

Results from scenario with 2 diploids, 3 tetraploids, 1-3 hybridizations

Graham Jones

2012-11-05

1 Scenarios

Scenarios shown in Fig 1. All genes have length 500. Population sizes are 100,000 at tips, and at rootward ends of branches, and 200,000 at tipward ends of internal branches and at the root. The root height is 0.012.

Strict clock branch rates assumed in simulation and BEAST.

HKY substitution model assumed in simulation and BEAST. In simulations, the Seq-Gen parameters are kappa=3, frequencies .3 A and T, .2 C and G. (Seq-Gen called with `-t3.0 -f0.3,0.2,0.2,0.3`. These are estimated in BEAST.

No site rate heterogeneity assumed within genes in simulation and BEAST.

Genes have the same mutation rate in simulations. Rates are estimated in BEAST.

Priors on population size scaling factor η , relative mutation rates of genes, and the ‘event rate’ (something like a diversification rate) λ for the diploid and tetraploid trees in the network, are all diffuse lognormals.

$G=3,9$. $N=1,3$. $T=8 \times 10^{-8}$, 1.6×10^{-7} .

When $T = 8 \times 10^{-8}$, then root height is $.012/(8 \times 10^{-8}) = 150,000$ generations. When $T = 1.6 \times 10^{-7}$, then root height is 75,000. The expected time for a coalescence between two genes in the same population of size 100,000 (100,000 gene copies per diploid genome) is 100,000 generations.

MCMC chains are 3,000,000 generations with first 1,000,000 discarded as burnin.

Fig 2 shows the simulated gene trees for scenario D, with $G=9$, $N=1$, $T=8 \times 10^{-8}$, replicate 1.

2 Results

Fig 3 shows the posterior probabilities for the number of hybridizations being equal to the correct value. Each graph shows ten replicates.

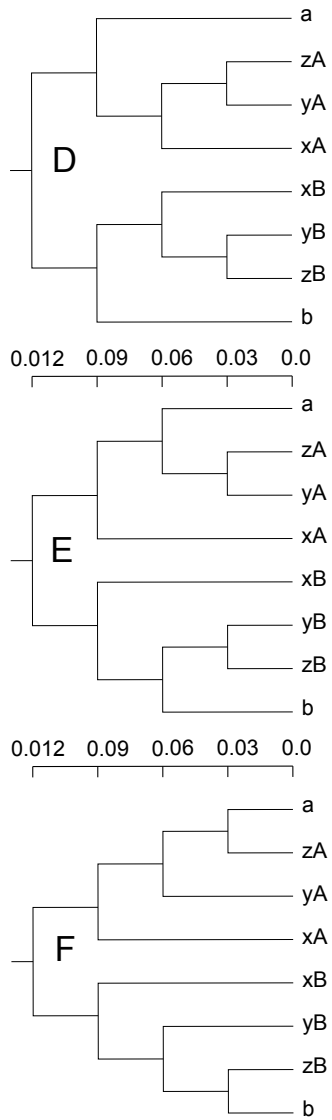


Figure 1: Scenario D,E,F: the true MUL-trees

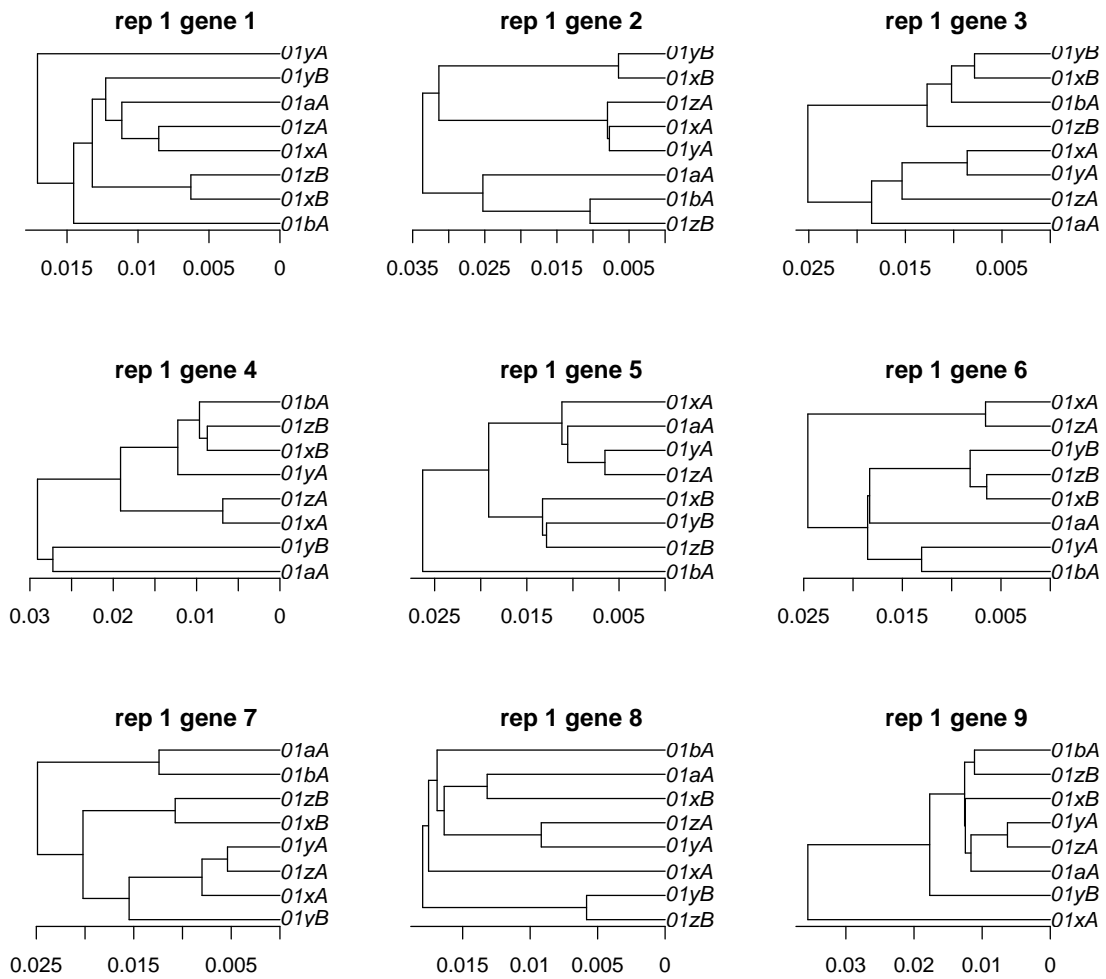


Figure 2: Simulated gene trees for scenario D, with $G=9$, $N=1$, $T=8e-8$, replicate 1.

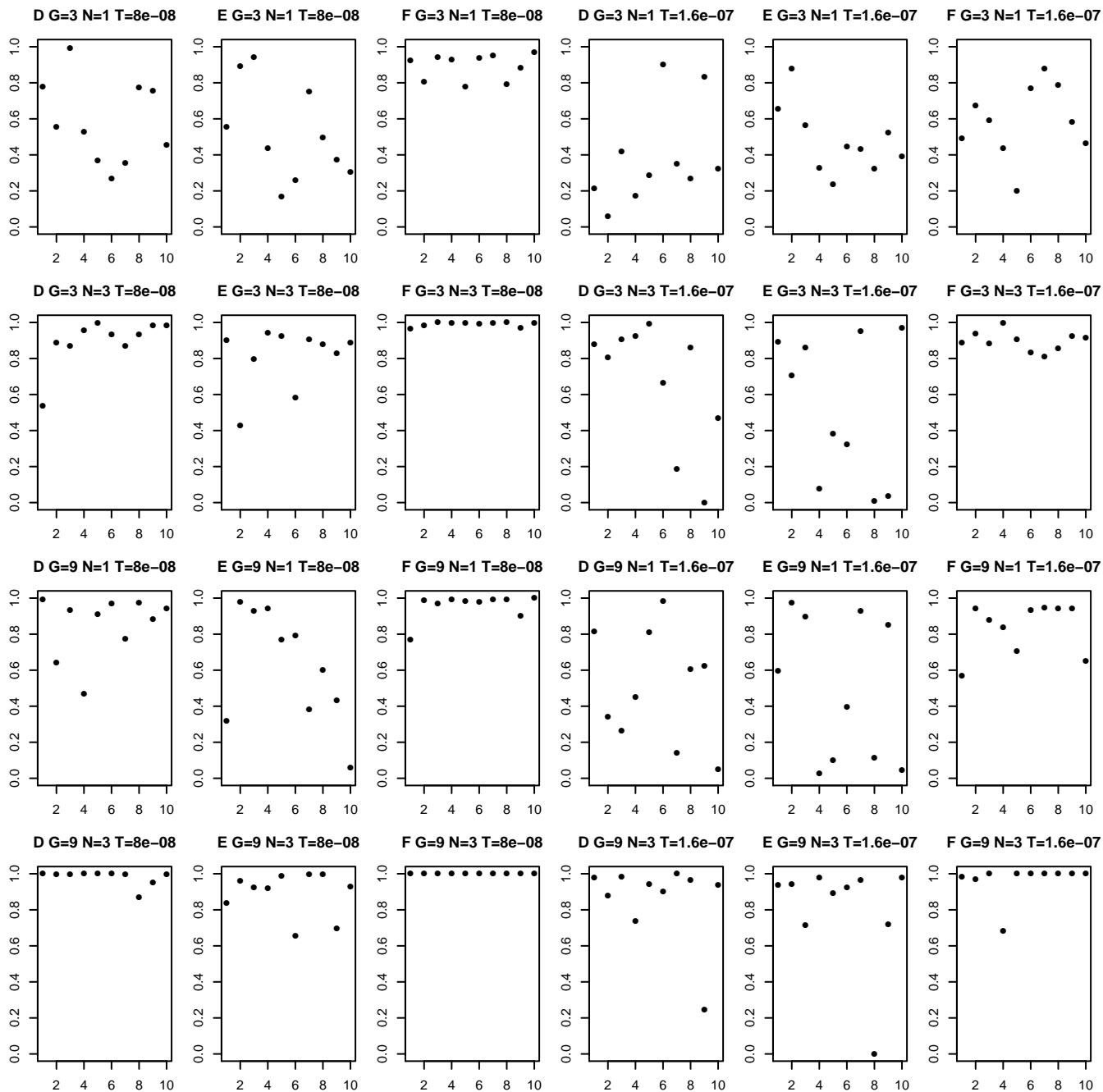


Figure 3: Posterior probabilities for the correct number of hybridizations

3 Configuration files

D - one hybridization

```
PROGRAMPARAMETERS
beastseed 652989819
beastchainlength 3000000
beastscreen.logevery 1000
beastparams.logevery 1000
beastgtrees.logevery 1000
beastmultree.logevery 1000
beastdbugtune.logevery 10000
treeannburnin 1001
SCENARIO
nofGNRT 4
3 1 10 8e-8 1.6e-7
3 3 10 8e-8 1.6e-7
9 1 10 8e-8 1.6e-7
9 3 10 8e-8 1.6e-7
genelength 500
nofdips 2
noftets 3
aA hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
bA hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
zA hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
zB hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
yA hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
yB hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
xA hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
xB hgt 0 hybhgt * tipp 100000 rootp 100000 hybp *
(zA,yA) hgt 0.003 hybhgt * tipp * rootp 100000 hybp *
(zB,yB) hgt 0.003 hybhgt * tipp * rootp 100000 hybp *
((zA,yA),xA) hgt 0.006 hybhgt 0.008 tipp * rootp 100000 hybp 100000
((zB,yB),xB) hgt 0.006 hybhgt 0.008 tipp * rootp 100000 hybp 100000
(aA,((zA,yA),xA)) hgt 0.009 hybhgt * tipp * rootp 100000 hybp *
(bA,((zB,yB),xB)) hgt 0.009 hybhgt * tipp * rootp 100000 hybp *
((aA,((zA,yA),xA)),(bA,((zB,yB),xB))) hgt 0.012 hybhgt * tipp * rootp 100000 hybp *
```

E - two hybridizations

```

PROGRAMPARAMETERS
beastseed 652989819
beastchainlength 3000000
beastscreen.logevery 10000
beastparams.logevery 1000
beastgtrees.logevery 1000
beastmultree.logevery 1000
beastdbgtune.logevery 10000
treeannburnin 1001

```

SCENARIO

```

nofGNRT 4
3 1 10 8e-8 1.6e-7
3 3 10 8e-8 1.6e-7
9 1 10 8e-8 1.6e-7
9 3 10 8e-8 1.6e-7
genelength 500
nofdips 2
noftets 3

```

aA	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
bA	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
zA	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
zB	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
yA	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
yB	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
xA	hgt 0	hybhgt 0.008	tipp 100000	rootp 100000	hybp 100000
xB	hgt 0	hybhgt 0.008	tipp 100000	rootp 100000	hybp 100000
(zA,yA)	hgt 0.003	hybhgt 0.005	tipp *	rootp 100000	hybp 100000
(zB,yB)	hgt 0.003	hybhgt 0.005	tipp *	rootp 100000	hybp 100000
(aA,(zA,yA))	hgt 0.006	hybhgt *	tipp *	rootp 100000	hybp *
(bA,(zB,yB))	hgt 0.006	hybhgt *	tipp *	rootp 100000	hybp *
(xA,(aA,(zA,yA)))	hgt 0.009	hybhgt *	tipp *	rootp 100000	hybp *
(xB,(bA,(zB,yB)))	hgt 0.009	hybhgt *	tipp *	rootp 100000	hybp *
((xA,(aA,(zA,yA))),(xB,(bA,(zB,yB))))	hgt 0.012	hybhgt *	tipp *	rootp *	hybp *

F - three hybridizations

PROGRAMPARAMETERS

beastseed 652989819
 beastchainlength 3000000
 beastscreen.logevery 10000
 beastparams.logevery 1000
 beastgtrees.logevery 1000
 bestmultree.logevery 1000
 bestdbugtune.logevery 10000
 treeannburnin 1001

SCENARIO

nofGNRT 4
 3 1 10 8e-8 1.6e-7
 3 3 10 8e-8 1.6e-7
 9 1 10 8e-8 1.6e-7
 9 3 10 8e-8 1.6e-7
 genelength 500
 nofdips 2
 noftets 3

aA	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
bA	hgt 0	hybhgt *	tipp 100000	rootp 100000	hybp *
zA	hgt 0	hybhgt 0.002	tipp 100000	rootp 100000	hybp 100000
zB	hgt 0	hybhgt 0.002	tipp 100000	rootp 100000	hybp 100000
yA	hgt 0	hybhgt 0.005	tipp 100000	rootp 100000	hybp 100000
yB	hgt 0	hybhgt 0.005	tipp 100000	rootp 100000	hybp 100000
xA	hgt 0	hybhgt 0.008	tipp 100000	rootp 100000	hybp 100000
xB	hgt 0	hybhgt 0.008	tipp 100000	rootp 100000	hybp 100000
(zA, aA)	hgt 0.003	hybhgt *	tipp *	rootp 100000	hybp *
(zB, bA)	hgt 0.003	hybhgt *	tipp *	rootp 100000	hybp *
(yA, (zA, aA))	hgt 0.006	hybhgt *	tipp *	rootp 100000	hybp *
(yB, (zB, bA))	hgt 0.006	hybhgt *	tipp *	rootp 100000	hybp *
(xA, (yA, (zA, aA)))	hgt 0.009	hybhgt *	tipp *	rootp 100000	hybp *
(xB, (yB, (zB, bA)))	hgt 0.009	hybhgt *	tipp *	rootp 100000	hybp *
((xA, (yA, (zA, aA))), (xB, (yB, (zB, bA))))	hgt 0.012	hybhgt *	tipp *	rootp *	hybp *