Assumptions of WF model

parmiete (random mating)

Simultaneous coalercent events do NOT occur

N= effective population size constant

Exchangeability of lineages

Any 2 lineages equally probable to coclerce.

time = number of gent before The present non-overlapping

Kingman's Coclescent Model will be a limit of this or a continuous time approximation

Assum- parmietre population, time measured in "coalescent units" so that · the rate of coalesce for 2 lineages is 1

Using uso for time in coalescent units and h(u) = probability 2 lineages have not coalesced at time

then the pertinent diff. eq is

$$\frac{d}{du} f(u) = - f(u)$$

23/ + u

with solution

Let P(u) = probability Z lineages did coalerce before time u

$$P(u) = 1 - e^{-u}$$
 Compared to $P(c_2 \le uN) = 1 - (1 - ||_N)$
 VF
 VF
 VF
 VF
 VF
 VF

Under the Kingman Cossescent Model, the expected time to coalerce is

$$E(2 \text{ lineages coalesce}) = \int_{0}^{\infty} u e^{-u} du du du = u \text{ in coalescence units!}$$

$$= (1-u)e^{-u} \Big|_{0}^{\infty} = 0 - (1-0)e^{-0} = 1! \text{ as preordained}$$

Some important comments freterations lete.

Important comments of consequence of using COALESCENCE UNITS
$$\Delta u \propto \Delta t$$

$$\Delta u \propto \Delta t$$

$$N$$
WF.

2) Bottlenecks:

Henecks:

Slower
$$\mathbb{E}_{WF}(C_2) > 3 = N_e$$

Some $\mathbb{E}_{WF}(C_2) > 3 = N_e$

Specd Up in time to coelescence

 $\mathbb{E}_{WF}(C_2) = N_e = 3$

The time to coalercence of 2 lineages depends on Ne(t), the current population cize. However, in coalescent units (i.e. Kingman coalescent population cize. However, in coalescent units (i.e. Kingman coalescent model) $IE(G_2)=1$ always. This is because if, say $N=N_c$ doubles model) $IE(G_2)=1$ always. This is because if, say $N=N_c$ doubles original

$$u = 1 = \frac{\Delta t}{N} = \frac{N \text{ gens}}{N \text{ indiv.}} = \frac{2N \text{ generators}}{2N \text{ individuals}}$$

i.e. you can rescale generation time and population size and still have well confescent units essentially hide any estimation fassumption/knowledge of generation time and population size (Both a plus and a minus.)

Coalescent und

under the Kingman coalescent we can also compute the expected time that n lineages coclesce into n-1

The crucial assumption is the iid.

which makes the rate $\binom{n}{2}$

all pairs equally likely to

coalerce

n= 4

In coalescence units

if K(u) = probability that in lineager remained time u, then

$$\frac{d}{du} K(u) = -\binom{n}{2} K(u)$$

 $K(0) = \Delta$

 $= \binom{n}{2} u$ and K(u) = e

There are lots of ways to determine

(E. (time to coalescente for n lineager to n-1)

* Exponental dist with rate
$$(2) = \overline{2}$$

* cdf: P(n | lineages coalesce to n-1) = 1-k(u) = 1-e

$$= |-e| - (2)u - (2)u - (2)u - (2)u - (2)u$$

$$= |-e| - (2)u - (2)u - (2)u - (2)u$$

$$= -e| (-2) = (2)e - (2)e$$

Again exponental ...

· More formally,

$$|E(n | lineager coclere)| = \int_{0}^{\infty} u(2)e du = \frac{1}{(2)} = \frac{2}{n(n-1)}$$

Theorem: Under the kingman contenent model, given a lineager in

a population, the expected time to the next coclescent event is

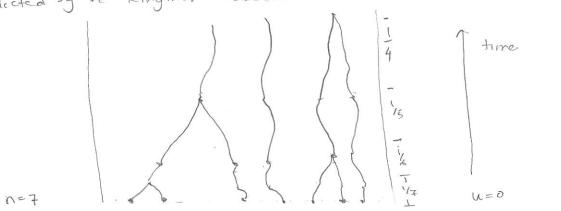
$$= \frac{2}{n(n-1)} \left[\frac{1}{3} \right] \frac{1}{6} \frac{1}{10} \frac{1}{15} \frac{1}{21} \text{ etc.}$$

$$= \frac{2}{n(n-1)} \left[\frac{1}{3} \right] \frac{1}{6} \frac{1}{10} \frac{1}{15} \frac{1}{21} \text{ etc.}$$

$$= \frac{1331}{14(41)}$$

This has a profound effect on the types of gene trees

predicted by the Kingman cockscent model.



Le. Coalescence events are more numerous near the present.

R example.

Finally, given in lineages then the expected time for them to coalesce down

F (gene tree w/ n lineage formed) = $\sum_{i=2}^{n} \frac{1}{i} = \sum_{i=2}^{n} \frac{1}{(i-1)} = 2\sum_{i=2}^{n} \frac{1$ to one lineage is

$$= 2\left(1-\frac{1}{n}\right) \quad \text{and} \quad \lim_{n\to\infty} z\left(1-\frac{1}{n}\right) = 2 \quad \text{confirmed points}$$