

2018-02-27_eclosion_summaries

Eclosion data

```
library(data.table)
library(ggplot2)
library(plyr)
```

```
##grabbing data if git cloned
#eclosions<-fread("../Data/2018-01-26_rhagoletis_masterdata_data_slice.csv")
#grabbing from online
eclosions<-fread("https://raw.githubusercontent.com/adnguyen/Circadian_rhythm_runs_seasonal_timing/master/Data/2018-01-26_rhagoletis_masterdata_data_slice.csv")
```

```
## Warning in fread("https://raw.githubusercontent.com/adnguyen/Circadian_rhythm_runs_seasonal_timing/master/Data/2018-01-26_rhagoletis_masterdata_data_slice.csv"): Bumped column 41 to
## type character on data row 530, field contains 'change food 2017-10-31
## 21:50'. Coercing previously read values in this column from logical,
## integer or numeric back to character which may not be lossless; e.g., if
## '00' and '000' occurred before they will now be just '0', and there may
## be inconsistencies with treatment of ',', ' and ',NA,' too (if they occurred
## in this column before the bump). If this matters please rerun and set
## 'colClasses' to 'character' for this column. Please note that column type
## detection uses a sample of 1,000 rows (100 rows at 10 points) so hopefully
## this message should be very rare. If reporting to datatable-help, please
## rerun and include the output from verbose=TRUE.
```

```
head(eclosions,10)
```

```
##      Ind_ID tape Site_name mass_day10 purge_time_1 purgel collection_date
## 1:      1 blue      OG      6.938      13:38      NA      2017-08-21
## 2:      2 blue  Ferris     11.175     13:39      NA      2017-08-21
## 3:      3 blue  Ferris     6.719     13:39      NA      2017-08-21
## 4:      4 blue  Ferris    10.719     13:40      NA      2017-08-21
## 5:      5 blue      OG     3.848     13:41      NA      2017-08-21
## 6:      6 blue      OG     7.576     13:37      NA      2017-08-21
## 7:      7 blue      OG     6.413     13:35      NA      2017-08-21
## 8:      8 blue      OG     9.365     13:57      NA      2017-08-21
## 9:      9 blue  Ferris     7.978     13:41      NA      2017-08-21
## 10:     10 blue      OG     4.895     13:36      NA      2017-08-21
##      day10 cohort_date cohort_day Host Li-cor_1 resp_time_1
## 1: 2017-09-04 2017-08-25          2 Apple      6262
## 2: 2017-09-04 2017-08-25          2 Apple      6262 10:13:15
## 3: 2017-09-04 2017-08-25          2 Apple      6262 10:14:47
## 4: 2017-09-04 2017-08-25          2 Apple      6262 10:16:43
## 5: 2017-09-04 2017-08-25          2 Apple      6262 10:18:41
## 6: 2017-09-04 2017-08-25          2 Apple      6262 10:20:19
## 7: 2017-09-04 2017-08-25          2 Apple      6262 10:22:07
## 8: 2017-09-04 2017-08-25          2 Apple      6262 10:23:42
## 9: 2017-09-04 2017-08-25          2 Apple      6262 10:25:39
## 10: 2017-09-04 2017-08-25          2 Apple      6262 10:27:14
##      resp_day11 mass_day14 purge_time_2 resp_time_2 resp_day15 Li_cor2
```

## 1:	NA	6.187	10:51	10:13:48	0.1432514	6262	
## 2:	0.2941100	9.967		10:16:22	0.1702350	6262	
## 3:	0.1052925	6.118		10:18:10	0.1076286	6262	
## 4:	1.3445380	9.539		10:19:38	1.3873900	6262	
## 5:	0.1492267	3.479		10:21:23	0.1182286	6262	
## 6:	1.4202160	6.824		10:22:55	1.4689120	6262	
## 7:	0.1365561	5.738		10:24:36	0.1601623	6262	
## 8:	1.3082910	8.401		10:25:57	1.4328830	6262	
## 9:	0.1591373	7.323		10:27:35	0.0809000	6262	
## 10:	0.1027624	4.349		10:28:53	0.0642000	6262	
##	notes	Resp_code	treatment	uniqueID	eclosion_date	eclosion_days	well_id
## 1:		0	SO	2b1		NA	A1
## 2:		1	GC	2b2		NA	
## 3:		1	RT	2b3	2017-10-29	64	A1
## 4:		1	GC	2b4		NA	
## 5:		1	RT	2b5	2017-10-20	56	A2
## 6:		1	SO	2b6		NA	A2
## 7:		1	SO	2b7		NA	A3
## 8:		1	RT	2b8	2017-09-26	32	A3
## 9:		1	SO	2b9		NA	A4
## 10:		1	GC	2b10		NA	
##	organism	Trikinetics_position		Trikinetics_monitor			
## 1:			NA		NA		
## 2:			NA		NA		
## 3:	fly		8		1		
## 4:			NA		NA		
## 5:	fly		15		2		
## 6:			NA		NA		
## 7:			NA		NA		
## 8:	fly		30		2		
## 9:			NA		NA		
## 10:			NA		NA		
##	Trikinetics_entry_LD_time		Trikinetic_exit_date				
## 1:							
## 2:							
## 3:		3:34		2107-11-05			
## 4:							
## 5:		13:03		2017-10-25			
## 6:							
## 7:							
## 8:		16:08		2017-10-05			
## 9:							
## 10:							
##	Trikinetics_exit_LD_time		notes_2				
## 1:							
## 2:							
## 3:		20:33	changed water	2017-11-03	11:56		
## 4:							
## 5:		9:58					
## 6:							
## 7:							
## 8:		12:45					
## 9:							
## 10:							

```
##      Free_run_monitor Free_run_position Free_run_entry_date
## 1:      NA              NA
## 2:      NA              NA
## 3:      4              11      2017-11-05
## 4:      NA              NA
## 5:      NA              NA
## 6:      NA              NA
## 7:      NA              NA
## 8:      NA              NA
## 9:      NA              NA
## 10:     NA              NA
##      Free_run_entry_time Free_run_exit_date Free_run_exit_time
## 1:
## 2:
## 3:      20:34      2017-12-09      20:00
## 4:
## 5:
## 6:
## 7:
## 8:
## 9:
## 10:
##
##      notes_3 V41
## 1:
## 2:
## 3: changed water 2017-11-28
## 4:
## 5:
## 6:
## 7:
## 8:
## 9:
## 10:
```

```
tail(eclosions,10)
```

```
##      Ind_ID  tape Site_name mass_day10 purge_time_1 purge1 collection_date
## 1:      8 white   Ferris    11.578      NA      NA      2017-09-19
## 2:      9 white   Ferris     6.663      NA      NA      2017-09-19
## 3:     10 white   Ferris     9.641      NA      NA      2017-09-19
## 4:     11 white    Blank      NA      NA      NA      2017-09-19
## 5:     12 white    Blank      NA      12:28    NA      2017-09-19
## 6:      1 white   Ferris    12.129      12:20    NA      2017-09-19
## 7:      2 white   Ferris    11.226      NA      NA      2017-09-19
## 8:      3 white   Ferris     7.681      NA      NA      2017-09-19
## 9:      4 white    Blank      NA      NA      NA      2017-09-19
## 10:     5 white    Blank      NA      12:21    NA      2017-09-19
##      day10 cohort_date cohort_day Host Li-cor_1 resp_time_1 resp_day11
## 1: 2017-10-12 2017-10-02      11 Haw    6262    10:40:24  0.3168173
## 2: 2017-10-12 2017-10-02      11 Haw    6262    10:42:06  0.1840791
## 3: 2017-10-12 2017-10-02      11 Haw    6262    10:43:56  1.1545140
## 4: 2017-10-12 2017-10-02      11 Haw    6262    10:46:05  0.0120000
## 5: 2017-10-12 2017-10-02      11 Haw    6262      NA
## 6: 2017-10-13 2017-10-03      12 Haw    7000    10:44:39  0.1462170
## 7: 2017-10-13 2017-10-03      12 Haw    7000    10:45:41  1.0436650
```

##	8:	2017-10-13	2017-10-03	12	Haw	7000	10:46:57	0.0739000
##	9:	2017-10-13	2017-10-03	12	Haw	7000	10:48:01	0.0090700
##	10:	2017-10-13	2017-10-03	12	Haw	7000	10:48:55	0.0367000
##		mass_day14	purge_time_2	resp_time_2	resp_day15	Li_cor2	notes	Resp_code
##	1:	10.750		14:30:24	0.1589825	7000		1
##	2:	6.126		14:31:35	0.0881000	7000		1
##	3:	8.644		14:32:41	0.1198599	7000		1
##	4:	NA		14:33:40	0.0112000	7000		1
##	5:	NA	10:43	14:34:27	0.0104000	7000		0
##	6:	11.012	14:44		NA	NA		NA
##	7:	8.514			NA	NA		NA
##	8:	7.011			NA	NA		NA
##	9:	NA			NA	NA		NA
##	10:	NA	14:45		NA	NA		NA
##		treatment	uniqueID	eclosion_date	eclosion_days	well_id	organism	
##	1:	RT	h11w8	2017-12-04		64	D2	fly
##	2:		h11w9			NA		
##	3:		h11w10			NA		
##	4:		h11w11			NA		
##	5:		h11w12			NA		
##	6:		h12w1			NA		
##	7:		h12w2			NA		
##	8:		h12w3			NA		
##	9:		h12w4			NA		
##	10:		h12w5			NA		
##		Trikinetics_position	Trikinetics_monitor	Trikinetics_entry_LD_time				
##	1:		3	2			10:38	
##	2:		NA	NA				
##	3:		NA	NA				
##	4:		NA	NA				
##	5:		NA	NA				
##	6:		NA	NA				
##	7:		NA	NA				
##	8:		NA	NA				
##	9:		NA	NA				
##	10:		NA	NA				
##		Trikinetic_exit_date	Trikinetics_exit_LD_time	notes_2	Free_run_monitor			
##	1:	2017-12-12		21:24			4	
##	2:						NA	
##	3:						NA	
##	4:						NA	
##	5:						NA	
##	6:						NA	
##	7:						NA	
##	8:						NA	
##	9:						NA	
##	10:						NA	
##		Free_run_position	Free_run_entry_date	Free_run_entry_time				
##	1:	16	2017-12-12	21:24				
##	2:	NA						
##	3:	NA						
##	4:	NA						
##	5:	NA						
##	6:	NA						

```
## 7: NA
## 8: NA
## 9: NA
## 10: NA
## Free_run_exit_date Free_run_exit_time notes_3
## 1: changed water 2017-12-21 22:20
## 2:
## 3:
## 4:
## 5:
## 6:
## 7:
## 8:
## 9:
## 10:
## V41
## 1:
## 2:
## 3:
## 4:
## 5:
## 6:
## 7:
## 8:
## 9:
## 10:
```

```
### getting numbers for eclosions without simulated overwintering, so these pupae were left at rearing +
sub<-subset(eclosions,eclosion_days>1)
t1<-ddply(sub,. (Host,organism),summarize,counts=length(eclosion_days))
knitr::kable(t1)
```

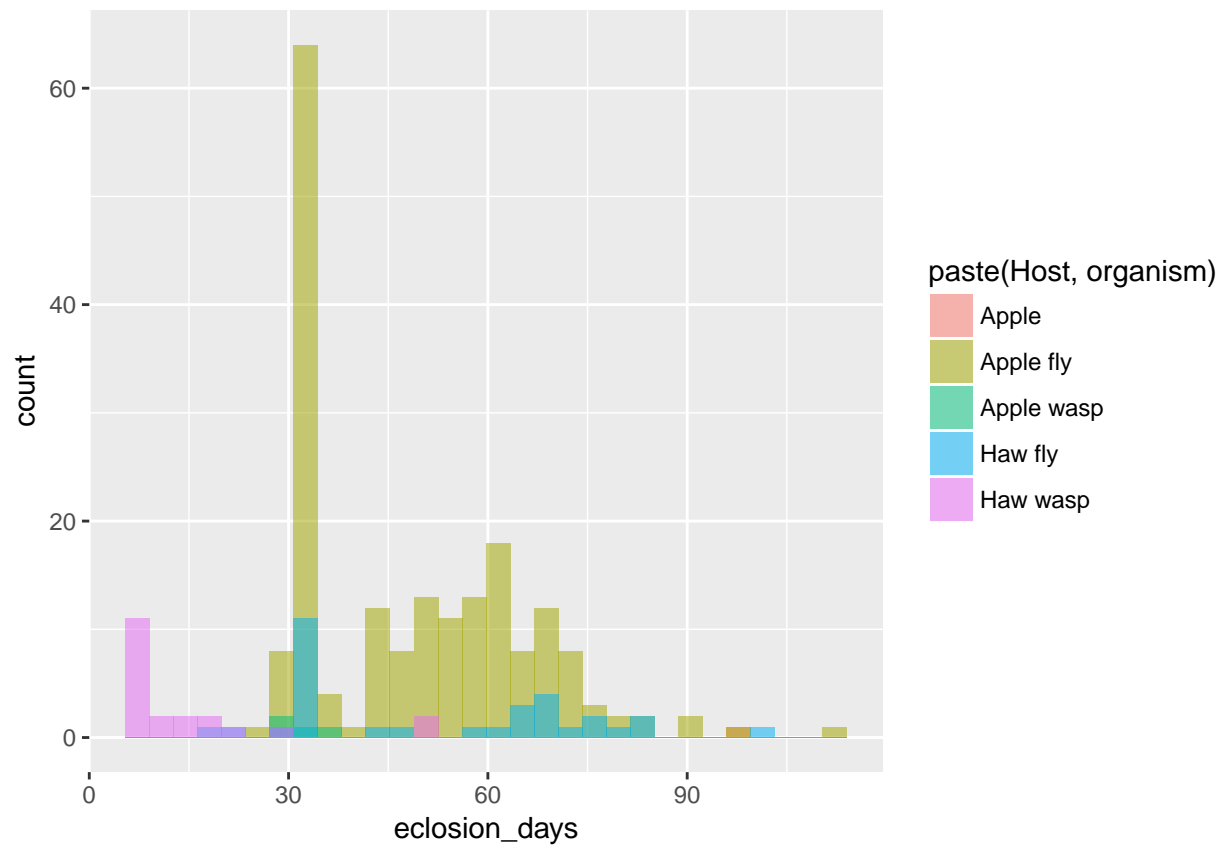
Host	organism	counts
Apple		1
Apple	fly	192
Apple	wasp	4
Haw	fly	32
Haw	wasp	21

```
# plotting distributions
```

```
ggplot(eclosions,aes(x=eclosion_days,fill=paste(Host,organism)))+geom_histogram(position="identity",alpha=0.5)
```

```
## `stat_bin()` using `bins = 30`. Pick better value with `binwidth`.
```

```
## Warning: Removed 1659 rows containing non-finite values (stat_bin).
```



```
flies<-subset(eclosions,organism=="fly")
ggplot(flies,aes(x=eclosion_days,fill=paste(Host)))+geom_density(position="identity",alpha=.52)

## Warning: Removed 1 rows containing non-finite values (stat_density).
```

