Transition Rates

Beatriz Willink

Jan 2021

```
Read packages
```

```
require(ggplot2)
require(tidyr)
require(wesanderson)
require(gridExtra)
```

Create file with the posterior samples for each run

Read and combine runs

```
run1 <- read.table("./run1.txt", header = T, sep = "\t")[,c(1,7:18)]
run2 <- read.table("./run2.txt", header = T, sep = "\t")[,c(1,7:18)]

posterior.dat <- rbind(run1, run2)
rm(run1, run2)
n = nrow(posterior.dat)</pre>
```

Reshape datframe and rescale transition rates

```
dat <-gather(posterior.dat, transition, rate, q01:q32, factor_key=TRUE)
dat$rate <- dat$rate/100</pre>
```

Transition rates for gains of female polymorphism

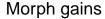
```
facet_wrap(~facet, scales = "free", nrow = 1)+
xlim(-0.01,0.3)+
theme_minimal(base_size = 11) +
labs(title = "Morph_gains", x = "Transition_rate", y = "Posterior_
    density", fill = "Transition") +
theme(legend.position = "none") +
theme(strip.text.x = element_text(angle = 0, hjust = 0)) +
scale_fill_manual(values = wes_palette("IsleofDogs1")[c(1,2,5,3,4)])
```

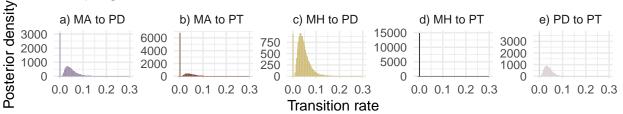
Transition rates for losses of female polymorphism

```
losses <- (dat[which(dat$transition == "q10" | dat$transition == "q12" |</pre>
                         dat$transition == "q30" | dat$transition == "q32" |
                        dat$transition == "q31"),])
losses facet \leftarrow rep(c("f)_{\sqcup}PD_{\sqcup}to_{\sqcup}MA", "h)_{\sqcup}PD_{\sqcup}to_{\sqcup}MH", "g)_{\sqcup}PT_{\sqcup}to_{\sqcup}MA", "j)_{\sqcup}PT_{\sqcup}
   to_{\square}PD", "i)_{\square}PT_{\square}to_{\square}MH"), each = n)
B <- ggplot (data = losses, aes(x = rate, fill = transition)) +
  geom_histogram(position = "identity", bins = 100, colour = NA, alpha =
      0.8) +
  facet_wrap(~facet, scales = "free", nrow = 1)+
  xlim(-0.01,0.3) +
  theme minimal(base size = 11) +
  theme(legend.position = "none") +
  theme(strip.text.x = element_text(angle = 0, hjust = 0))+
  labs(title = "Morph_{\sqcup}losses" , x = "Transition_{\sqcup}rate", y = "Posterior_{\sqcup}
      density", fill = "Transition") +
  scale_fill_manual(values = wes_palette("IsleofDogs1")[c(1,3,2,4,5)])
```

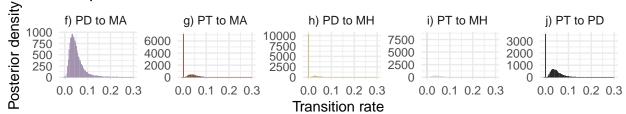
Transitions between sexual dimorphism and monomorphism with monomorphic females

Plot everything

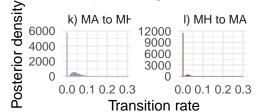




Morph losses



Morph changes



How common are non-zero rates for each transition

```
##
       transition
                     type prob_nonzero
##
                               0.8078750
  1
               q01
                      gain
##
  2
               q02 change
                               0.6420625
                               0.5778750
##
  3
               q03
                      gain
               q10
##
                      loss
                               0.9975000
##
   5
               q12
                      loss
                               0.3517500
               q13
   6
                               0.7675625
                      gain
##
   7
                               0.2815625
               q20 change
##
   8
                               0.9422500
               q21
                      gain
##
  9
                               0.0670000
               q23
                      gain
## 10
               q30
                      loss
                               0.5537500
```

## 11	q31	loss	0.7762500
## 12	a32	loss	0.4630000