uptake_exp

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```r
head(d)
```
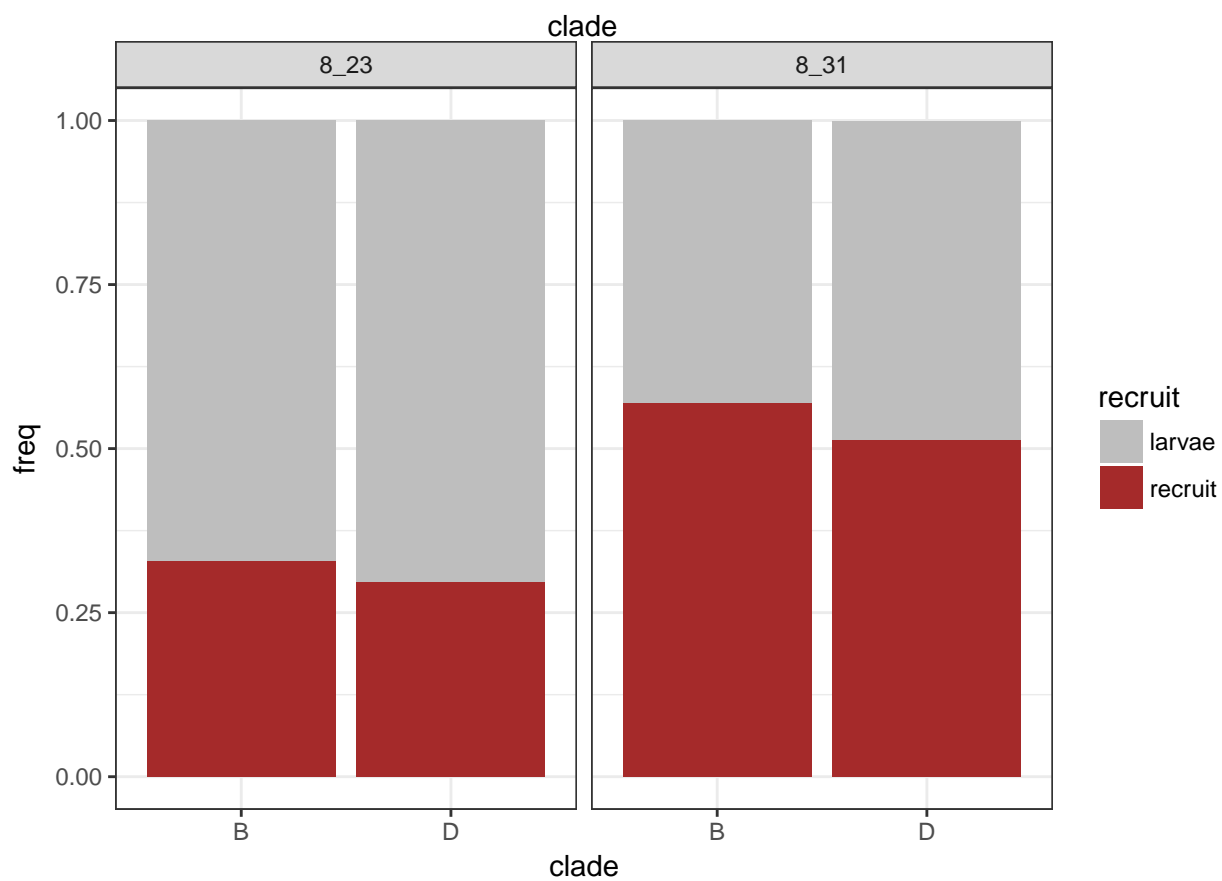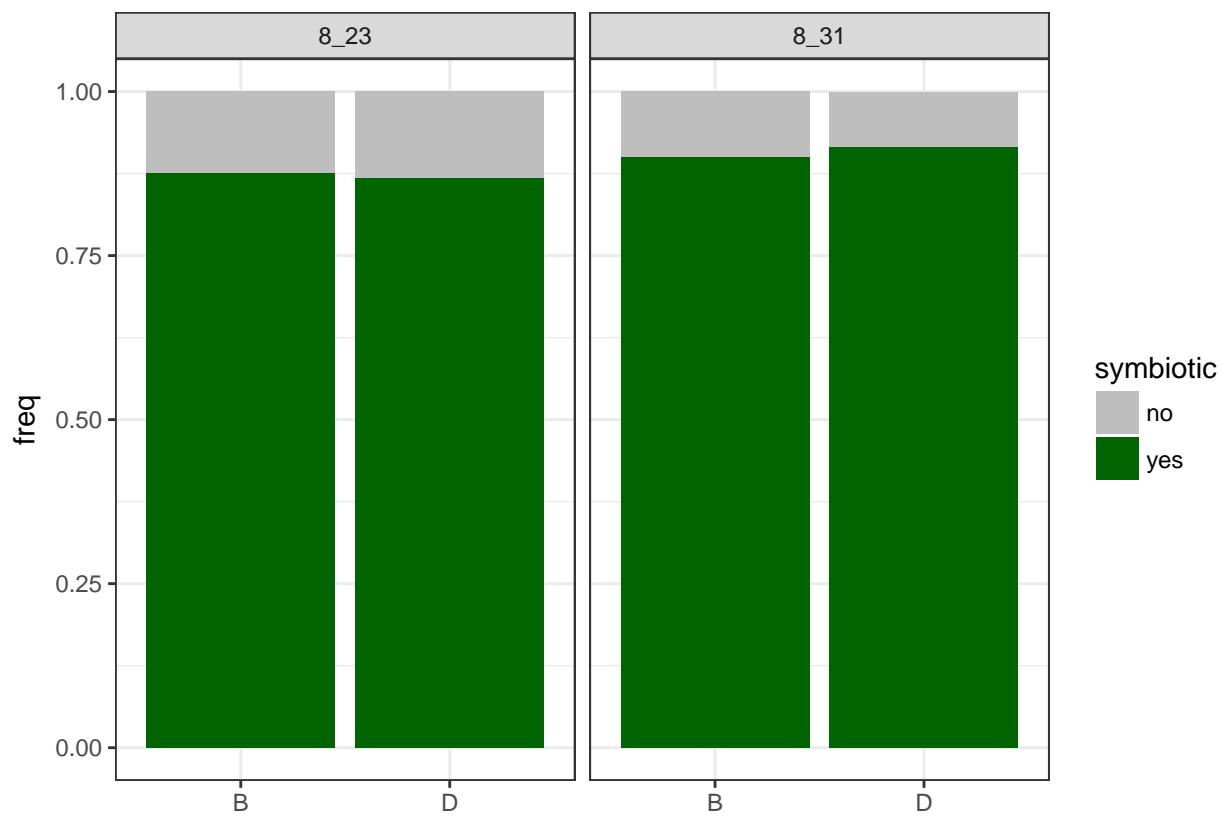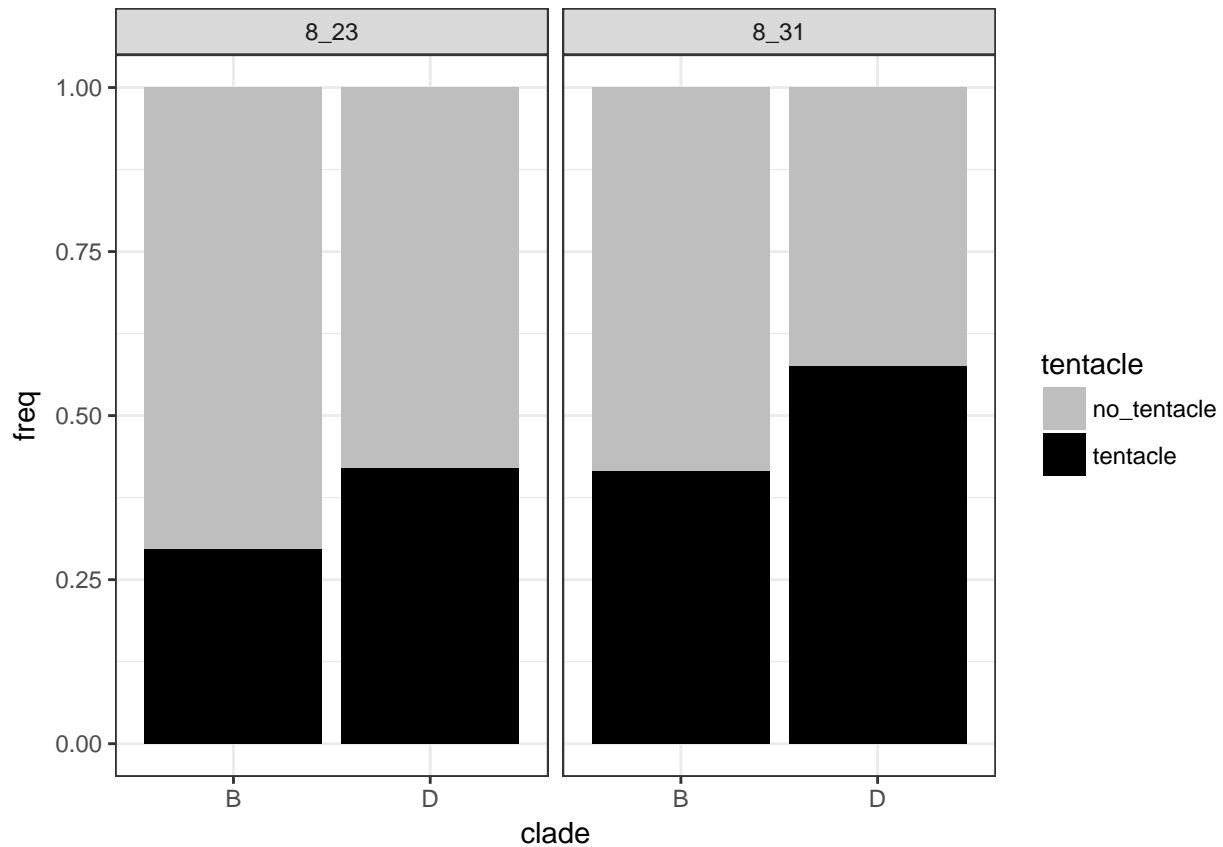
```
## # A tibble: 6 x 11
##    clade plate  well type  count sym   swim  day   tentacle     symbiotic
##    <chr> <chr> <int> <chr> <int> <chr> <chr> <chr> <fct>        <fct>
## 1 D     p1        1 slt       3 s     l     8_23  tentacle     yes
## 2 D     p1        1 sl        4 s     l     8_23  no_tentacle  yes
## 3 D     p1        1 al        3 a     l     8_23  no_tentacle  no
## 4 D     p1        2 ss        1 s     s     8_23  no_tentacle  yes
## 5 D     p1        2 slt       4 s     l     8_23  tentacle     yes
## 6 D     p1        2 sl        4 s     l     8_23  no_tentacle  yes
## # ... with 1 more variable: recruit <fct>
```

```r
d %>% ungroup() %>% group_by(day, clade,symbiotic) %>%
  summarize(sum(count))
```

```
## # A tibble: 9 x 4
## # Groups:   day, clade [?]
##   day   clade symbiotic `sum(count)`
##   <chr> <chr> <fct>            <int>
## 1 8_23  B     no                  49
## 2 8_23  B     yes                346
## 3 8_23  D     no                  45
## 4 8_23  D     yes                295
## 5 8_31  B     no                  20
## 6 8_31  B     yes                180
## 7 8_31  D     no                  15
## 8 8_31  D     yes                164
## 9 8_31  NoSym no                 163
```

```
summary(mcmcSymbiotic)
```

```
##
##  Iterations = 3001:12991
##  Thinning interval  = 10
##  Sample size  = 1000
##
##  DIC: 306.5741
##
##  G-structure:  ~plate
##
##       post.mean  l-95% CI u-95% CI eff.samp
## plate     0.386 5.695e-17    1.526    79.53
##
##  R-structure:  ~units
##
##       post.mean l-95% CI u-95% CI eff.samp
## units     1.393   0.6831    2.238    73.72
##
##  Location effects: symbiotic_count ~ clade * day
##
##                post.mean l-95% CI u-95% CI eff.samp  pMCMC
## (Intercept)      2.66795  1.96568  3.55106   1095.1 <0.001 ***
## cladeD          -0.05927 -0.94323  0.84665   1000.0  0.886
## day8_31         -0.56422 -1.38773  0.39873   1000.0  0.234
## cladeD:day8_31   0.18714 -1.16861  1.54020    795.8  0.774
## ---
```

```
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```
```r
summary(mcmcRecruit)
```
```
##
##  Iterations = 3001:12991
##  Thinning interval  = 10
##  Sample size  = 1000
##
##  DIC: 334.4549
##
##  G-structure:  ~plate
##
##         post.mean l-95% CI u-95% CI eff.samp
## plate      1.011   0.1597    2.562     1000
##
##  R-structure:  ~units
##
##         post.mean l-95% CI u-95% CI eff.samp
## units     0.2444   0.1189   0.3932    865.7
##
##  Location effects: recruit_count ~ clade * day
##
##                post.mean l-95% CI u-95% CI eff.samp  pMCMC
## (Intercept)       3.0324   2.2739   3.8568     1000 <0.001 ***
## cladeD           -0.2711  -0.6737   0.1681     1000  0.222
## day8_31          -0.6878  -1.0918  -0.2522     1000  0.006 **
## cladeD:day8_31    0.1654  -0.4187   0.7709     1000  0.600
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```
```r
summary(mcmcTentacle)
```
```
##
##  Iterations = 3001:12991
##  Thinning interval  = 10
##  Sample size  = 1000
##
##  DIC: 322.3711
##
##  G-structure:  ~plate
##
##         post.mean l-95% CI u-95% CI eff.samp
## plate    0.7512   0.1136    1.877     1000
##
##  R-structure:  ~units
##
##         post.mean l-95% CI u-95% CI eff.samp
## units    0.1608  0.07458   0.2644    753.3
##
##  Location effects: tentacle_count ~ clade * day
##
##                post.mean l-95% CI u-95% CI eff.samp  pMCMC
## (Intercept)     3.10855  2.41597  3.81769     1000 <0.001 ***
## cladeD         -0.15743 -0.51574  0.18780     1000  0.376
```

```
## day8_31       -0.66744 -1.06329 -0.30175     1000 <0.001 ***
## cladeD:day8_31  0.02943 -0.44643  0.58529     1000  0.914
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```