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Spatio-temporal regulatory landscape of sex-determination



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CNAG-CRG · ICREA

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<http://marciuslab.org>
<http://3DGenomes.org>
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Sex determination: a 3,000 year-old enigma

Mythology



Hermaphrodite primeval men

Plato's symposium, 385-370 BC

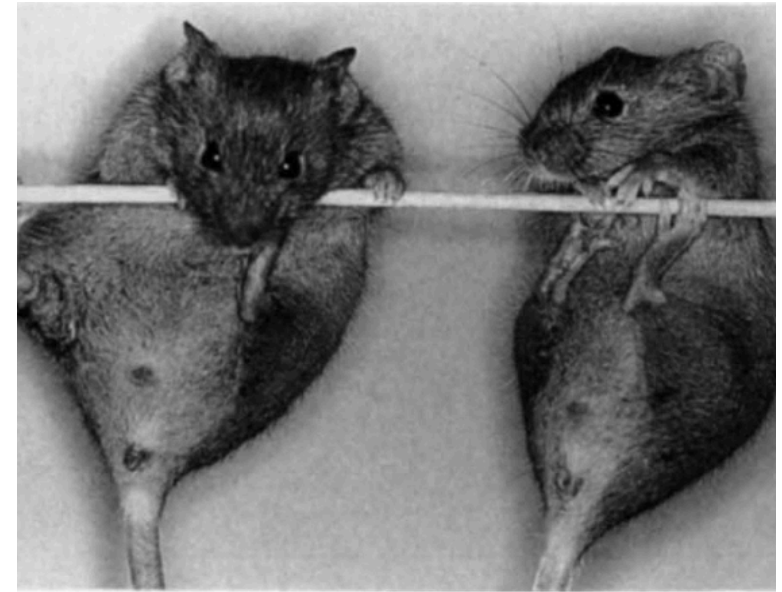
Theories



Left-right theory

Alexandrian manuscripts, 1st cent. BC

Genetics



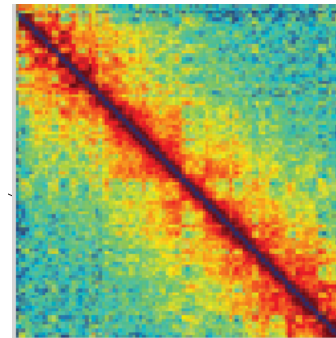
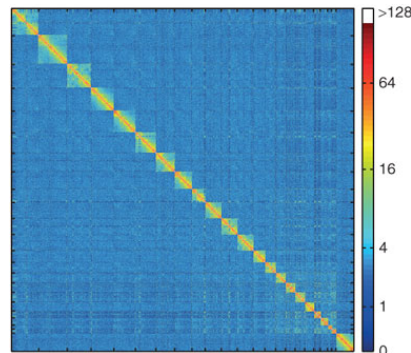
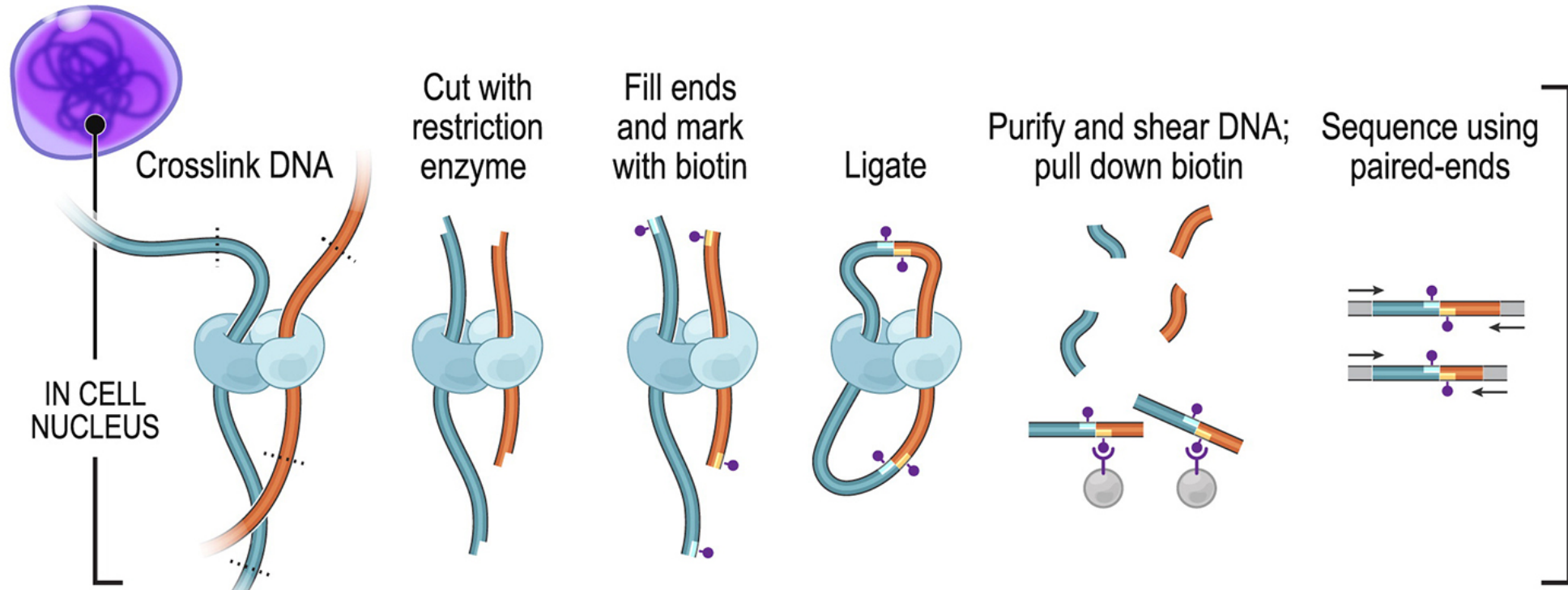
Discovery of Sry gene

*Koopman et al., Nature, 1991
(Goodfellow & Lovell Badge labs)*

Chromosome Conformation Capture

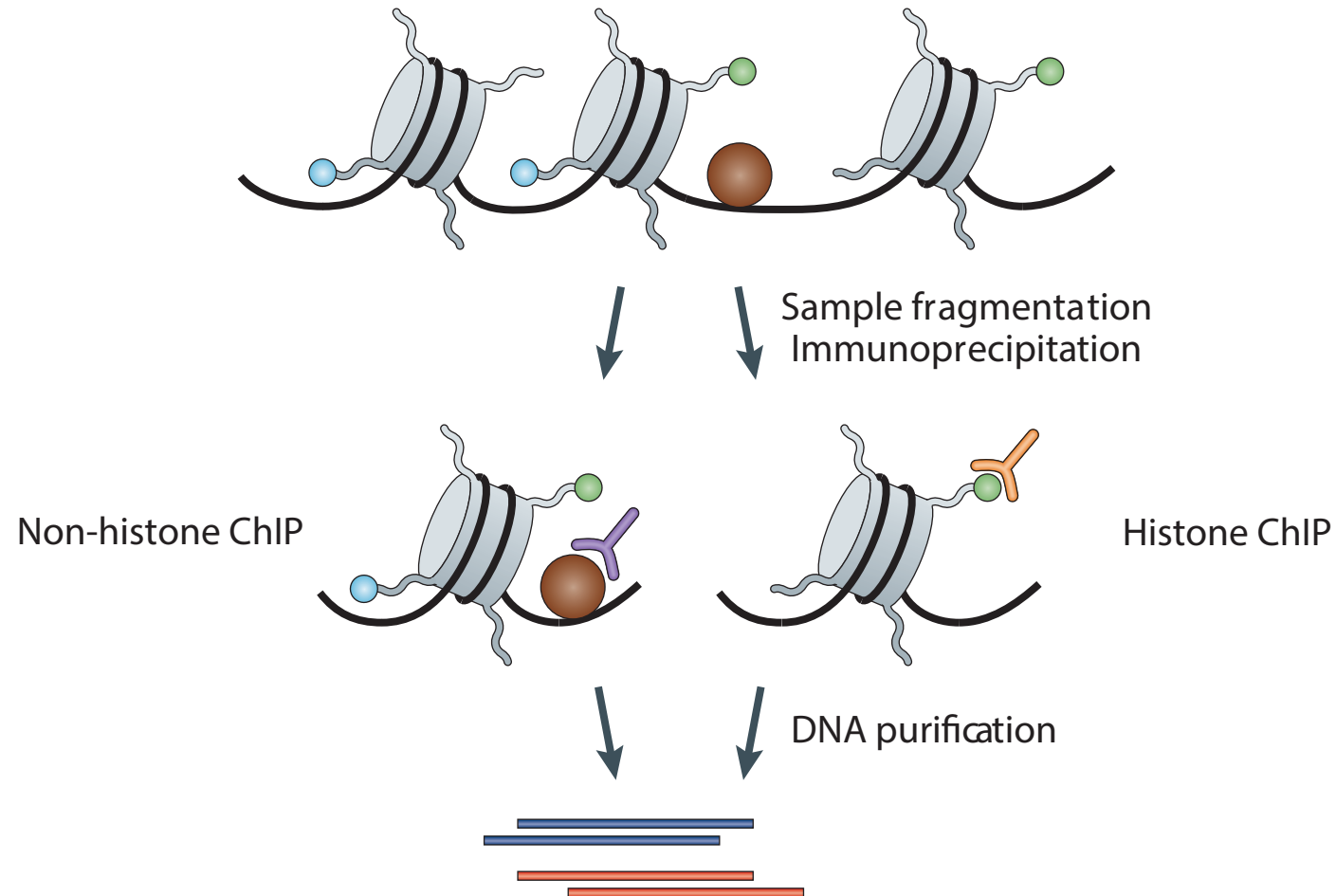
Dekker, J., Rippe, K., Dekker, M., & Kleckner, N. (2002). *Science*, 295(5558), 1306–1311.

Lieberman-Aiden, E., et al. (2009). *Science*, 326(5950), 289–293.

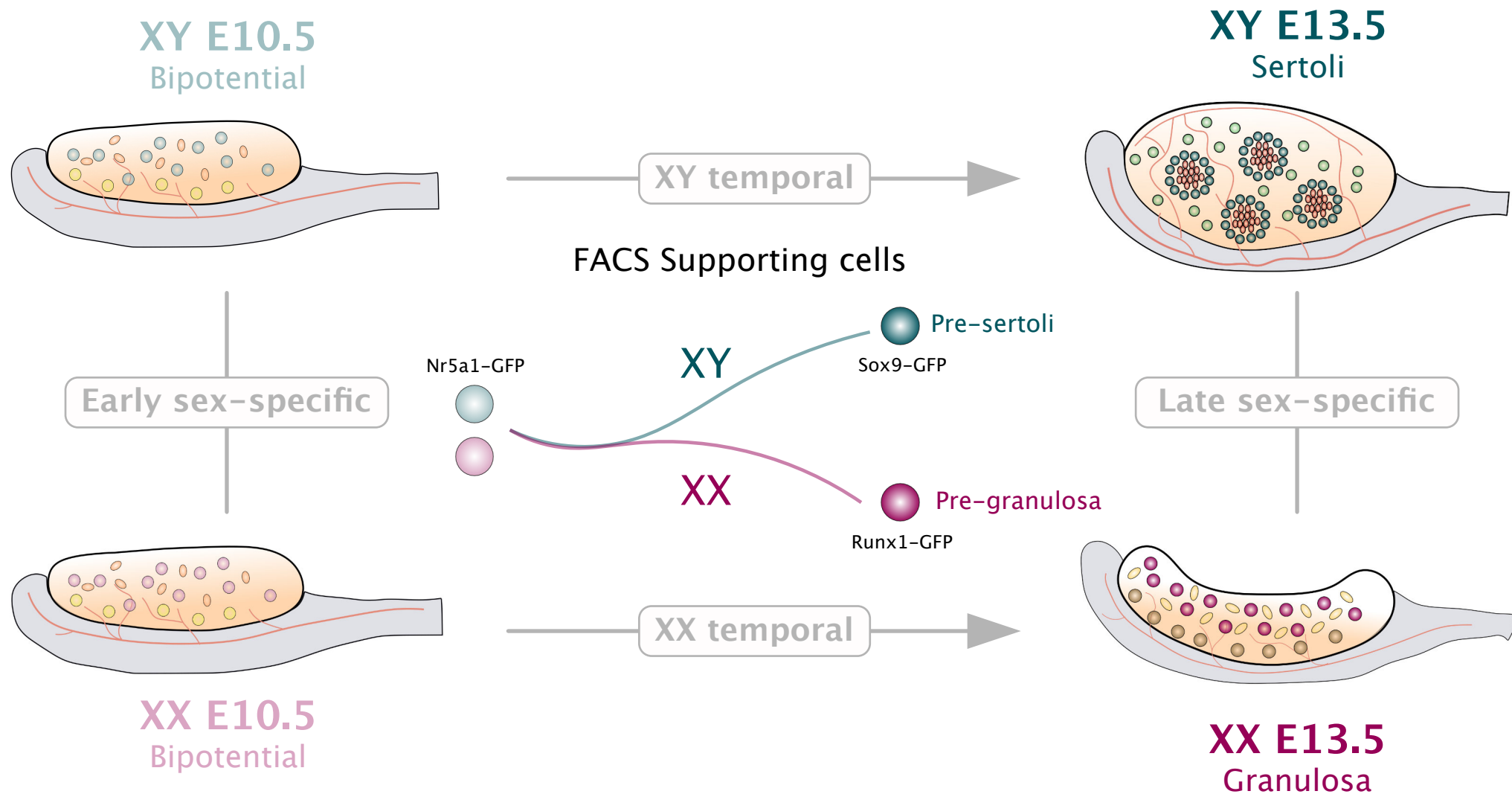


Chromatin Immunoprecipitation (ChIP)

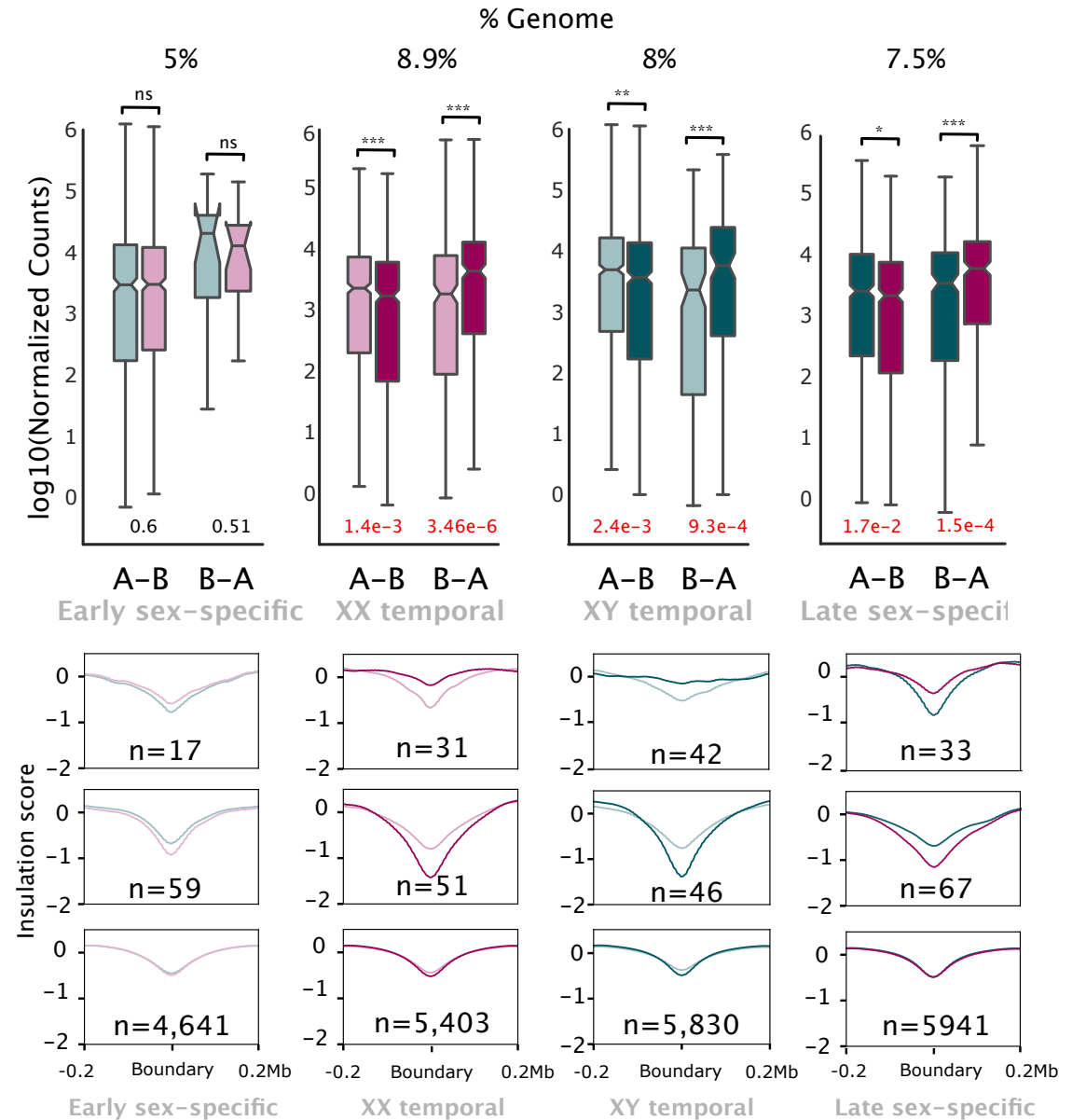
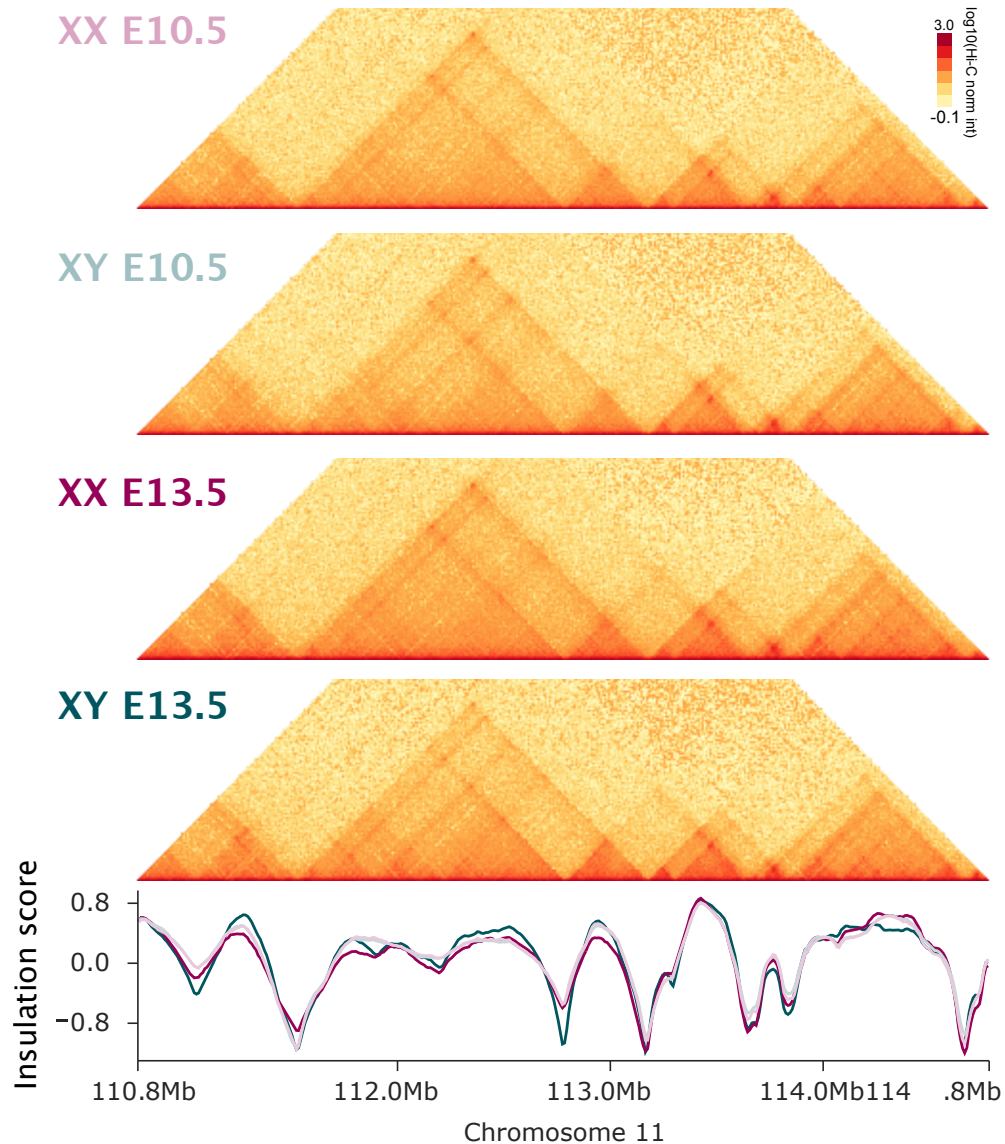
Solomon, M. J., Larsen, P. L. & Varshavsky, A. (1988) Cell 53, 937–947.
Park, P.J. (2009) Nature Reviews Genetics 10, 669–680.

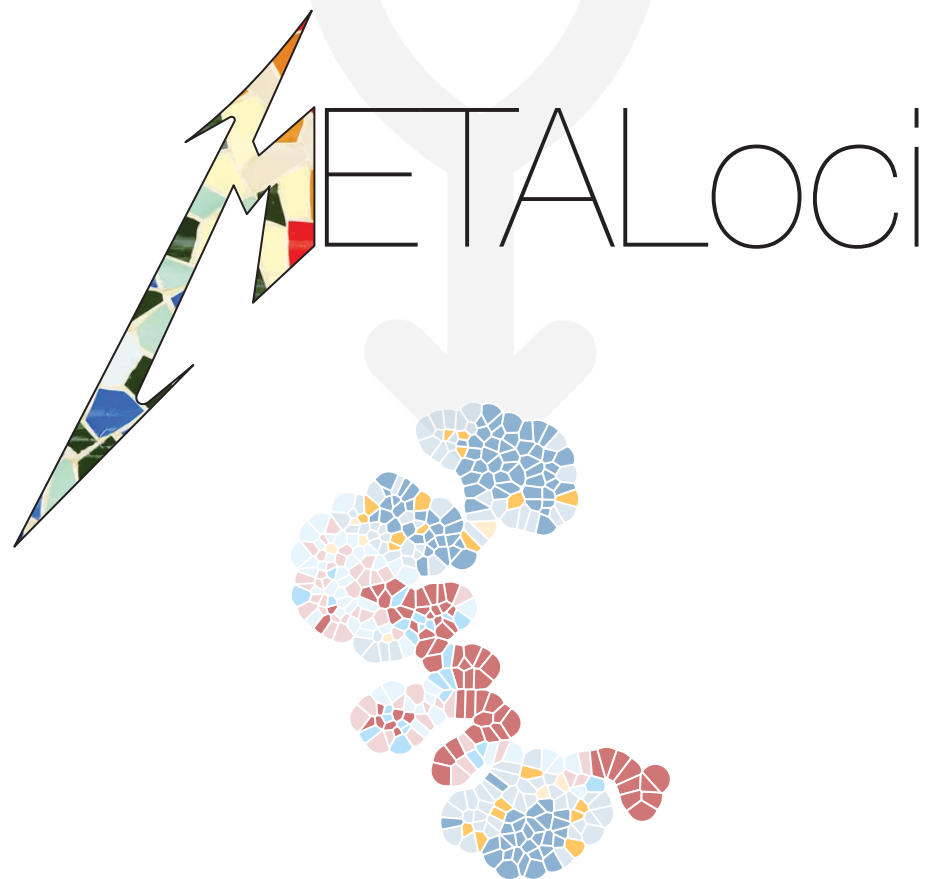


Sex-determination as a model for “bipotential” commitment



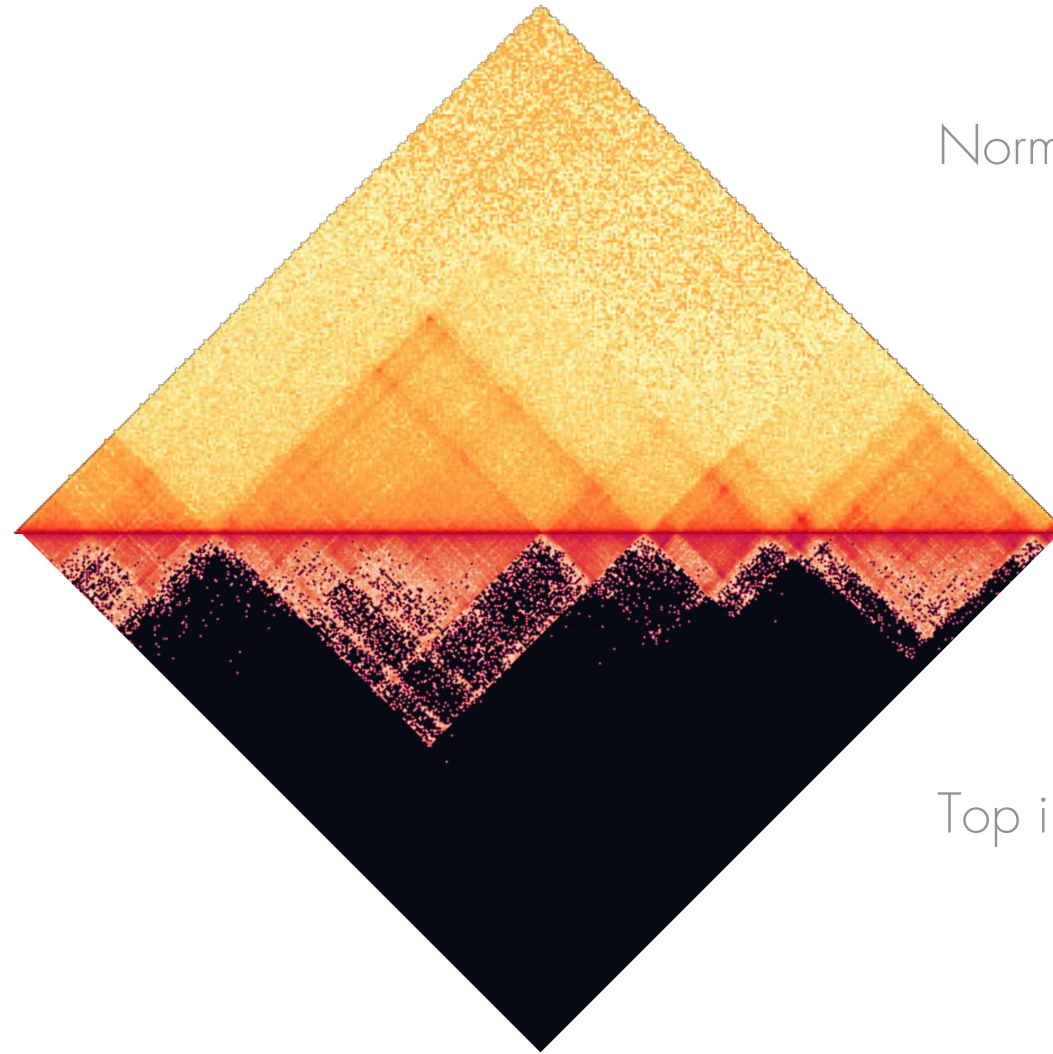
No major structural (apparent) differences





Hi-C normalization and interaction selection

chr11:110780000-114770000

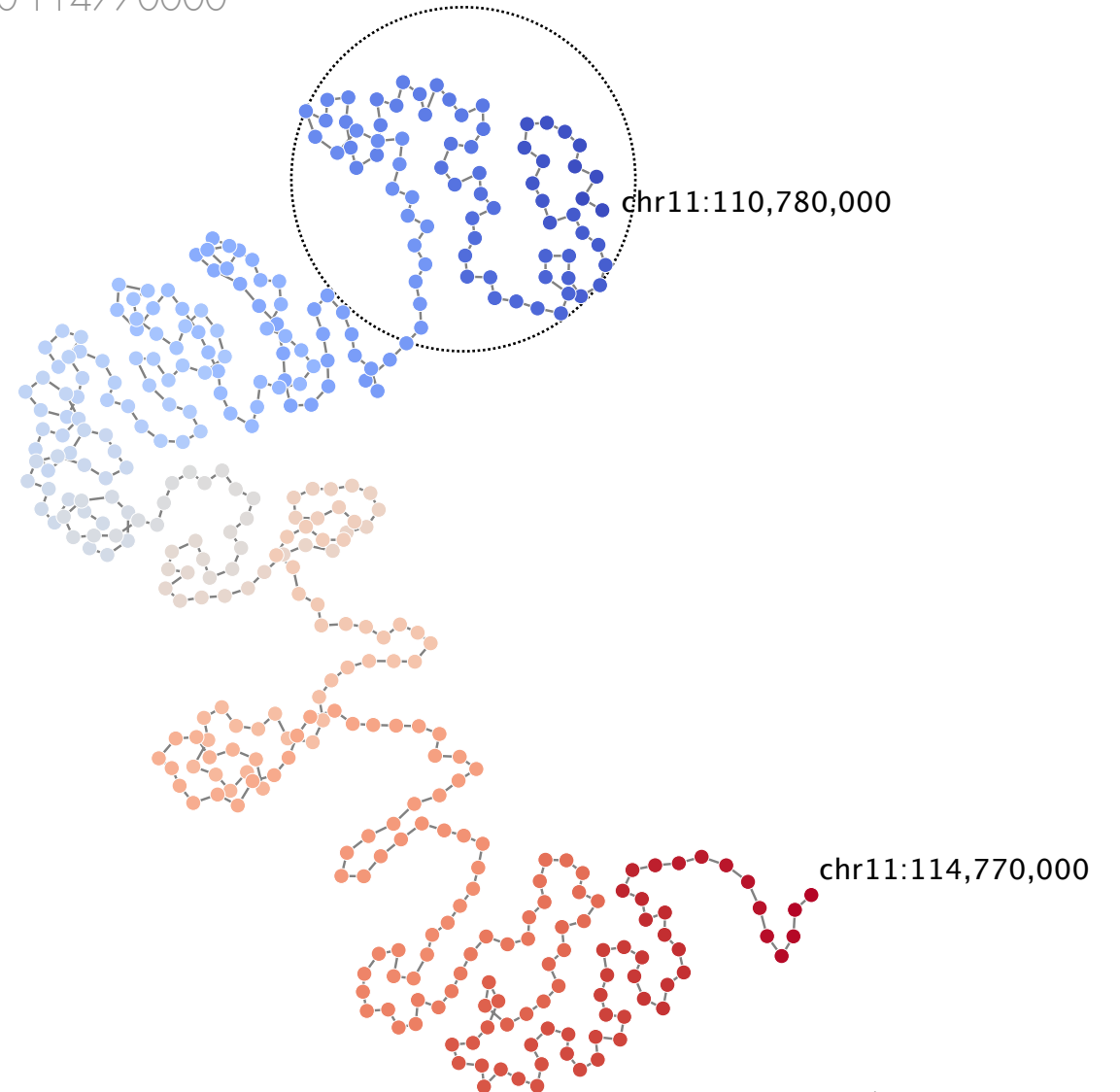
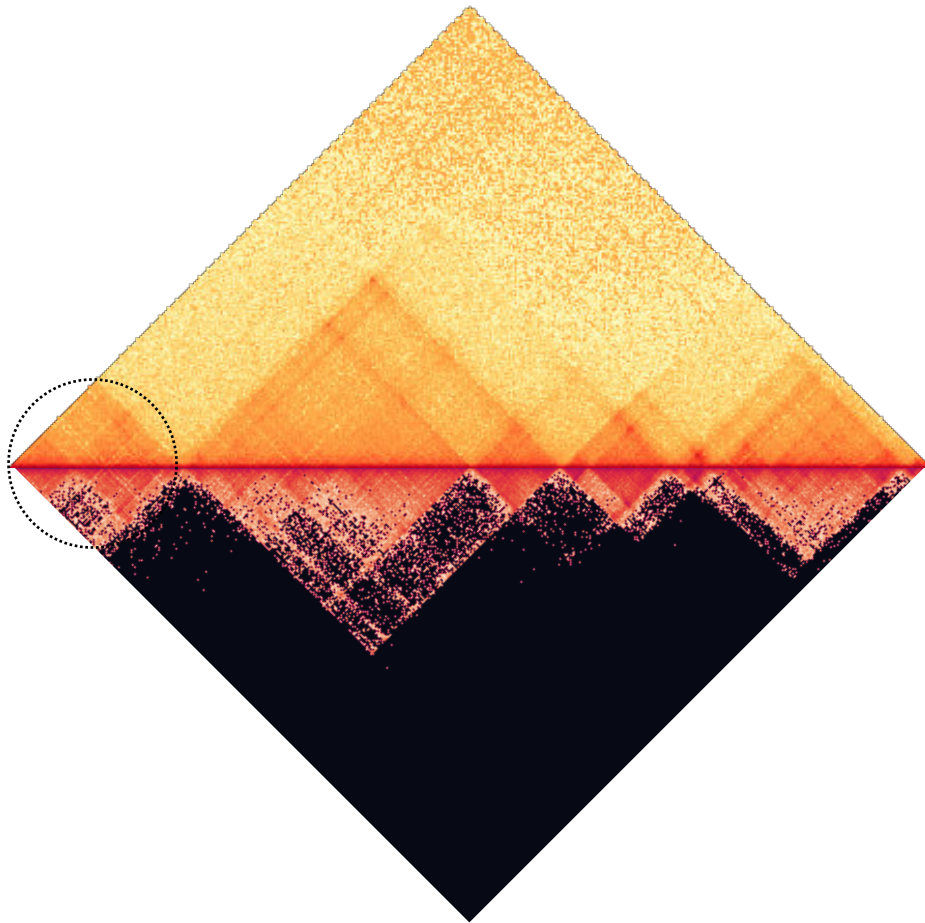


Normalized Hi-C

Top interactions

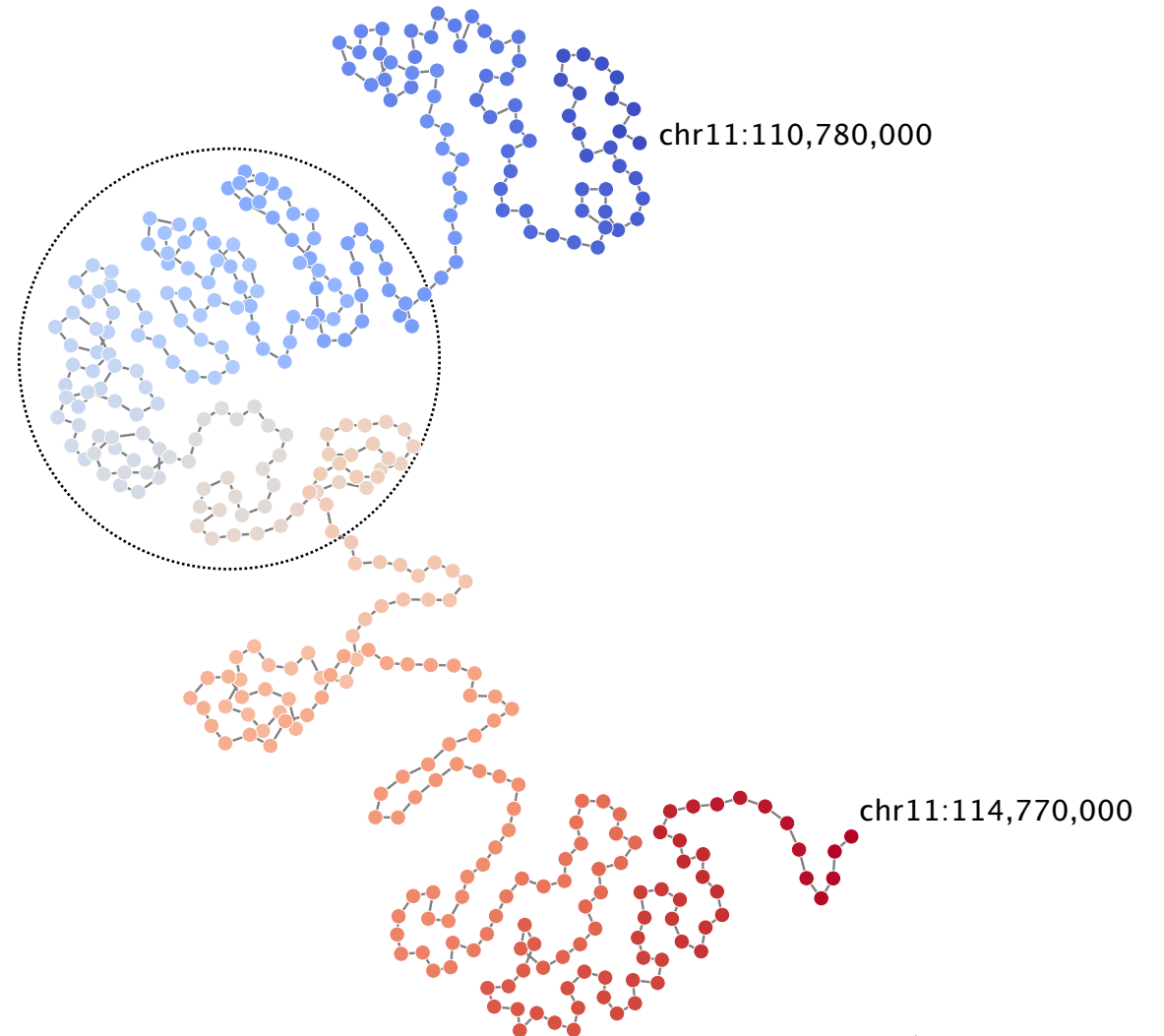
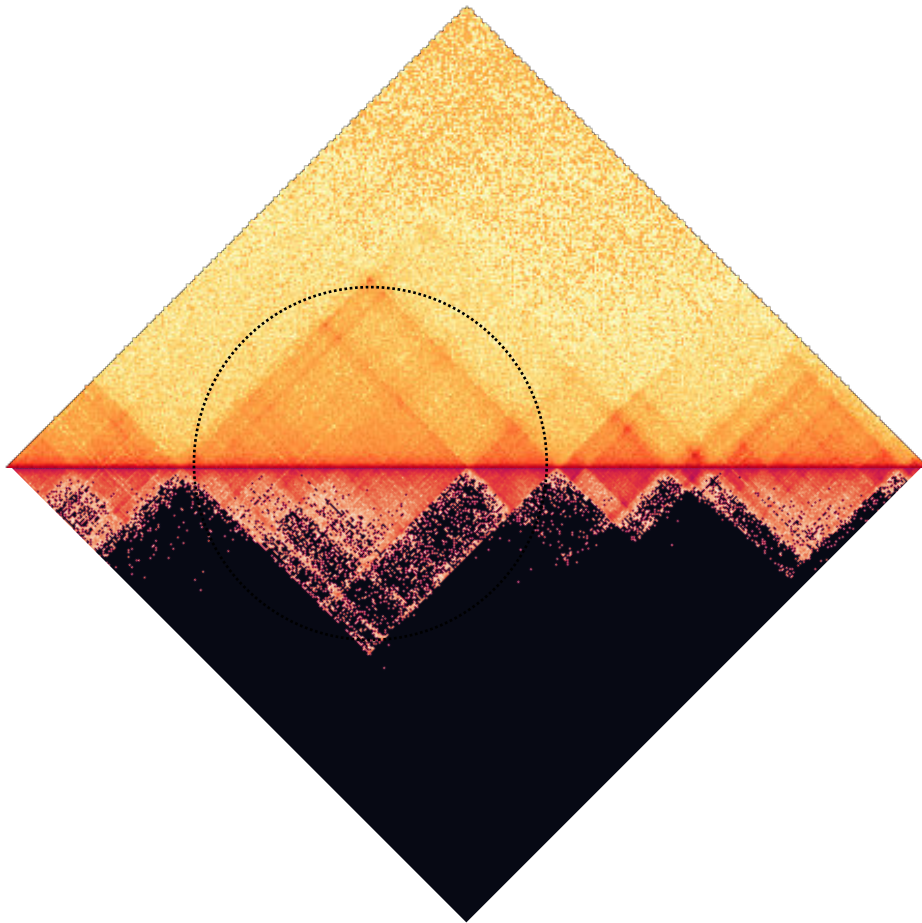
Spatial lay-out of significant interactions

chr11:110780000-114770000



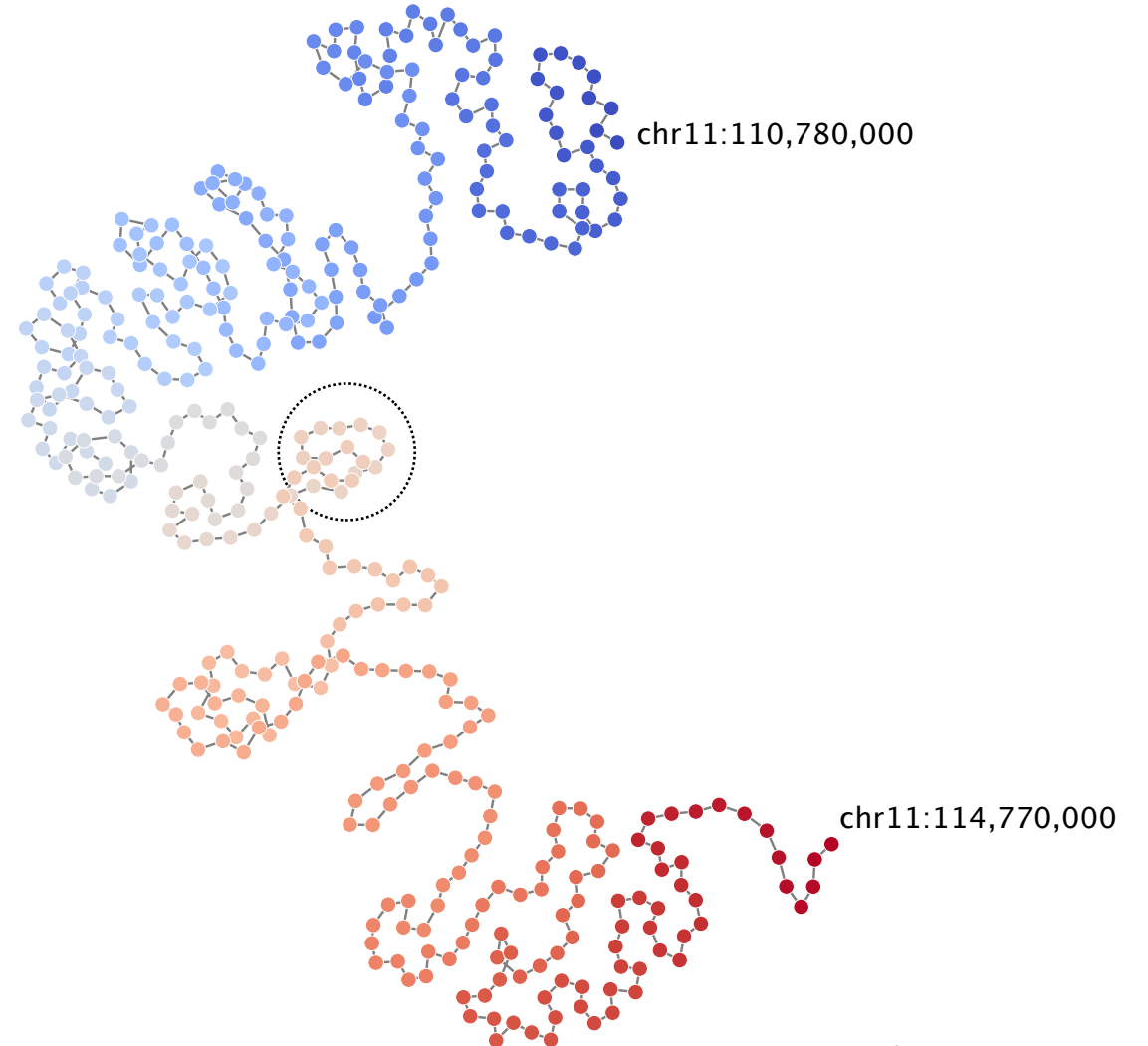
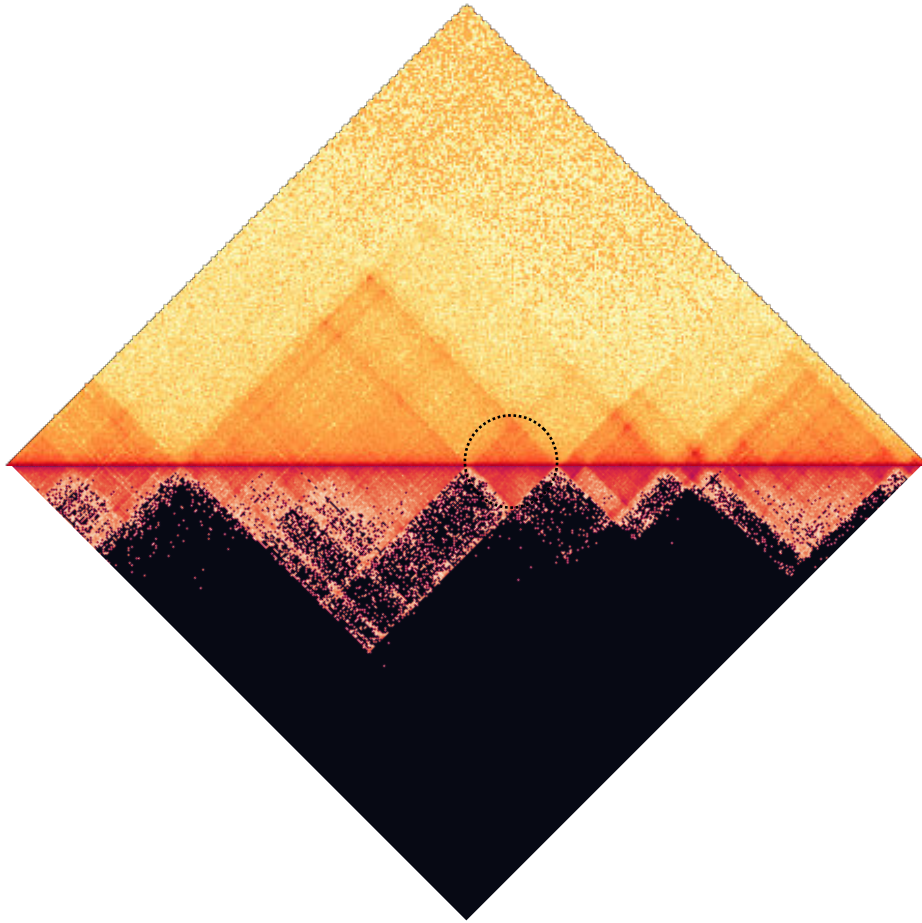
Spatial lay-out of significant interactions

chr11:110780000-114770000



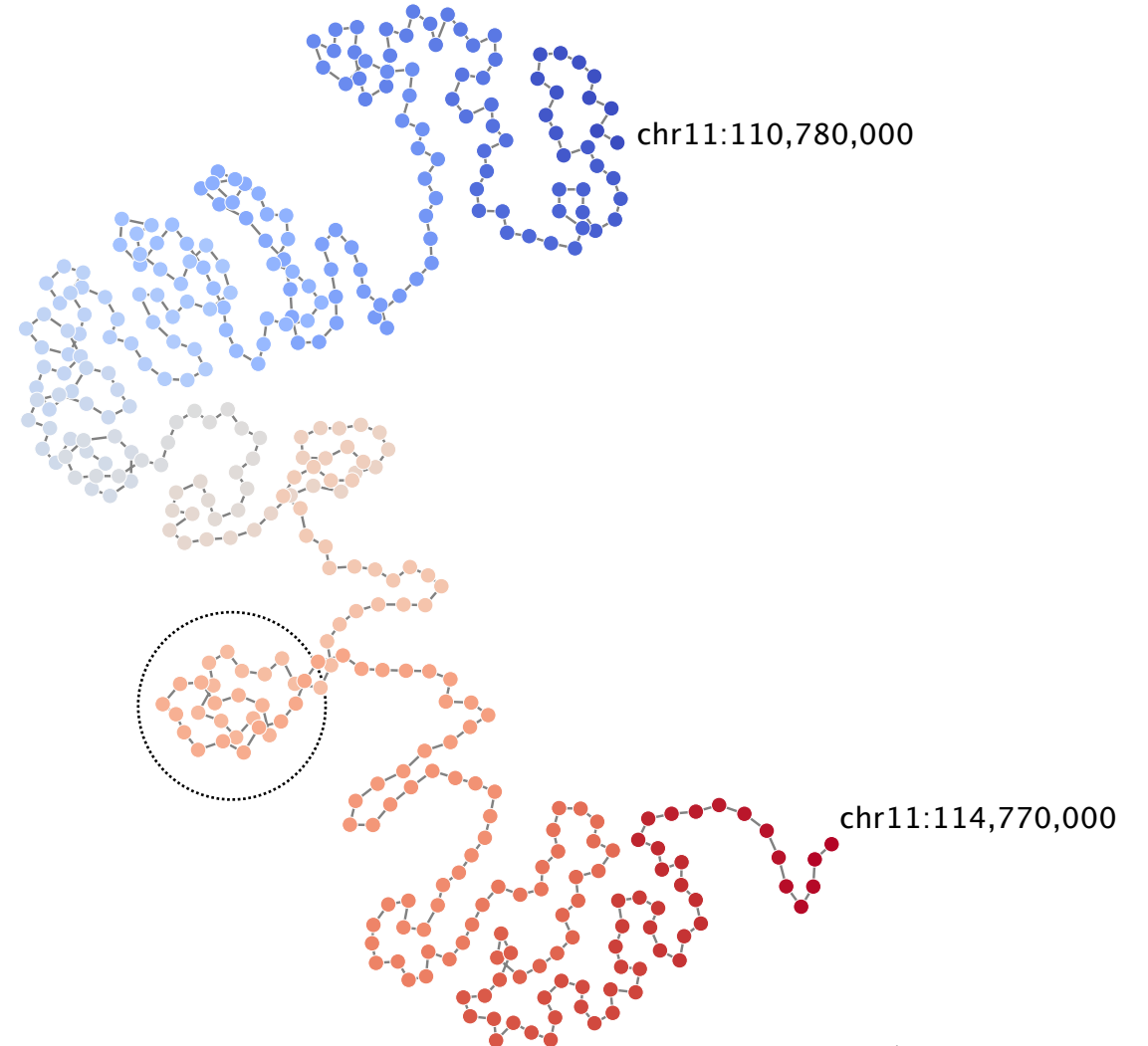
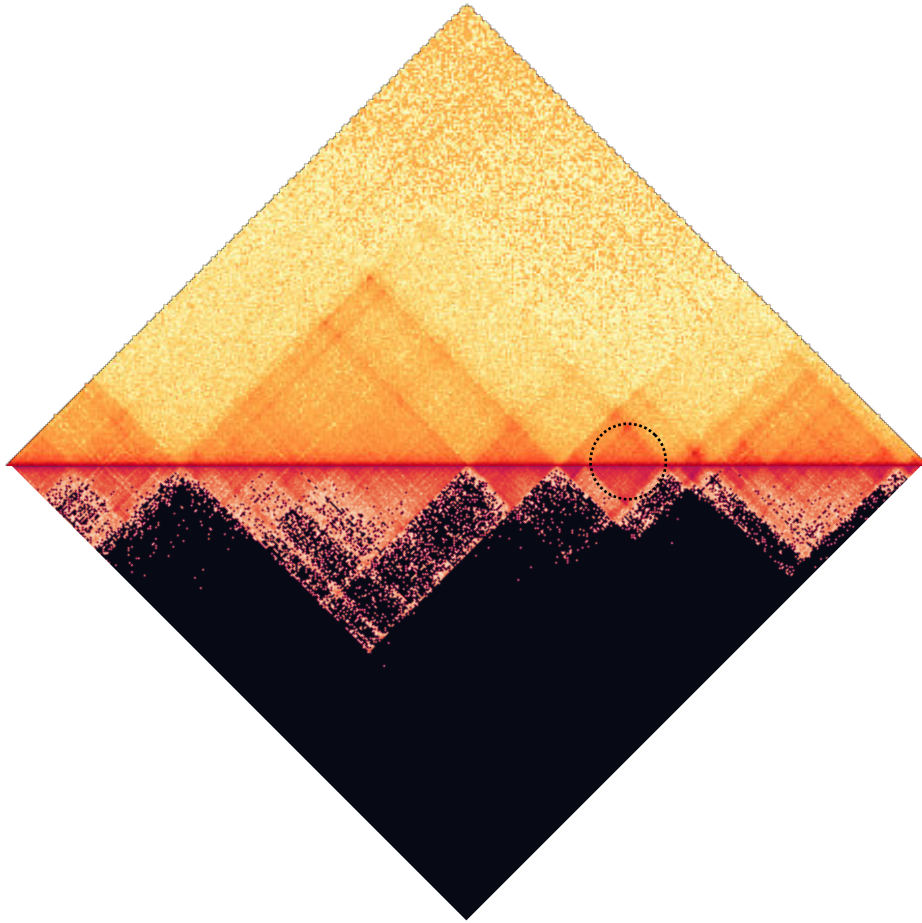
Spatial lay-out of significant interactions

chr11:110780000-114770000



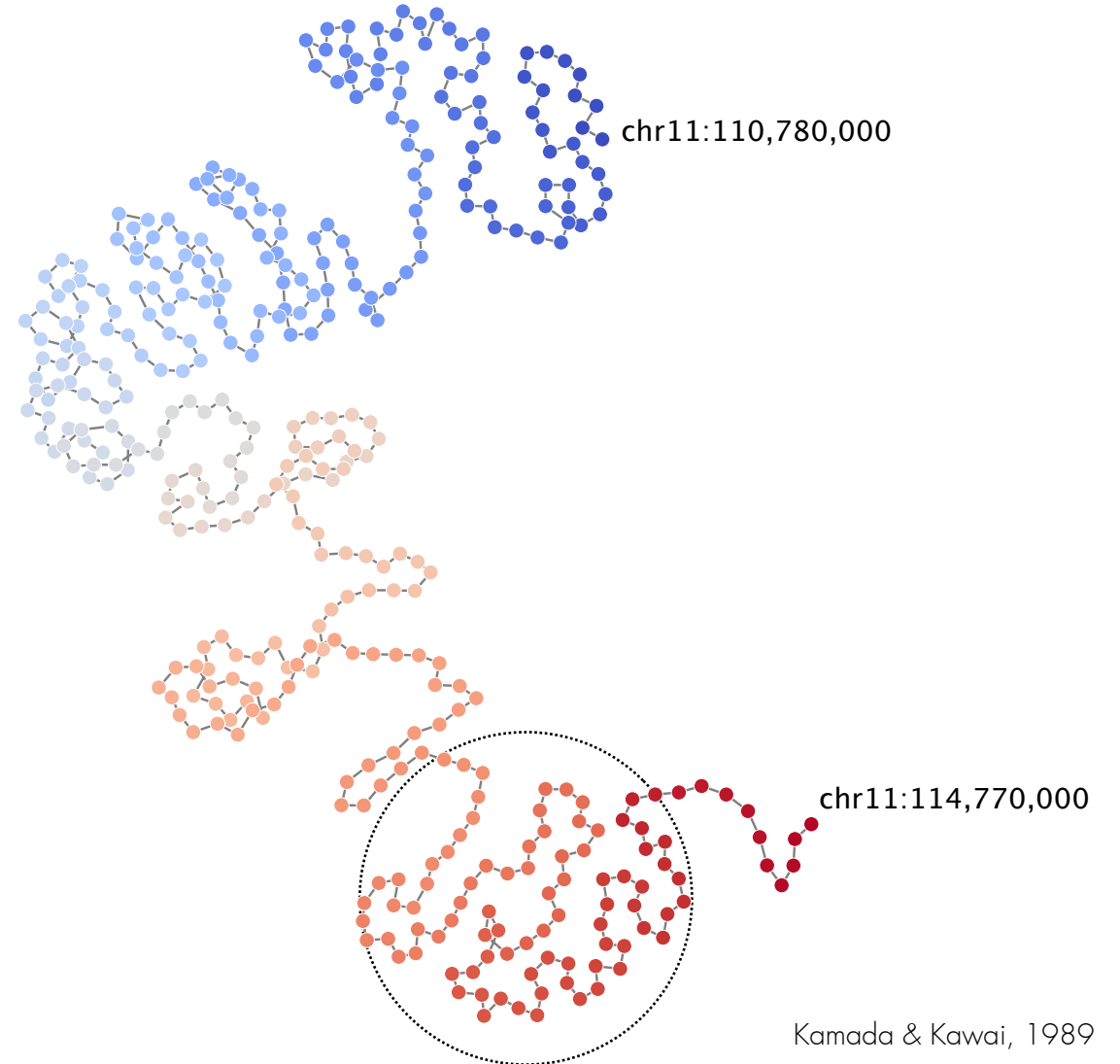
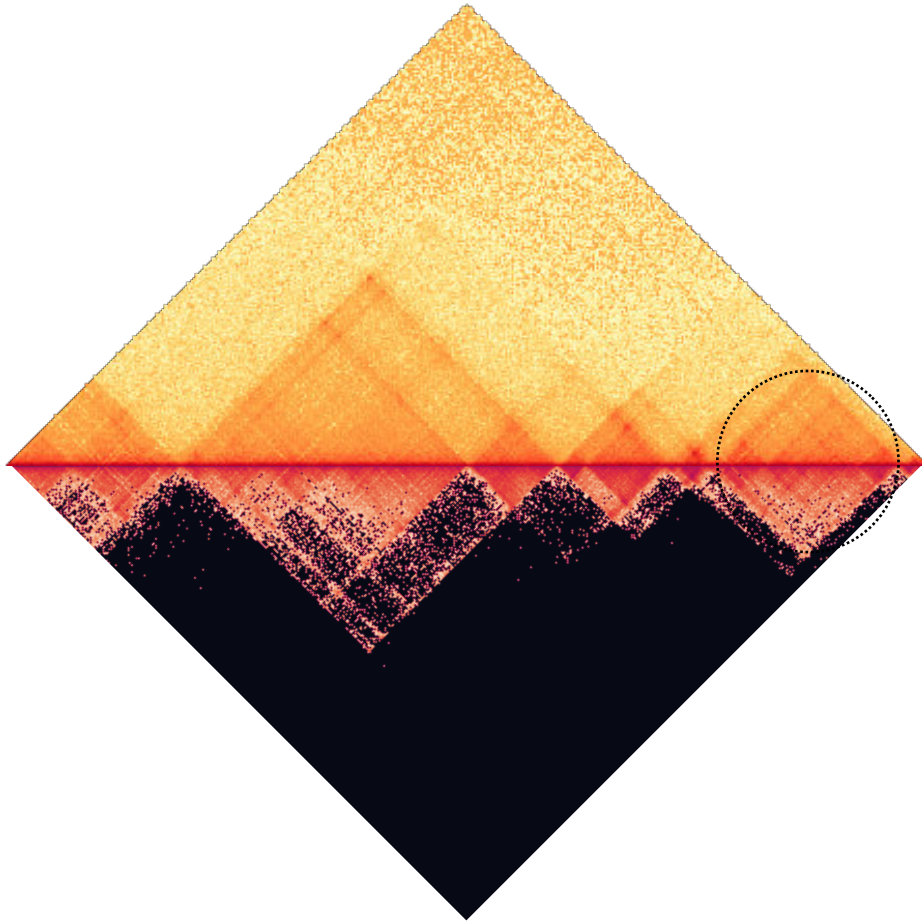
Spatial lay-out of significant interactions

chr11:110780000-114770000



Spatial lay-out of significant interactions

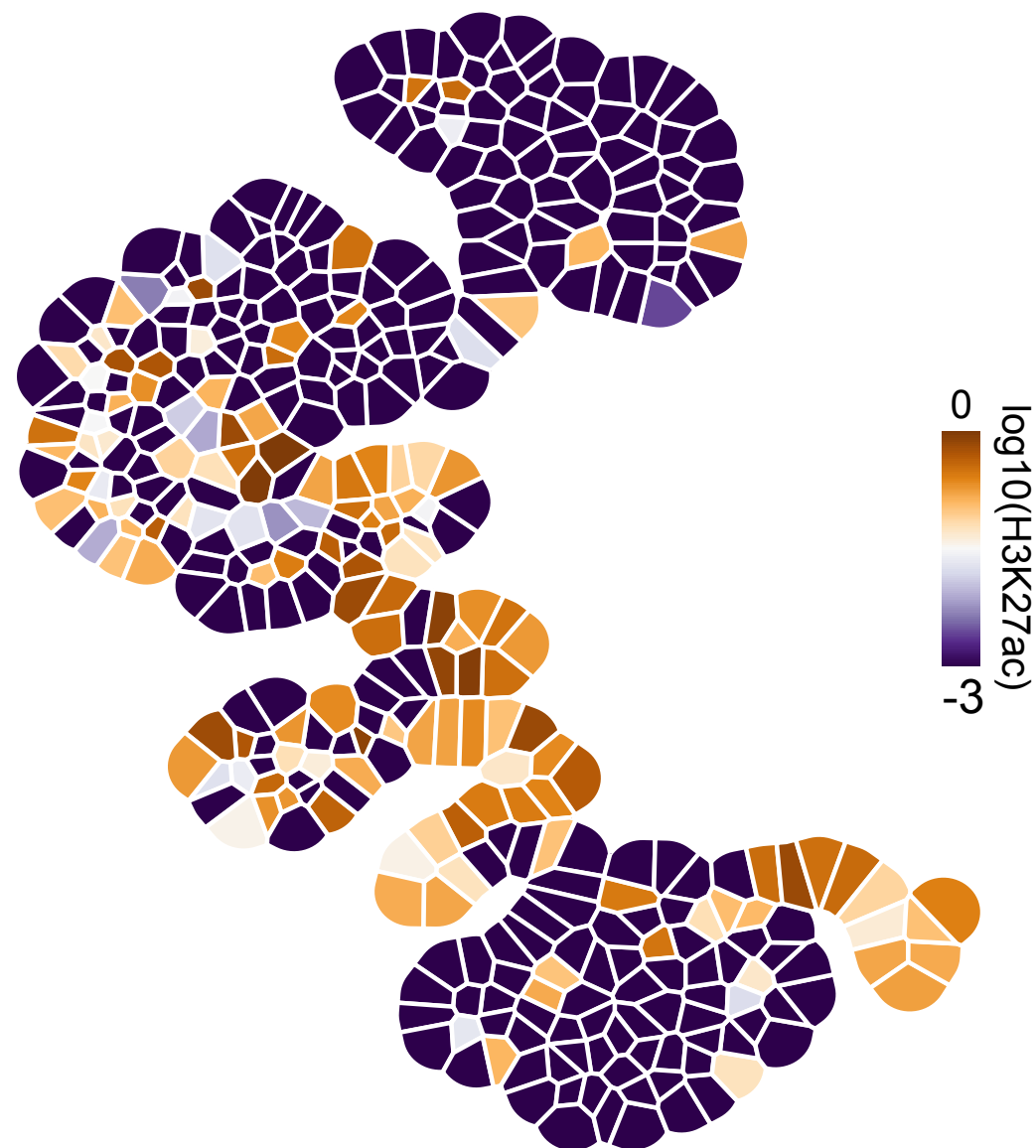
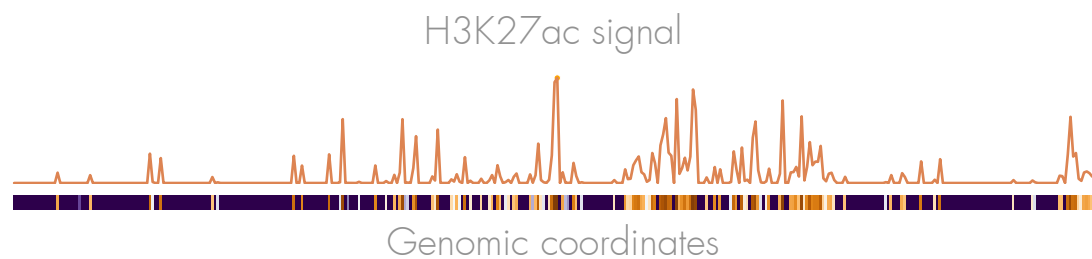
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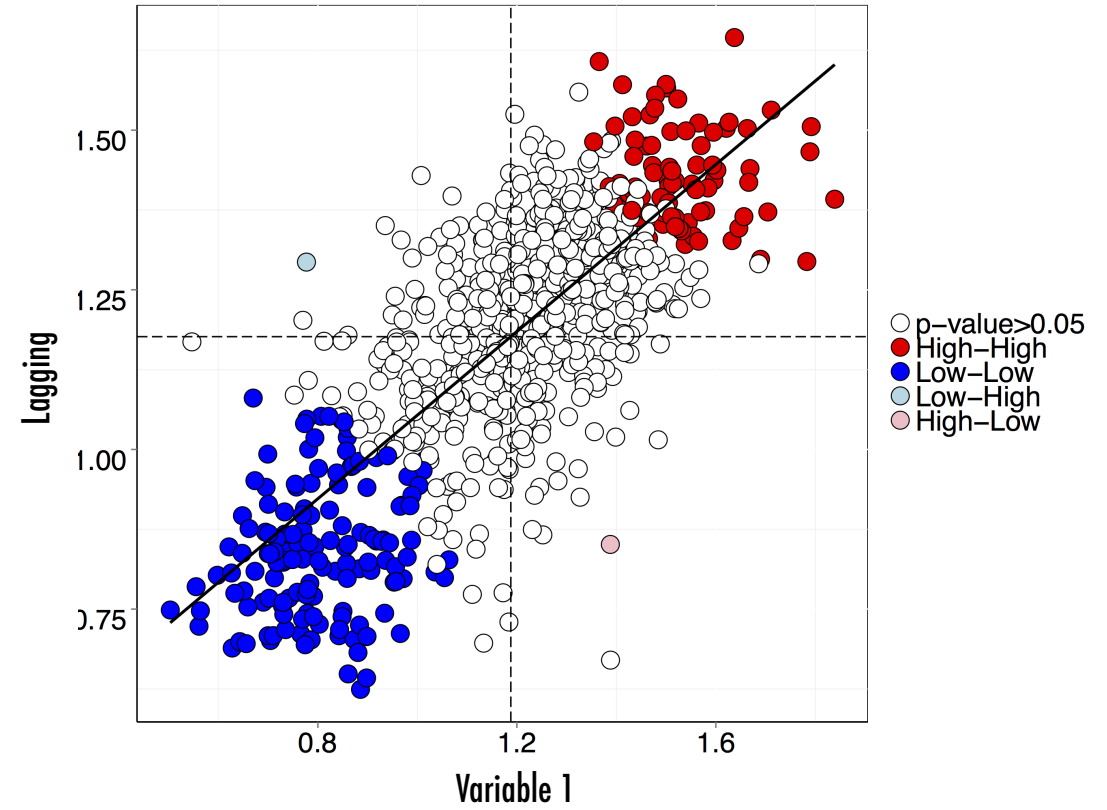
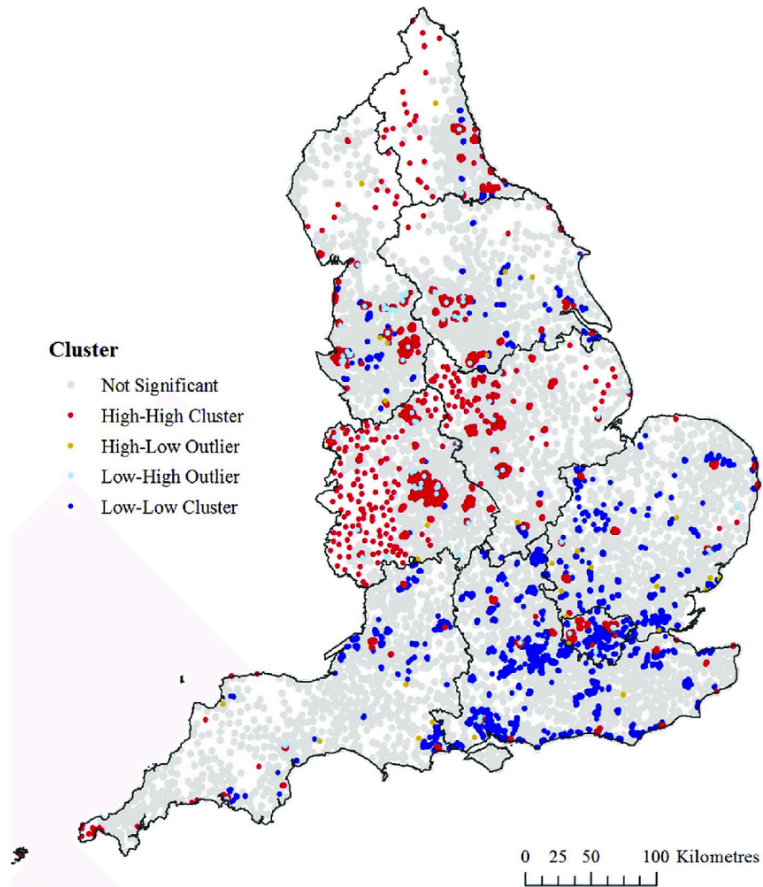
Kamada & Kawai, 1989

Marker (H3K27ac) into 2D mapping

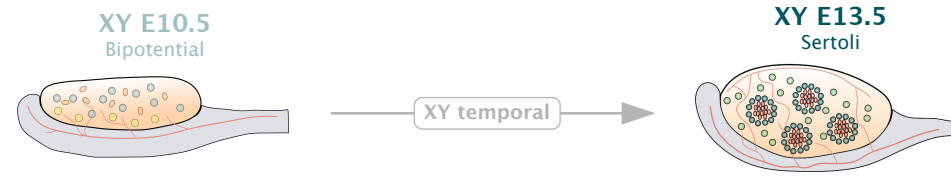
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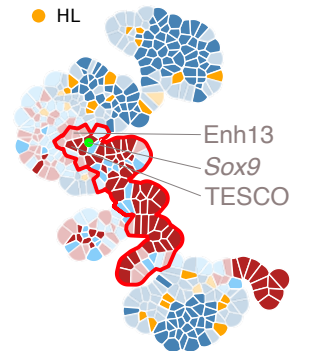
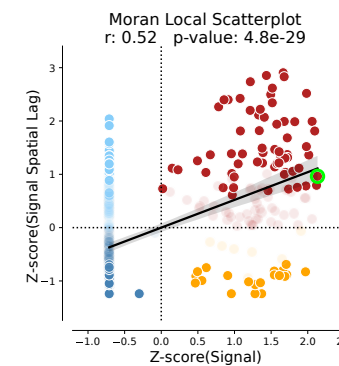
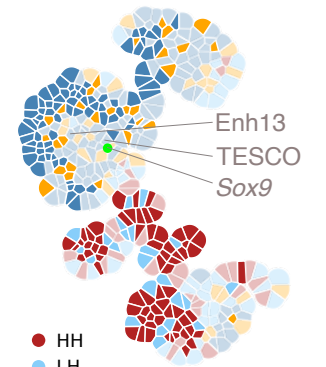
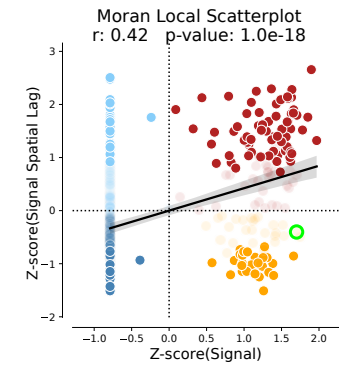
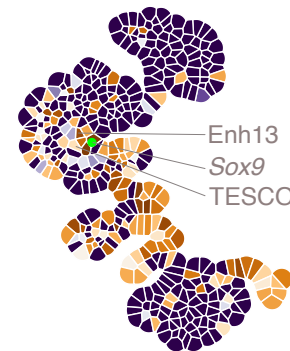
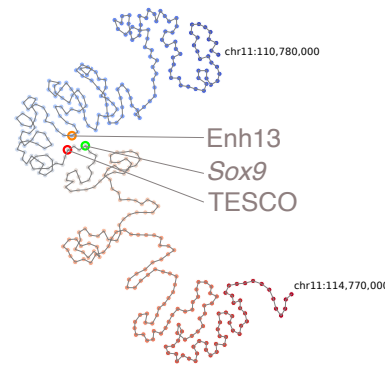
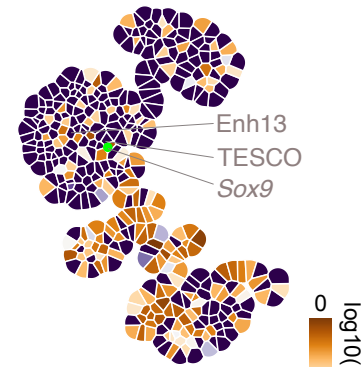
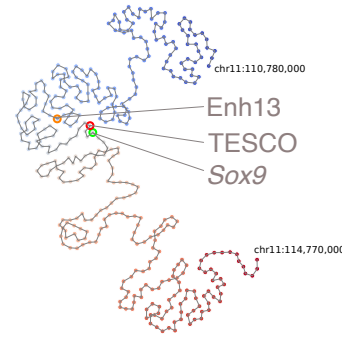
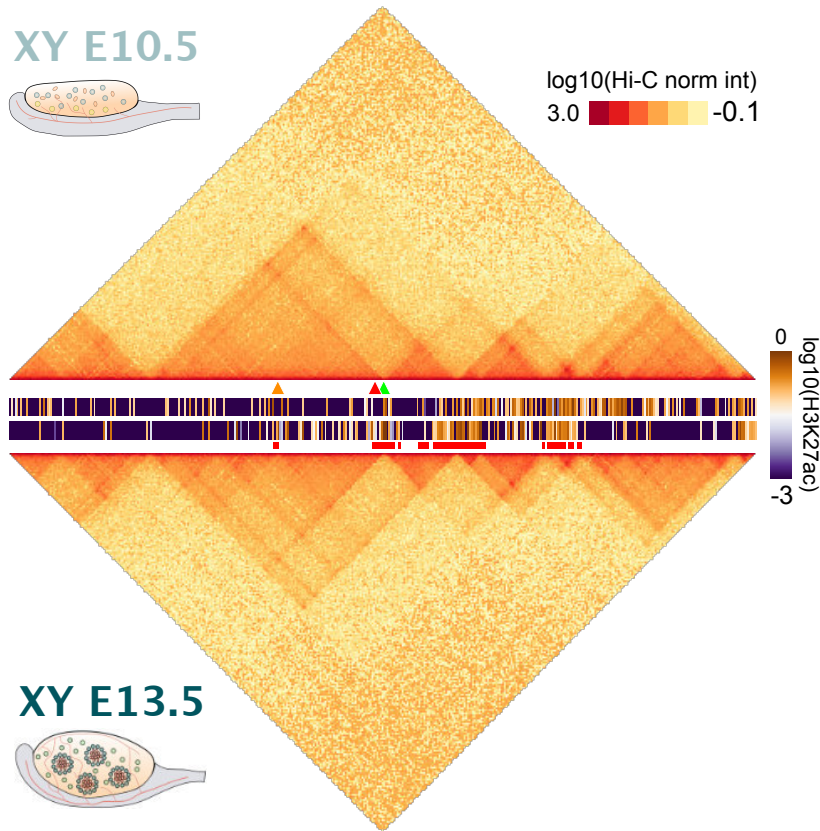
Local Moran Index



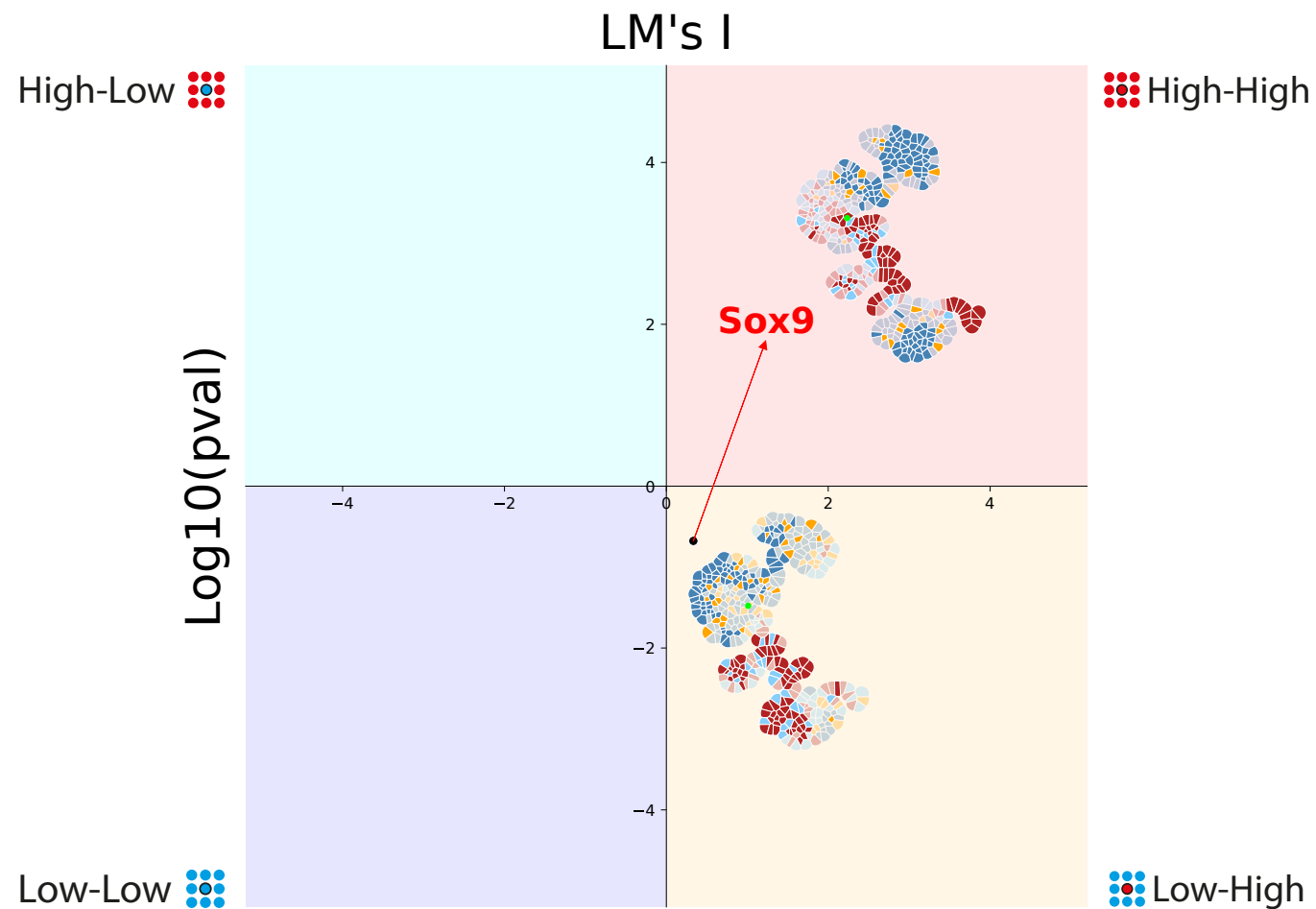
Quantifying regulatory environments bin by bin



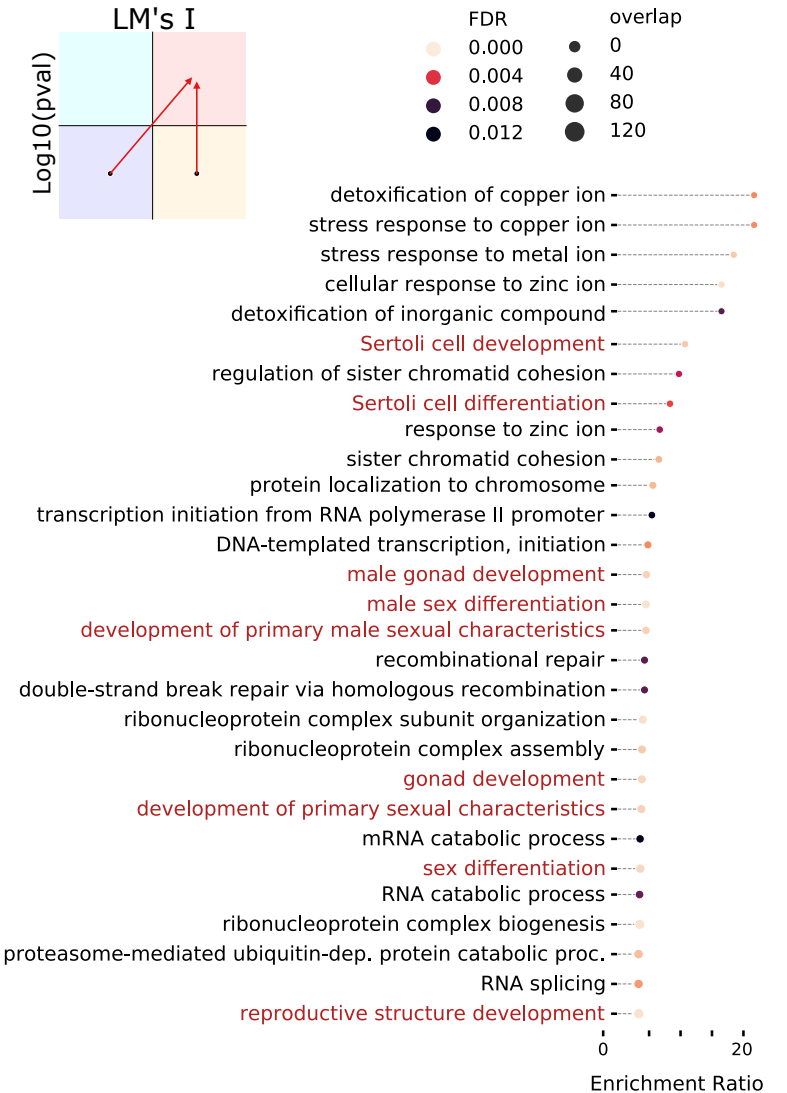
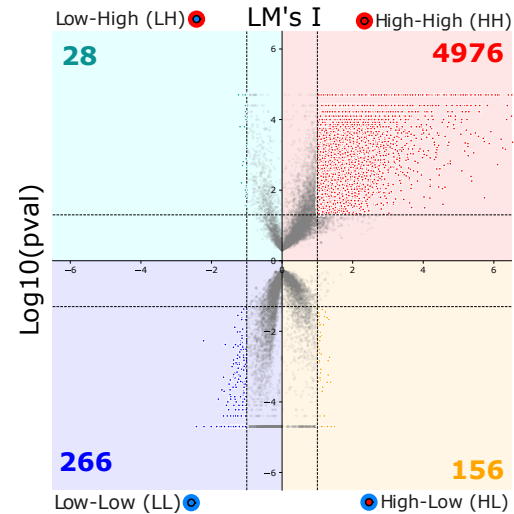
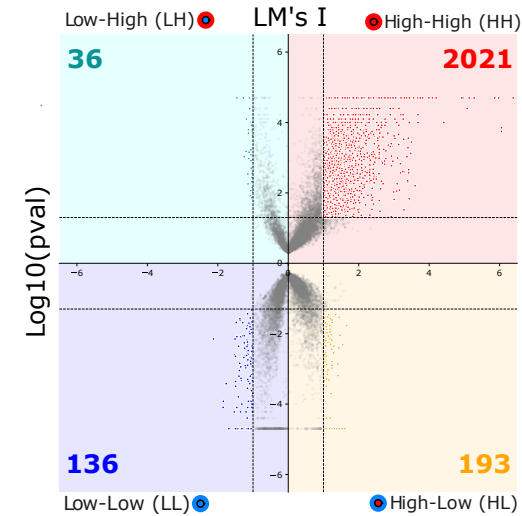
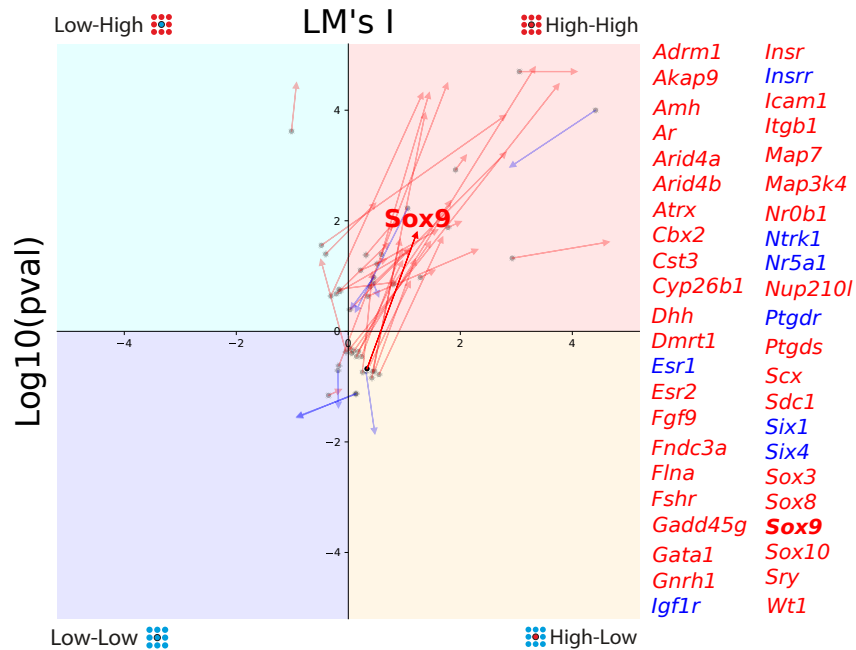
Sox9 locus chr11:110,780,000-114,770,000



LMI Trip for Sox9 gene

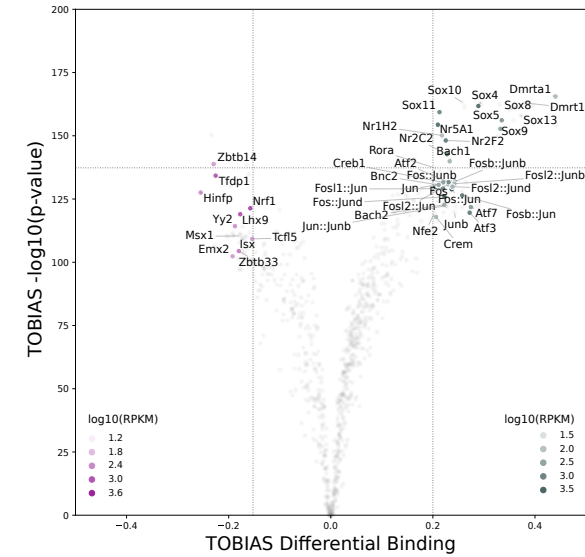
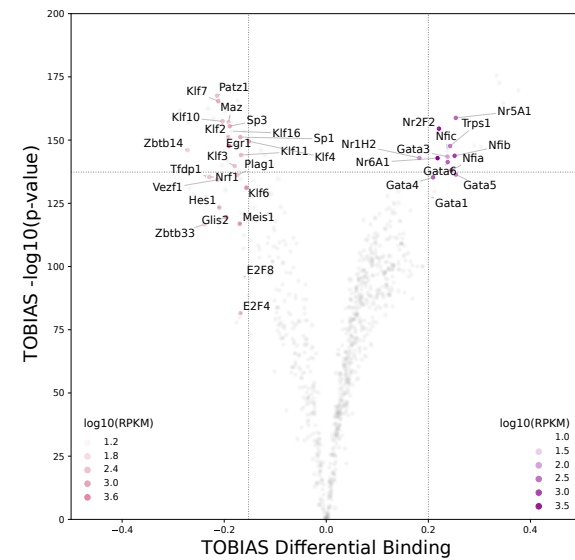
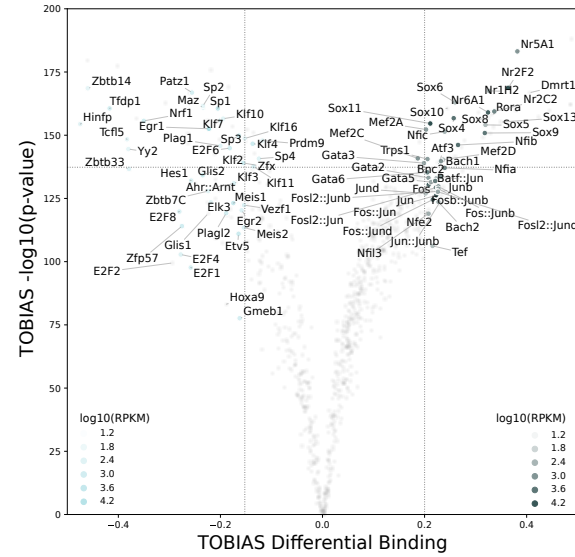
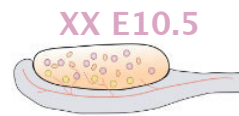
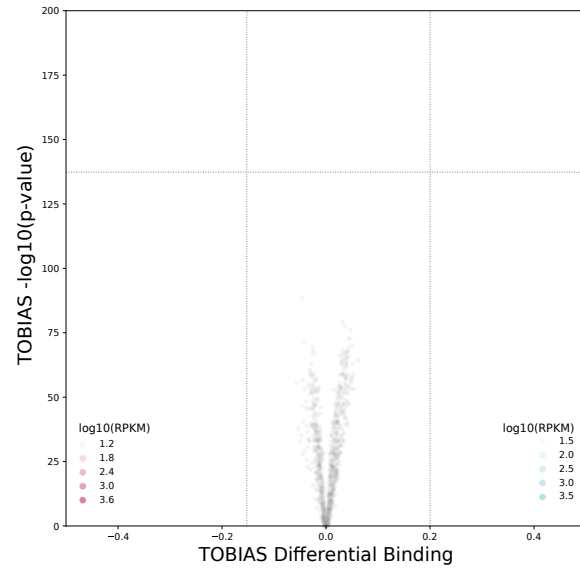
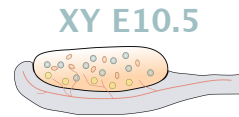


All genes LMI Trip



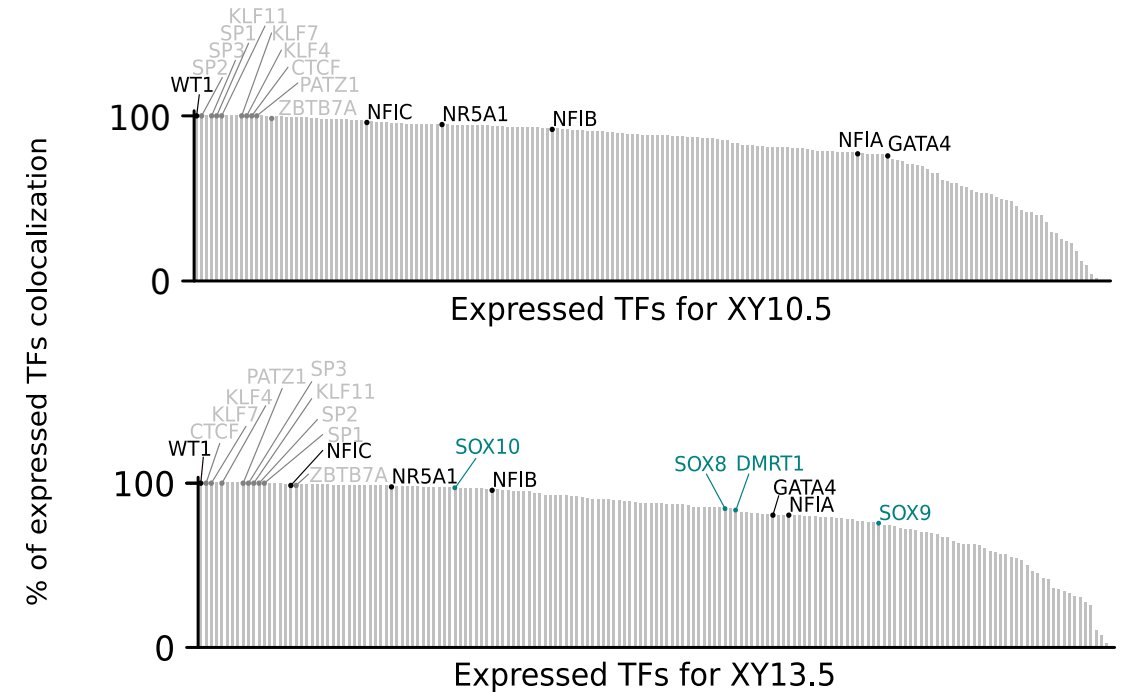
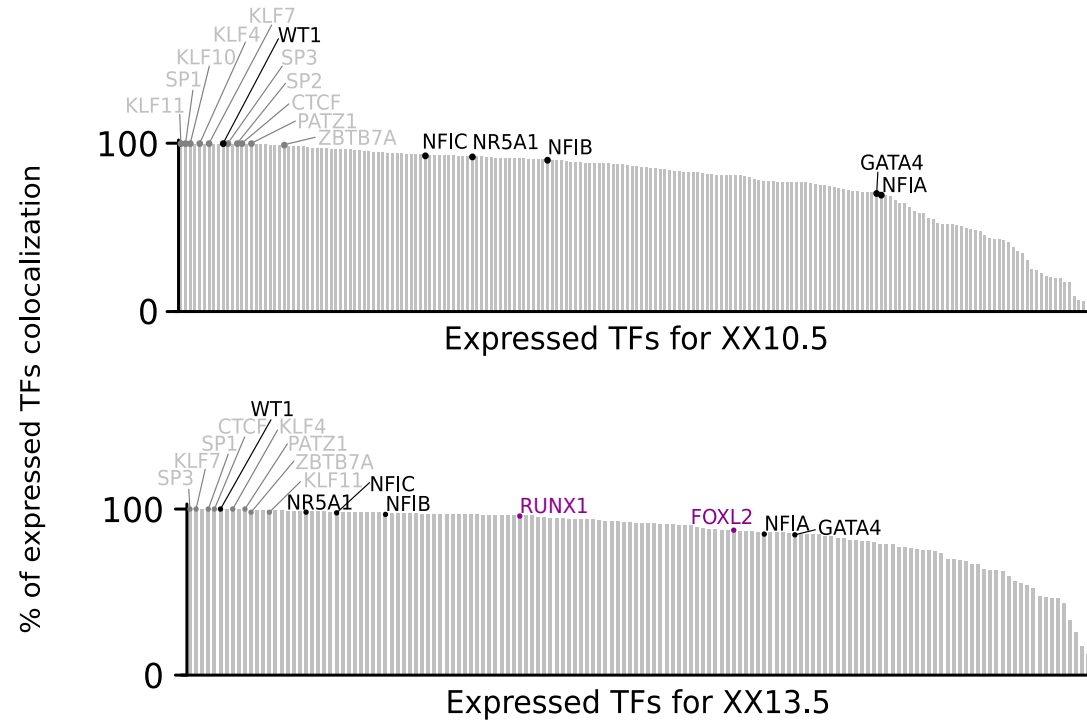


TF footprints





"Stripe" TFs

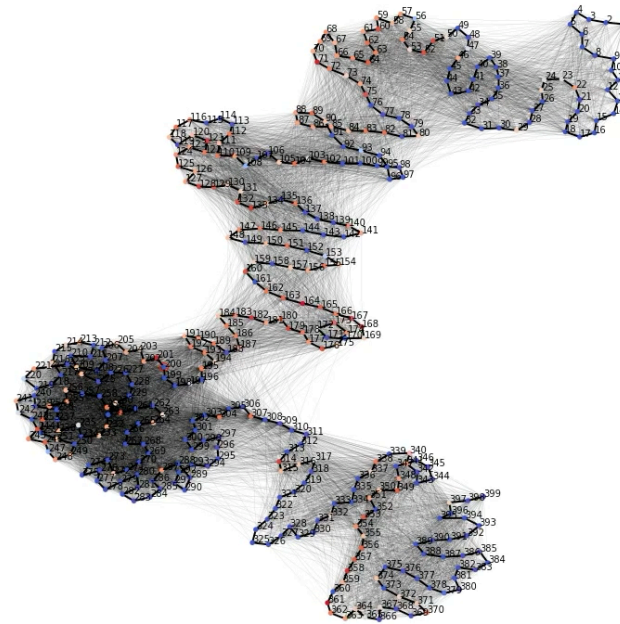


Now that we know the genes....

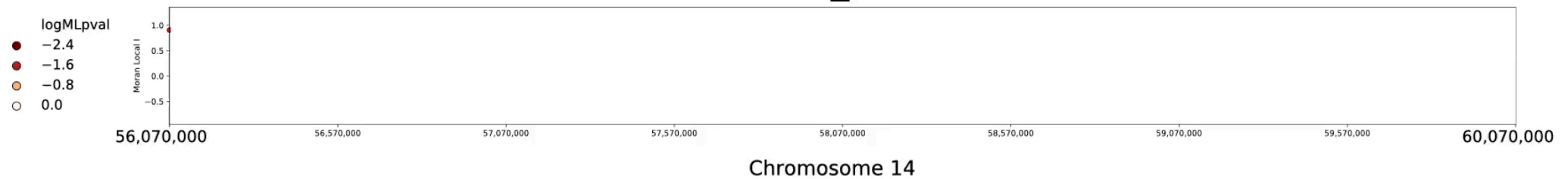
Can we identify regulatory elements using  METALoci

METALoci predictive mode

Fgf9 locus chr14:56,070,000-60,070,000



Fgf9

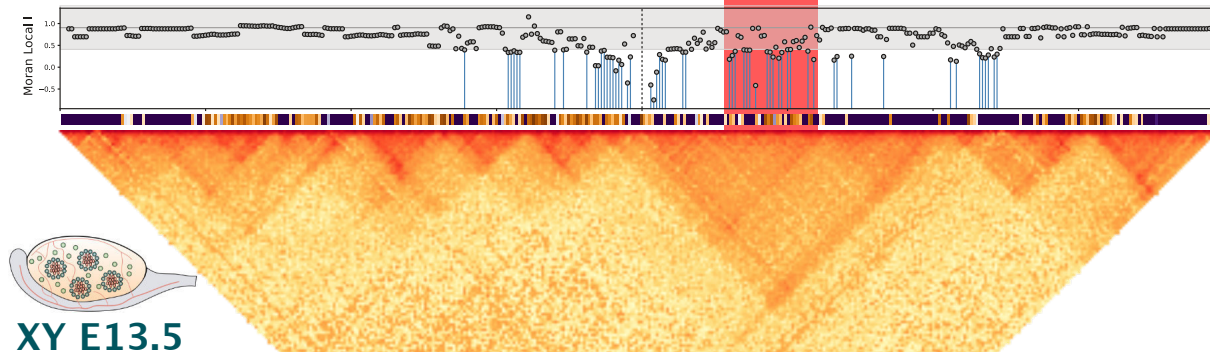
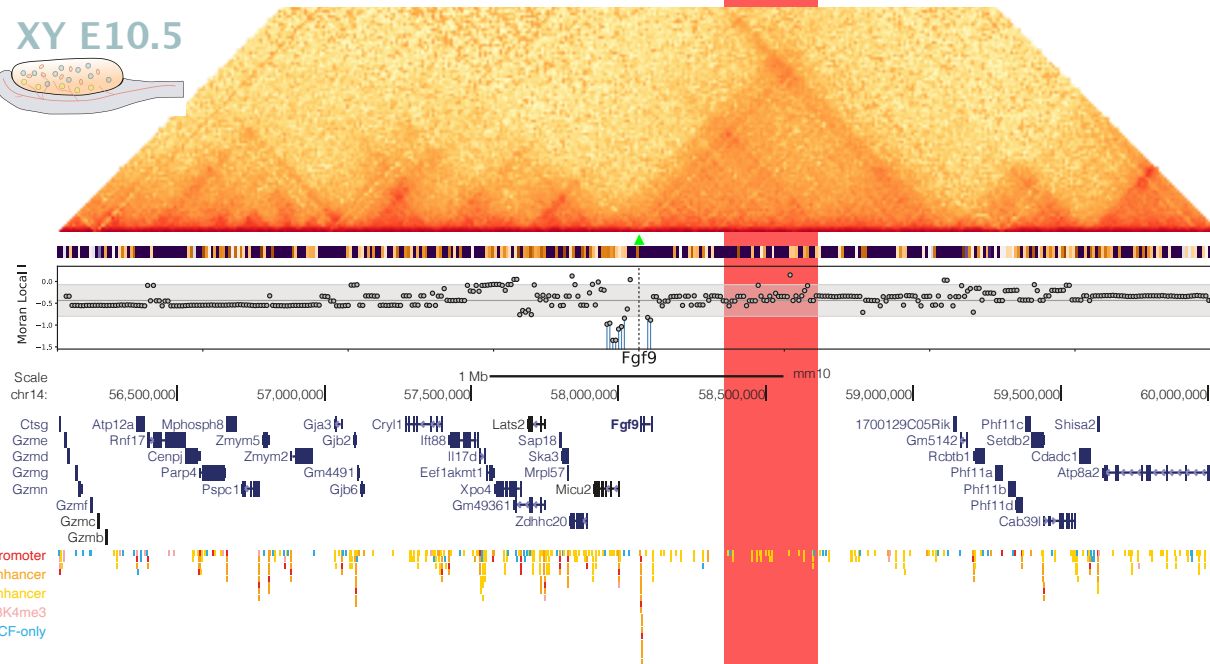
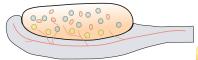


METALoci predictive mode

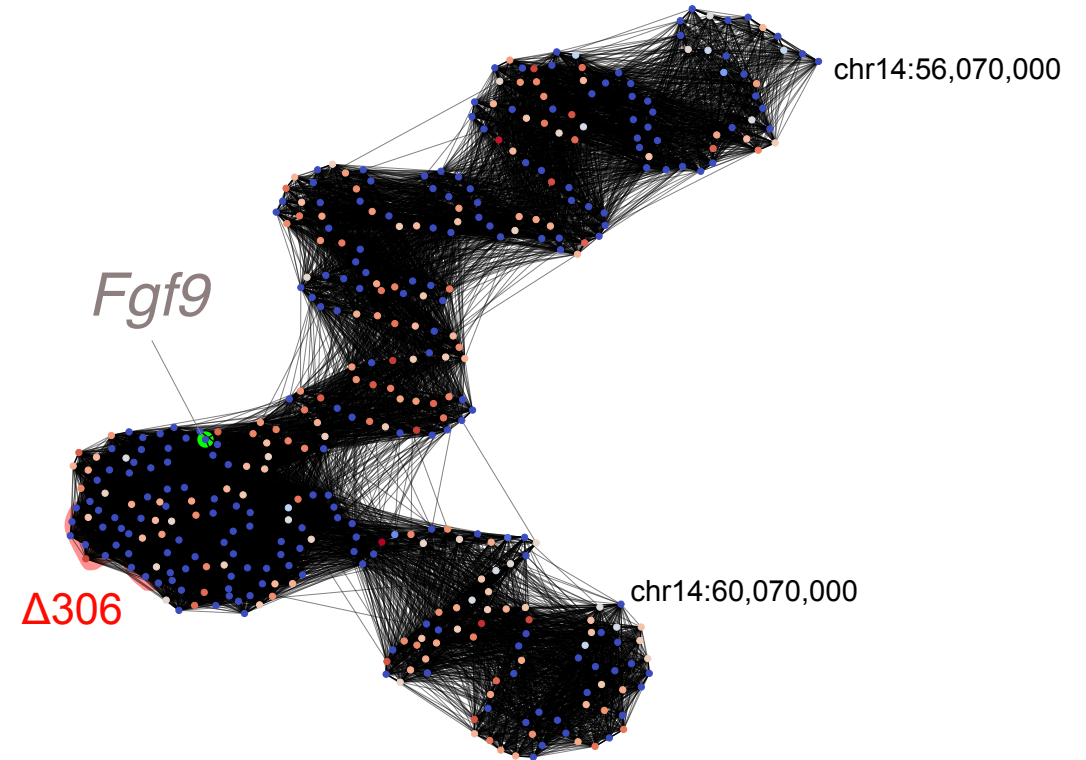
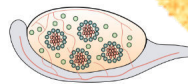
Fgf9 locus chr14:56,070,000-60,070,000

$\Delta 306$

XY E10.5



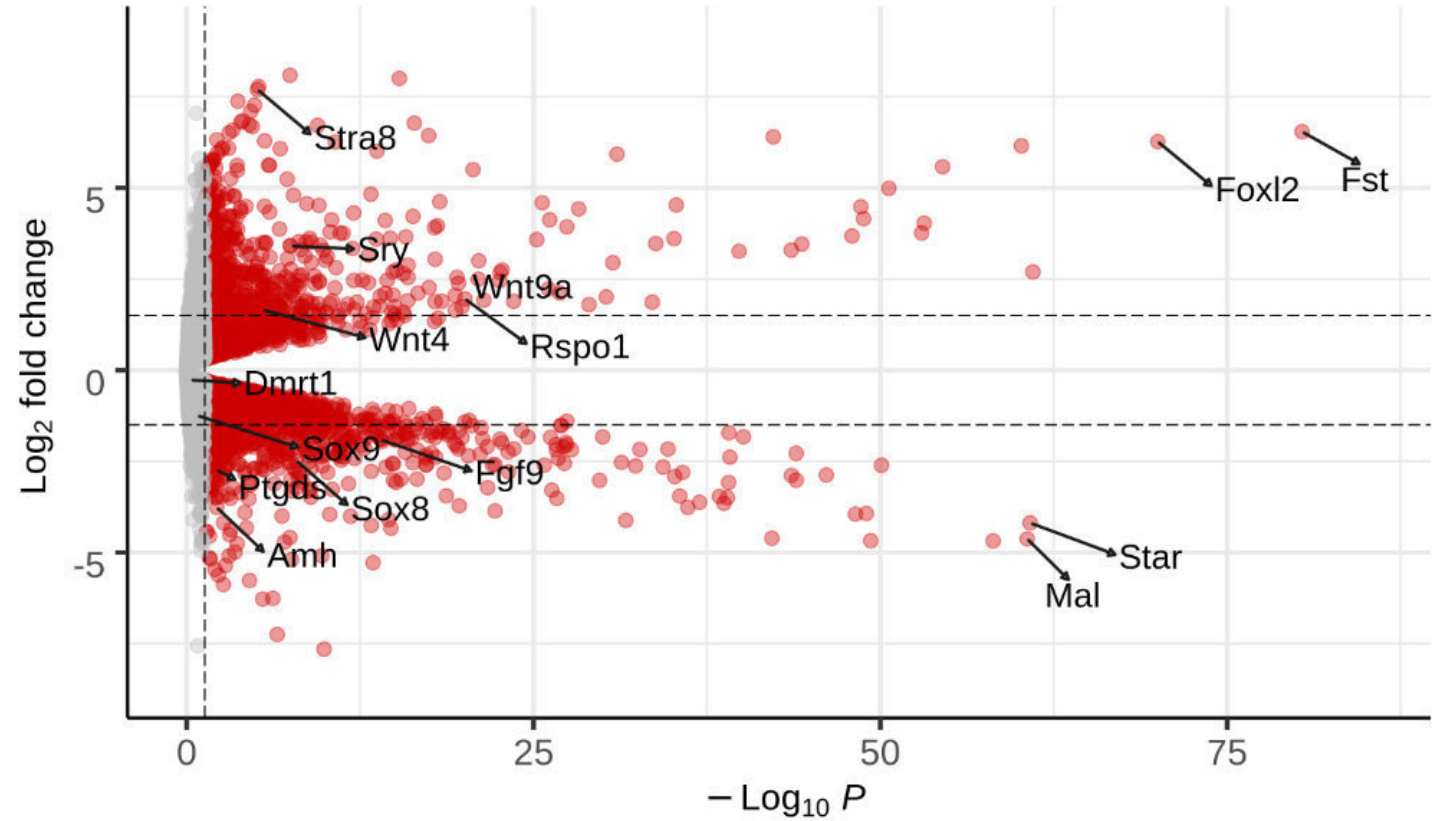
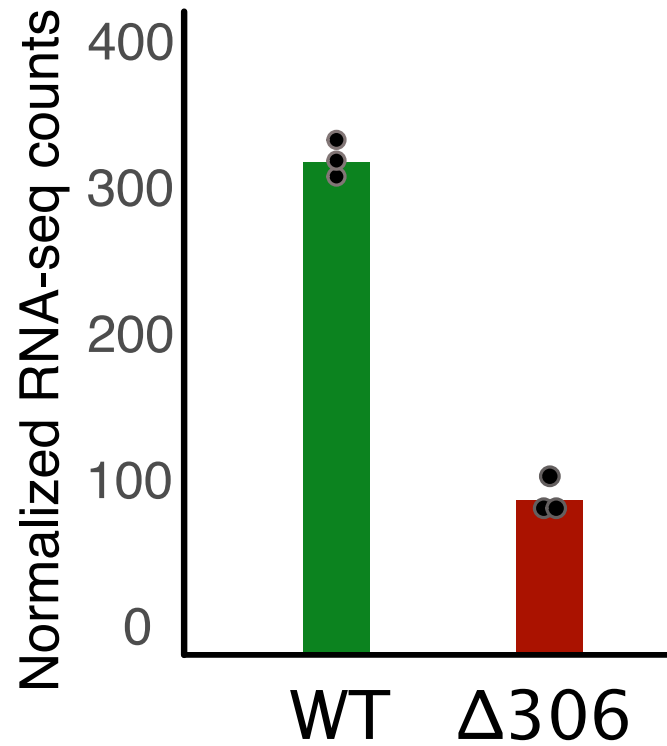
XY E13.5



$\Delta 306$

METALoci predictive mode

Fgf9 XY $\Delta 306$ mutant



METALoci predictive mode

Fgf9 XY $\Delta 306$ mutant

XY Wildtype



Testis

XY $\Delta 306$



Ovotestis



Ovary-like

XX Wildtype

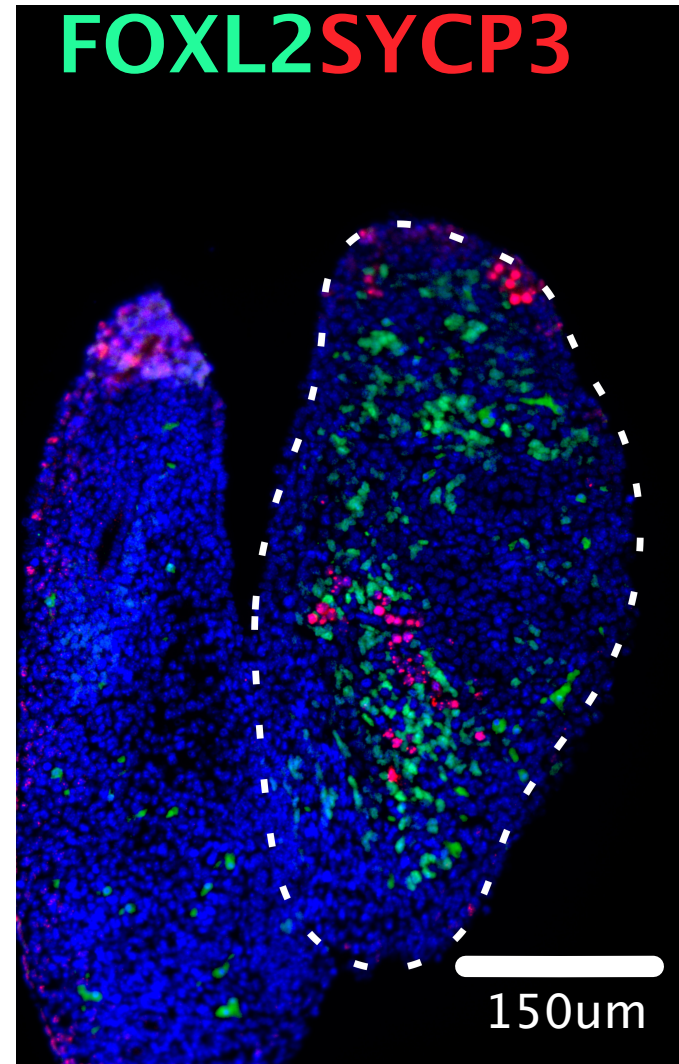
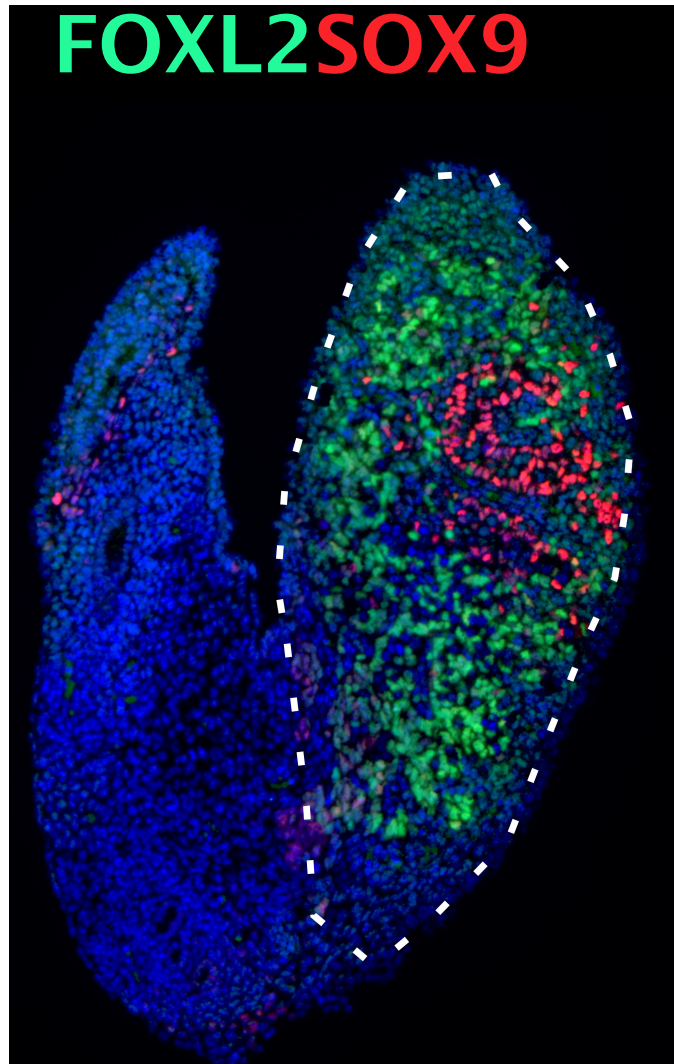


Ovary

250um

METALoci predictive mode

Fgf9 XY $\Delta 306$ mutant



Take home messages:

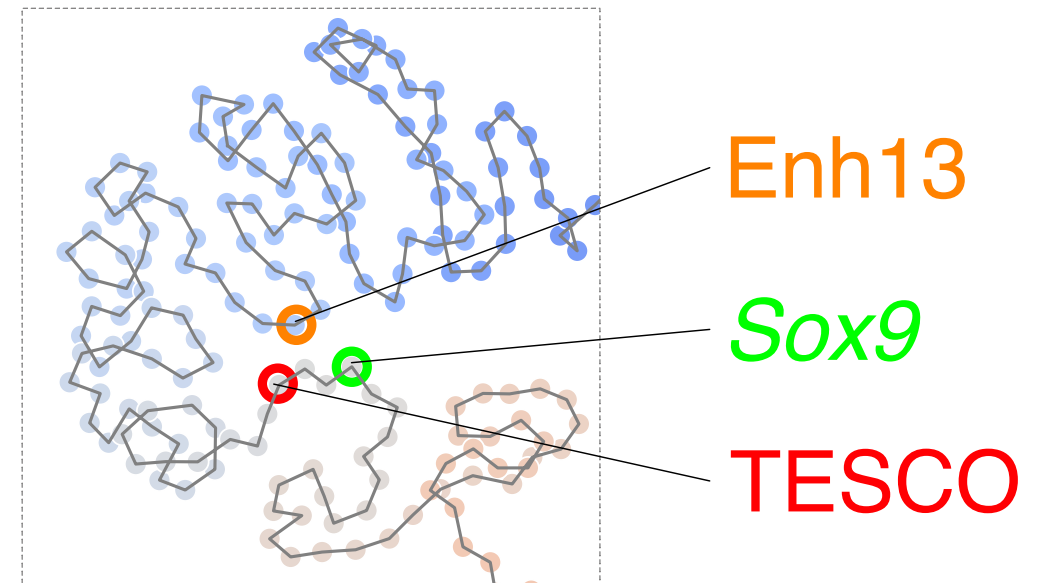
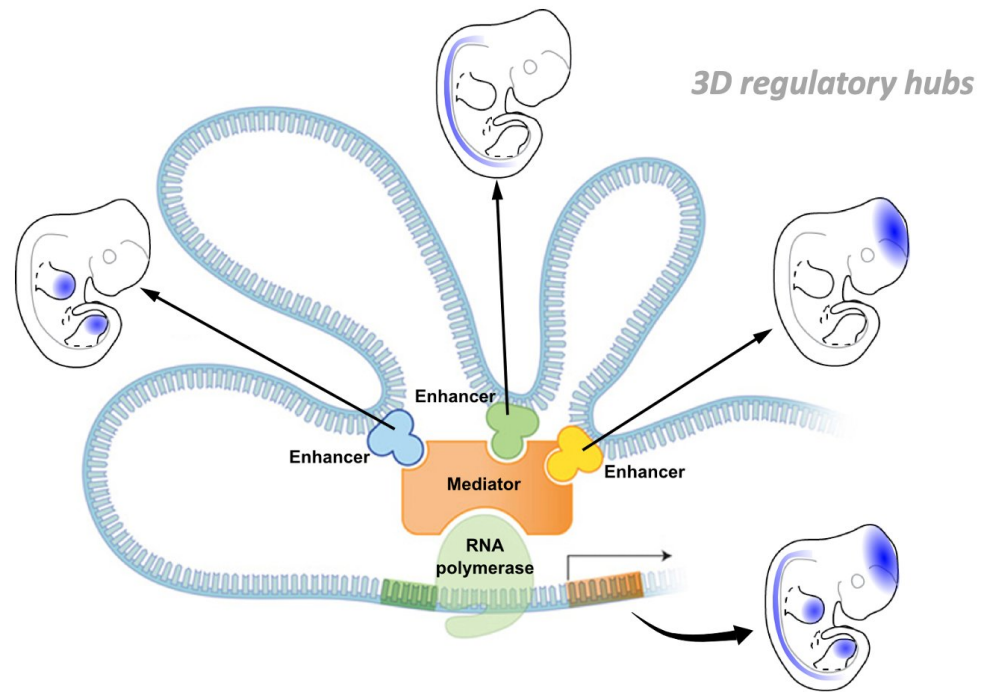
First characterization of the 3D regulatory landscape of sex determination

METALoci is an unbiased approach to quantify gene regulatory activity

METALoci is a predictive tool to identify critical regulatory loci

Discovery of a novel non-coding region controlling sex determination

Take home messages:



<http://marciuslab.org>
<http://3DGenomes.org>

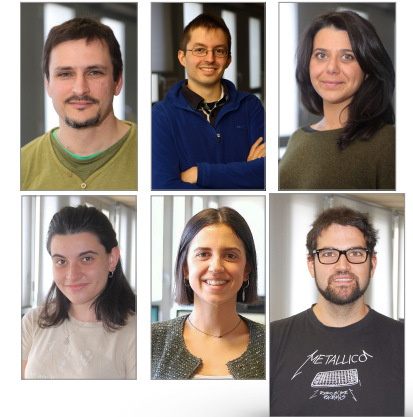
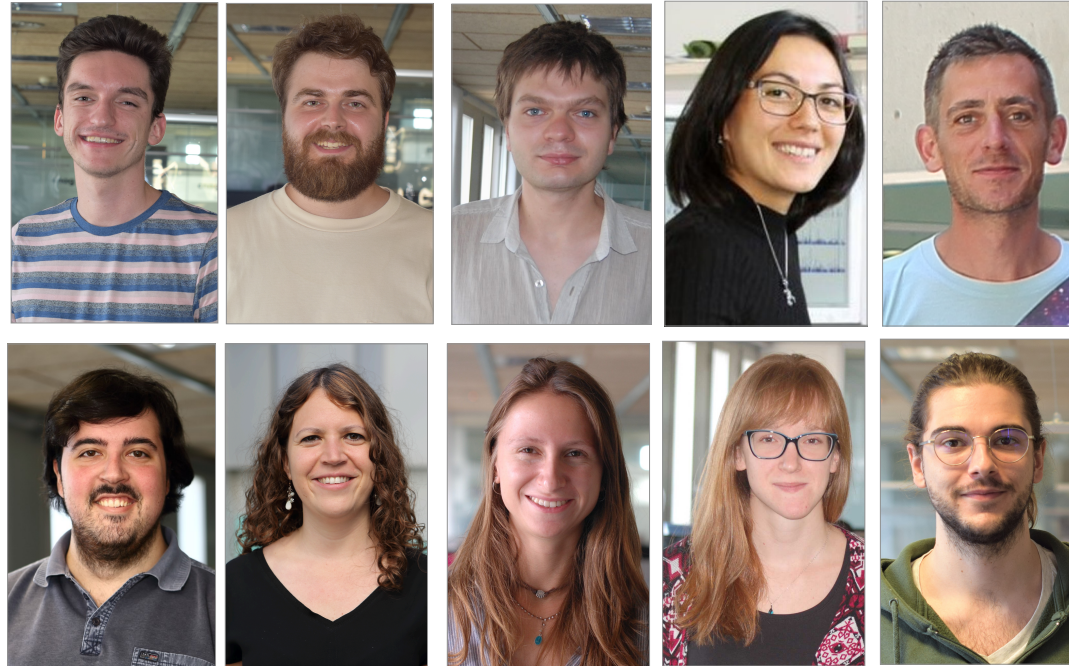
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∴ Our current sponsors ∴



∴ Conflict of Interest Statement ∴

Marc A. Marti-Renom serves as a consultant to Acuity Spatial Genomics, Inc., and receives compensation for these services.