 12.10.1-12.10.19,2007

### FLIM-FRET principle



## FRET resolution in the range of the nucleosomal compaction



## Nucleosome-nucleosome interactions contribute to the FRET readout

FRET (%) map



#### Localisation & Quantification of chromatin domains compaction in living cells

HDAC inhibitor (TSA)



Chromatin domain compaction is controlled by histone modifications

Llères et al. JCB. 2009

### Maintenance of chromatin domains along mitosis



HeLa H2B-GFP

HeLa H2B-GFP/mCherry-H2B









#### Llères et al. JCB. 2009

Measuring chromatin compaction in an organism

#### Caenorhabditis elegans as an organism model



High degree of conservation

#### Hermaphrodites c. elegans 2FPs-H2B



#### Frøkjær-Jensen et al, Nat Methods. 2012

#### FLIM-FRET measurements in C.elegans gonads



### Highest chromatin compaction in meiosis

**Fluorescence Lifetime** 





C.elegans 2FPs-H2B

Fluorescence Intensity GFP::H2B mCherry::H2B 2.0 2.8  $\tau$  (ns) **Fluorescence Lifetime** 

 $\tau$  (ns)

2.8

2.0



Llères D, Bailly A et al. Cell reports. 2017

## Distinct chromatin domain structures in pachytene-stage chromosomes



FRET efficiency (%)



**Relative FRET Fraction** 140 Sub-low 120 100 sub-Low Pixels Frequency 100 Low FRET 80-Pixels Frequency (%) Intermediate FRET 80 📕 High FRET 60 60-40 20 40-Intermediate Hig OW 0 20-60 80 20 FRET efficiency (%)

0%

Ongoing work: Modeling Chromatin domain structures in pachytene-stage chromosomes



- Low FRET interactions
- High FRET interactions

- Fraction of donor interacting with acceptor



Coarse grained modeling



Open loops

#### What do these compaction states correspond to ?

#### How are they controlled ?

#### HP1 homolog is essential for heterochromatin compaction



Llères D, Bailly A et al. Cell reports. 2017

Silenced repeated sequences acquired structurally compacted chromatin controlled by HP1



#### Structural regulators of chromosomes: Condensin complexes



Condensin I

Controls chromosomes in early meiosis

#### Condensin II



Restructure meiotic chromosomes after pachytene exit

Adapted from Strome S et al. CSHLP. 2014.

#### Condensin complexes differentially regulate compaction



#### Relationship between Condensin complexes and lampbrush meiotic architecture



Chromosome length (pixels)

#### Summary



#### Conclusions

- Meiotic chromosomes in living C. *elegans* display heterogeneous chromatin domain structures
- Heterochromatin architecture revealed a highly compacted nanoscale organisation *in vivo* controlled by HP1
- Tandem repeat-enriched ectopic chromosomes acquire heterochromatic structure
- Condensin I and II show differential effects on chromosome structure

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