



**Dra. Kasia Oktaba**

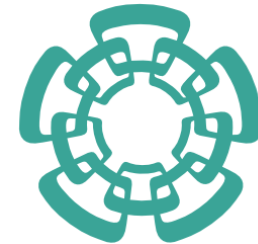
## Laboratorio de Regulación y Topología del Genoma

Departamento de Ingeniería Genética

Cinvestav-Unidad Irapuato

Edificio E3 Laboratorio 3

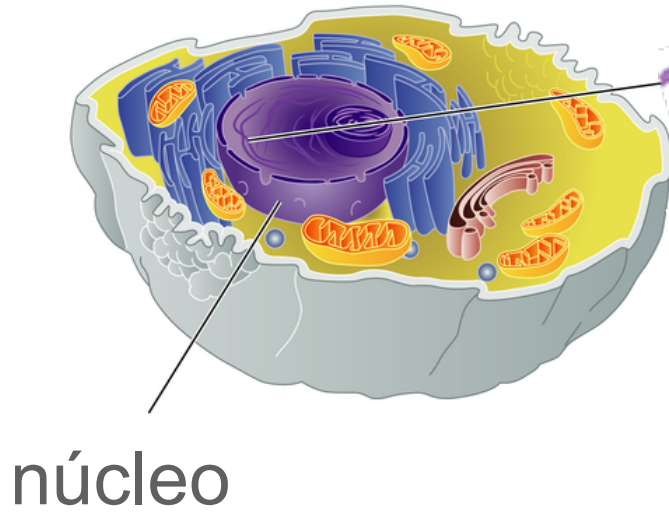
[k.oktaba@cinvestav.mx](mailto:k.oktaba@cinvestav.mx)



**Cinvestav**  
Unidad Irapuato

# Regulación y topología del genoma

## Parte II: Topología



## Parte I: Regulación

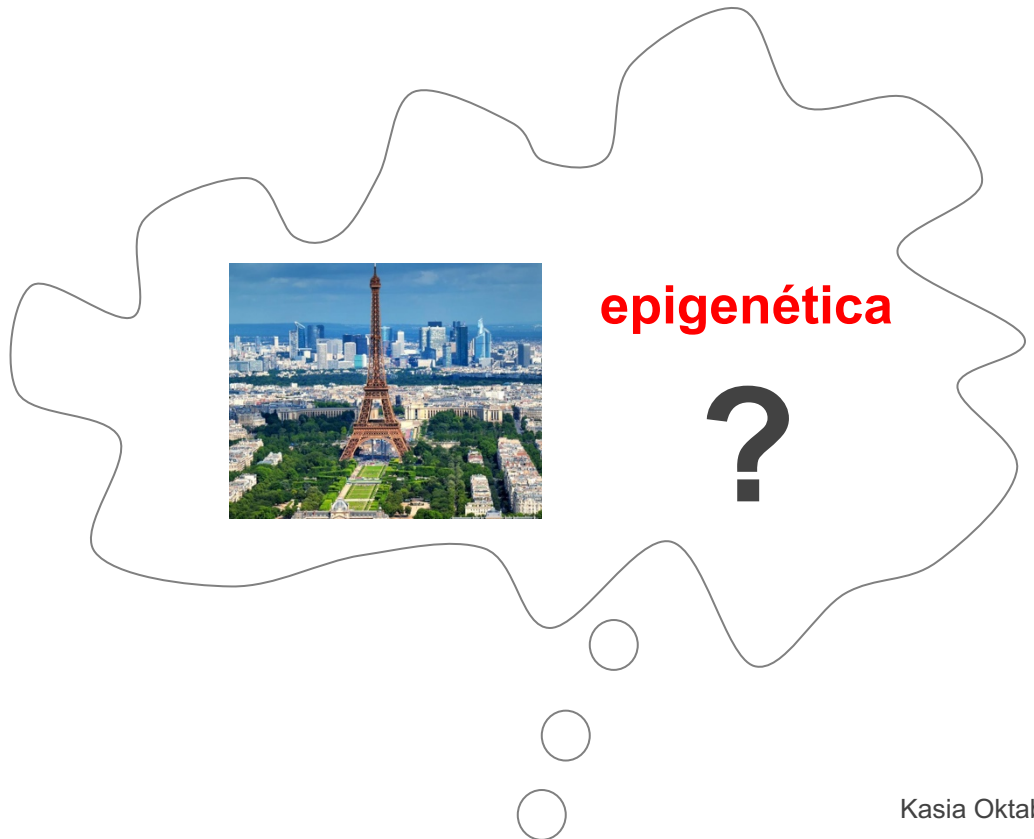
DNA



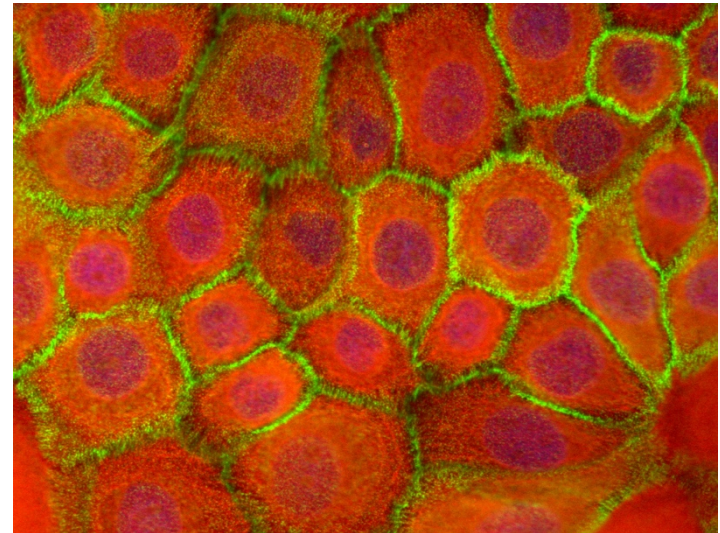
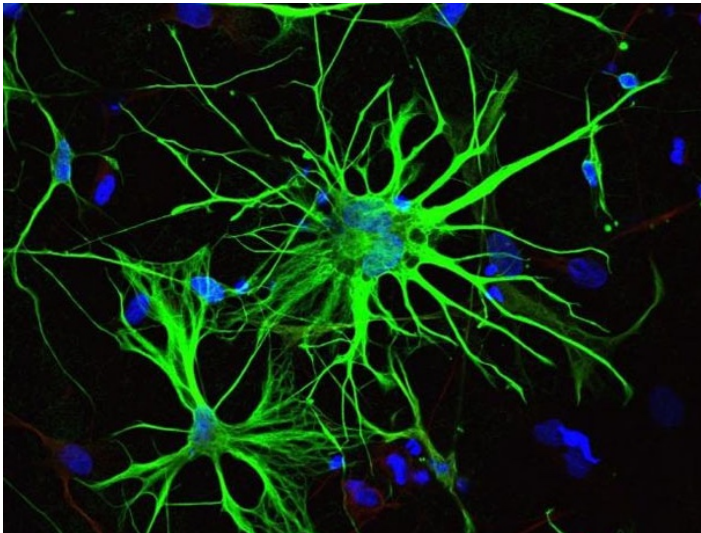
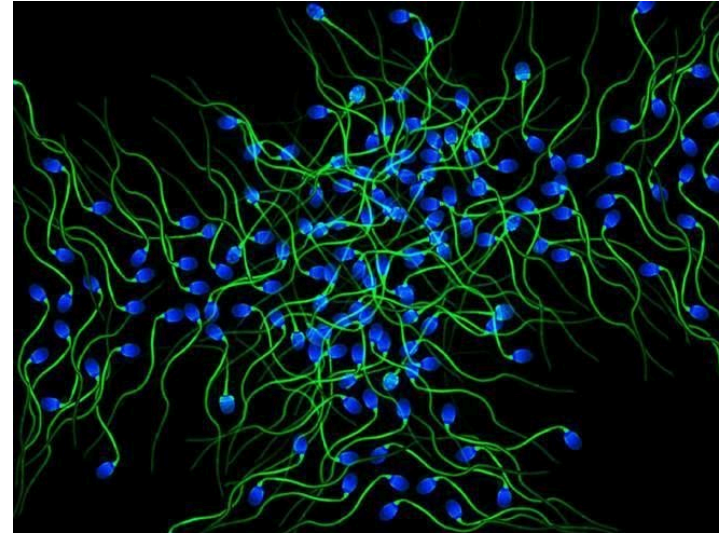
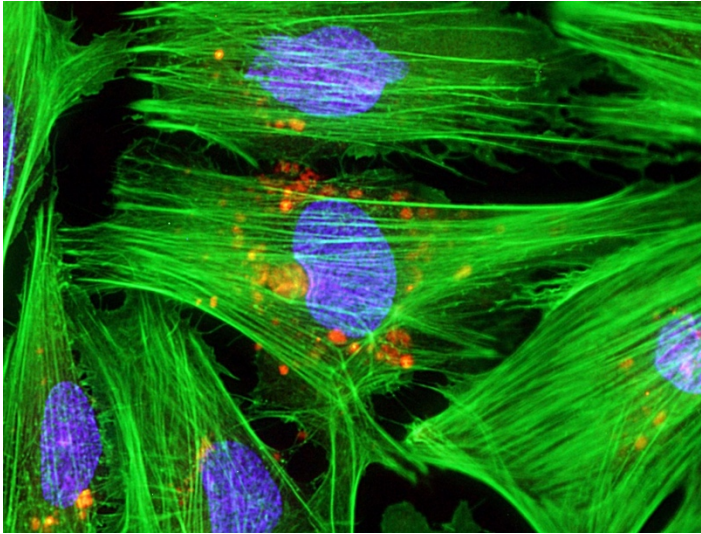
# ...finales de 2002



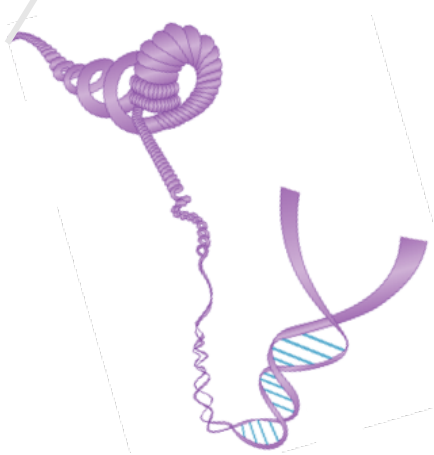
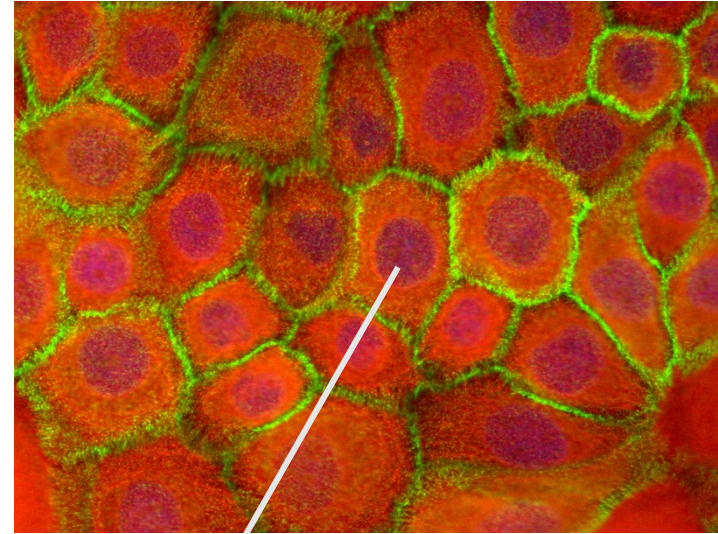
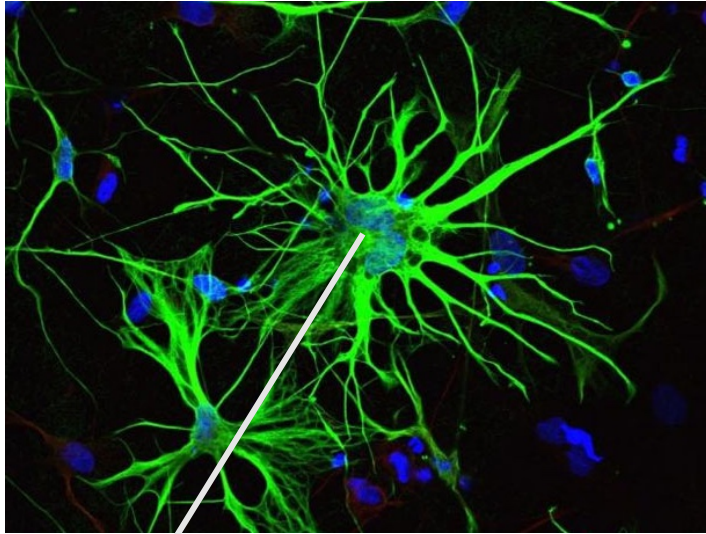
¿Quieres ir a Paris a hacer un proyecto de epigenética?



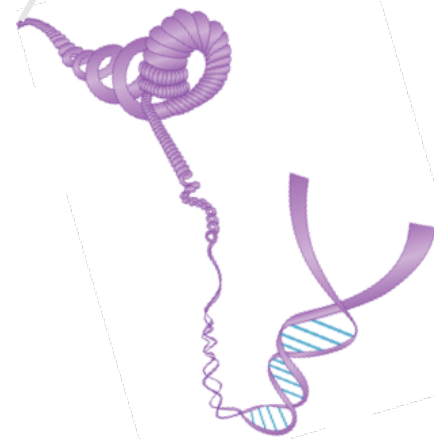
# Regulación del genoma



# Regulación del genoma

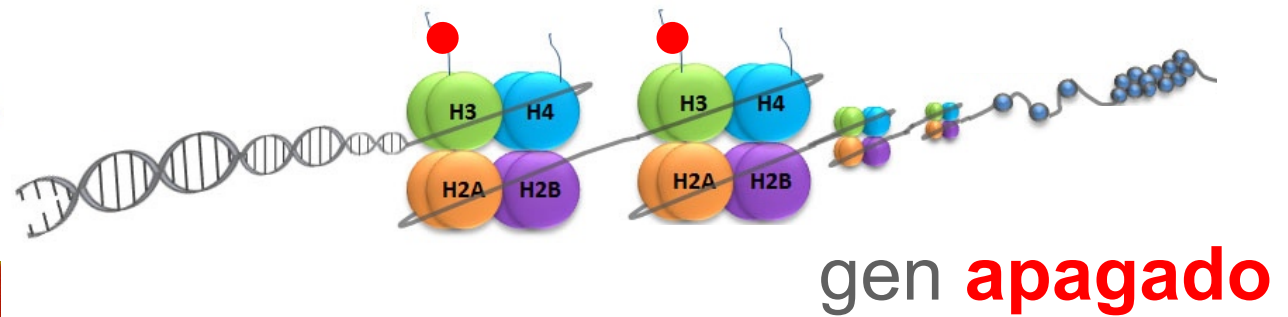
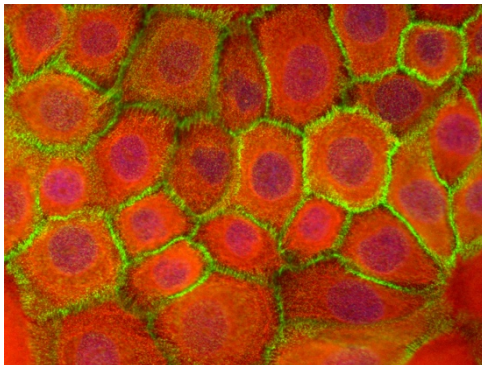
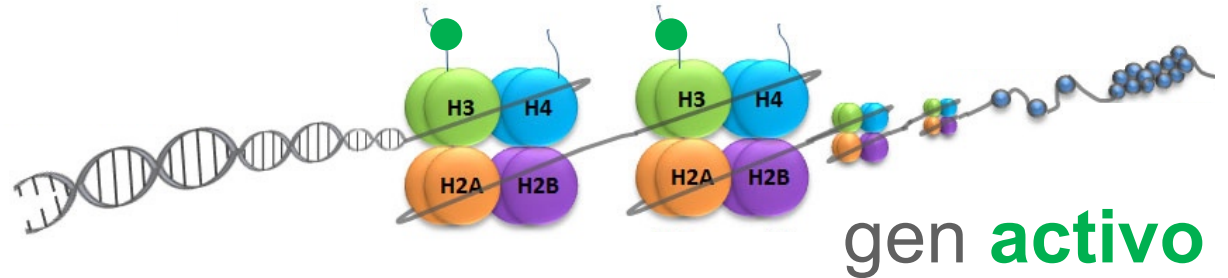
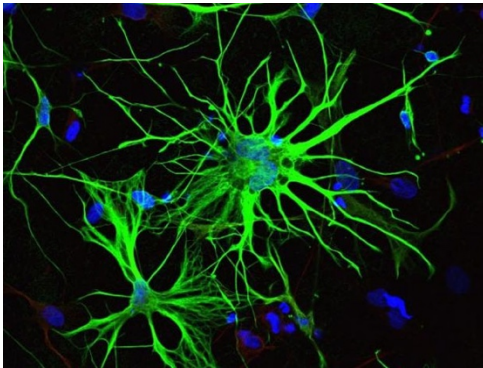


?



# Regulación del genoma: epigenética

marcas epigenéticas en colas de histonas: código



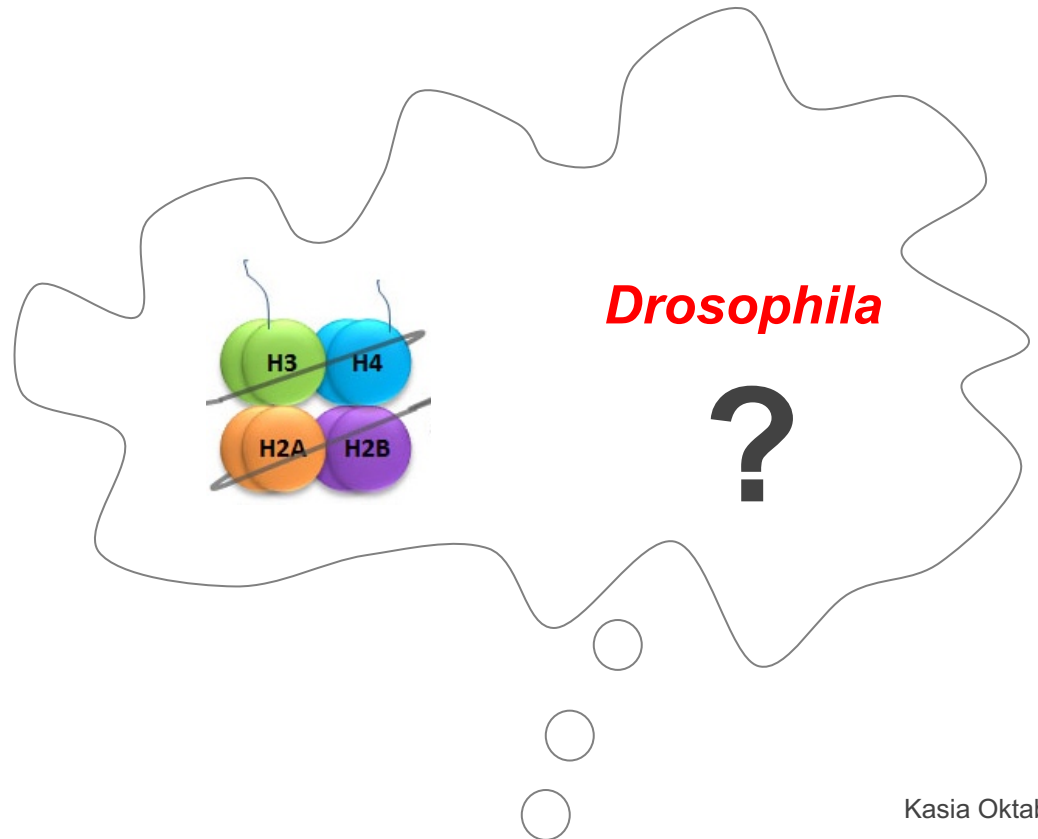
# 2004-2011 en Alemania



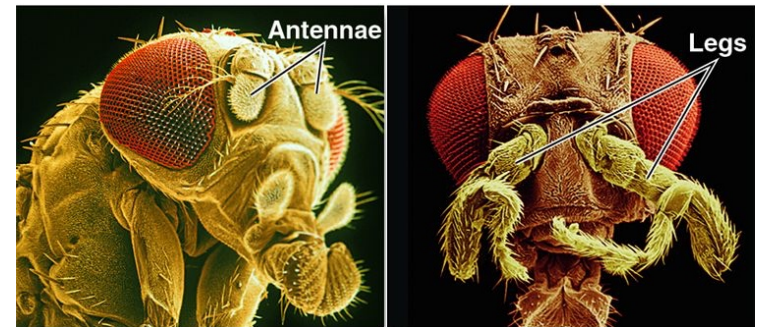
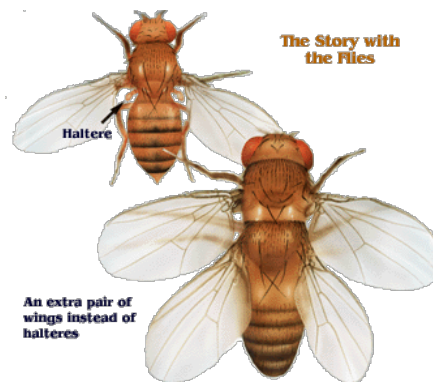
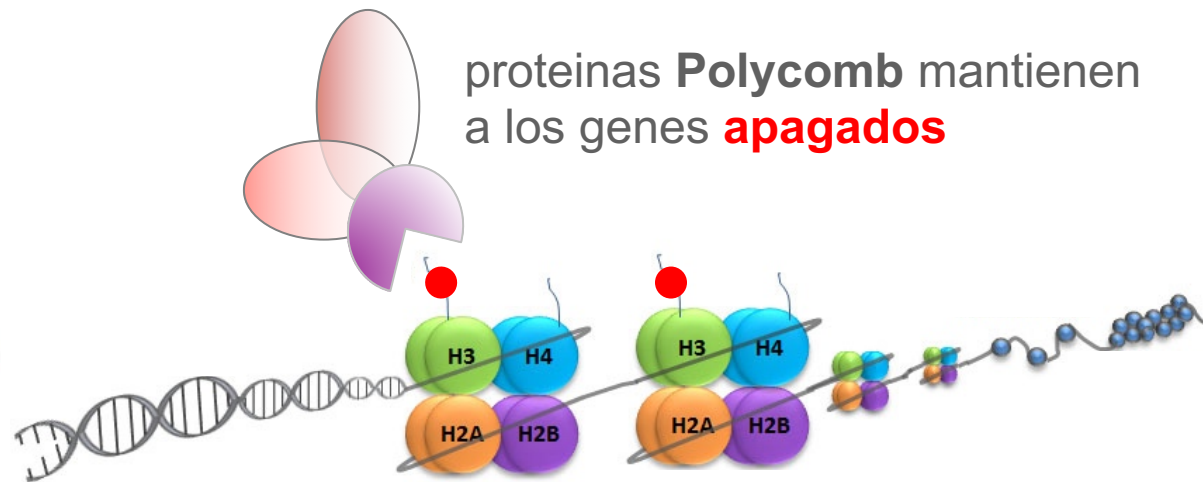
Jürg  
Müller



¿Epigenética? está bien  
pero usando como modelo  
la mosca de la fruta.



# Regulación del genoma: en *Drosophila*



(a) Normal fly

(b) Mutant fly



# 2011-2016 en EUA



Mike  
Levine

Berkeley  
UNIVERSITY OF CALIFORNIA

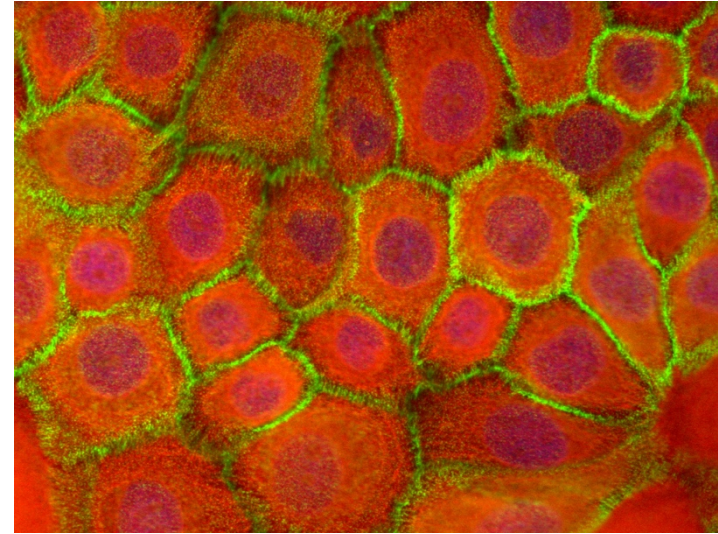
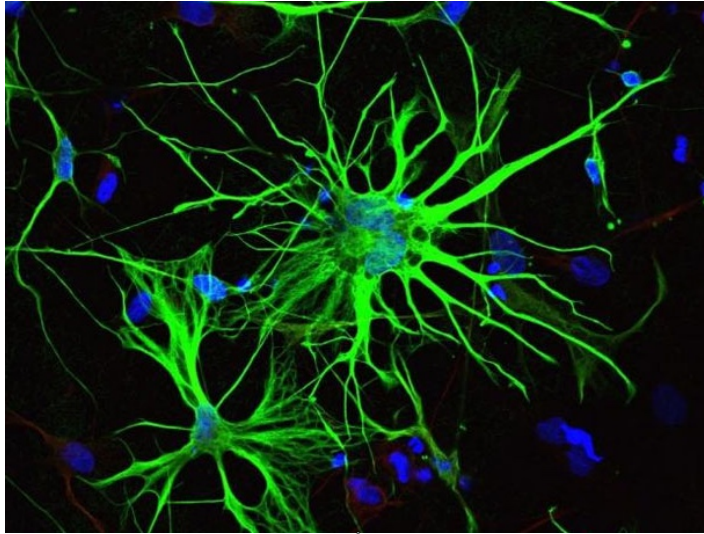
¿Y si buscamos todos los  
'enhancers' en el genoma  
de *Drosophila*?



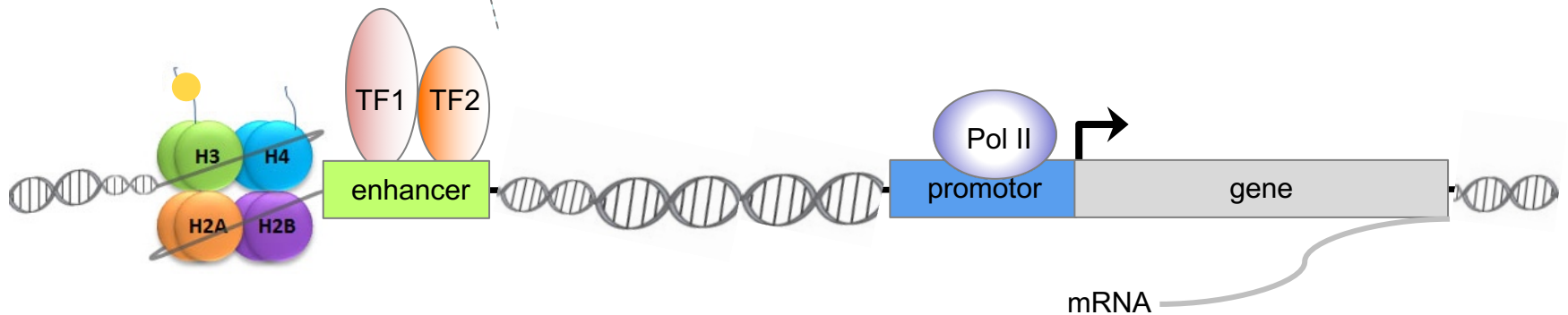
'enhancers'

?

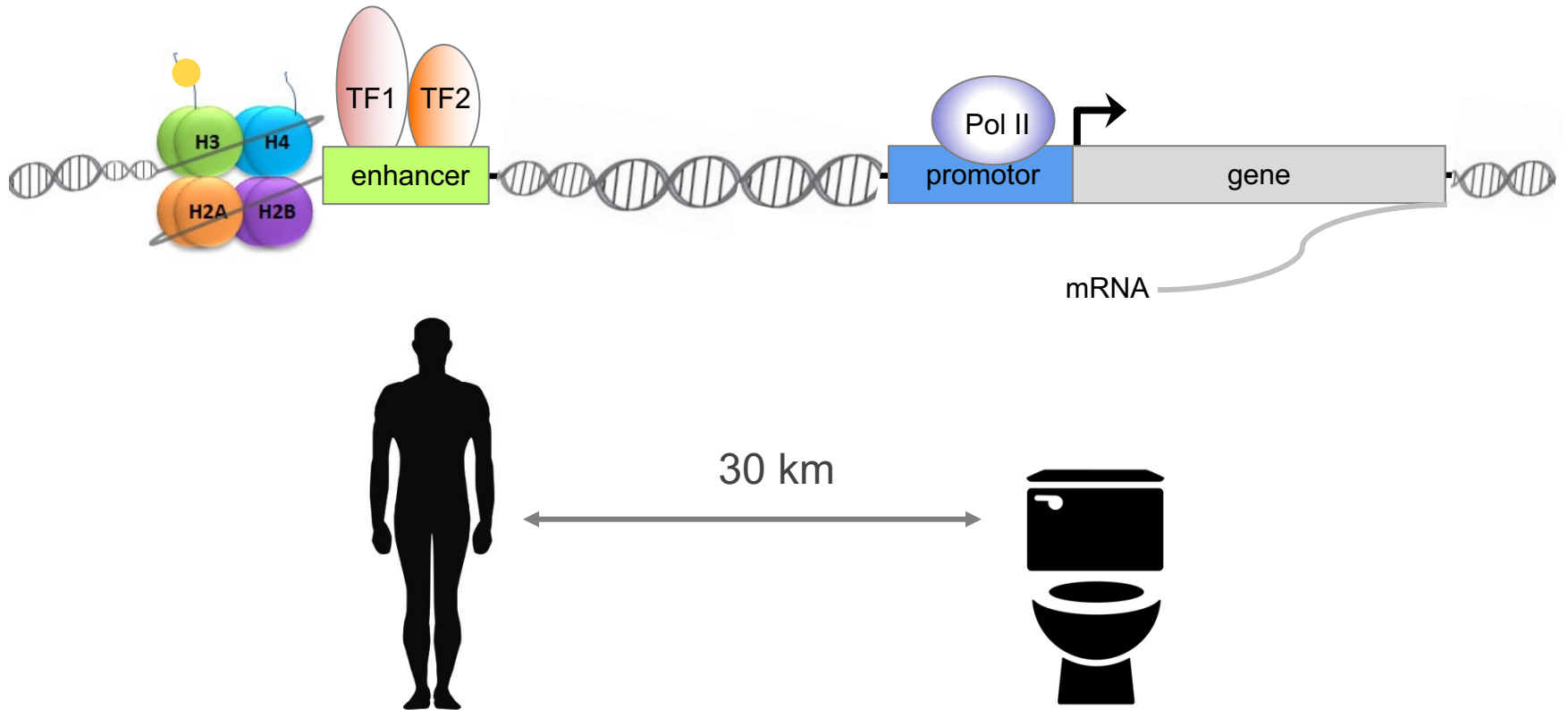
# Regulación del genoma: 'enhancers'



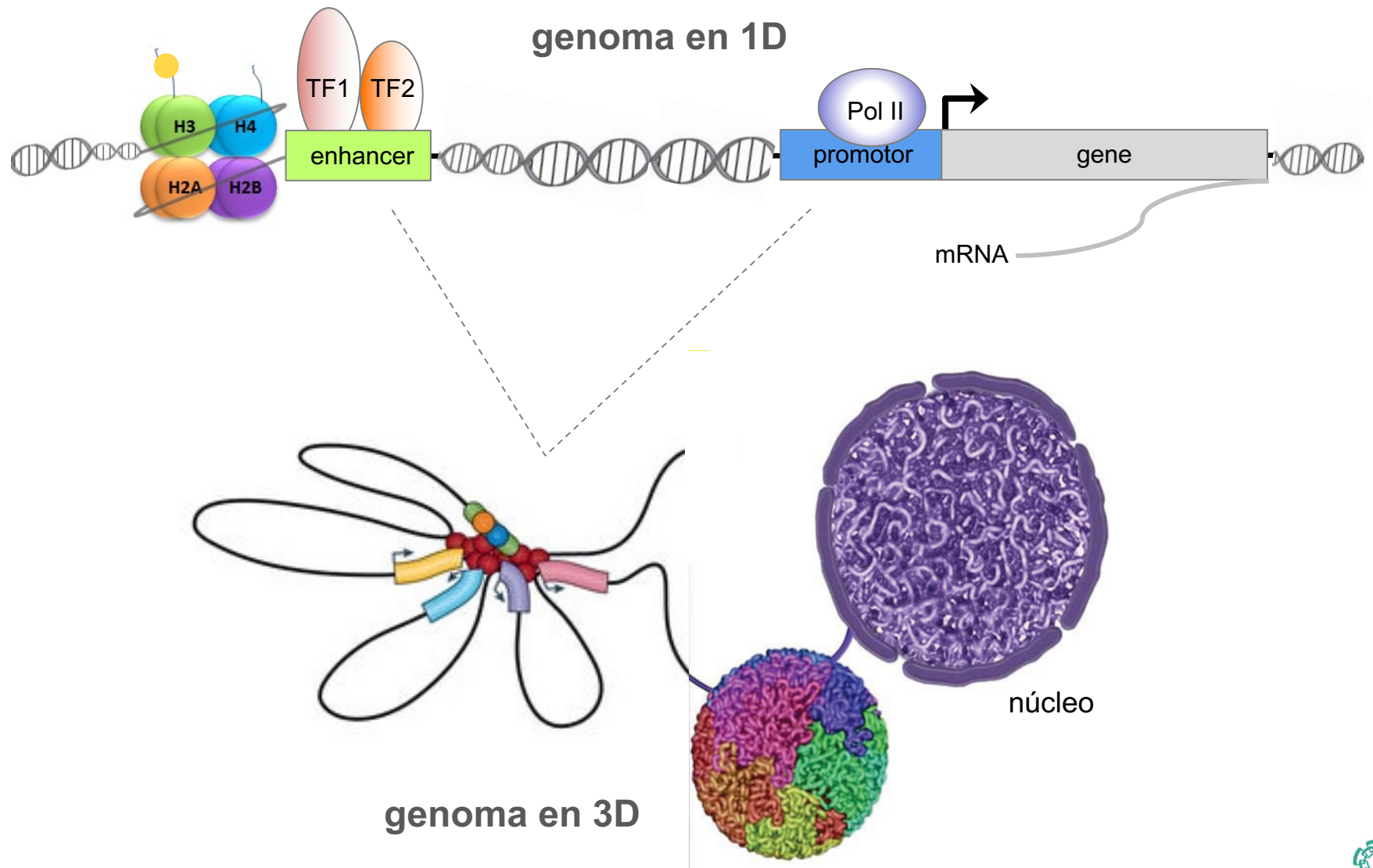
factores de transcripción  
**activan** o **apagan** genes



# Problema no. 1



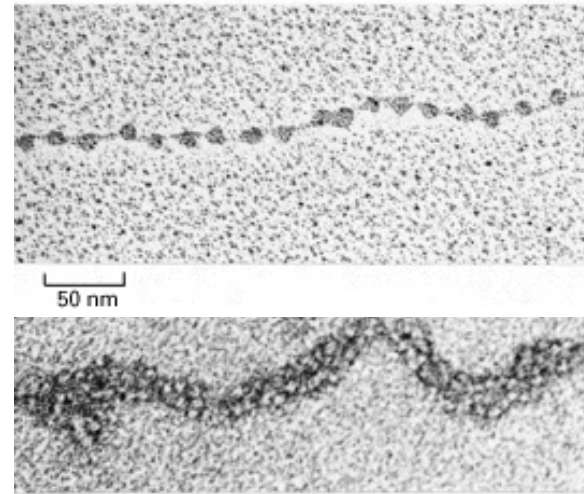
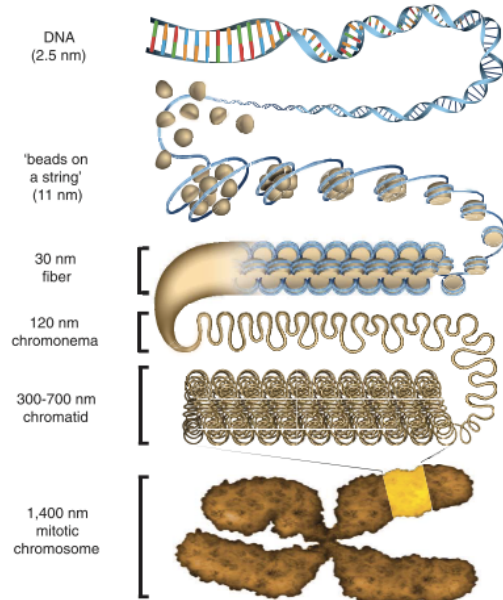
# Topología del genoma



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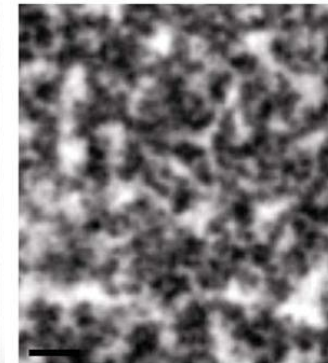
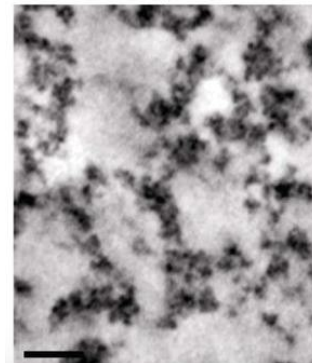


## disordered 5-24 nm diameter chain



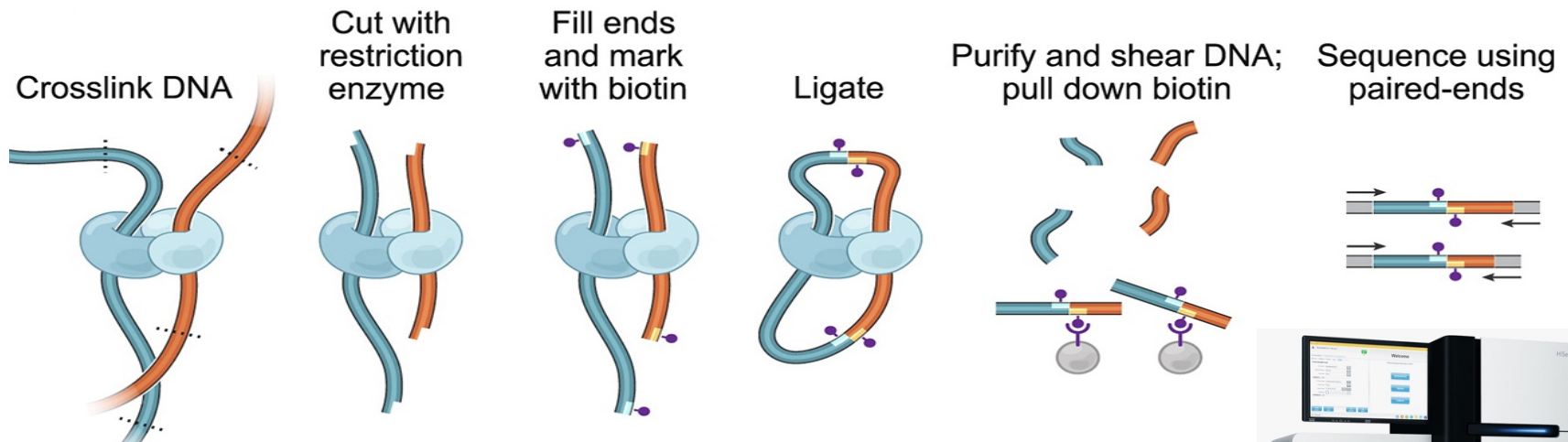
interphase

mitotic

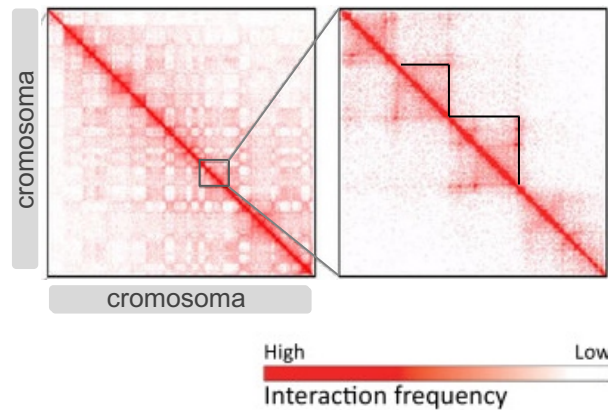


# desde finales de 2016 en Irapuato

## biología molecular: método HiC

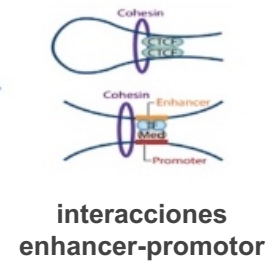
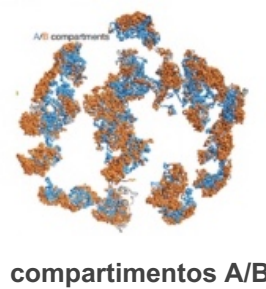
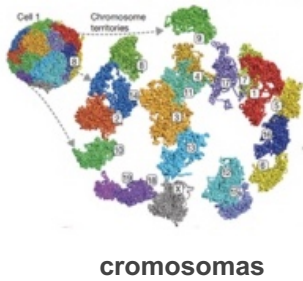
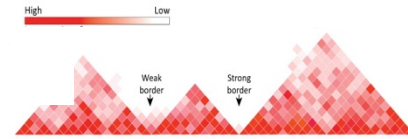
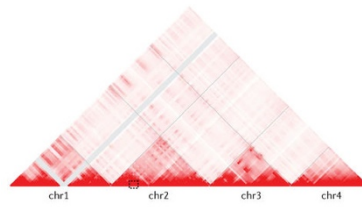
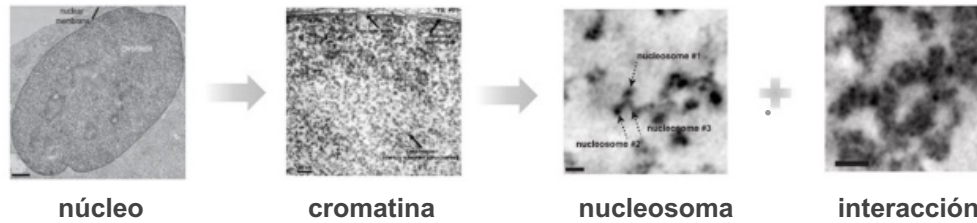


análisis de datos de secuenciación



# Topología del genoma

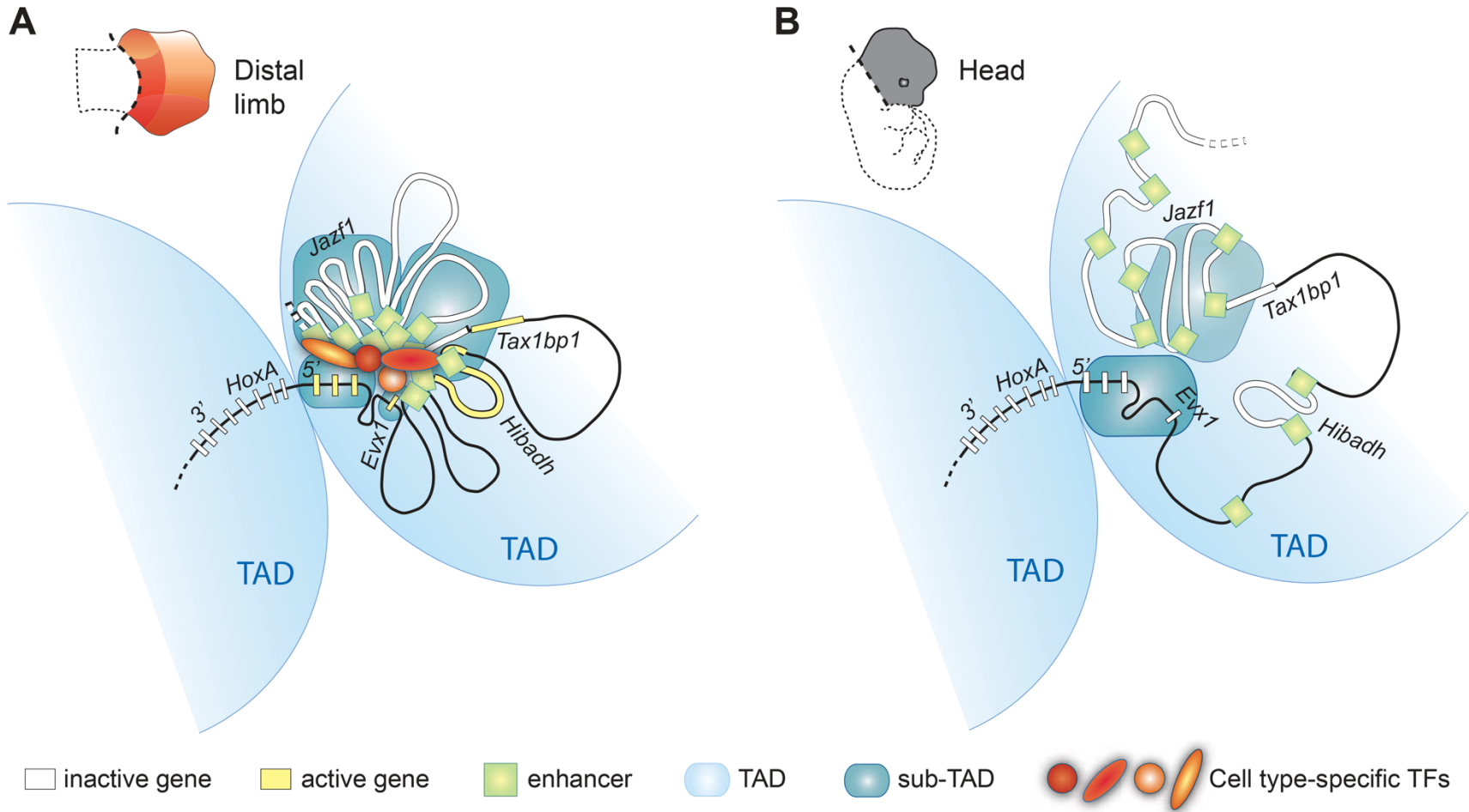
## microscopía



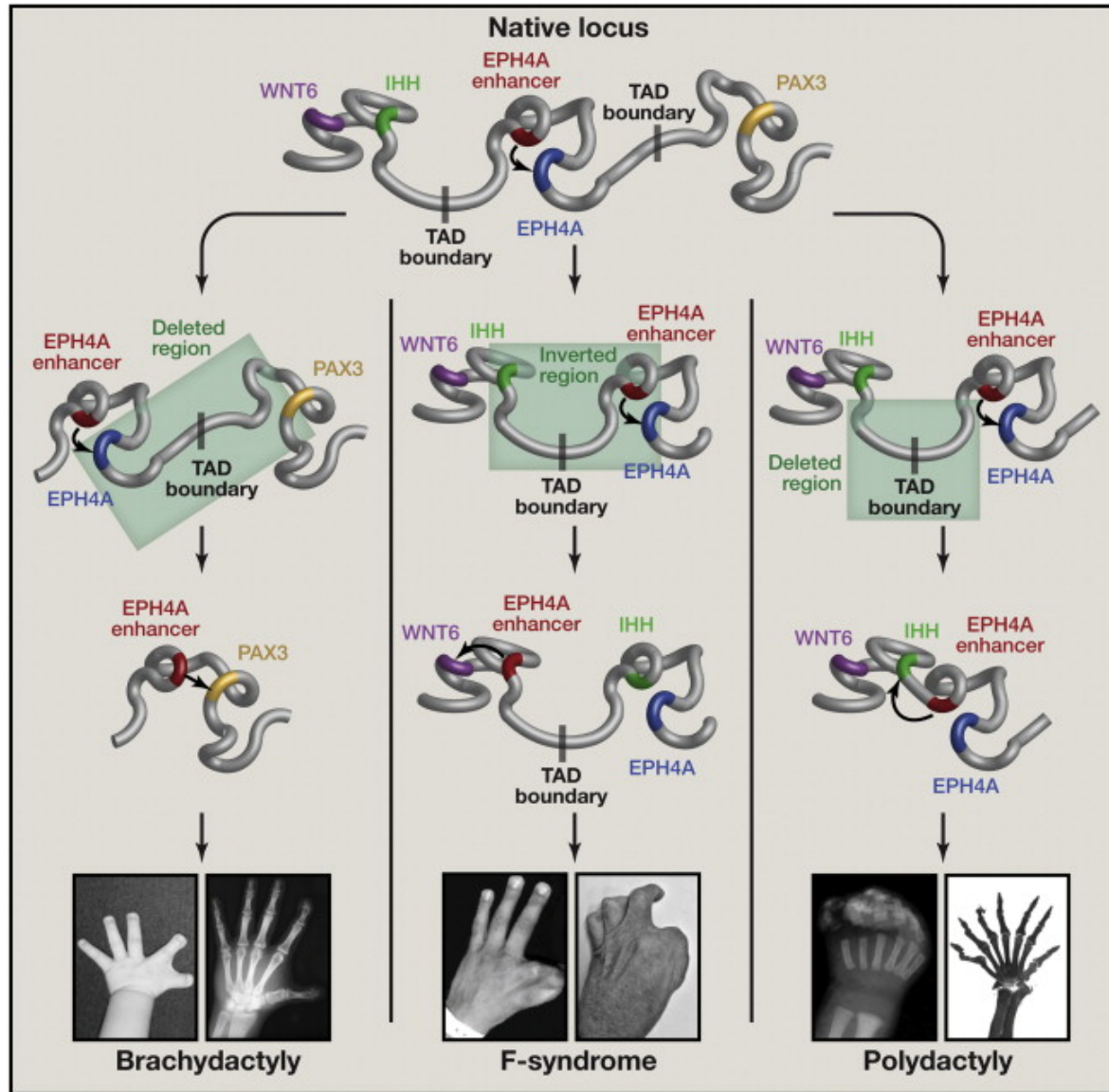
## método HiC



# Topología del genoma: tejido específica?



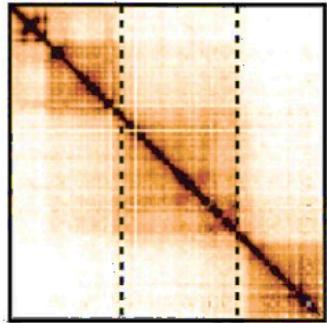
# Topología del genoma: relevante?



# Topología del genoma en distintas especies



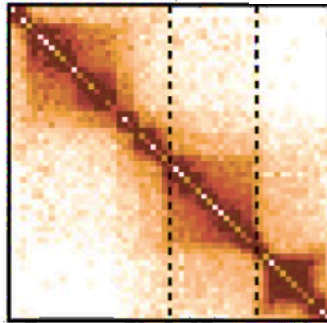
CTCF, Cohesin  
promoters, tRNA



1 MB



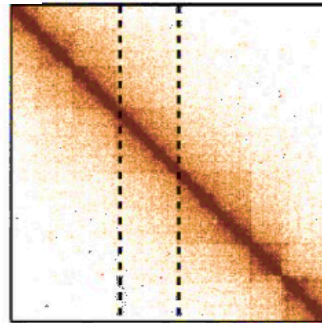
CTCF, BEAF, CP190  
Su(Hw), promoters



0.25 MB



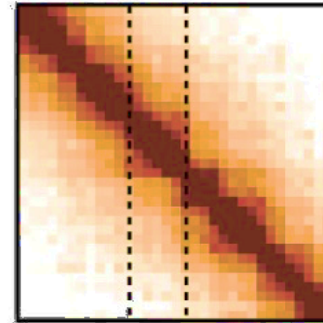
DCC



3 MB



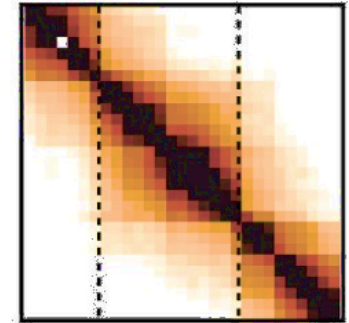
Convergent genes  
Cohesin



0.30 MB



Active genes



0.25 MB

Dekker & Heard (2015) *FEBS Letters*

¿qué mecanismos?  
¿cómo evoluciona?

# Laboratorio de Regulación y Topología del Genoma



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Kasia Oktaba

