

```
getwd()

## [1] "C:/Users/naszh/Google Drive (shulai@iu.edu)/IUB/Shiffrin Lab/mix_experiment/analysis"

knitr::opts_knit$set(root.dir = "/home/shulai/Documents/exp1_data0")
knitr::opts_knit$set(root.dir = "C:/Users/naszh/Google Drive (shulai@iu.edu)/IUB/Shiffrin Lab/mix_experiment")

rn=c("Block", "Trial", "Condition", "OldNew", "Setsize", "Serpos", "Lag",
      "Response", "Correctness", "RT", "WhichProbe",
      paste("prob", seq(1,6), sep=""), "surveyq")
# read.table("cmvman1048.txt", head=FALSE, sep="", fill = TRUE, col.names = rn)
```

Basic

```
subnum=16+5
basen="cmvmspec1_"
```

Functions

load data

```
ld=function(basen, subnum){
  for (i in 1:subnum){
    if (i==1){
      d=read.table(paste(basen,i, ".txt", sep=""), head=FALSE, sep="", fill = TRUE, col.names = rn)
      d=d %>% mutate(Subnum=i) #>% replace_na(list(surveyq=surveyq[length(surveyq)]))
    } else{
      d0=read.table(paste(basen,i, ".txt", sep=""), head=FALSE, sep="", fill = TRUE, col.names = rn)
      d0=d0 %>% mutate(Subnum=i) #>% replace_na(list(surveyq=surveyq[length(surveyq)]))
      d=rbind(d, d0)
    }
    d=d%>% select(Subnum, everything()) %>% arrange(Subnum) %>% filter(RT>0)
  }
  return(d)
}
```

plot RT

```
plot_rtall=function(d, tit){
  dase=summarySEwithin(data=d, measurevar = "RT", withinvars = c("Setsize", "Correctness", "OldNew", "Condition"),
    # print(dase)
    dalse=subset(dase, Correctness==1)
  p1=ggplot(data=dalse, aes(Setsize, RT))+
    geom_point(aes(color=as.factor(OldNew), shape=as.factor(Condition), group=OldNew), size=5)+
    geom_line(aes(color=as.factor(OldNew), linetype=as.factor(Condition),
      group=interaction(OldNew, Condition)))+
    scale_color_manual(name="Old-New", #breaks=c(1,0),
      labels=c(`1`="Old", `2`="New"),
      values=c("#F23005", "#FFBE73"))+
    scale_shape_discrete(name="Condition",
      labels = c(`1`="CMpos_s", `2`="CMpos_reg", `3`="CMneg_h", `4`="CMneg_l", `5`="vm"))+
    # theme_minimal()
  print(p1)
}
```

```

scale_linetype_discrete(name="Condition",
                        labels = c(`1`="CMpos_s",`2`="CMpos_reg",`3`="CMneg_h",`4`="CMneg_l",`5`="vm"))+
ggtitle(paste("Mean CorrectRT - Set Size",tit))
return(p1)
}

```

plot Correct

```

plot_error=function(d,tit){
  da_p=summarySEwithin(data=d,measurevar = "Correctness",withinvars = c("Setsize","OldNew","Condition"))
  # da_p
  p1=ggplot(data=da_p,aes(Setsize,1-Correctness))+
    geom_point(aes(color=as.factor(OldNew),shape=as.factor(Condition),group=OldNew),size=5)+
    geom_line(aes(color=as.factor(OldNew),linetype=as.factor(Condition),
                  group=interaction(OldNew,Condition)))+
    scale_color_manual(name="Old-New",#breaks=c(1,0),
                      labels=c(`1`="Old", `2`="New"),
                      values=c("#F23005", "#FFBE73"))+
    scale_shape_discrete(name="Condition",
                        labels = c(`1`="CMpos_s",`2`="CMpos_reg",`3`="CMneg_h",`4`="CMneg_l",`5`="vm"))+
    scale_linetype_discrete(name="Condition",
                          labels = c(`1`="CMpos_s",`2`="CMpos_reg",`3`="CMneg_h",`4`="CMneg_l",`5`="vm"))+
    ggtitle(paste("Error Rate- Set Size - ",tit))+
    scale_y_continuous("Probability of Error")
  return(p1)
}

```

mutate new condition

```

creatnewc=function(dc){
  d0=dc %>% mutate(Condition=as.numeric(Condition)) %>%
  mutate(specondi=case_when( Condition %in% as.factor(1:4)~ Condition,
                             Condition==5&OldNew==1 ~ 5, #c5=vm-old
                             Condition==5&OldNew==2 ~ 6) ) %>% #6~vm-new
  mutate(specondi=as.factor(specondi))
  return(d0)
}

```

plot rt blockwise

```

plot_rtbblockwise = function(d,tit){
  # d=dc
  dp3=creatnewc(d)%>% filter(Correctness==1)
  dp3=summarySEwithin(data=dp3, measurevar = "RT", withinvars = c("Block","Setsize","specondi"))

  Condition_n=c("CMpos_s","CMpos_reg","CMneg_h","CMneg_l","vm_old","vm_new");names(Condition_n)=as.character(1:6)

  p=ggplot(dp3,aes(Block,RT))+
    geom_point(aes(color=Setsize))+
    geom_line(aes(color=Setsize,group=interaction(Setsize)))+
    ggtitle(paste("RT accross blocks -"),tit)+
    facet_wrap(specondi~.,

```

```

        labeller = labeller(specondi=Condition_n) ,ncol=3)
        # labeller = label_both)
    return(p)

}

plot_rtbblockwise2=function(d,tit){
  dp3=creatnewc(d) %>% filter(Correctness==1)
  dp34=summarySEwithin(data=dp3, measurevar = "RT", withinvars =
                        c("Block", "specondi"))
  Condition_n=c("CMpos_s", "CMpos_reg", "CMneg_h", "CMneg_l", "vm_old", "vm_new");names(Condition_n)=as.ch
  p=ggplot(dp34,aes(Block,RT))+
    geom_point(aes(color=specondi))+
    geom_line(aes(color=specondi,
                  group=specondi))+
    scale_color_discrete(labels = Condition_n )+
    ggtitle(paste("RT accross blocks"),tit)
  return(p)
}

```

participant wise

```

plot_rtbblockpar=function(d,tit){
  dp3=creatnewc(dc) %>% filter(Correctness==1)
  dp34=summarySEwithin(data=dp3, measurevar = "RT", withinvars =
                        c("Block", "specondi", "Subnum"))
  Condition_n=c("CMpos_s", "CMpos_reg", "CMneg_h", "CMneg_l", "vm_old", "vm_new");names(Condition_n)=as.ch
  p=ggplot(dp34,aes(Block,RT))+
    geom_point(aes(color=specondi))+
    geom_line(aes(color=specondi,
                  group=specondi))+
    scale_color_discrete(labels = Condition_n )+
    ggtitle(paste("RT accross blocks across participant"),tit)+
    facet_wrap(Subnum~.,ncol=3)
  return(p)
}

```

load data

```

d=ld(basen,subnum)
d=d%>%mutate_at(c("Setsize", "OldNew", "Condition"),as.factor)

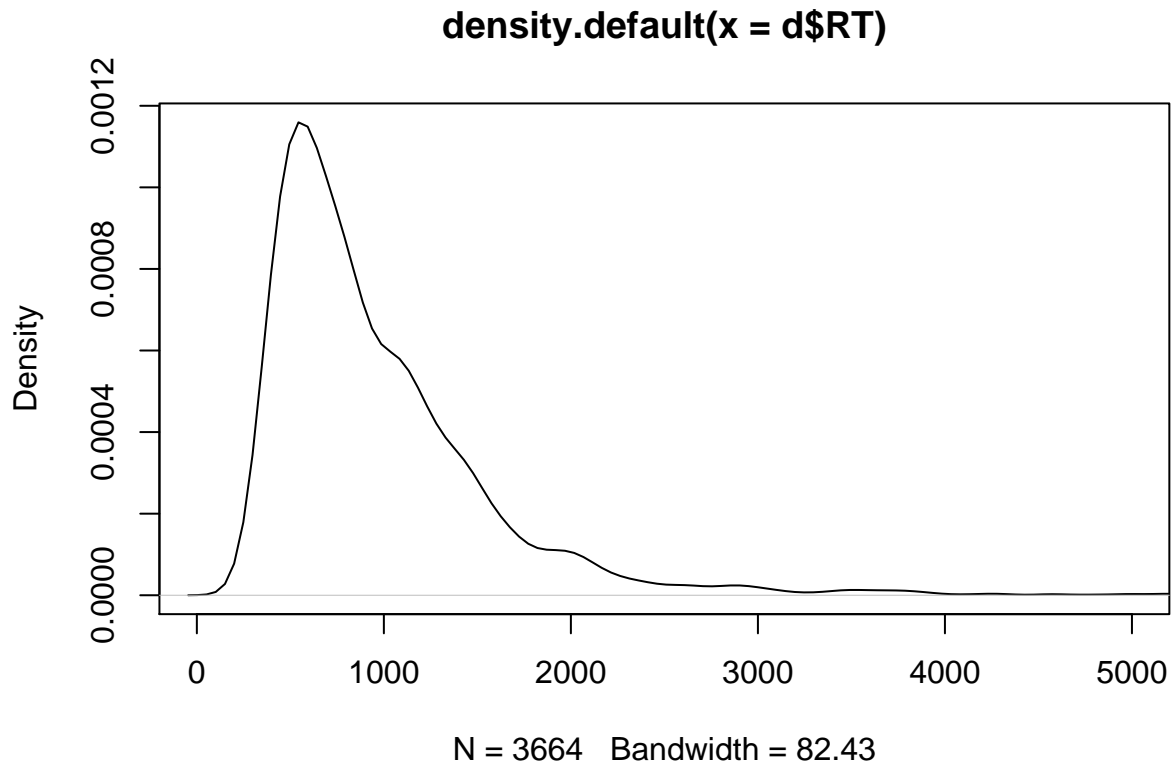
```

clean data

```

plot(density(d$RT),xlim=c(0,5000))

```



```
# d$Trial[d$RT>4000]
#
# d[d$Trial<10,]

dc=d%>%filter(RT<5000&RT!=0)
```

Manipulation check

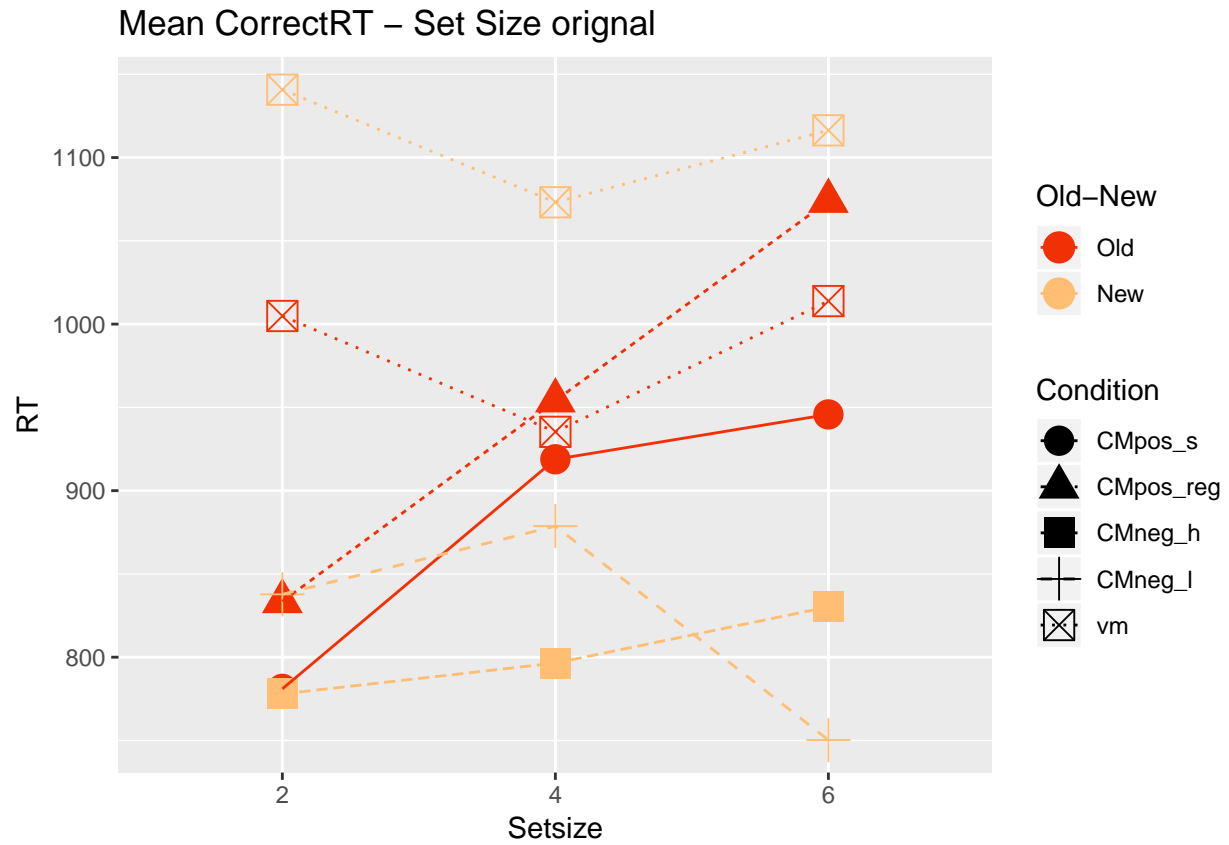
```
a=array()
for(i in 1:10){
  a[i]=sum(d$WhichProbe==i)/sum(length(d$Block))
}
a

## [1] 0.10725983 0.09989083 0.14546943 0.06741266 0.09579694 0.09497817
## [7] 0.09497817 0.10343886 0.10289301 0.08788210
```

Plor rt for correct response

```
plot_rtall(d,"original")

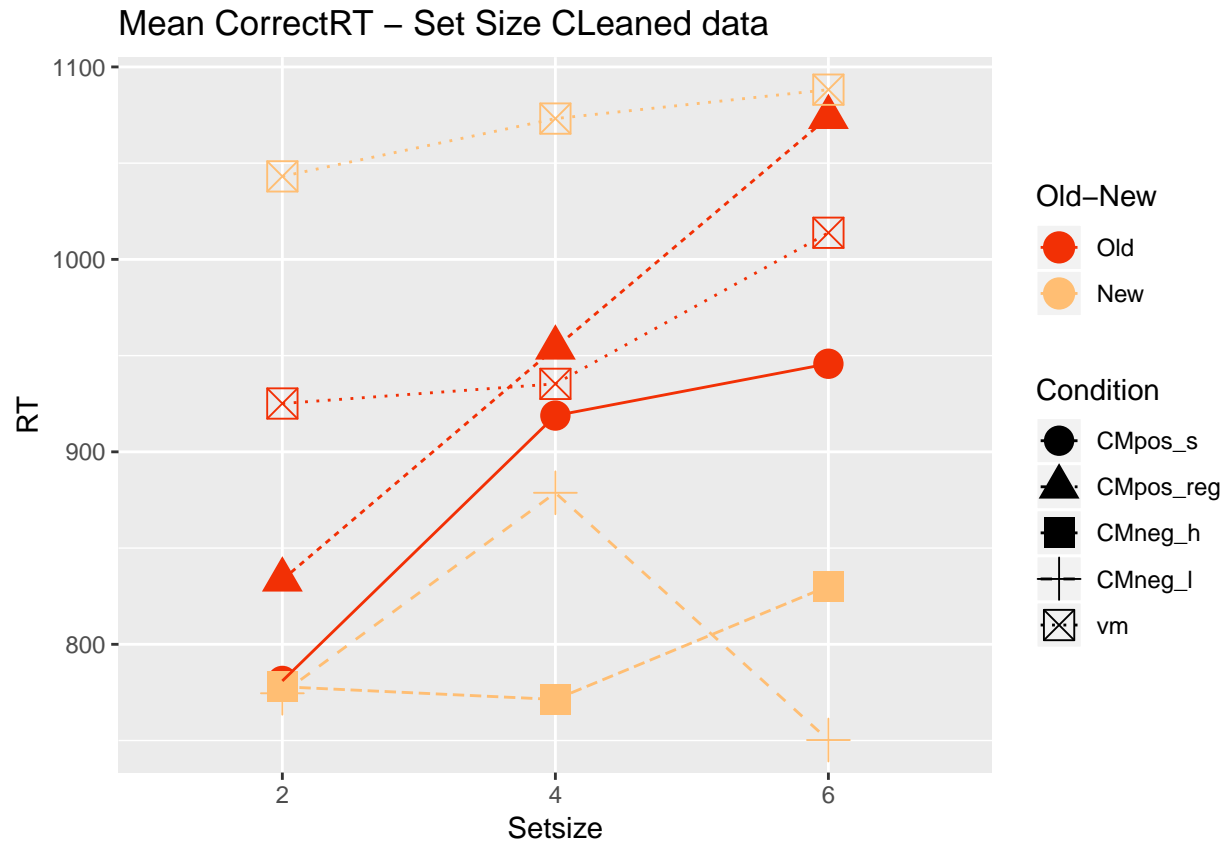
## Automatically converting the following non-factors to factors: Correctness
## Warning in qt(conf.interval/2 + 0.5, datac$N - 1): NaNs produced
```



```
plot_rtall(dc, "Cleaned data")
```

```
## Automatically converting the following non-factors to factors: Correctness
```

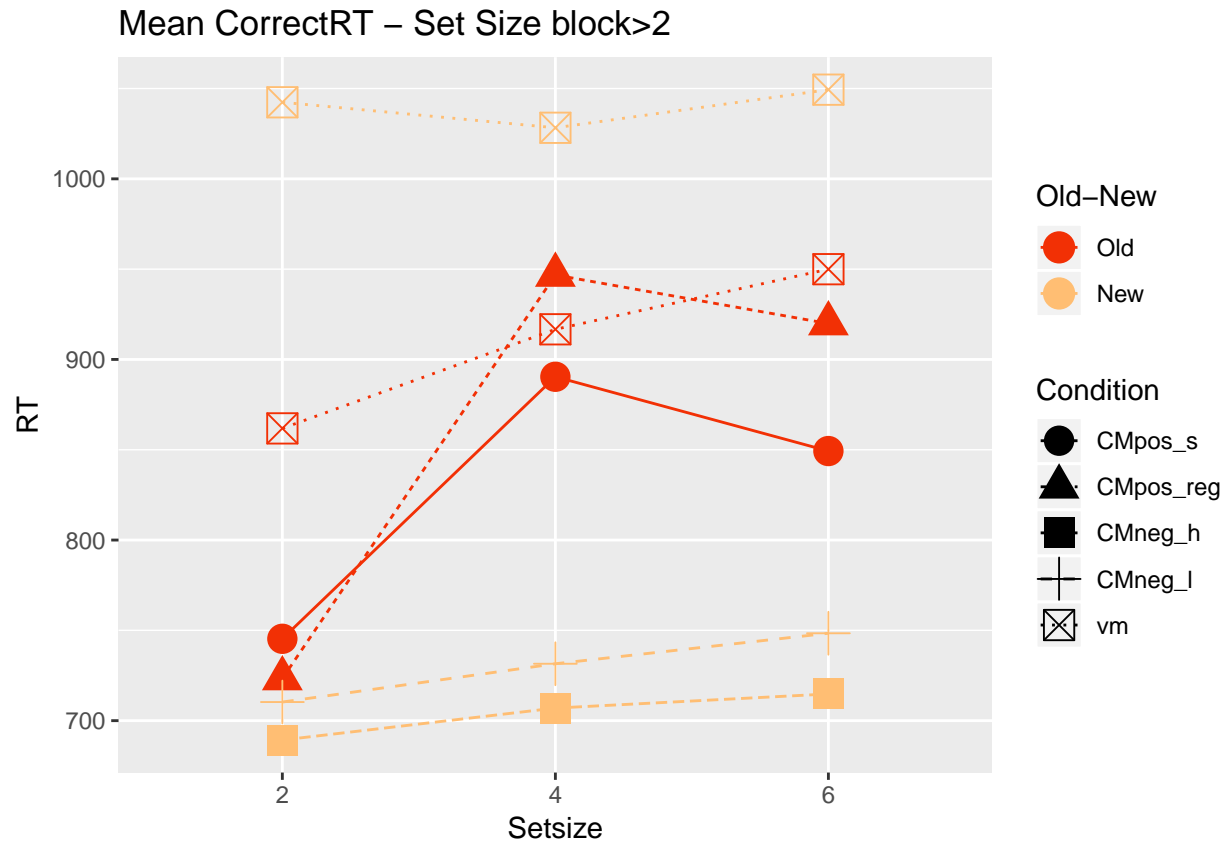
```
## Warning in qt(conf.interval/2 + 0.5, datac$N - 1): NaNs produced
```



```
plot_rtall(filter(dc,Block>2),"block>2")
```

```
## Automatically converting the following non-factors to factors: Correctness
```

```
## Warning in qt(conf.interval/2 + 0.5, datac$N - 1): NaNs produced
```



VM-New is weird, let's see what if block>2

frequent inference

Want: 1-2 cmpos vs cm neg 1-5 cmpos vs vm pos 2-5 cmneg vs vm pos

#make new condition variable for every conditions

```
dcmut=creatnewc(dc) %>% filter(Correctness==1)
```

Want: 1-2 cmpos vs cm neg

1-5 cmpos vs vm pos

2-5 cmneg vs vm pos

```
TukeyHSD(data=dcmut,aov(RT~Condition,data=dc))
```

```
## Tukey multiple comparisons of means
```

```
## 95% family-wise confidence level
```

```
##
```

```
## Fit: aov(formula = RT ~ Condition, data = dc)
```

```
##
```

```
## $Condition
```

```
## diff lwr upr p adj
```

```
## 2-1 54.2289868 -58.63958 167.097557 0.6842350
```

```
## 3-1 -112.4581922 -215.80754 -9.108847 0.0249903
```

```
## 4-1 -113.0883211 -239.56777 13.391131 0.1049962
```

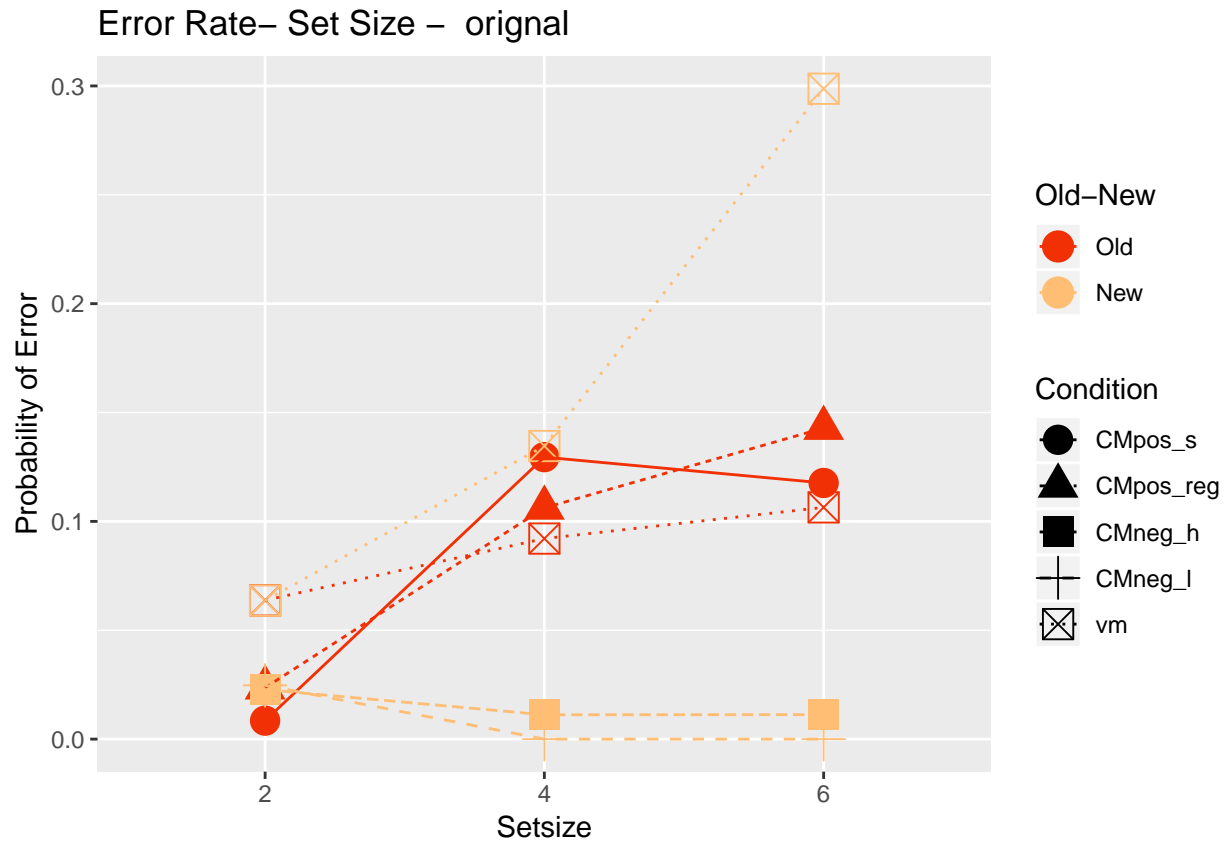
```
## 5-1 107.8900971 22.52787 193.252321 0.0051475
```

```
## 3-2 -166.6871790 -272.20620 -61.168156 0.0001621
```

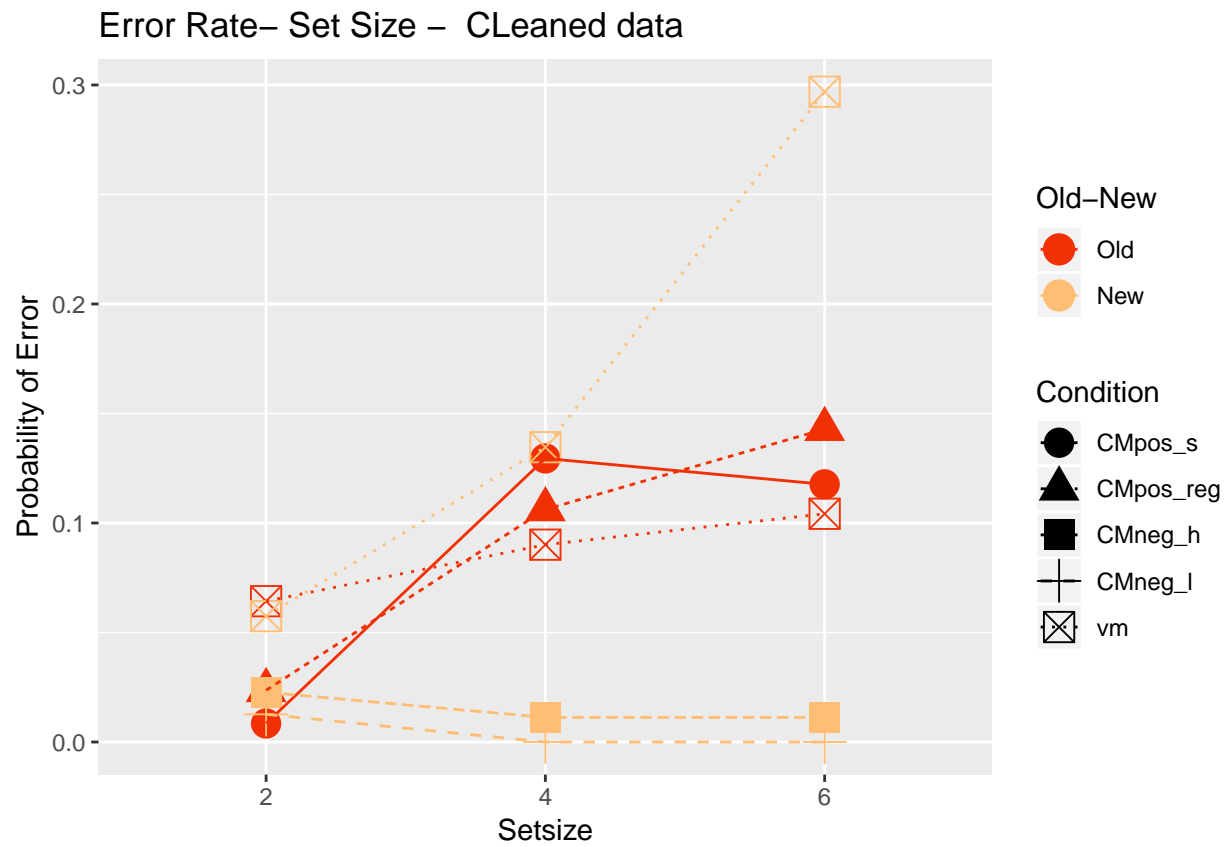
```
## 4-2 -167.3173079 -295.57575 -39.058862 0.0034464
## 5-2 53.6611103 -34.31551 141.637735 0.4561557
## 4-3 -0.6301289 -120.59680 119.336541 1.0000000
## 5-3 220.3482893 144.97151 295.725073 0.0000000
## 5-4 220.9784182 116.10886 325.847976 0.0000001
```

plot correctness

```
plot_error(d,"original")
```

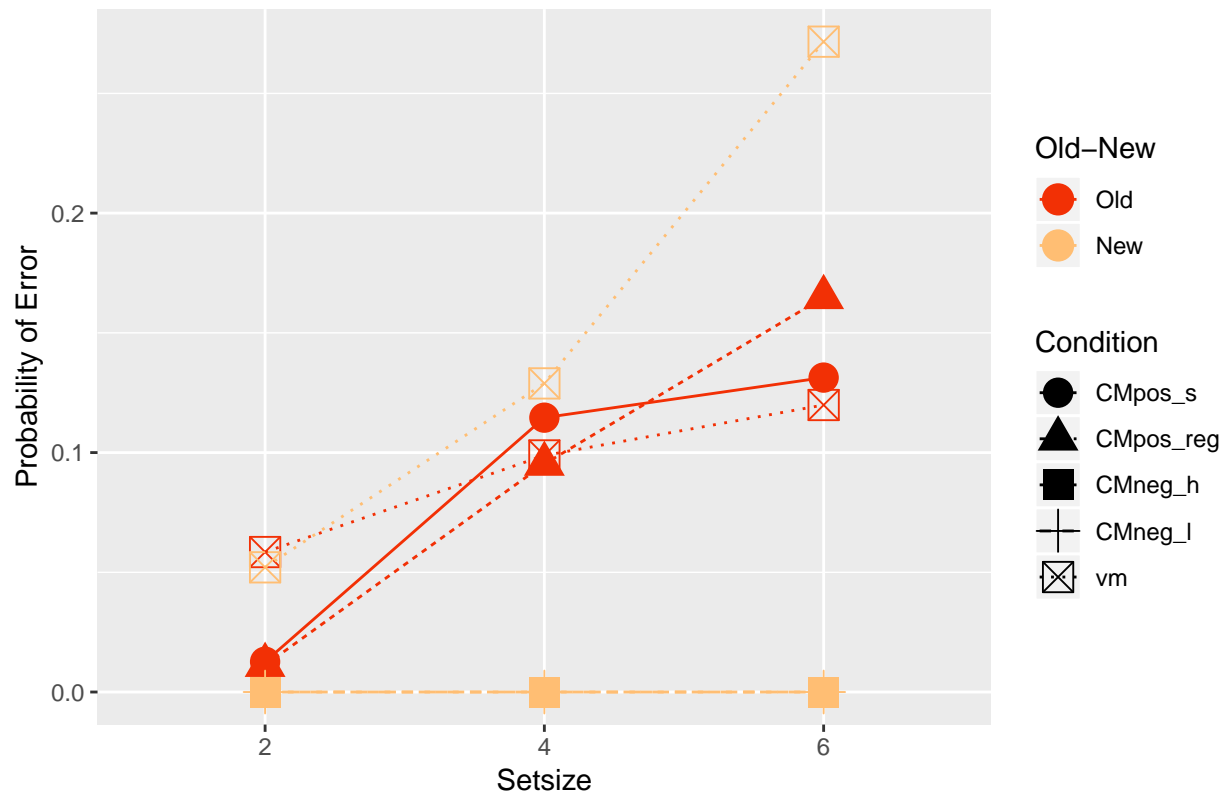


```
plot_error(dc,"CLEaned data")
```

```
plot_error(filter(dc,Block>2),"block>2")
```

Error Rate– Set Size – block>2



##freq Want: 1-2 cmpos vs cm neg
 1-5 cmpos vs vm pos
 2-5 cmneg vs vm pos

```
TukeyHSD(data=dcmut,aov(Correctness~Condition,data=dc))
```

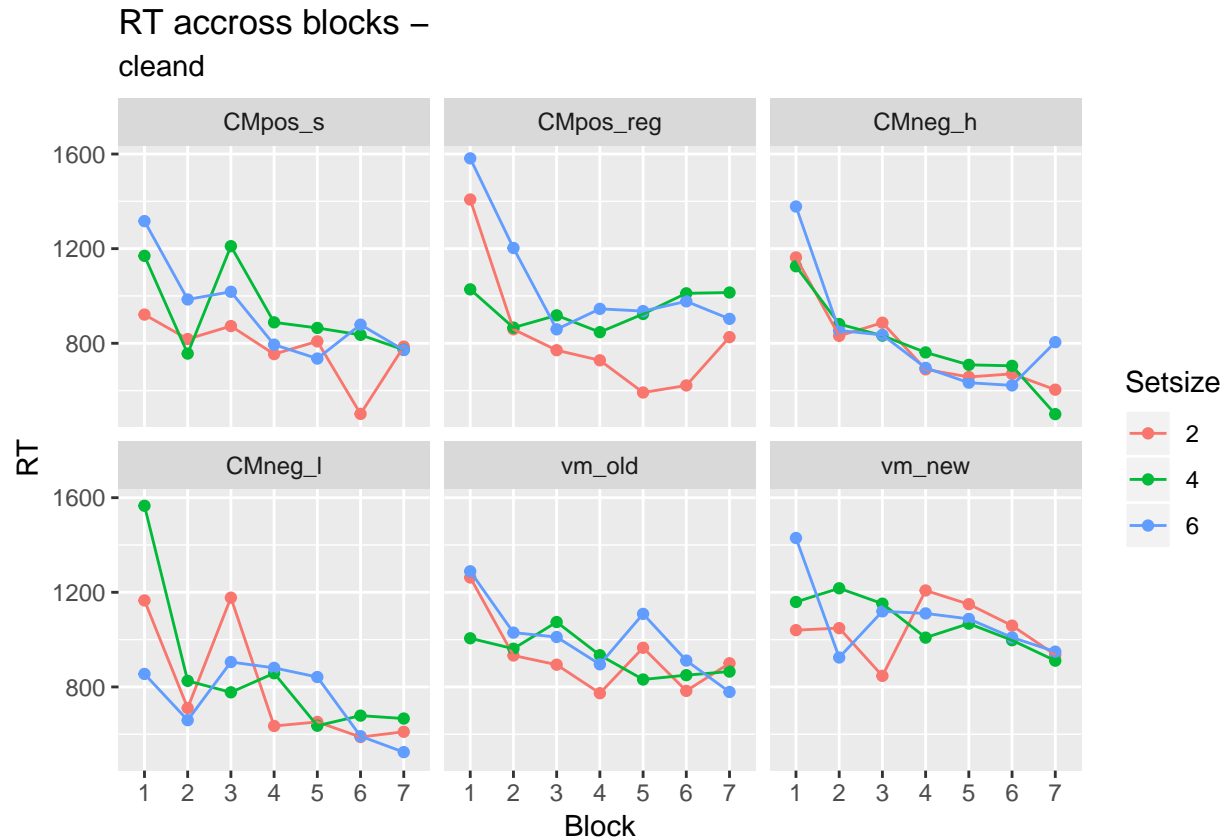
```
## Tukey multiple comparisons of means
## 95% family-wise confidence level
##
## Fit: aov(formula = Correctness ~ Condition, data = dc)
##
## $Condition
##      diff      lwr      upr    p adj
## 2-1 -0.00110541 -0.05618985 0.05397903 0.9999980
## 3-1  0.07402093  0.02358226 0.12445960 0.0006044
## 4-1  0.08497689  0.02324979 0.14670399 0.0016368
## 5-1 -0.02321054 -0.06487077 0.01844968 0.5491046
## 3-2  0.07512634  0.02362878 0.12662390 0.0006668
## 4-2  0.08608230  0.02348698 0.14867762 0.0016619
## 5-2 -0.02210513 -0.06504129 0.02083103 0.6244228
## 4-3  0.01095596 -0.04759264 0.06950456 0.9863335
## 5-3 -0.09723147 -0.13401840 -0.06044455 0.0000000
## 5-4 -0.10818743 -0.15936803 -0.05700684 0.0000001
```

plot RT- Blockwise

```
# head(d)
```

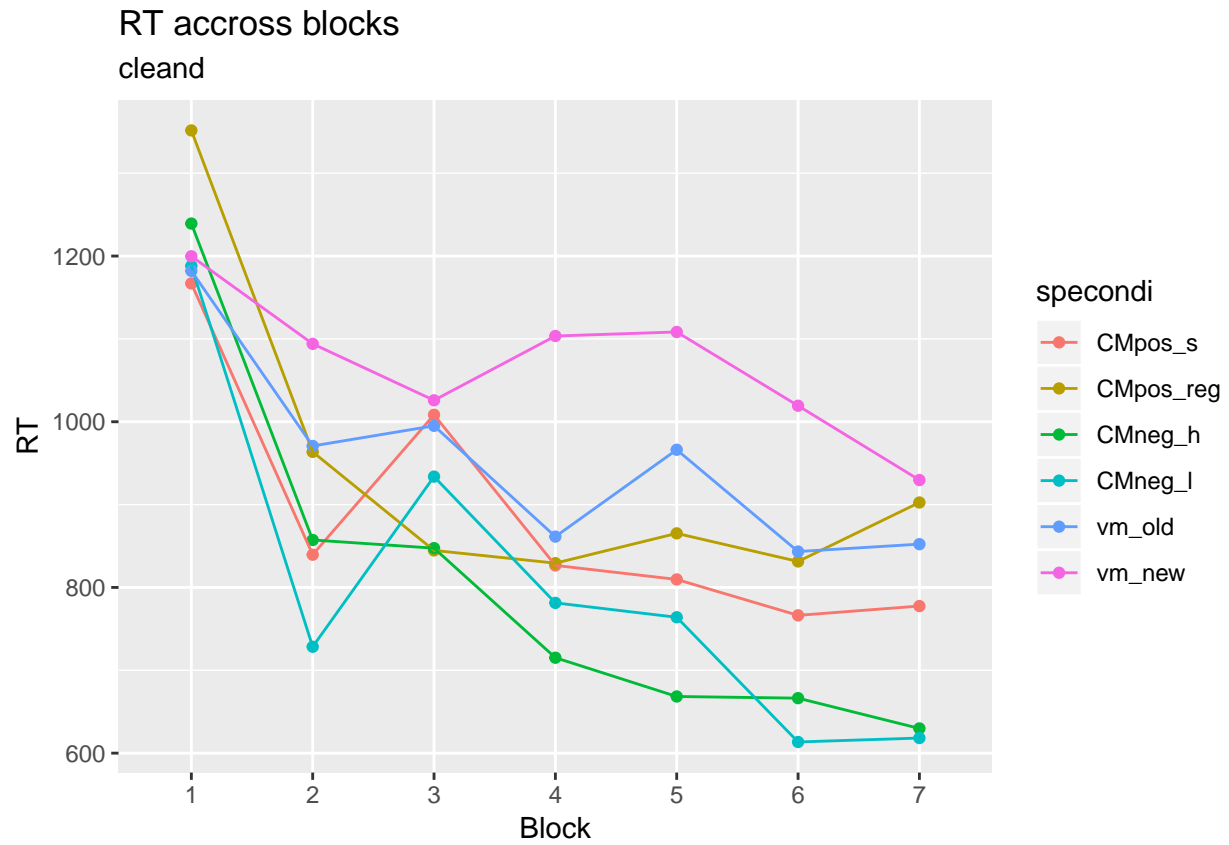
```
plot_rtblockwise(dc,"cleand")
```

```
## Automatically converting the following non-factors to factors: Block
```



```
plot_rtblockwise2(dc,"cleand")
```

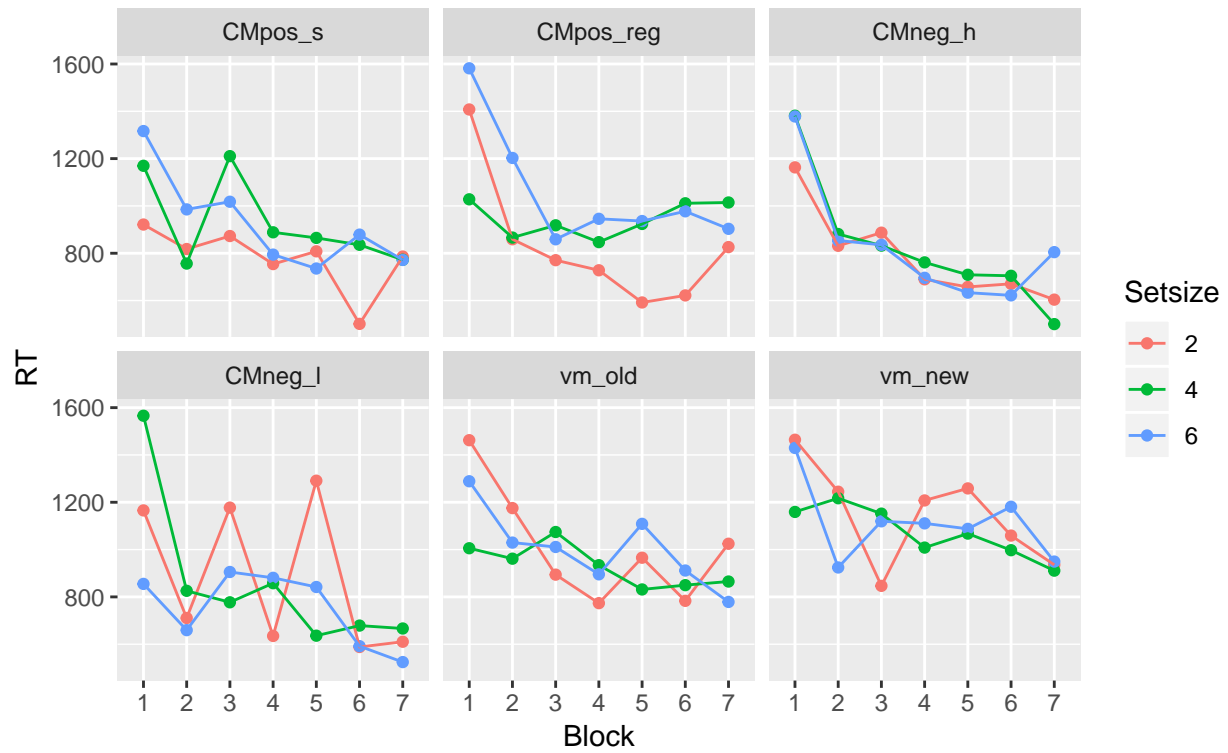
```
## Automatically converting the following non-factors to factors: Block
```



```
plot_rtblockwise(d,"orgid")
```

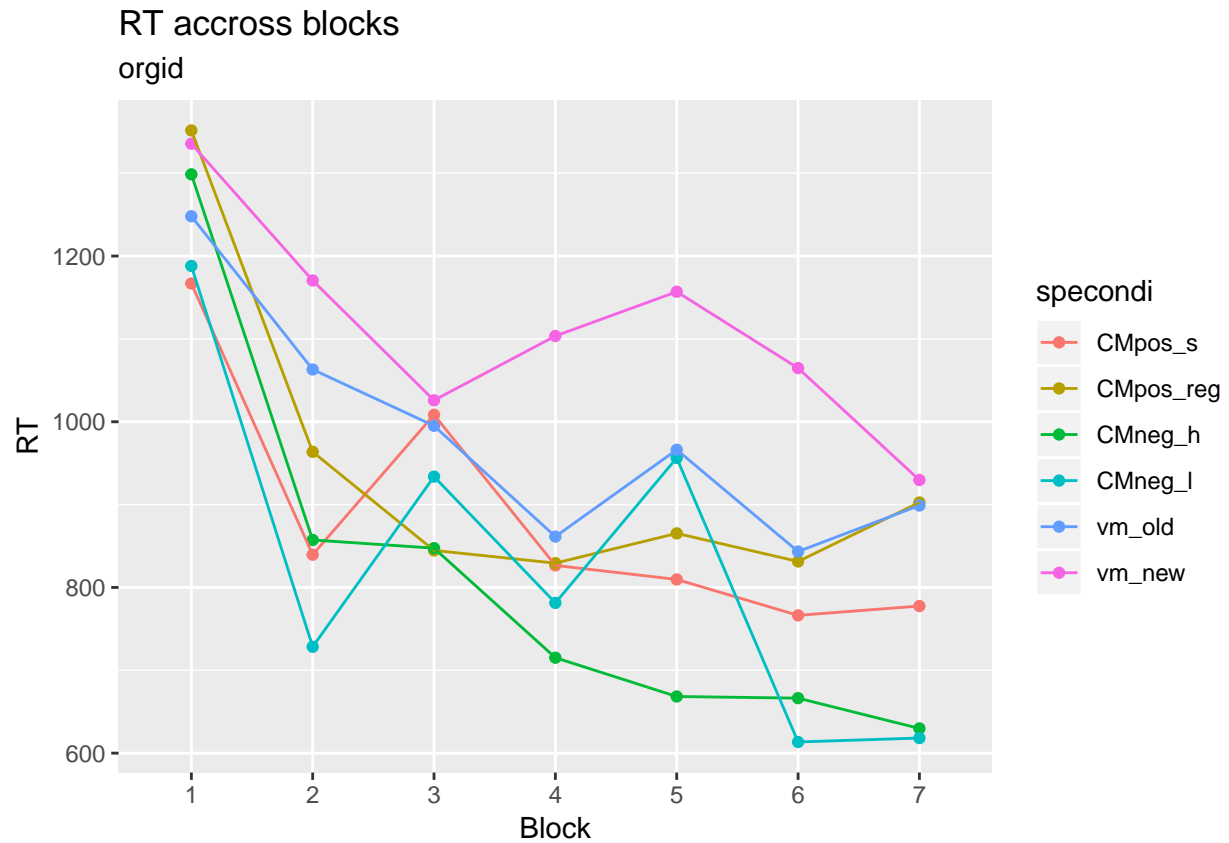
```
## Automatically converting the following non-factors to factors: Block
```

RT accross blocks – orgid



```
plot_rtbblockwise2(d,"orgid")
```

```
## Automatically converting the following non-factors to factors: Block
```



p-participant wise

```
plot_rtblockpar(dc, "cleand")
```

```
## Automatically converting the following non-factors to factors: Block, Subnum
```

```
## Warning in qt(conf.interval/2 + 0.5, datac$N - 1): NaNs produced
```

RT accross blocks across participant
cleand

