The scaling limit of the minimum spanning tree of the complete graph

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Joint work with Louigi Addario-Berry (McGill), Nicolas Broutin (INRIA Rocquencourt) and Grégory Miermont (ENS Lyon).

Minimum spanning trees

Suppose that G = (V(G), E(G)) is a finite connected graph. To each edge $e \in E(G)$, associate a positive weight w(e).

Problem: find the connected subgraph T of G which has the same vertex set as G and minimises

$$\sum_{e\in E(T)}w(e).$$

Since the weights are positive, the minimum is necessarily attained by a tree. If the weights are all distinct, this tree is unique and is the so-called minimum spanning tree (MST).





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- ► Start from a forest of isolated vertices. List the edges as e₁, e₂,... in increasing order of weight.
- ▶ At step *i*, add edge *e_i* as long as it does not create a cycle.
- Stop when all vertices are connected.

At every stage, we have a forest.















































- ► Start with all edges present. List the edges as e₁, e₂,... in decreasing order of weight.
- ► At step *i*, remove *e_i* as long as it does not disconnect the graph (i.e. as long as it lies in a cycle).
- Stop when no cycles remain.




































The minimum spanning tree of the complete graph

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Question (Aldous (1990))

Does the MST of the complete graph on n vertices possess a scaling limit?



[Picture by Louigi Addario-Berry]

Graphs as metric spaces



We don't really care about the vertex labels or the weights.

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So we will think of the tree as a metric space using the graph distance for the metric. We will also want to be able to pick points uniformly at random from the tree, so we equip it with the uniform measure (call this a measured metric space).

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Again, think of T_n as a measured metric space by using the graph distance and assigning each vertex weight 1/n.

Theorem. (Aldous (1993); Le Gall (2006)) As $n \to \infty$,

$$\frac{1}{\sqrt{n}}T_n \stackrel{d}{\to} \mathcal{T},$$

in the sense of the Gromov-Hausdorff-Prokohorov distance d_{GHP} between compact measured metric spaces, where ${\cal T}$ is the Brownian continuum random tree.

Let $h: [0,1] \to \mathbb{R}^+$ be an excursion, that is a continuous function such that h(0) = h(1) = 0 and h(x) > 0 for $x \in (0,1)$.



















Now put glue on the underside of the excursion and push the two sides together to get a real tree.



The Brownian CRT



[Picture by Grégory Miermont]

The Brownian continuum random tree \mathcal{T} is the random real tree we obtain by doing this gluing procedure to the function 2e, where $(e(x), 0 \le x \le 1)$ a standard Brownian excursion. Properties of the Brownian CRT

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 \mathcal{T} comes naturally equipped with a uniform measure μ , which is the probability measure induced on \mathcal{T} from the Lebesgue measure on [0, 1]. μ is concentrated on the leaves of the tree.

The scaling limit of the UST

Consider T_n , the UST of the complete graph on *n* vertices. Think of T_n as a measured metric space by using the graph distance and assigning each vertex weight 1/n.

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The distortion of R is

$$\mathsf{dis}(R) = \sup\{|d(x,y) - d'(x',y')| : (x,x'), (y,y') \in R\}.$$

 (X, d, μ) and (X', d', μ') are close in the Gromov-Hausdorff-Prokhorov distance if:

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► R has small distortion (i.e. d(x, y) and d'(x', y') are close whenever x ∈ X corresponds to x' ∈ X' and y ∈ X corresponds to y' ∈ X');

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- $\pi(R^c)$ is small.

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Back to the original question: what about the MST?



Does the MST of the complete graph on n vertices possess a scaling limit?

Hold on, isn't it the same?!





a UST on 3000 vertices

an MST on 3000 vertices

[Pictures by Karl-Friedrich Israel]

The scaling limit of the MST

Let M_n be the MST of the complete graph on n vertices, considered as a metric space using the graph distance, and endowed with the uniform measure on its vertices.

Theorem. (Addario-Berry, Broutin, G. & Miermont) There exists a random compact measured metric space \mathcal{M} such that

$$\frac{1}{n^{1/3}}M_n \stackrel{d}{\to} \mathcal{M}$$

as $n \to \infty$, in the sense of d_{GHP}.

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The key to understanding this result is a connection to the Erdős-Rényi random graph.

Reminder: the Erdős-Rényi random graph

Take *n* vertices labelled by $\{1, 2, ..., n\}$ and put an edge between any pair independently with probability *p*. Call the resulting model $\mathbb{G}(n, p)$. (Bond percolation on the complete graph.)

Example: n = 10, p = 0.4 (vertex labels omitted).



The Erdős-Rényi random graph process

Take the obvious coupling of $(\mathbb{G}(n, p), p \in [0, 1])$: simply assign each edge e an independent random edge-weight W(e) with uniform distribution on [0, 1] and keep all edges with weight $W(e) \leq p$.

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It's then natural to think of p as a time and the evolving graph as a process.

Kruskal and Erdős-Rényi

By using the same weights, we can think of (a continuous-time version of) Kruskal's algorithm as "sitting inside" the Erdős-Rényi process: we call this the Kruskal process. When an edge appears in the Erdős-Rényi process, it gets included in the Kruskal process as long as it doesn't create a cycle.

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Indeed, the Kruskal components are the MST's of the Erdős-Rényi components.

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So the action is happening a bit earlier, in the critical window for the Erdős-Rényi phase transition.

The critical random graph

The critical window: $p = \frac{1}{n} + \frac{\lambda}{n^{4/3}}$, where $\lambda \in \mathbb{R}$. For such p, the largest components have size $\Theta(n^{2/3})$.

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Define the surplus of a component to be the number of edges more than a tree that it has.

A component with surplus 3:



Convergence of the sizes and surpluses

Fix $\lambda \in \mathbb{R}$. Let $C_{\lambda}^{n,1}, C_{\lambda}^{n,2}, \ldots$ be the sequence of component sizes of $\mathbb{G}(n, n^{-1} + \lambda n^{-4/3})$ in decreasing order, and let $S_{\lambda}^{n,1}, S_{\lambda}^{n,2}, \ldots$ be their surpluses.

Write $\mathbf{C}_{\lambda}^{n} = (C_{\lambda}^{n,1}, C_{\lambda}^{n,1}, \ldots)$ and $\mathbf{S}_{\lambda}^{n} = (S_{\lambda}^{n,1}, S_{\lambda}^{n,2}, \ldots)$.

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Theorem (Aldous (1997)). For fixed $\lambda \in \mathbb{R}$, as $n \to \infty$,

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(Convergence for the first co-ordinate takes place in

$$\ell^2_{\searrow} := \left\{ \mathbf{x} = (x_1, x_2, \ldots) : x_1 \ge x_2 \ge \ldots \ge 0, \sum_{i=1}^{\infty} x_i^2 < \infty \right\}$$

and for the second in the sense of finite-dimensional distributions.)

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Let $B_{\lambda}(t) = W_{\lambda}(t) - \min_{0 \le s \le t} W_{\lambda}(s)$ be the process reflected at its minimum.



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Decorate the picture with the points of a rate one Poisson process which fall above the *x*-axis and below the graph.

 \mathbf{C}_{λ} is the sequence of excursion-lengths of this process, in decreasing order.

 \mathbf{S}_{λ} is the sequence of numbers of points falling in the corresponding excursions.

The critical Erdős-Rényi random graph

Let $\mathbf{G}_{\lambda}^{n} = (G_{\lambda}^{n,1}, G_{\lambda}^{n,2}, \ldots)$ be the sequence of components of $\mathbb{G}(n, n^{-1} + \lambda n^{-4/3})$ in decreasing order of size, each considered as a metric space with the graph distance. Put mass $n^{-2/3}$ on each vertex.
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where $\mathscr{G}_{\lambda} = (\mathscr{G}_{\lambda}^{1}, \mathscr{G}_{\lambda}^{2}, ...)$ is a sequence of measured real trees with vertex-identifications encoded by Aldous' marked limit process in decreasing order of length.

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Convergence is with respect to the distance

$$\mathsf{dist}^4_{\mathsf{GHP}}(\mathcal{A},\mathcal{B}) := \left(\sum_{i=1}^\infty \mathsf{d}_{\mathsf{GHP}}(\mathcal{A}_i,\mathcal{B}_i)^4\right)^{1/4}$$

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- The kth-longest excursion has length C^k_λ and so the corresponding tree is endowed with a uniform measure of that total mass.
- ► In the limit, surplus edges correspond to vertex-identifications.
- In each excursion, the points of the Poisson process tell us where these vertex-identifications should occur.

Spanning tree and vertex identifications



To get from the state of the Erdős-Rényi process at a fixed time $p \in [0, 1]$ to the state of the Kruskal process, we need to break the cycles at high-weight edges.

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For fixed λ and k, we can construct $T_{\lambda}^{n,k}$ from $G_{\lambda}^{n,k}$ by running the cycle-breaking algorithm on the latter.

Reminder: cycle-breaking

- ► Start with all edges present. List them as e₁, e₂,... in decreasing order of weight.
- At step *i*, remove *e_i* as long as it does not disconnect the graph (i.e. as long as it lies in a cycle).
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This tells us what the limiting analogue of the cycle-breaking procedure should be: repeatedly remove points chosen according to the uniform (Lebesgue) measure on the cycles in the metric space, until no cycles remain.

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(It's actually a little delicate to check that the cycle-breaking procedure passes nicely to the limit ...)

A limit for the Kruskal process

Let $\mathbf{T}_{\lambda}^{n} = (\mathcal{T}_{\lambda}^{n,1}, \mathcal{T}_{\lambda}^{n,2}, \ldots)$ be the components of the Kruskal process, each endowed with the measure which assigns mass $n^{-2/3}$ to each vertex, and let $\mathscr{T}_{\lambda} = (\mathscr{T}_{\lambda}^{1}, \mathscr{T}_{\lambda}^{2}, \ldots)$ be the sequence of spaces obtained from \mathscr{G}_{λ} by cycle-breaking.

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where convergence is with respect to the distance

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- As λ increases, we glue more and more little trees onto *T*¹_λ.
 However, its diameter remains bounded. Moreover, the additional mass coming from the little trees gets spread out essentially uniformly over the tree.
- The mass of 𝔅¹_λ diverges as λ → ∞, so we renormalise to get total mass measure 1 for each λ. If we do this, we obtain that there exists a limiting measured metric space 𝓜 as λ → ∞.

The scaling limit of the MST



We obtain

$$\frac{1}{n^{1/3}}M^n \stackrel{d}{\to} \mathscr{M}$$

as $n \to \infty$, in the sense of d_{GHP}.

References

The continuum limit of critical random graphs

L. Addario-Berry, N. Broutin and C. Goldschmidt *Probability Theory and Related Fields* 152, 3-4 (2012), pp.367-406.

Critical random graphs: limiting constructions and distributional properties

L. Addario-Berry, N. Broutin and C. Goldschmidt *Electronic Journal of Probability* 15 (2010), pp.741-775.

The scaling limit of the minimum spanning tree of the complete graph

L. Addario-Berry, N. Broutin, C. Goldschmidt and G. Miermont arXiv:1301.1664 (2013+).