

# Hidden Hamiltonian Cycle Recovery via Linear Programming

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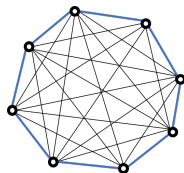
Joint work with  
Vivek Bagaria (Stanford), Jian Ding (Penn), David Tse (Stanford)  
and Yihong Wu (Yale)

Vilnius, July 5, 2018

# Mathematical problem: Hidden Hamiltonian cycle model

- Given a weighted undirected complete graph on  $n$  vertices
- Latent: a Hamiltonian cycle  $C^*$
- Edge weight

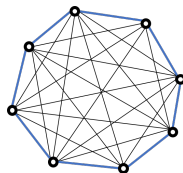
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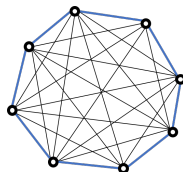


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## Remarks:

- For this talk,  $Q = N(0, 1)$  and  $P = N(\mu, 1)$ , so that

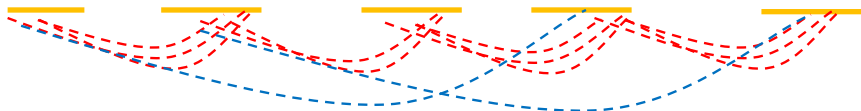
$$W = \mu \cdot \underbrace{\text{adj matrix of } C^*}_{\text{"signal"}} + \text{noise}$$

- Hidden Hamiltonian cycle planted in Erdős-Rényi graph  
[\[Broder-Frieze-Shamir '94\]](#)

# Motivation: Link information in Chicago datasets

- ① Reconstitute chromatin in vitro upon naked DNA
- ② Produce cross-links by fixing chromatin with formaldehyde

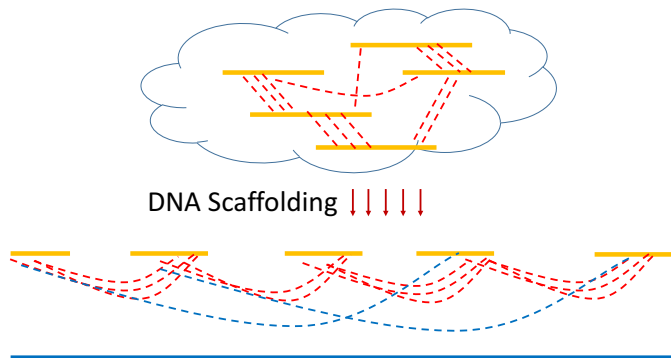
AGCTCGACTTGCAATTTCCGAGCTATGGCCAGTACTGCATACGGGCTTACGCGTAC



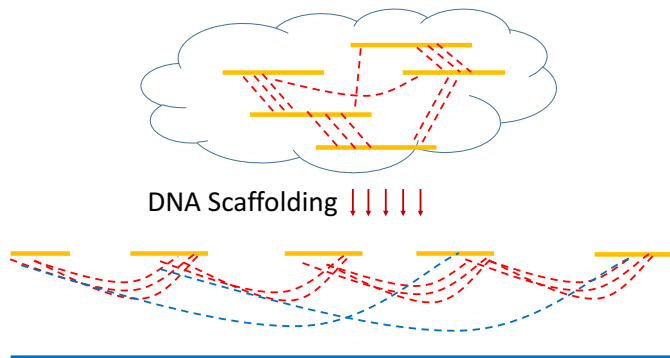
Chicago datasets generate cross-links among contigs [Putnam et al. '16]

On average **more** cross-links exist between **adjacent** contigs

# Ordering DNA contigs with Chicago cross-links



# Ordering DNA contigs with Chicago cross-links



Reduces to traveling salesman problem (TSP)

Find a path (tour) that visits every contig exactly once with the maximum number of cross-links

# Key challenges for DNA scaffolding with Chicago data

- Computational: TSP is NP-hard in the **worst-case**
- Statistical: spurious cross-links between contigs that are far apart



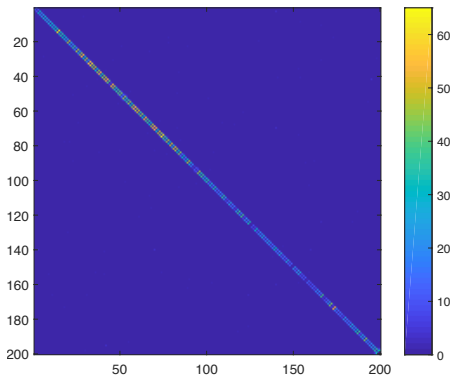
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## Key questions:

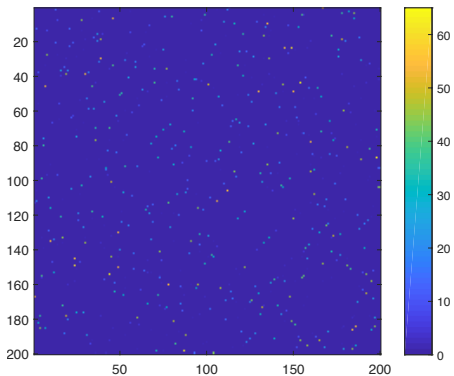
- How to **efficiently** order hundreds of thousands of contigs?
- How much **noise** can be tolerated for accurate DNA scaffolding?

# Mathematical model for DNA scaffolding



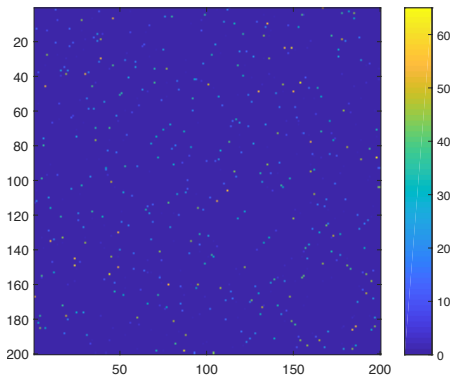
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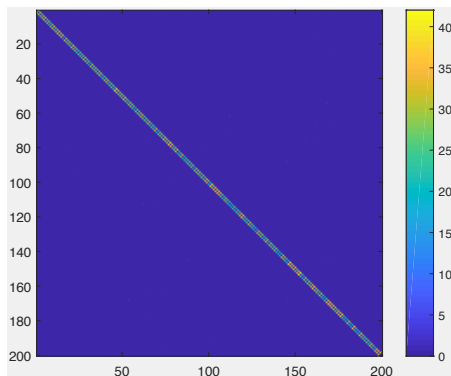


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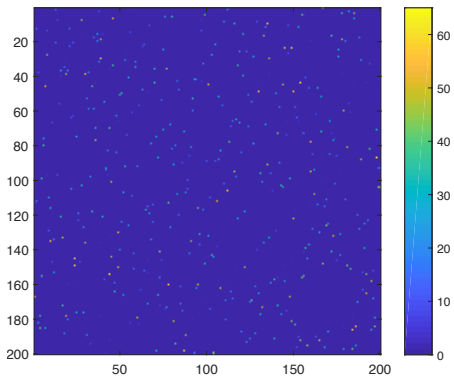


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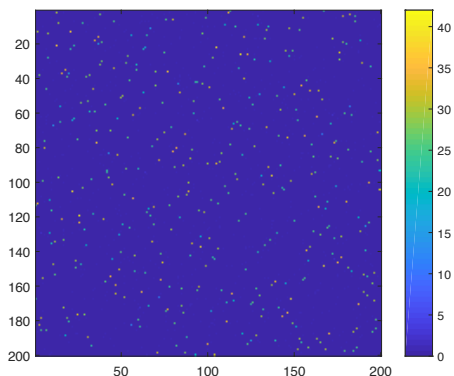


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# Mathematical model for DNA scaffolding



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# What is known information-theoretically

Maximum likelihood estimator reduces to TSP

$$\hat{X}_{\text{TSP}} = \arg \max_X \langle W, X \rangle$$

s.t.  $X$  is the adjacency matrix of some Hamiltonian cycle

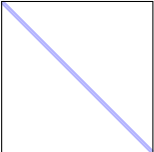
## Theorem (Sharp threshold)

*If  $\mu^2 < 4 \log n$ , exact recovery is information-theoretically impossible*

*If  $\mu^2 > 4 \log n$ , MLE succeeds in exact recovery*

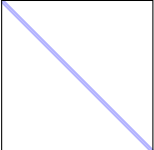
# What is known algorithmically

- **Spectral methods** fail miserably:
  - ▶  $\mu \gg n^{2.5}$  (spectral gap of cycle is too small)

$$W = \mu \begin{array}{|c|} \hline \square \\ \hline \end{array} + \text{Gaussian noise}$$


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- This talk: **linear programming** achieves sharp threshold

$$\begin{array}{l} \frac{\mu^2}{\log n} > 4 : \quad \text{LP succeeds} \\ \frac{\mu^2}{\log n} < 4 : \quad \text{Everything fails} \end{array}$$

Threshold determined by **Battacharyya distance** (a.k.a. Rényi divergence of order  $\frac{1}{2}$ ):

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LP succeeds when

$$B(P, Q) - \log n \rightarrow +\infty$$

**optimal** under mild assumptions

## Convex relaxations of TSP

$$\begin{aligned}\hat{X}_{\text{TSP}} &= \arg \max_X \langle W, X \rangle \\ \text{s.t.} \quad & \sum_j X_{ij} = 2, \quad \forall i \\ & X_{ij} \in \{0, 1\} \\ & \sum_{i \in I, j \notin I} X_{ij} \geq 2, \quad \forall \emptyset \neq I \subset [n]\end{aligned}$$

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- The last constraint: subtour elimination

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- Replacing the integrality constraint with box constraint: **SUBTOUR LP** relaxation [Dantzig-Fulkerson-Johnson '54, Held-Karp '70]
- Exponentially many linear constraints, nevertheless solvable using interior point method

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- How it performs in our random instance?

Theorem (Bagaria-Ding-Tse-Wu-X. '18)

*If  $\mu^2 - 4 \log n \rightarrow \infty$ , then  $\widehat{X}_{\text{F2F}} = X^*$  with high probability.*

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Remarks

- Achieving the IT-limit  $\mu^2 = 4 \log n$

## Max-Product Belief Propagation

$$m_{i \rightarrow j}(t) = w_{ij} - 2^{\text{nd}} \max_{\ell \neq j} \{m_{\ell \rightarrow i}(t-1)\}$$

$$m_{i \rightarrow j}(0) = w_{ij}$$

After  $T$  iterations, for each vertex  $i$ , keep the two largest incoming messages  $m_{\ell \rightarrow i}(T)$  and delete the rest.

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- BP is exact provided the optimal solution of F2F is integral  
[Bayati-Borgs-Chayes-Zecchina '11]
- It can be shown that  $T = O(n^2 \log n)$  whp



Theoretical analysis of convex relaxation

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- Primal argument:
  - ▶ No feasible solution other than the ground truth has a better objective value whp
  - ▶ Key: for LP, can restrict to **extremal points** (vertices of the feasible polytope)

- KKT conditions (Farkas' lemma):  $\widehat{X}_{\text{F2F}} = X^* \iff \exists u \in \mathbb{R}^n$  (dual certificate):

$$u_i + u_j \leq W_{ij}, \quad \text{for } i \sim j \text{ in } C^*$$

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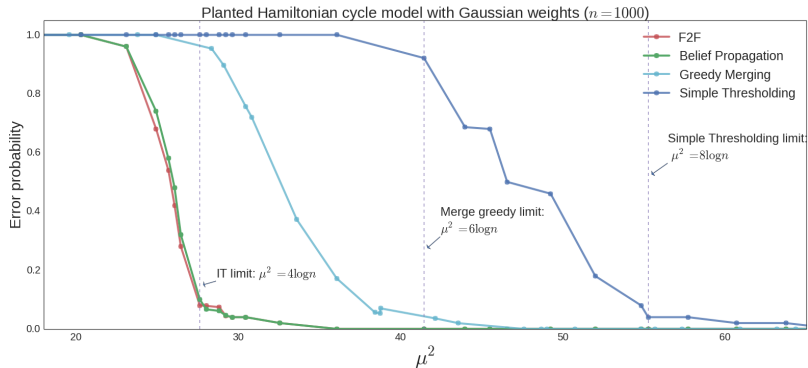
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- This certificate shows correctness if  $\mu^2 > 6 \log n$  (same as greedy merging)

# Synthetic data experiment



- Show whp for all extremal points  $X \neq X^*$ :

$$\langle W, X \rangle < \langle W, X^* \rangle$$

- F2F polytope:

$$\left\{ X \in [0, 1]^{n \times n} : \sum_{j=1}^n X_{ij} = 2 \right\}$$

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Proof of correctness for F2F LP



- 1 Encode the solution: for any extremal point  $X$ , represent  $2(X - X^*)$  as a **bicolored multigraph**  $G_X$

$$w(G_X) = \langle W, 2(X - X^*) \rangle$$

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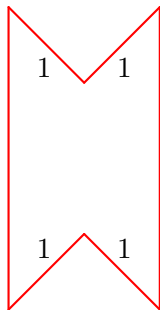
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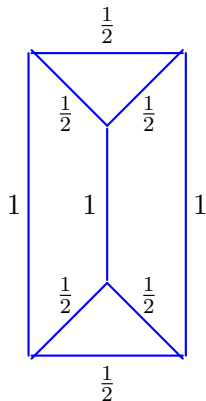
- ③ Counting: Show that whp  $w(F) < 0$  for all  $F \in \mathcal{F}$

# Step 1: Bicolored multigraph representation



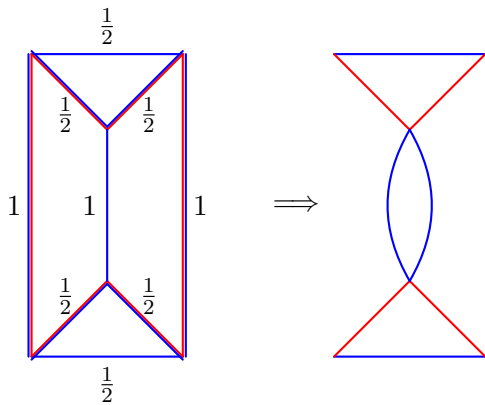
$X^*$ : true cycle

## Step 1: Bicolored multigraph representation



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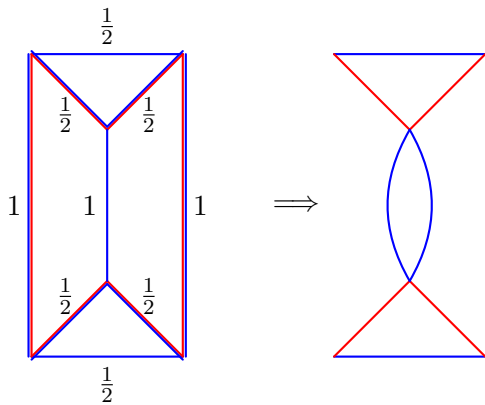
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key observation

$G_X$  is always balanced: red degree = blue degree

## Step 2: Edge decomposition

Theorem (Kotzig '68)

*Every connected balanced bicolored multigraph has an **alternating Eulerian circuit**.*



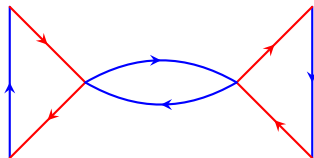
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### Remarks

- An Eulerian circuit may traverse a double edge twice

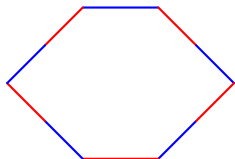


“Dumbbell” structure

## Step 2: Edge decomposition

$\mathcal{U}$ : collection of graphs recursively constructed

- 1 Start with an even cycle in alternating colors
- 2 **Blossoming procedure**: At each step, contract an edge in any cycle and attach a **flower** (alternating path of double edges followed by an alternating odd cycle)

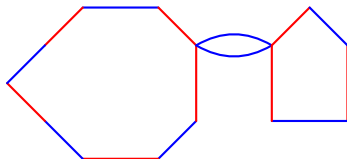


Obtained by starting with a 10-cycle and blossoming 3 times

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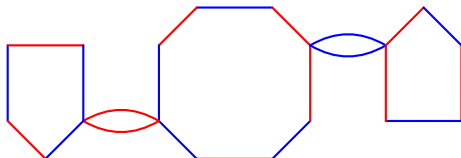


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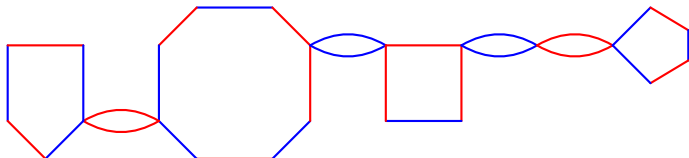


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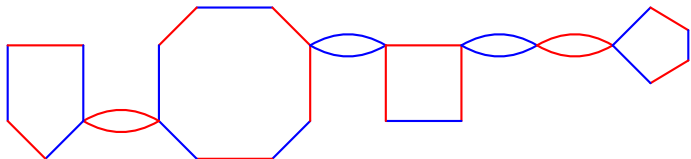


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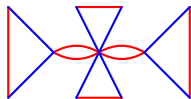
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- 1 Start with an even cycle in alternating colors
- 2 **Blossoming procedure**: At each step, contract an edge in any cycle and attach a **flower** (alternating path of double edges followed by an alternating odd cycle)

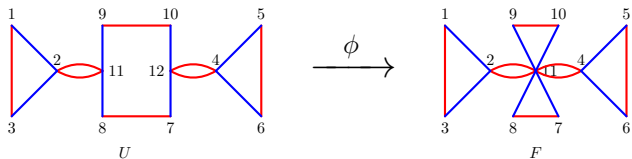


Obtained by starting with a 10-cycle and blossoming 3 times

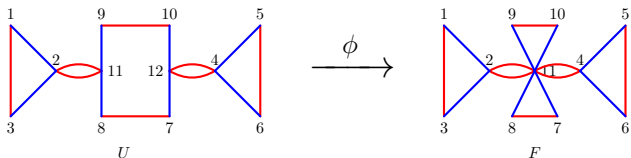
However, not every  $G_X$  is of this form...



- Graph homomorphism  $\phi : U \rightarrow F$  is a vertex map that preserves edges and edge multiplicity



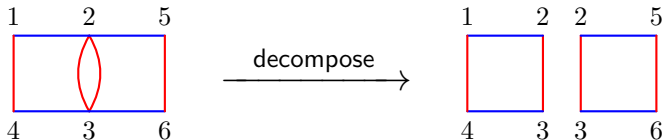
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### Lemma (Decomposition)

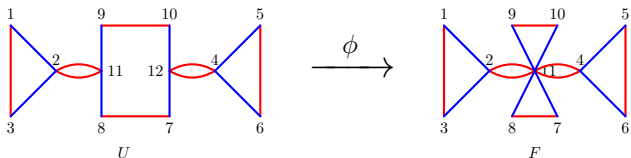
Every balanced bicolored multigraph  $G$  with edge multiplicity at most 2 can be decomposed as an edge-disjoint union of graphs in

$$\mathcal{F} = \{F : U \rightarrow F \text{ for some } U \in \mathcal{U}\}$$





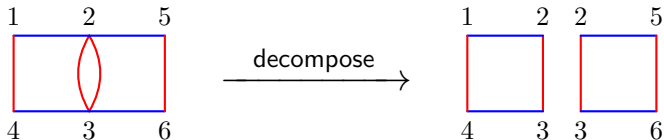
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- It remains to show  $\min_{F \in \mathcal{F}} w(F) < 0$  whp

## Step 3: Counting and large deviation arguments

$$\mathcal{F}_{k,\ell} = \{F \in \mathcal{F} : E(F) \text{ consists of } k \text{ double edges and } \ell \text{ single edges} \}$$

### Lemma

For any  $k \geq 0$  and  $\ell \geq 3$ . With probability at least  $1 - n^{-\Theta(k+\ell)}$ ,

$$\max_{F \in \mathcal{F}_{k,\ell}} (w(F) - \mathbb{E}[w(F)]) \leq (1 + \epsilon) (2k + \ell) \sqrt{\log n}$$

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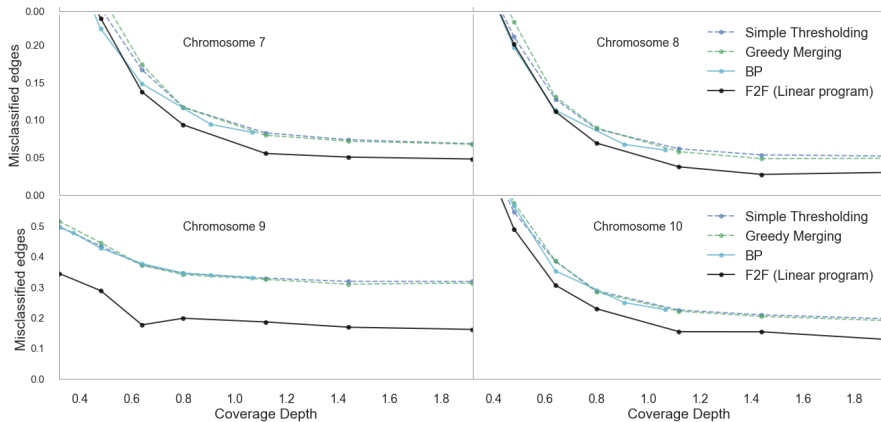
### Remarks

- Total:  $2k + \ell$  edges, half red half blue.
- Weights on red edges  $\sim \mathcal{N}(-\mu, 1)$ ; weights on blue edges  $\sim \mathcal{N}(0, 1)$

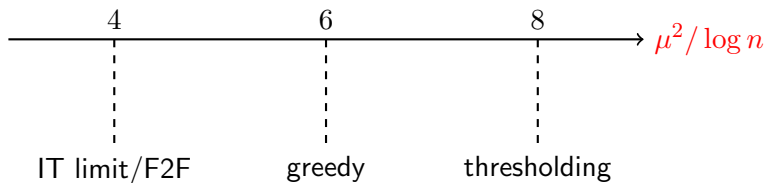
$$w(F) \sim \mathcal{N}\left(-\frac{2k + \ell}{2}\mu, 4k + \ell\right)$$

- 1000 DNA contigs of size 100 kbps
- 0.45 million Chicago cross-links
- Subsample each cross-link with probability  $p$

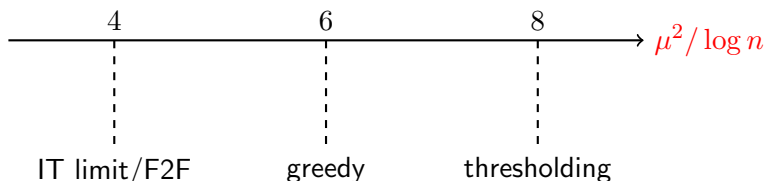
# Homosapiens [Putnam et al 16, Genome Research]



## Conclusion and remarks



# Conclusion and remarks



## References

- Vivek Bagaria, Jian Ding, David Tse, Yihong Wu & X. (2018). *Hidden Hamiltonian Cycle Recovery via Linear Programming*, <https://arxiv.org/abs/1804.05436>