

The total external length in the evolving Kingman coalescent

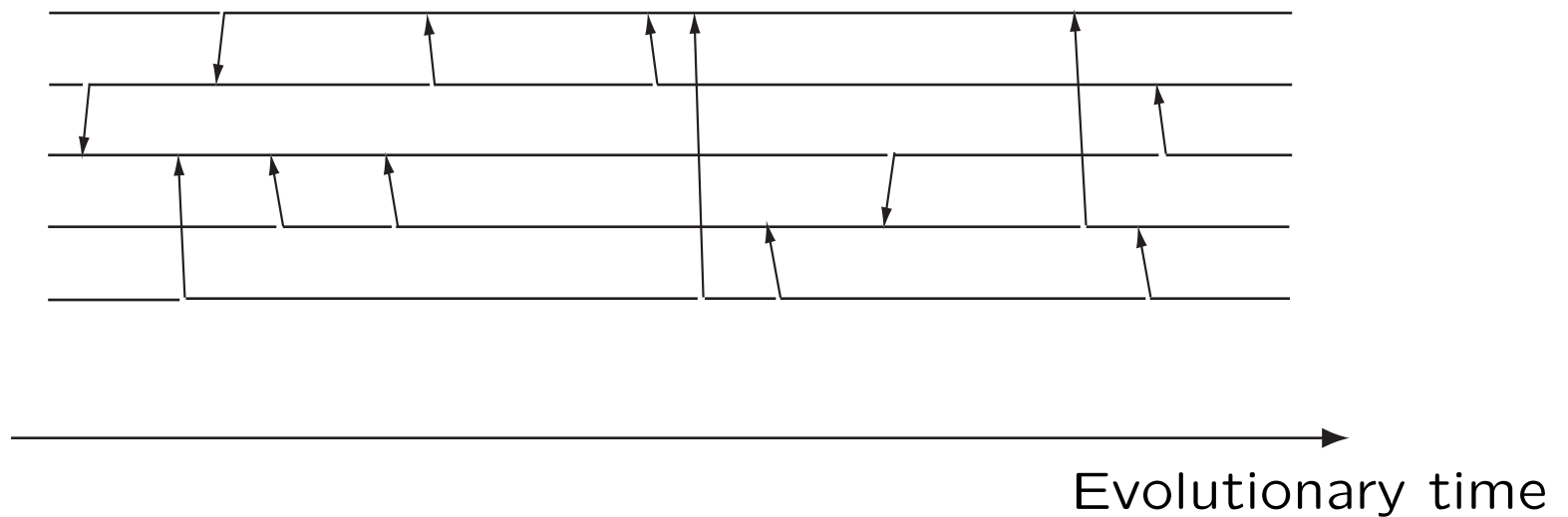
Götz Kersting and Iulia Stanciu

Goethe Universität, Frankfurt am Main

CIRM, Luminy
June 11–15 2012



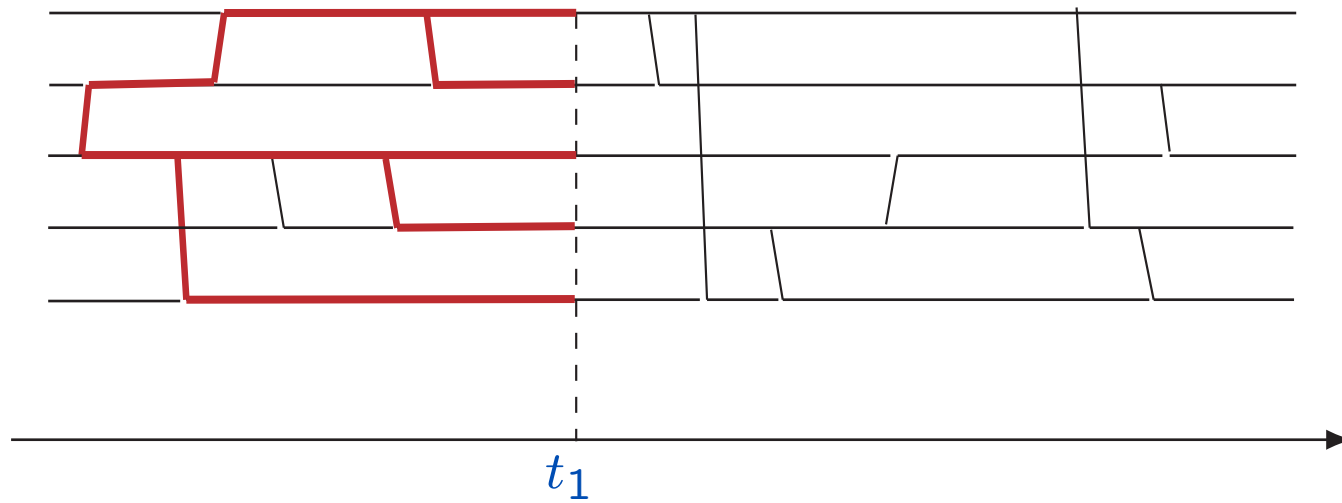
The evolving Kingman N -coalescent ($N = 5$):



Moran's model with time $-\infty < t < \infty$:

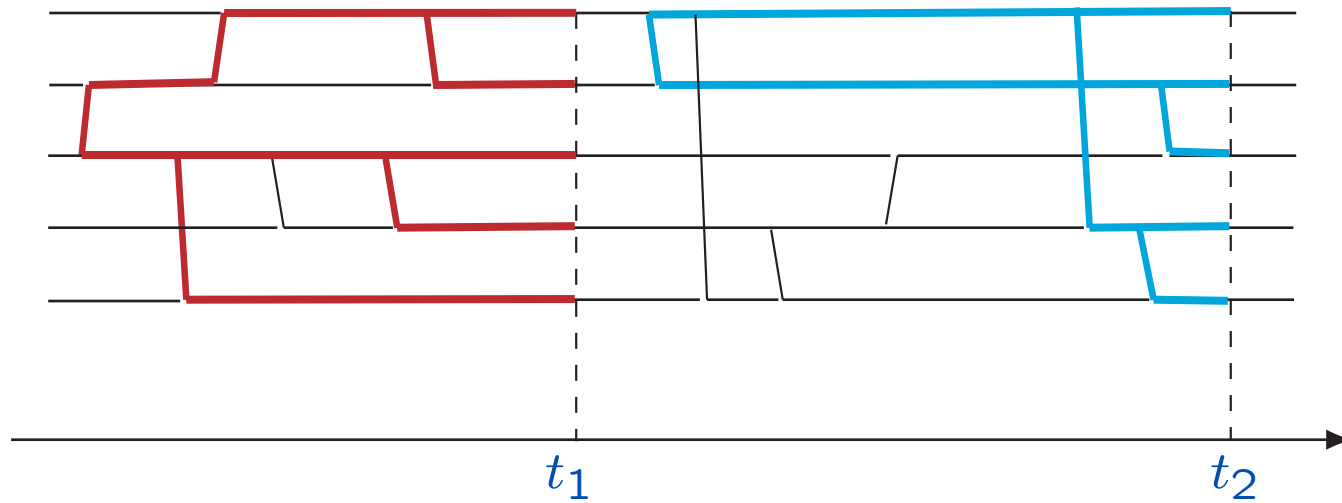
Links between pairs of lines appear at rate 1, independent between the different pairs.

The evolving Kingman N -coalescent:



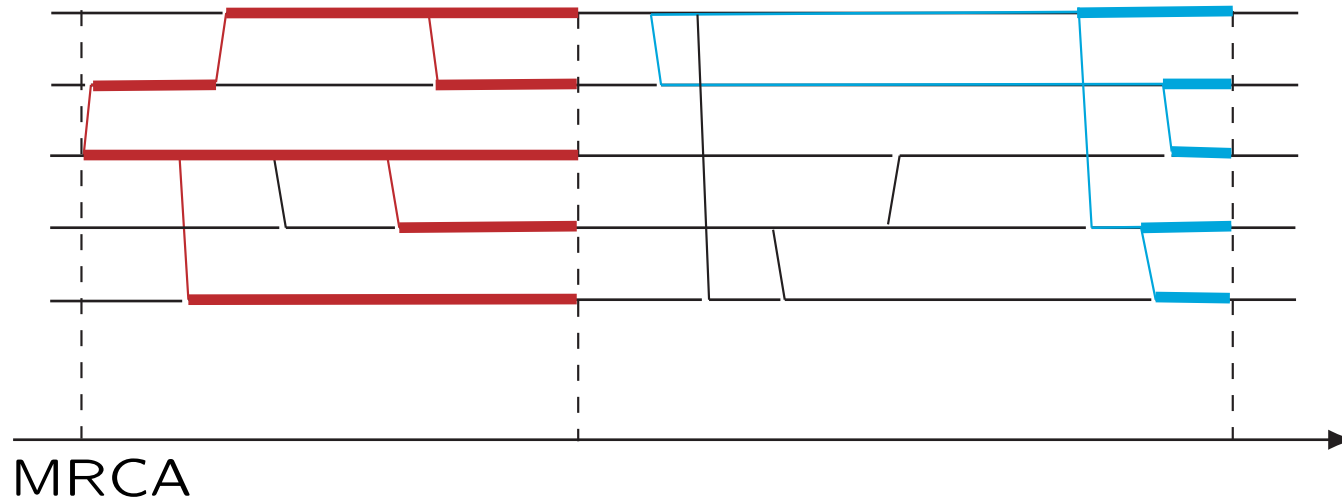
Kingman's coalescent at time t_1

The evolving Kingman N -coalescent:



The coalescent tree evolves in time.

The evolving Kingman N -coalescent:



Evolving time to MRCA
tree topology
total length
total external length

These results are rather different in nature, none covering any other.

Interesting aspects:

- limiting processes with
a.s. continuous paths versus
a.s. (compensated) pure jump paths
- different time scalings

A BASKET OF RESULTS

Theorem: (Pfaffelhuber/Wakolbinger , Donnelly/Kurtz, 2006)

Let $A_N(t)$ be the time to the MRCA of the evolving Kingman N -coalescent at time $t \in \mathbb{R}$. Then, as $N \rightarrow \infty$,

$$\left(A_N(t) \right)_{t \in \mathbb{R}} \xrightarrow{d} A ,$$

where the limiting process $A = (A_t)_{t \in \mathbb{R}}$ is stationary, a.s. pure jump, non-Markovian.

For a related result on the two oldest families in the genealogy see Delmas, Dhersin, and Siri-Jegousse (2010).

Theorem: (Greven, Pfaffelhuber, Winter, 2009, 2010)

Let $T_N(t)$ be the tree, induced by the evolving Kingman N -coalescent at time $t \in \mathbb{R}$ in the space of real trees furnished with the Gromov-weak topology. Then, as $N \rightarrow \infty$,

$$\left(T_N(t)\right)_{t \in \mathbb{R}} \xrightarrow{d} T ,$$

where the limiting tree-valued process $T = (T_t)_{t \in \mathbb{R}}$ is stationary, a.s. continuous, and unique solution of a martingale problem.

Theorem: (Pfaffelhuber, Wakolbinger, Weisshaupt, 2011)

Let $L'_N(t)$ be the total length of the evolving Kingman N -coalescent at time $t \in \mathbb{R}$. Then, as $N \rightarrow \infty$,

$$\left(L'_N(t) - 2 \log N \right)_{t \in \mathbb{R}} \xrightarrow{d} L' ,$$

where the limiting process $L' = (L'_t)_{t \in \mathbb{R}}$ is stationary, a.s. pure jump, non-Markovian.

L' has *infinite quadratic variation* (Knobloch, Stanciu, Wakolbinger, 2011), thus fails to be a semimartingale.

Theorem: (Schweinsberg, 2011)

Let $L''_N(t)$ be the total length of the evolving *Bolthausen-Sznitman* N -coalescent at time $t \in \mathbb{R}$. Then, as $N \rightarrow \infty$,

$$\left(\frac{(\log N)^2}{N} L''_N\left(\frac{t}{\log N}\right) - \log N - \log \log N \right)_{t \in \mathbb{R}} \xrightarrow{d} L'' ,$$

where the stationary limiting process $L'' = (L''_t)_{t \in \mathbb{R}}$ solves the SDE

$$dL'' = -L'' dt + dY$$

for a certain Lévy-process Y of index 1.

Theorem: (K., Stanciu, 2012, ongoing work)

Let $L_N(t)$ be the total *external length* of the evolving Kingman N -coalescent. Then

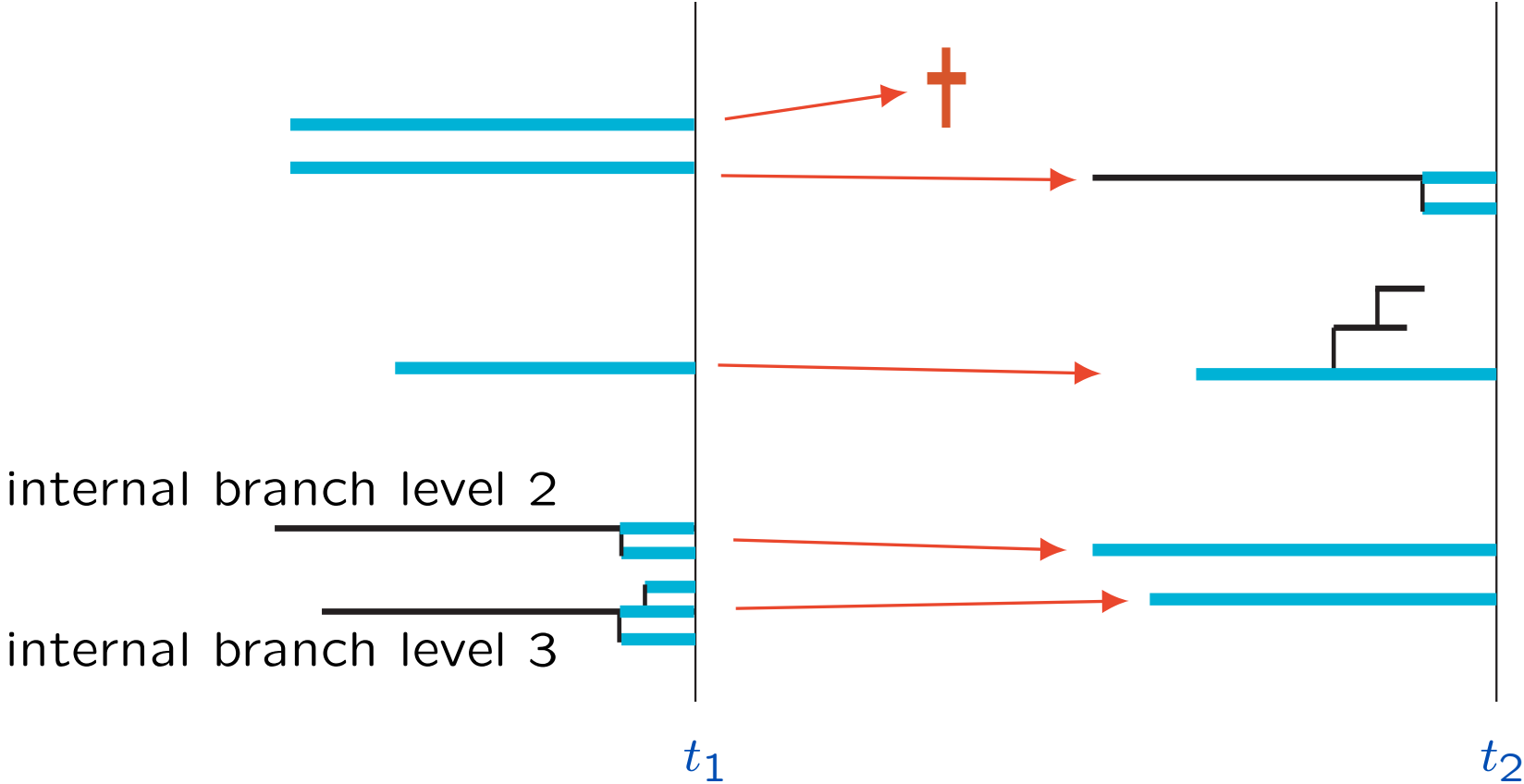
$$\left(\sqrt{\frac{N}{4 \log N}} \left(L_N\left(\frac{t}{N}\right) - 2 \right) \right)_{t \in \mathbb{R}} \xrightarrow{d} L ,$$

where L is a stationary, Gaussian, a.s. continuous, with covariance function

$$\mathbf{Cov}(L_s, L_t) = \left(\frac{1}{1 + |t - s|} \right)^2 .$$

Note the different scaling of time (*real* instead of *evolutionary* periods).

The dynamics of the external lengths:



So let for the *static* Kingman N -coalescent (at time $t = 0$)

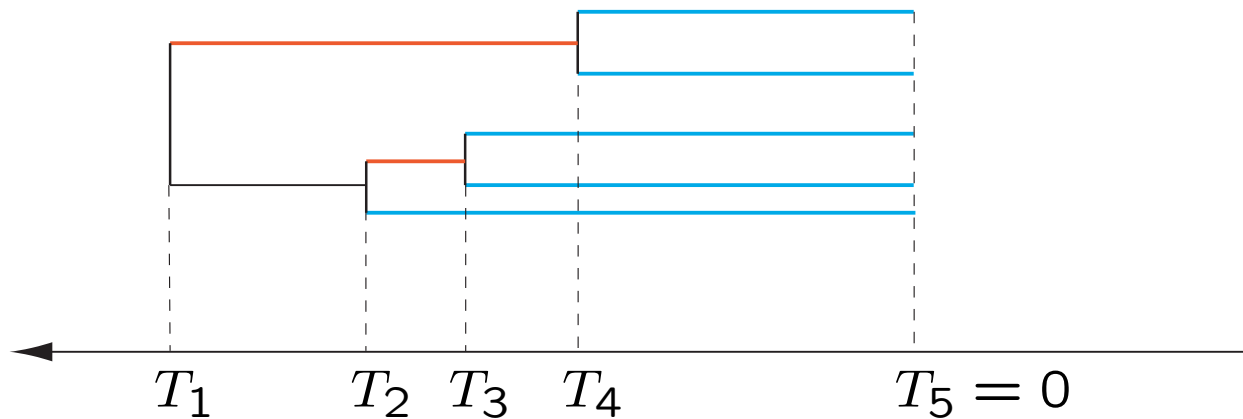
$L_N^i =$ total internal branch length of level i ,

in particular for $i = 1$

$L_N^1 =$ total external branch length .

How do we gain access to these quantities?

Branch numbers V_N, \dots, V_2 and W_N, \dots, W_2 .



$$V_2 = 0$$

$$W_2 = 1$$

$$V_5 = 5$$

$$W_5 = 0$$

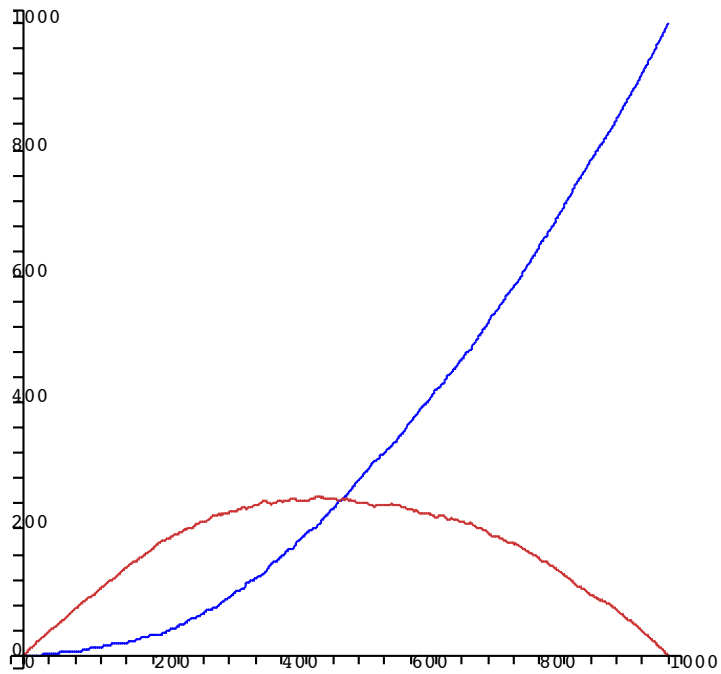
$$L_N^1 = \sum_{i=2}^N V_i (T_{i-1} - T_i) , \quad L_N^2 = \sum_{i=2}^N W_i (T_{i-1} - T_i)$$

$$\begin{aligned} L_N^1 &= \sum_{i=2}^N V_i(T_{i-1} - T_i) \\ &= \sum_{k=1}^{N-1} T_k(V_{k+1} - V_k) + T_1 V_2 \\ &\approx \sum_{k=1}^{N-1} \frac{2}{k} \Delta V_k \end{aligned}$$

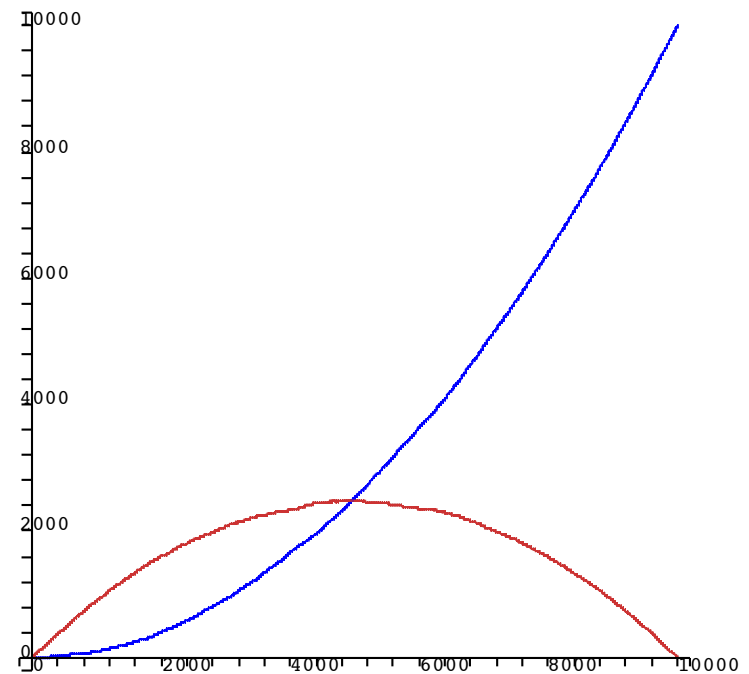
ΔV_k is easy to handle for k close to N .

External branch numbers $0 = V_1, V_2, \dots, V_N$, and
(total) internal branch numbers

$$U_1 = 1 - V_1, U_2 = 2 - V_2, \dots, U_N = 1 - V_N$$



$N = 1000$



$N = 10000$

Let us look at the randomness within

$$V_N - N \quad (= 0)$$

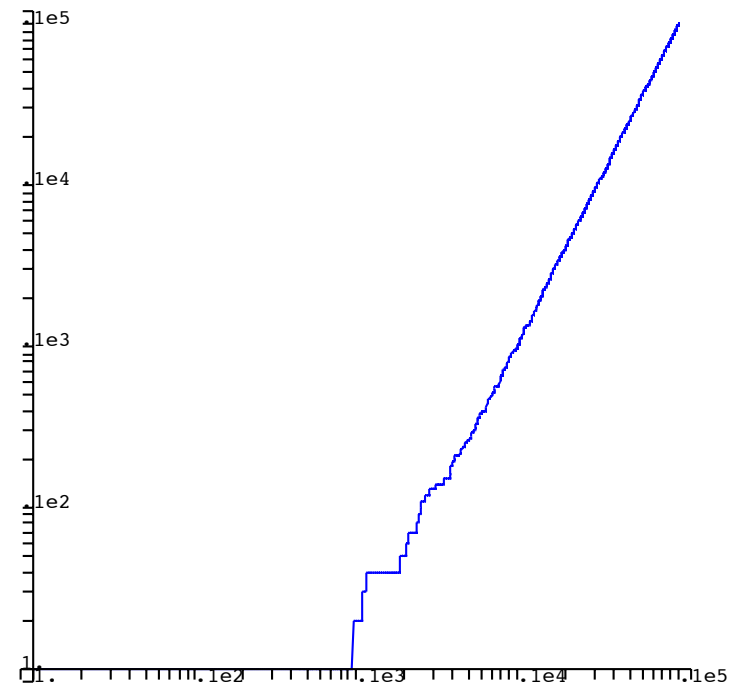
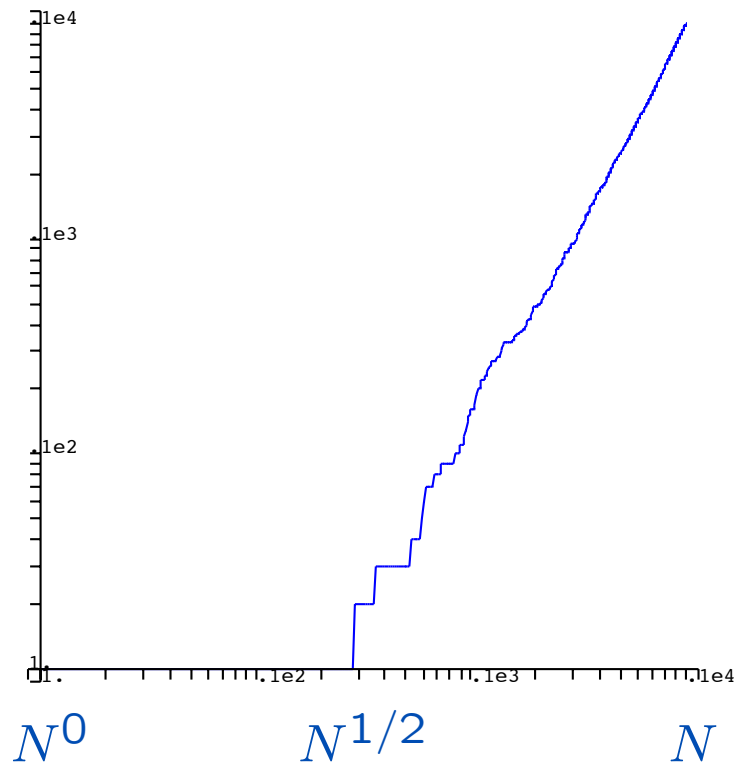
$$V_{N-1} - N + 2$$

⋮

$$V_{N-i} - N + 2i$$

⋮

Randomness within (V_i) close to N :
 $V_N - N + 1, V_{N-1} - N + 3, \dots$



Randomness enters (almost) independently.

However, since we consider

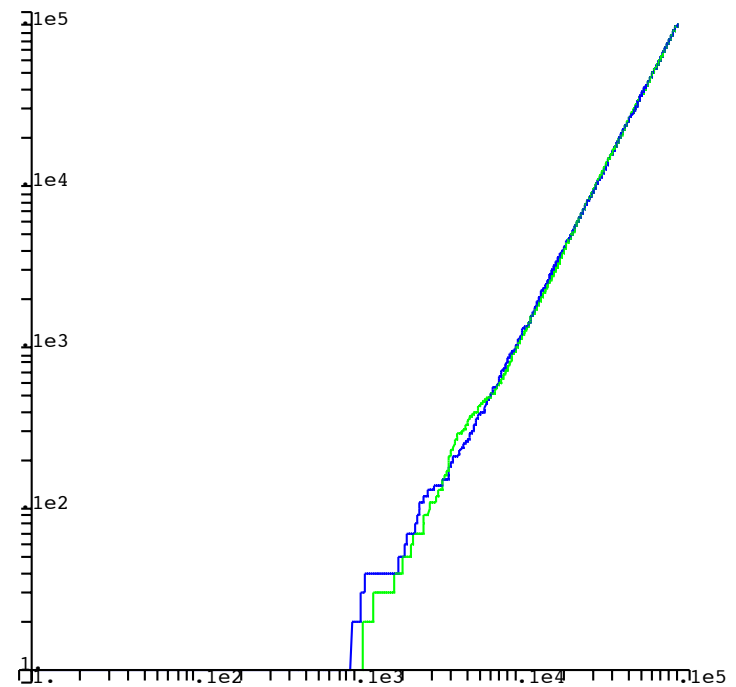
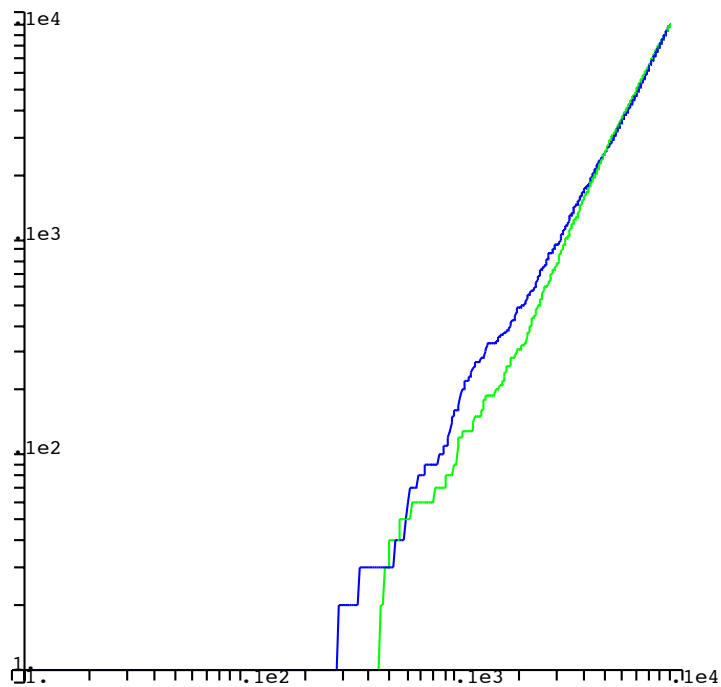
$$L_N^1 \approx \sum_{k=1}^{N-1} \frac{2}{k} \Delta V_k$$

we have to understand randomness at the beginning of V_2, \dots, V_N .
So let us compare

$$V_N - N, V_{N-1} - N + 2, \dots \quad \text{to} \quad V_1, V_2, \dots$$

Randomness within (V_i) close to N and 1 :

$V_N - N + 1, V_{N-1} - N + 3, \dots$ versus $V_1 + 1, V_2 + 1, \dots$



For the (total) internal numbers $U_1 = 1 - V_1, \dots, U_N = N - V_N$ we have reversibility:

Theorem: (Janson, K. 2011)

$$(U_1, \dots, U_{N-1}) \stackrel{d}{=} (U_{N-1}, \dots, U_1) .$$

For another proof see Knobloch, Stanciu, Wakolbinger (2011).

Theorem: (Janson, K. 2011)

$$\sqrt{\frac{N}{4 \log N}} (L_N^1 - 2) \xrightarrow{d} N(0, 1) .$$

The representation of the (total) internal numbers U_1, \dots, U_N as diminishing urn:

- Take urn with *blue* balls, altogether N balls.
- Remove them stepwise:
Successively remove a random pair of balls and replace it by one *orange* ball.
- If i balls are left,
let U_i the number of orange balls among them and V_i the number of blue balls.

Note:

$$V_{N-i} - N + 2i = i - U_{N-i} \quad , \quad V_i = i - U_i \quad .$$

Now recall

V_N, \dots, V_2 external branch numbers

W_N, \dots, W_2 internal branch numbers level 2

⋮

Note:

V_N, V_{N-1}, \dots, V_2 is a *Markov chain* (inhomogeneous in time).

$(V_N, W_N), (V_{N-1}, W_{N-1}), \dots, (V_2, W_2)$ is a Markov chain, or

W_N, \dots, W_2 is a Markov chain,
given the *random environment* V_N, \dots, V_2 .

The transition probabilities:

$$P_{v,w}^k(v', w') = \mathbf{P}(V_{k-1} = v', W_{k-1} = w' \mid V_k = v, W_k = w)$$

$$P_{v,w}^k(v-2, w+1) = \frac{\binom{v}{2}}{\binom{k}{2}}$$

$$P_{v,w}^k(v, w-2) = \frac{\binom{w}{2}}{\binom{k}{2}}$$

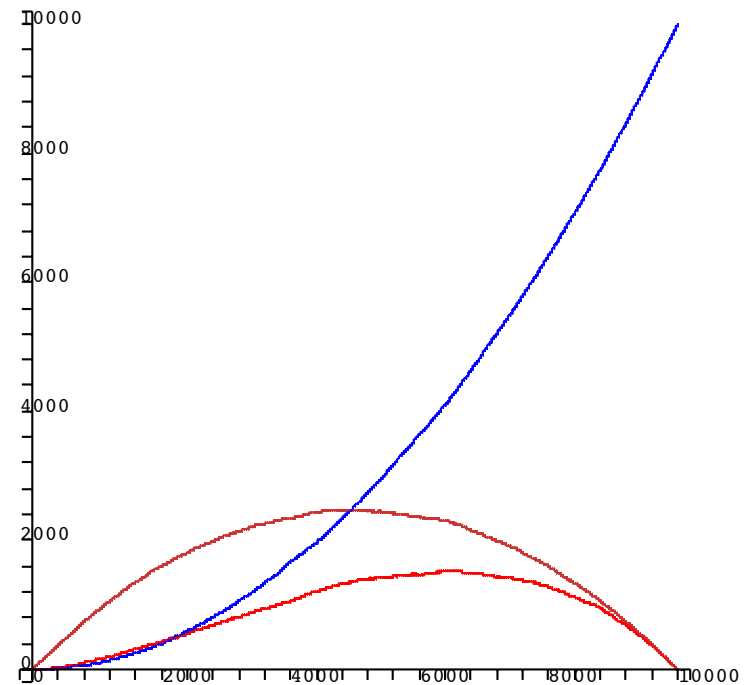
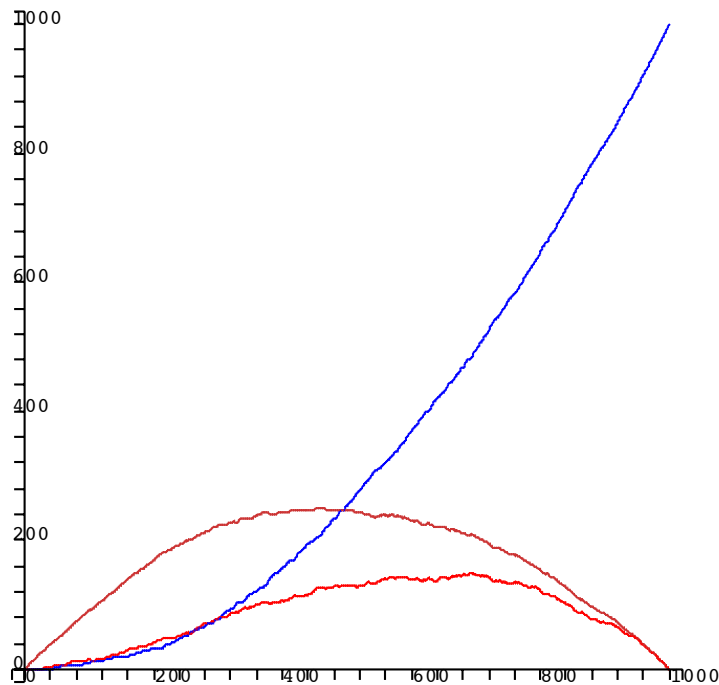
$$P_{v,w}^k(v, w) = \frac{\binom{k-v-w}{2}}{\binom{k}{2}}$$

$$P_{v,w}^k(v-1, w-1) = \frac{vw}{\binom{k}{2}}$$

$$P_{v,w}^k(v-1, w) = \frac{v(k-v-w)}{\binom{k}{2}}$$

$$P_{v,w}^k(v, w-1) = \frac{w(k-v-w)}{\binom{k}{2}}$$

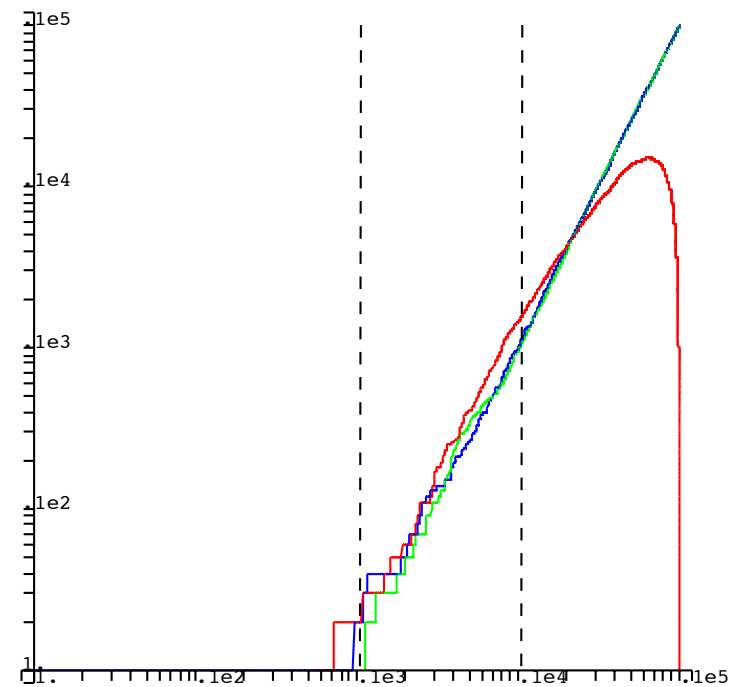
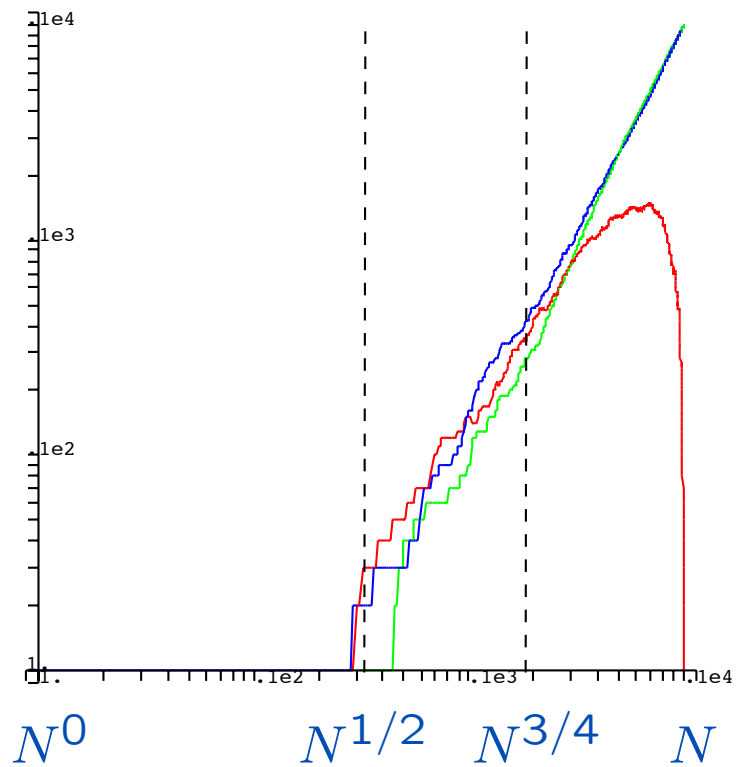
External branch numbers V_1, \dots, V_N , and
(total) internal branch numbers U_1, \dots, U_N and
internal branch numbers level 2 W_1, \dots, W_N .



Randomness within (V_i) and (W_i) :

$V_N - N + 1, V_{N-1} - N + 3, \dots$ and $V_1 + 1, V_2 + 1, \dots$

and $W_1 + 1, W_2 + 1, \dots$



Theorem: (K., Stanciu 2012)

For every $k \in \mathbb{N}$

$$\sqrt{\frac{N}{4 \log N}} \left(L_N^1 - \mu_1, \dots, L_N^k - \mu_k \right) \xrightarrow{d} N(0, I_k) .$$

with the $k \times k$ identity matrix I_k and

$$\mu_i = \frac{2}{i} .$$

Idea of proof:

Reversing time seems no longer feasible.

We couple the Markov chain

$$(V_N, W_N), \dots, (V_2, W_2)$$

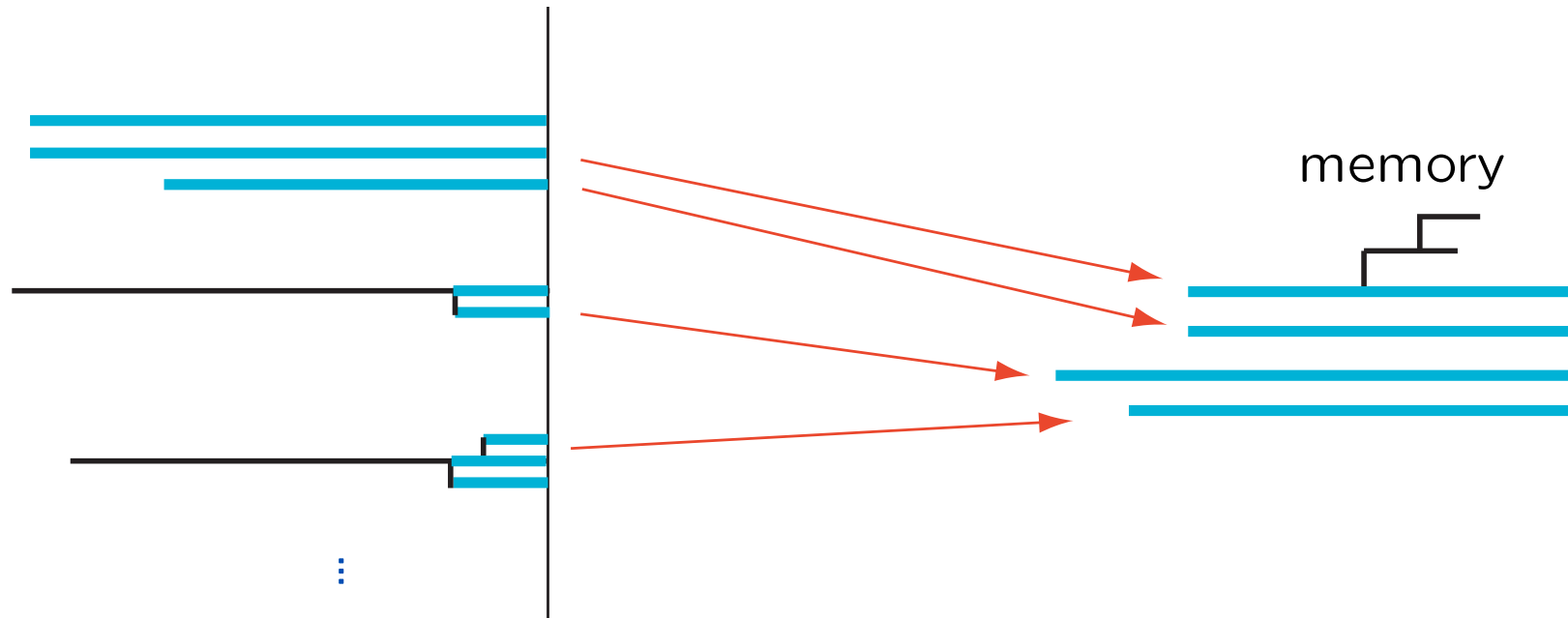
with two independent urns, i.e. with

$$(V_N, \tilde{V}_N), \dots, (V_2, \tilde{V}_2) ,$$

where $(\tilde{V}_N, \dots, \tilde{V}_2)$ is an independent copy of (V_N, \dots, V_2) .

Now the urns can be reversed.

Back to the evolving coalescent:



$$\text{Cov}(L_s, L_t) =$$

probability that a critical binary branching process consists of exactly 1 individual at time $|t - s|$.

