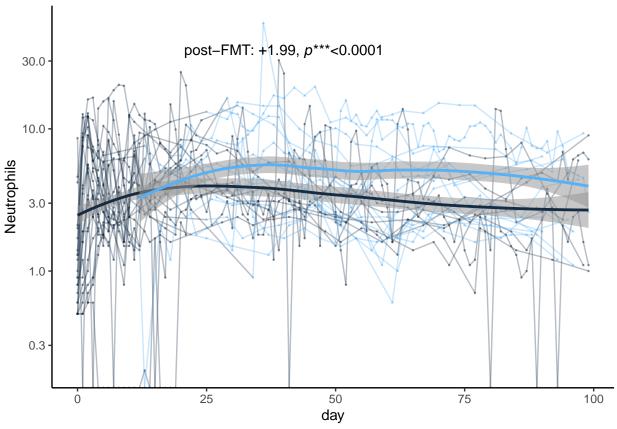
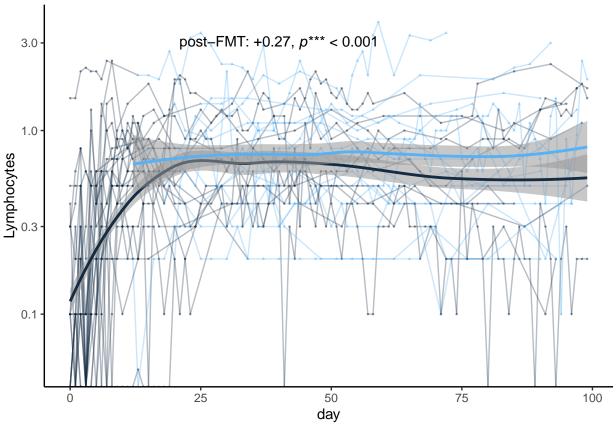
Model of FMT effect on white blood cell counts.

```
library(nlme)
library(dplyr)
library(tidyverse)
library(tidyr)
library(ggplot2)
d <- read.csv('../tidy_WBC_FMT_data_4r.csv', header = TRUE, )</pre>
length(unique(d$pid))
## [1] 24
library(effects)
require(lmerTest)
subsetd <- subset(d, d$wbc%in%c('neutrophils'))</pre>
subsetd <- subset(subsetd, subsetd$day<100)</pre>
# armpost: 1 only iff post randomization and treated with FMT
subsetd$armpost <- subsetd$arm * subsetd$postrandomization</pre>
subsetd$dayfactor <- as.factor(subsetd$day)</pre>
subsetd$patientfactor <- as.factor(subsetd$patientid)</pre>
fitn <- lme(fixed=value~armpost,random=list(~1|dayfactor, ~1|patientfactor),data=subsetd)
fitl <- lm(value~armpost,data=subsetd)</pre>
               numDF denDF F-value p-value
## (Intercept)
                1 820 1183.5514 <.0001
## armpost
                   1 820 44.8281 <.0001
## Approximate 95% confidence intervals
##
##
  Fixed effects:
                  lower
                            est.
                                     upper
## (Intercept) 3.912569 4.252921 4.593273
              1.406547 1.989926 2.573306
## armpost
## attr(,"label")
## [1] "Fixed effects:"
```



```
library(effects)
require(lmerTest)
subsetd <- subset(d, d$wbc%in%c('lymphocytes'))</pre>
subsetd <- subset(subsetd, subsetd$day<100)</pre>
# armpost: 1 only iff post randomization and treated with FMT
subsetd$armpost <- subsetd$arm * subsetd$postrandomization</pre>
subsetd$dayfactor <- as.factor(subsetd$day)</pre>
subsetd$patientfactor <- as.factor(subsetd$patientid)</pre>
fitn <- lme(fixed=value~armpost,random=list(~1|dayfactor, ~1|patientfactor),data=subsetd)
##
               numDF denDF F-value p-value
## (Intercept)
                        820 958.4423 <.0001
## armpost
                    1
                        820 41.4363 <.0001
## Approximate 95% confidence intervals
##
    Fixed effects:
##
##
                    lower
                               est.
                                         upper
## (Intercept) 0.5658711 0.6196076 0.6733440
## armpost
               0.1848796 0.2659868 0.3470939
## attr(,"label")
## [1] "Fixed effects:"
```



```
library(effects)
require(lmerTest)
subsetd <- subset(d, d$wbc%in%c('monocytes'))</pre>
subsetd <- subset(subsetd, subsetd$day<100)</pre>
# armpost: 1 only iff post randomization and treated with FMT
subsetd$armpost <- subsetd$arm * subsetd$postrandomization</pre>
subsetd$dayfactor <- as.factor(subsetd$day)</pre>
subsetd$patientfactor <- as.factor(subsetd$patientid)</pre>
fitn <- lme(fixed=value~armpost,random=list(~1|dayfactor, ~1|patientfactor),data=subsetd)
#summary(fitn)
##
               numDF denDF F-value p-value
## (Intercept)
                        820 434.6799 <.0001
                        820 70.3229 <.0001
## armpost
## Approximate 95% confidence intervals
```

##

##

armpost

attr(,"label")
[1] "Fixed effects:"

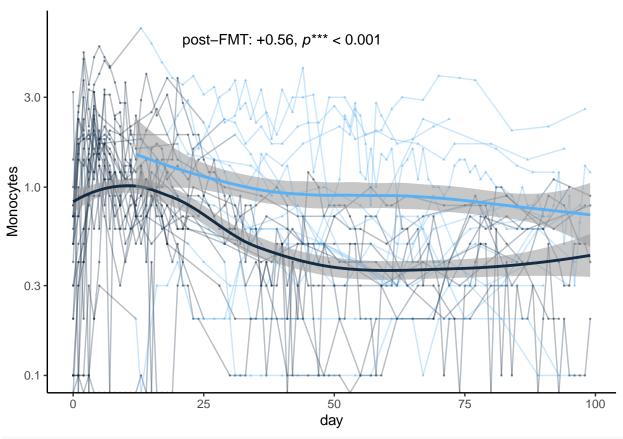
Fixed effects:

lower

(Intercept) 0.6400576 0.7424106 0.8447636

est.

0.4319588 0.5639645 0.6959703



```
library(effects)
require(lmerTest)
subsetd <- subset(d, d$wbc%in%c('wbctotal'))</pre>
subsetd <- subset(subsetd, subsetd$day<100)</pre>
# armpost: 1 only iff post randomization and treated with FMT
subsetd$armpost <- subsetd$arm * subsetd$postrandomization</pre>
subsetd$dayfactor <- as.factor(subsetd$day)</pre>
subsetd$patientfactor <- as.factor(subsetd$patientid)</pre>
fitn <- lme(fixed=value~armpost,random=list(~1|dayfactor, ~1|patientfactor),data=subsetd)
#summary(fitn)
##
               numDF denDF
                              F-value p-value
## (Intercept)
                        820 1562.6573 <.0001
                        820
                              57.0658 <.0001
## armpost
## Approximate 95% confidence intervals
##
##
    Fixed effects:
##
                   lower
                             est.
                                      upper
```

(Intercept) 5.576004 5.989349 6.402694

armpost

attr(,"label")
[1] "Fixed effects:"

1.952193 2.637519 3.322845

?intervals

Help on topic 'intervals' was found in the following packages:

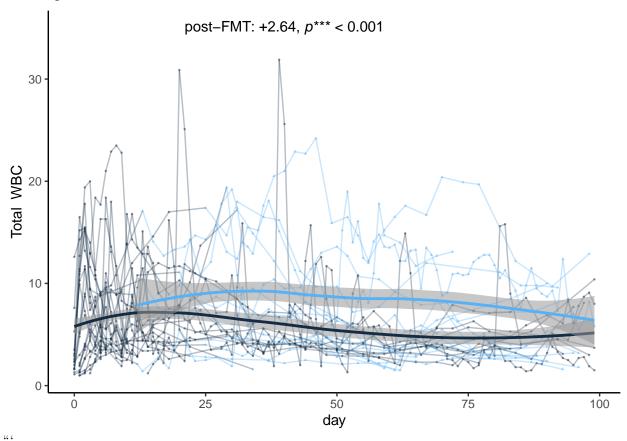
##

Package Library

lubridate /Library/Frameworks/R.framework/Versions/3.6/Resources/library
nlme /Library/Frameworks/R.framework/Versions/3.6/Resources/library

##

Using the first match \dots



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