

ON THE NUMBER OF SUBTREES ON THE FRINGE OF RANDOM TREES

(partly joint with Huilan Chang)

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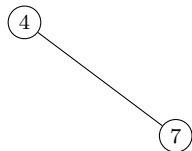
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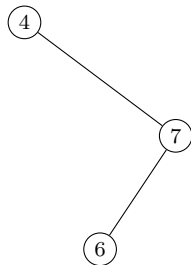
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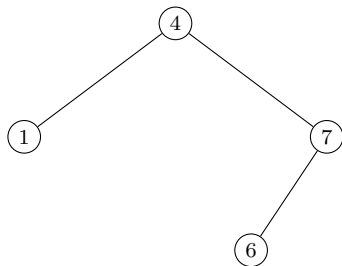
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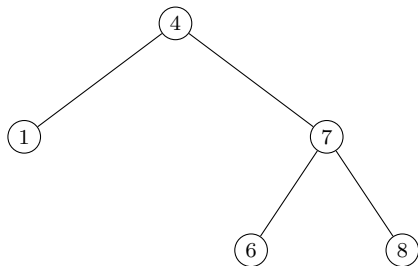
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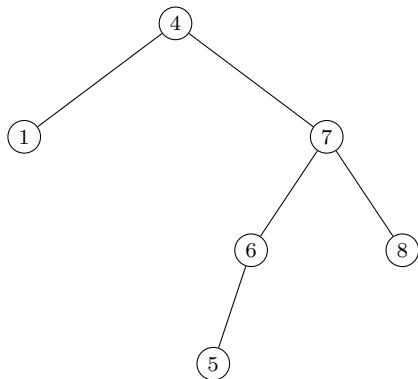
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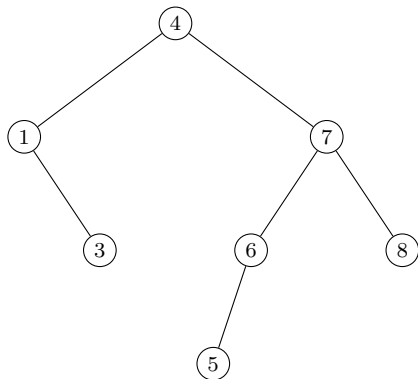
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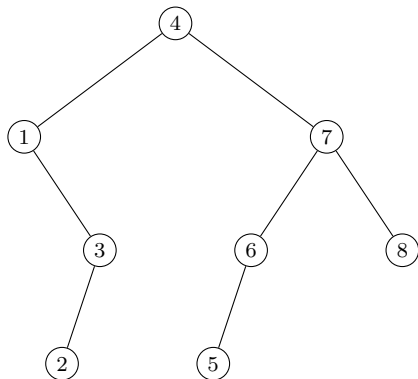
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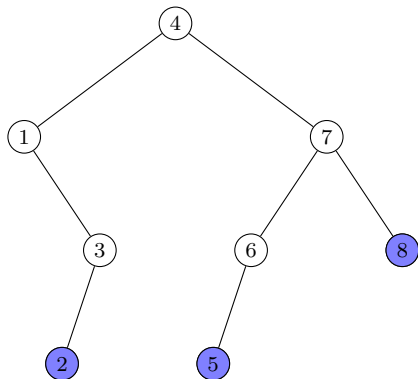
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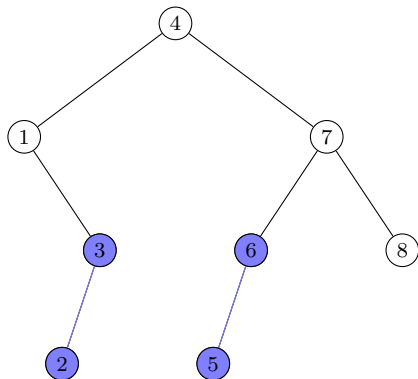


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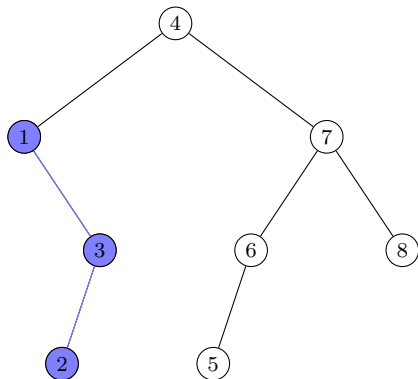
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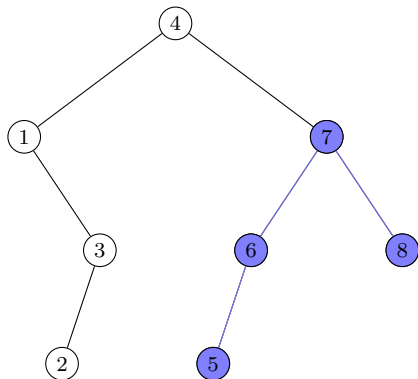
$$X_{8,2} = 2$$

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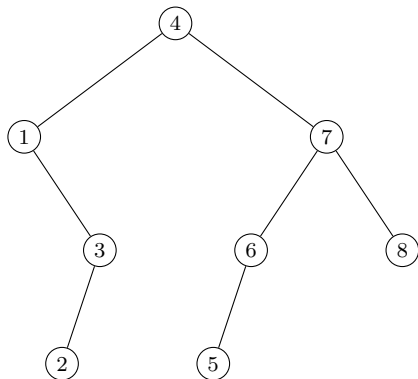
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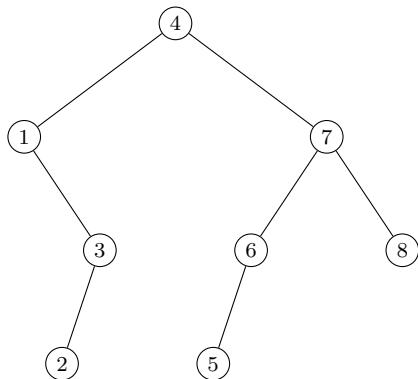
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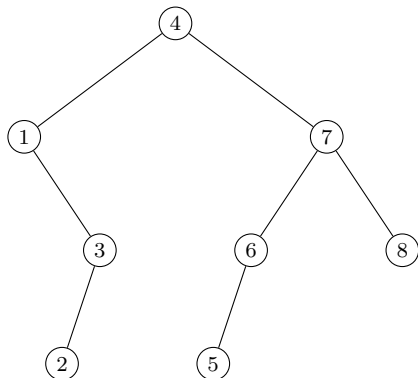
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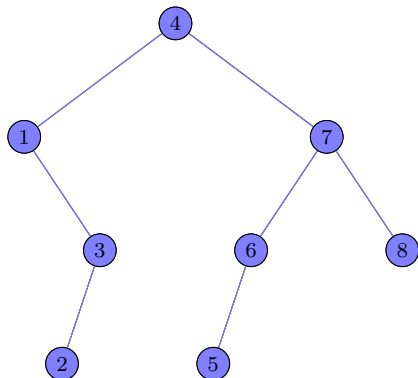
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$$X_{8,8} = 1$$

Mean value and variance

$X_{n,k}$ satisfies

$$X_{n,k} \stackrel{d}{=} X_{I_n,k} + X_{n-1-I_n,k}^*,$$

where $X_{k,k} = 1$, $X_{I_n,k}$ and $X_{n-1-I_n,k}^*$ are conditionally independent given I_n , and $I_n = \text{Unif}\{0, \dots, n-1\}$.

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This yields

$$\mu_{n,k} := \mathbf{E}(X_{n,k}) = \frac{2(n+1)}{(k+1)(k+2)}, \quad (n > k),$$

and

$$\sigma_{n,k}^2 := \text{Var}(X_{n,k}) = \frac{2k(4k^2 + 5k - 3)(n+1)}{(k+1)(k+2)^2(2k+1)(2k+3)}$$

for $n > 2k + 1$.

Some previous results

- Aldous (1991): Weak law of large numbers

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Central limit theorem with optimal Berry-Esseen bound and LLT

→ All the above results are for fixed k .

Results for $k = k_n$

Theorem (Feng, Mahmoud, Panholzer (2008))

(i) (Normal range) Let $k = o(\sqrt{n})$ and $k \rightarrow \infty$ as $n \rightarrow \infty$. Then,

$$\frac{X_{n,k} - \mu_{n,k}}{\sqrt{2n/k^2}} \xrightarrow{d} \mathcal{N}(0, 1).$$

(ii) (Poisson range) Let $k \sim c\sqrt{n}$ as $n \rightarrow \infty$. Then,

$$X_{n,k} \xrightarrow{d} \text{Poisson}(2c^{-2}).$$

(iii) (Degenerate range) Let $k < n$ and $\sqrt{n} = o(k)$ as $n \rightarrow \infty$. Then,

$$X_{n,k} \xrightarrow{L_1} 0.$$

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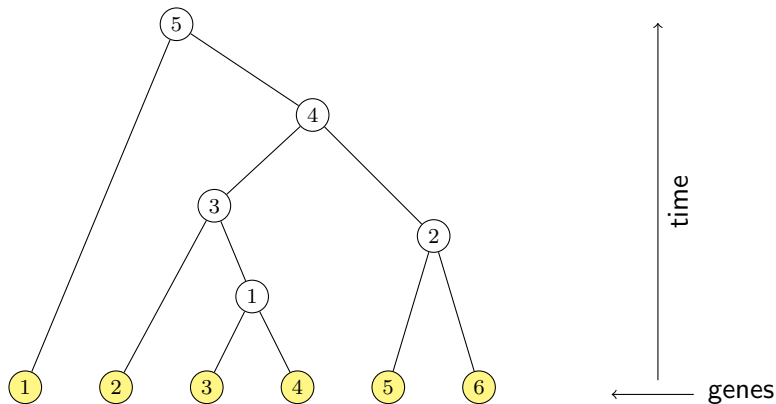
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- $X_{n,k}$ is related to parameters arising in genetics.

Yule generated random genealogical trees

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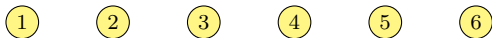


Yule generated random genealogical trees

Example:

Random model:

At every time point,
two yellow nodes
uniformly coalescent.

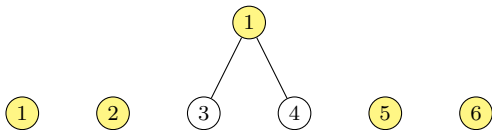


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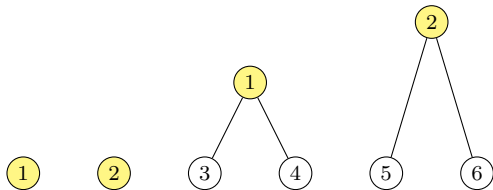
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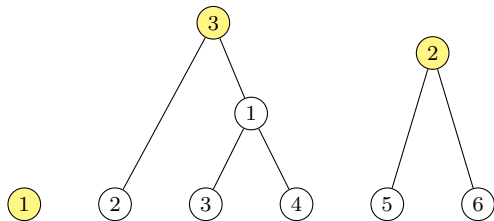


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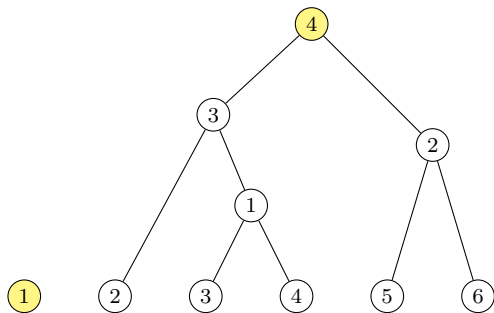


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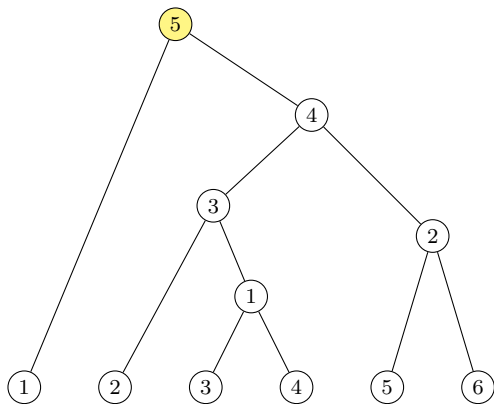


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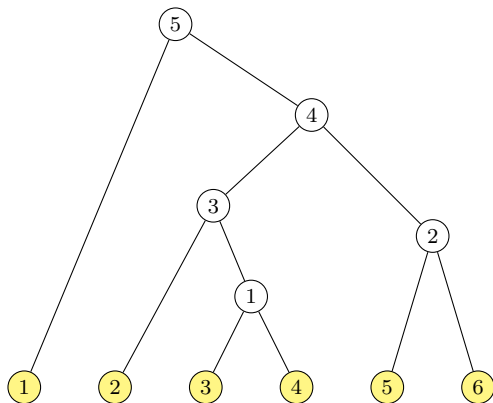


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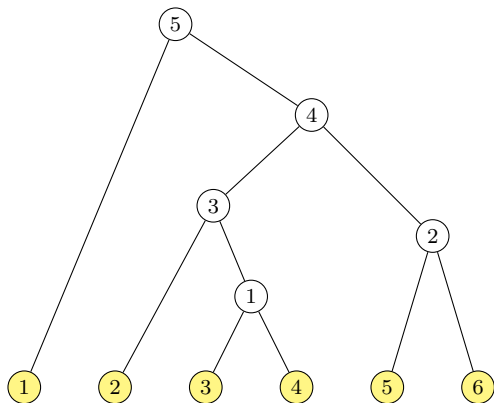


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Same model as
random binary
search tree model!

Shape parameters of genealogical trees

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- Nodes with minimal clade size k (Blum and François (2005)):

If $k \geq 3$, then they are internal nodes with induced subtree of size $k - 1$ and either an empty right subtree or empty left subtree.

Counting pattern in random binary search trees

Consider $X_{n,k}$ with

$$X_{n,k} \stackrel{d}{=} X_{I_n,k} + X_{n-1-I_n,k}^*,$$

where $X_{k,k} = \text{Bernoulli}(p_k)$, $X_{I_n,k}$ and $X_{n-1-I_n,k}^*$ are conditionally independent given I_n , and $I_n = \text{Unif}\{0, \dots, n-1\}$.

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Then,

| p_k | shape parameter |
|--------------|--|
| 1 | # of $k+1$ -pronged nodes |
| $2/k$ | # of nodes with minimal clade size $k+1$ |
| $2^{k-1}/k!$ | # of $k+1$ caterpillars |

Underlying recurrence and solution

All (centered or non-centered) moments satisfy

$$a_{n,k} = \frac{2}{n} \sum_{j=0}^{n-1} a_{j,k} + b_{n,k},$$

where $a_{k,k}$ is given and $a_{n,k} = 0$ for $n < k$.

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We have

$$a_{n,k} = \frac{2(n+1)}{(k+1)(k+2)} a_{k,k} + 2(n+1) \sum_{k < j < n} \frac{b_{j,k}}{(j+1)(j+2)} + b_{n,k},$$

where $n > k$.

Mean value and variance

We have

$$\mathbf{E}(X_{n,k}) = \frac{2(n+1)}{(k+1)(k+2)}p_k, \quad (n > k),$$

and

$$\text{Var}(X_{n,k}) = \frac{2p_k(4k^3 + 16k^2 + 19k + 6 - (11k^2 + 22k + 6)p_k)(n+1)}{(k+1)(k+2)^2(2k+1)(2k+3)}$$

for $n > 2k + 1$.

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Note that

$$\mathbf{E}(X_{n,k}) \sim \text{Var}(X_{n,k}) \sim \frac{2p_k}{k^2}n$$

for $n > 2k + 1$ and $k \rightarrow \infty$ as $n \rightarrow \infty$.

Higher moments

Denote by

$$A_{n,k}^{(m)} := \mathbf{E}(X_{n,k} - \mathbf{E}(X_{n,k}))^m.$$

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Then,

$$A_{n,k}^{(m)} = \frac{2}{n} \sum_{j=0}^{n-1} A_{j,k}^{(m)} + B_{n,k}^{(m)},$$

where

$$B_{n,k}^{(m)} := \sum_{\substack{i_1+i_2+i_3=m \\ 0 \leq i_1, i_2 < m}} \binom{m}{i_1, i_2, i_3} \frac{1}{n} \sum_{j=0}^{n-1} A_{j,k}^{(i_1)} A_{n-1-j,k}^{(i_2)} \Delta_{n,j,k}^{i_3}$$

and

$$\Delta_{n,j,k} = \mathbf{E}(X_{j,k}) + \mathbf{E}(X_{n-1-j,k}) - \mathbf{E}(X_{n,k}).$$

Normal range

Proposition

Uniformly for $n, k, m \geq 1$ and $n > k$

$$A_{n,k}^{(m)} = \mathcal{O} \left(\max \left\{ \frac{2p_k n}{k^2}, \left(\frac{2p_k n}{k^2} \right)^{m/2} \right\} \right).$$

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Proposition

For $\mathbf{E}(X_{n,k}) \rightarrow \infty$ as $n \rightarrow \infty$,

$$A_{n,k}^{(2m-1)} = o \left(\left(\frac{2p_k n}{k^2} \right)^{m-1/2} \right), \quad A_{n,k}^{(2m)} \sim g_m \left(\frac{2p_k n}{k^2} \right)^m,$$

where

$$g_m = (2m)! / (2^m m!).$$

Poisson range

Consider

$$\bar{A}_{n,k}^{(m)} = \mathbf{E}(X_{n,k}(X_{n,k} - 1) \cdots (X_{n,k} - m + 1)).$$

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Then, similarly as before:

Proposition

(i) *Uniformly for $n, k, m \geq 1$ and $n > k$*

$$\bar{A}_{n,k}^{(m)} = \mathcal{O} \left(\max \left\{ \frac{2p_k n}{k^2}, \left(\frac{2p_k n}{k^2} \right)^m \right\} \right).$$

(ii) *For $\mathbf{E}(X_{n,k}) \rightarrow c$ and $k < n$ as $n \rightarrow \infty$,*

$$\bar{A}_{n,k}^{(m)} \longrightarrow c^m.$$

The phase change

Theorem

(i) (Normal range) Let $\mathbf{E}(X_{n,k}) \rightarrow \infty$ and $k \rightarrow \infty$ as $n \rightarrow \infty$. Then,

$$\frac{X_{n,k} - \mathbf{E}(X_{n,k})}{\sqrt{2p_k n/k^2}} \xrightarrow{d} \mathcal{N}(0, 1).$$

(ii) (Poisson range) Let $\mathbf{E}(X_{n,k}) \rightarrow c > 0$ and $k < n$ as $n \rightarrow \infty$. Then,

$$X_{n,k} \xrightarrow{d} \text{Poisson}(c).$$

(iii) (Degenerate range) Let $\mathbf{E}(X_{n,k}) \rightarrow 0$ as $n \rightarrow \infty$. Then,

$$X_{n,k} \xrightarrow{L_1} 0.$$

A comparison of the phase change

For k -caterpillars, we have

$$\mathbf{E}(X_{n,k}) = \frac{2^{k-1}n}{(k+2)!}.$$

Note that either

$$\mathbf{E}(X_{n,k}) \rightarrow \infty \quad \text{or} \quad \mathbf{E}(X_{n,k}) \rightarrow 0.$$

So, there is no Poisson range.

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| shape parameter | location | phase change |
|------------------------|-----------------------|-------------------------------|
| k -pronged nodes | \sqrt{n} | normal - poisson - degenerate |
| minimal clade size k | $\sqrt[3]{n}$ | normal - poisson - degenerate |
| k -caterpillars | $\ln n / (\ln \ln n)$ | normal - degenerate |

Refined results (for # of subtrees)

Define

$$\phi_{n,k}(y) = e^{-\sigma_{n,k}^2 y^2 / 2} \mathbf{E} \left(e^{(X_{n,k} - \mu_{n,k})y} \right).$$

and

$$\phi_{n,k}^{(m)} = \left. \frac{d^m \phi_{n,k}(y)}{dy^m} \right|_{y=0}.$$

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Proposition

Uniformly for $n, k \geq 1$ and $m \geq 0$

$$|\phi_{n,k}^{(m)}| \leq m! A^m \max \left\{ \frac{n}{k^2}, \left(\frac{n}{k^2} \right)^{m/3} \right\}$$

for a suitable constant A .

Berry-Esseen bound and LLT for the normal range

Theorem (Rate of convergency)

For $1 \leq k = o(\sqrt{n})$ as $n \rightarrow \infty$,

$$\sup_{x \in \mathbb{R}} \left| P \left(\frac{X_{n,k} - \mu_{n,k}}{\sigma_{n,k}} < x \right) - \Phi(x) \right| = \mathcal{O} \left(\frac{k}{\sqrt{n}} \right).$$

Berry-Esseen bound and LLT for the normal range

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Theorem (LLT)

For $1 \leq k = o(\sqrt{n})$ as $n \rightarrow \infty$,

$$P(X_{n,k} = \lfloor \mu_{n,k} + x\sigma_{n,k} \rfloor) = \frac{e^{-x^2/2}}{\sqrt{2\pi}\sigma_{n,k}} \left(1 + \mathcal{O}\left((1 + |x|^3)\frac{k}{\sqrt{n}}\right)\right),$$

uniformly in $x = o(n^{1/6}/k^{1/3})$.

LLT for the Poisson range

Define

$$\bar{\phi}_{n,k}(y) = e^{-\mu_{n,k}(y-1)} \mathbf{E} \left(y^{X_{n,k}} \right).$$

and

$$\phi_{n,k}^{(m)} = \left. \frac{d^m \bar{\phi}_{n,k}(y)}{dy^m} \right|_{y=1}.$$

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Proposition

Uniformly for $n > k$ and $m \geq 0$

$$|\bar{\phi}_{n,k}^{(m)}| \leq m! A^m \left(\frac{n}{k^3} \right)^{m/2}$$

for a suitable constant A .

Poisson approximation

Theorem (LLT)

For $k < n$ and $n \rightarrow \infty$,

$$P(X_{n,k} = l) = e^{-\mu_{n,k}} \frac{(\mu_{n,k})^l}{l!} + \mathcal{O}\left(\frac{n}{k^3}\right)$$

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Theorem (Poisson approximation)

Let $k < n$ and $k \rightarrow \infty$ as $n \rightarrow \infty$. Then,

$$d_{TV}(X_{n,k}, \text{Poisson}(\mu_{n,k})) \rightarrow 0.$$

Remark: A rate can be given as well.

Other types of random trees

- Random recursive trees

Non-plane, labelled trees with every label sequence from the root to a leaf increasing; random model is the uniform model.

Methods works as well (with minor modifications) and similar results can be proved.

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- Plane-oriented recursive trees (PORTs)

Plane, labelled trees with every label sequence from the root to a leaf increasing; random model is the uniform model.

Method works as well, but details more involved.

Mean value and variance of PORTs

We have,

$$\mu_{n,k} := \mathbf{E}(X_{n,k}) = \frac{2n-1}{4k^2-1}, \quad (n > k).$$

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$$\mu_{n,k} := \mathbf{E}(X_{n,k}) = \frac{2n-1}{4k^2-1}, \quad (n > k).$$

Moreover, for fixed k as $n \rightarrow \infty$,

$$\text{Var}(X_{n,k}) \sim c_k n,$$

where

$$c_k = \frac{8k^2 - 4k - 8}{(4k^2 - 1)^2} - \frac{((2k-3)!!)^2}{((k-1)!)^2 4^{k-1} k(2k+1)},$$

and, for $k < n$ and $k \rightarrow \infty$ as $n \rightarrow \infty$,

$$\mathbf{E}(X_{n,k}) \sim \text{Var}(X_{n,k}) \sim \frac{n}{2k^2}.$$

The phase change

Theorem

(i) (Normal range) Let $k = o(\sqrt{n})$ and $k \rightarrow \infty$ as $n \rightarrow \infty$. Then,

$$\frac{X_{n,k} - \mu_{n,k}}{\sqrt{n/(2k^2)}} \xrightarrow{d} \mathcal{N}(0, 1).$$

(ii) (Poisson range) Let $k \sim c\sqrt{n}$ as $n \rightarrow \infty$. Then,

$$X_{n,k} \xrightarrow{d} \text{Poisson}((2c^2)^{-1}).$$

(iii) (Degenerate range) Let $k < n$ and $\sqrt{n} = o(k)$ as $n \rightarrow \infty$. Then,

$$X_{n,k} \xrightarrow{L_1} 0.$$

More results and future research

- Parameters of genealogical trees under different random models

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Very simple classes of increasing trees and more general classes of increasing trees (polynomial varieties, mobile trees, etc.)

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Very simple classes of increasing trees and more general classes of increasing trees (polynomial varieties, mobile trees, etc.)

- Phase change results for the number of nodes with out-degree k

Important in computer science.

A phase change from normal to degenerate is expected (no Poisson range).