

# Wrap up

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## Prepare data

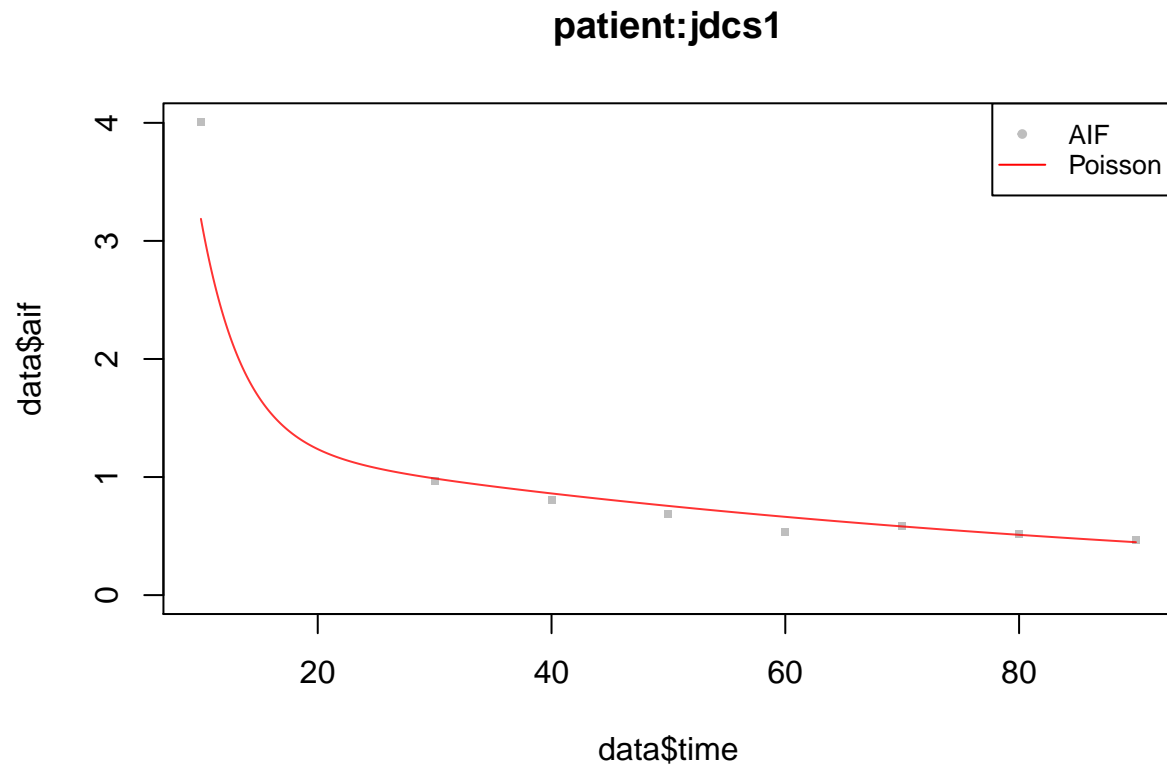
```
# load data
data = readdata()
# find tmax and slice the curve
data = data %>%
  group_by(patient) %>%
  mutate(count = map(count, ~findtmax(.x)),
         count_asc = map(count, ~slice_asc(.x)), # data before tmax
         count_dsc = map(count, ~slice_exp(.x))) # data after tmax
```

## Fit model

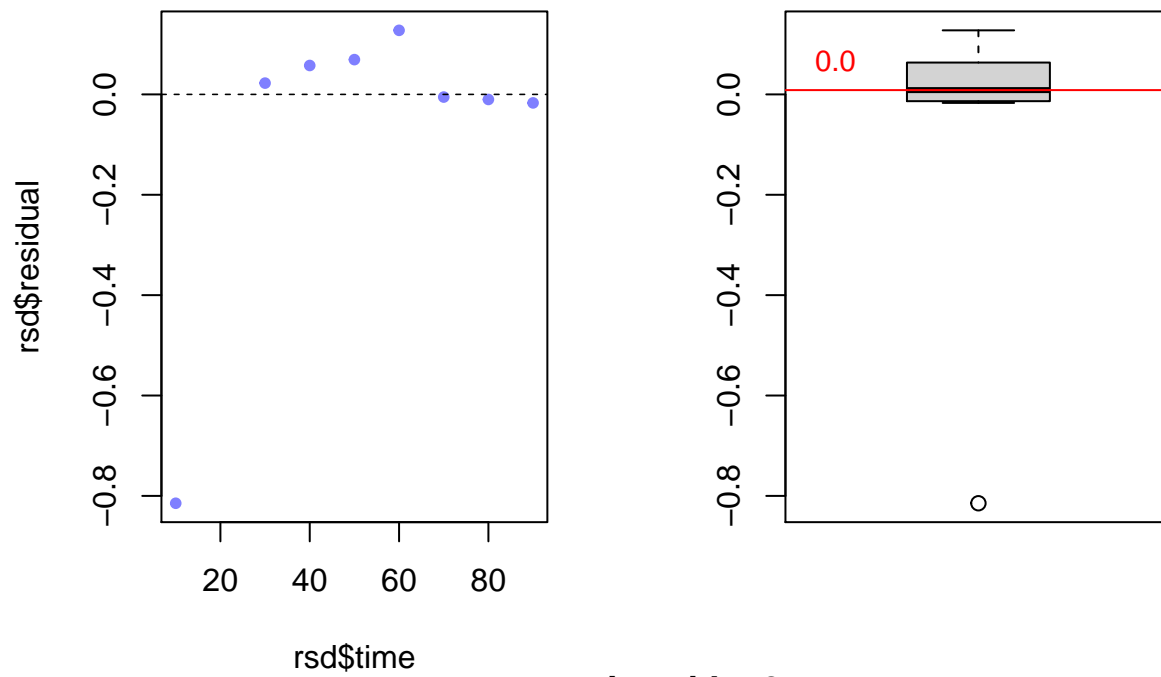
```
data = data %>%
  group_by(patient) %>%
  mutate(asc_res = map(count_asc, ~acs_inter(.x)), # detect t0 and interpolate aif between t0 and tmax
         count_asc = map(asc_res, ~.x$data), # add t0 to the data
         asc_mod = map(asc_res, ~.x$segmod), # save model that detecting t0
         dsc_mod = map(count_dsc, # fit nonlinear poisson regression for descending part
                       ~kinfitr_gnm(t = .x$time, # time since tmax
                                    t_G = .x$t_G, # time point put in gamma count since injection time
                                    y.sum = .x$count,
                                    delta = .x$delta, # time in the gamma counter
                                    vol = .x$vol,
                                    pf = .x$parentFraction,
                                    bpr = .x$bpr,
                                    disp = rep(1,nrow(.x))
                                   ),
         dsc_mod = map(dsc_mod, ~.x$result), # save model fit data after tmax
         asc_pred = map(asc_res, ~.x$pred), # get prediction before tmax
         # get prediction after tmax, contain interpolated aif
         dsc_pred = map2(dsc_mod, count_dsc, ~pred_aif(.x, .y)),
         pred = map2(asc_pred, dsc_pred, ~rbind(.x, y$rsd[-1,])), # combine predicted aif
         rsd = map2(count, pred, ~add_residual(.x, .y)) # calculate residuals
         ) %>%
  select(-asc_res)
```

plot

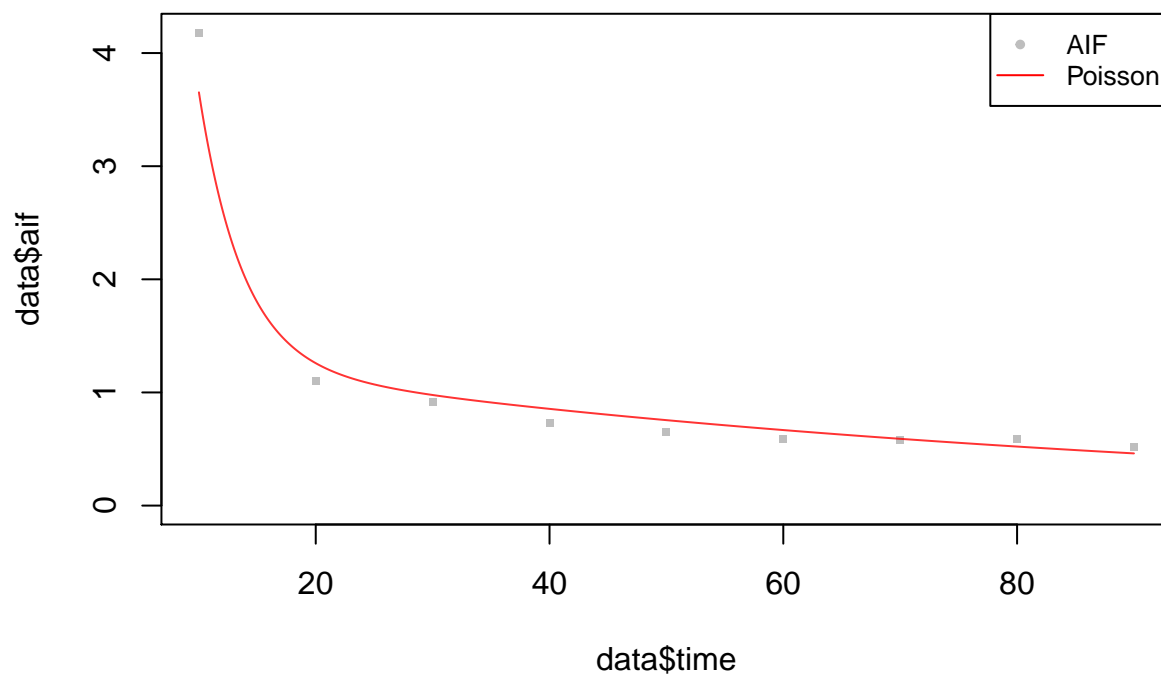
```
for (i in 1:nrow(data)){  
  par(mfrow=c(1,1))  
  # change range of time by using 2nd and 3rd arguments  
  scope_whole(data[i,],10,90) # plot aif and predicted aif  
  plot_rsd(data[i,],10,90) # plot residuals and boxplot with median  
}
```



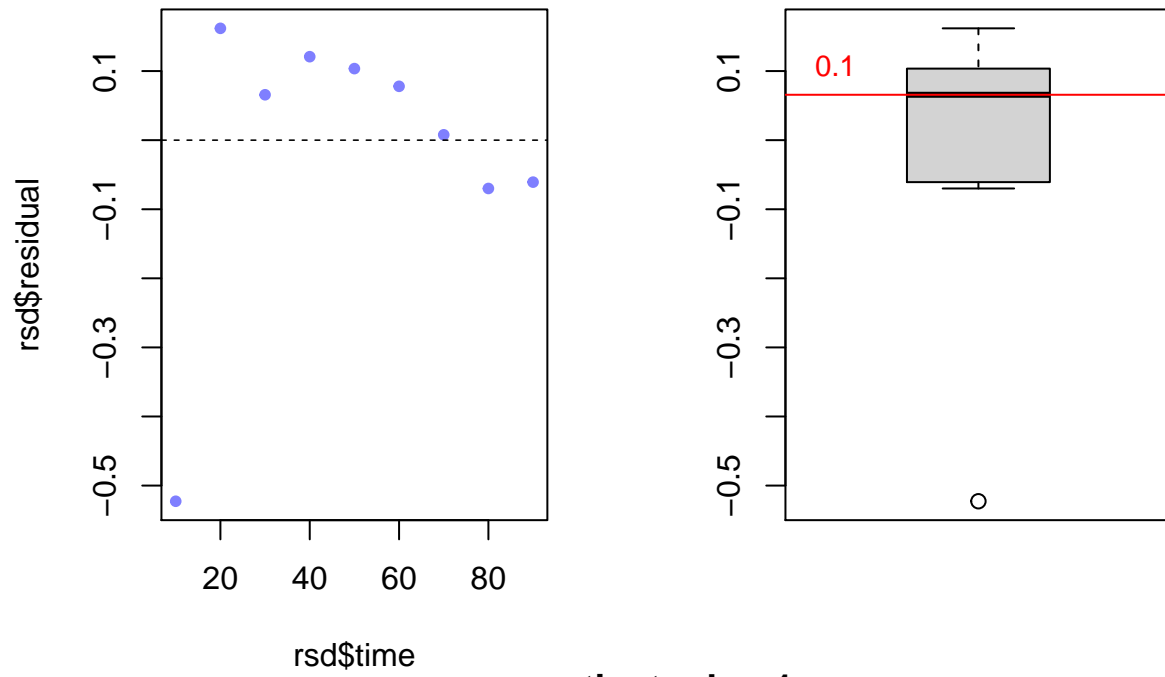
**patient:jdc1**



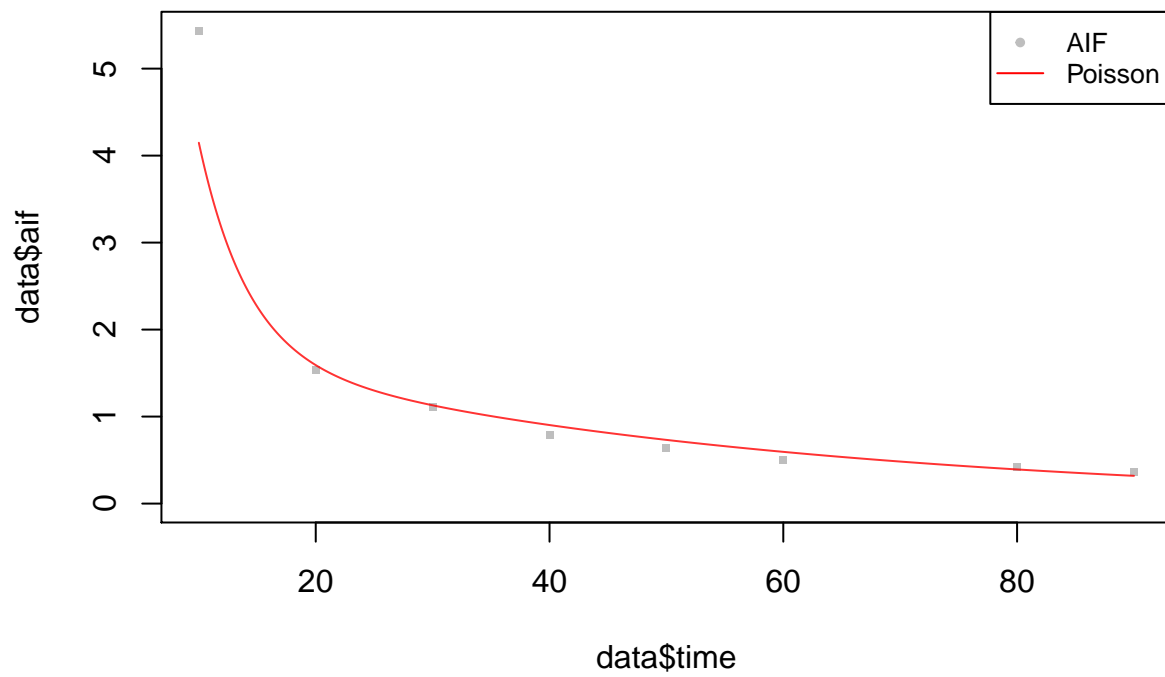
**patient:jdc2**



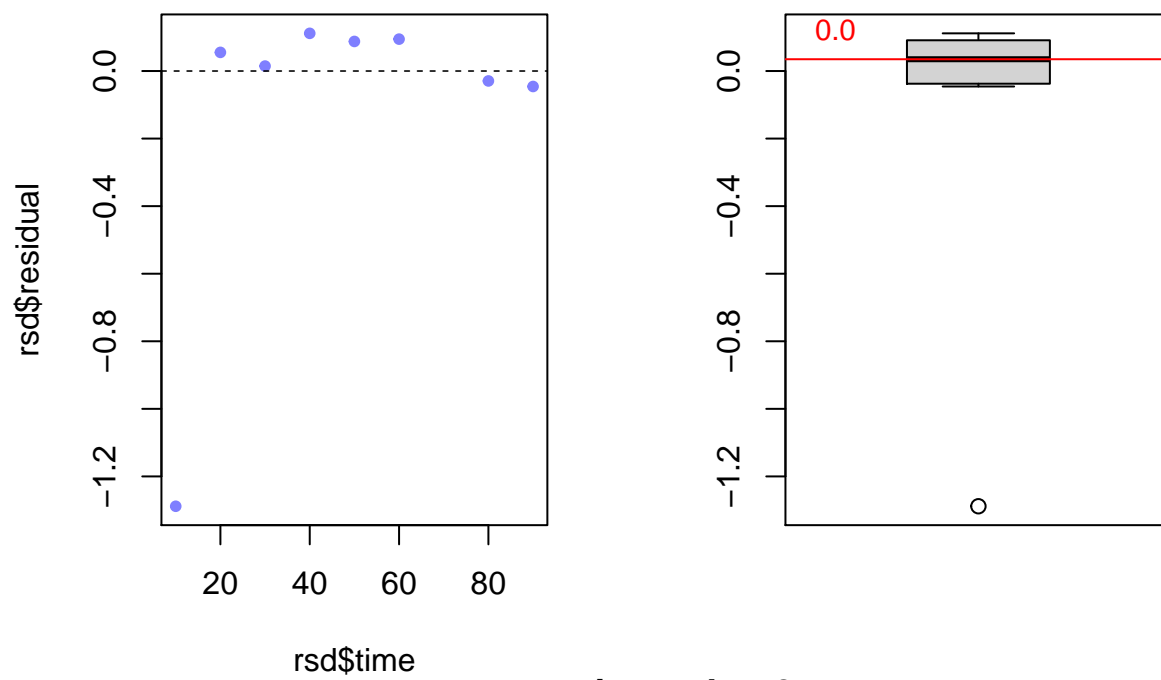
**patient:jdcs2**



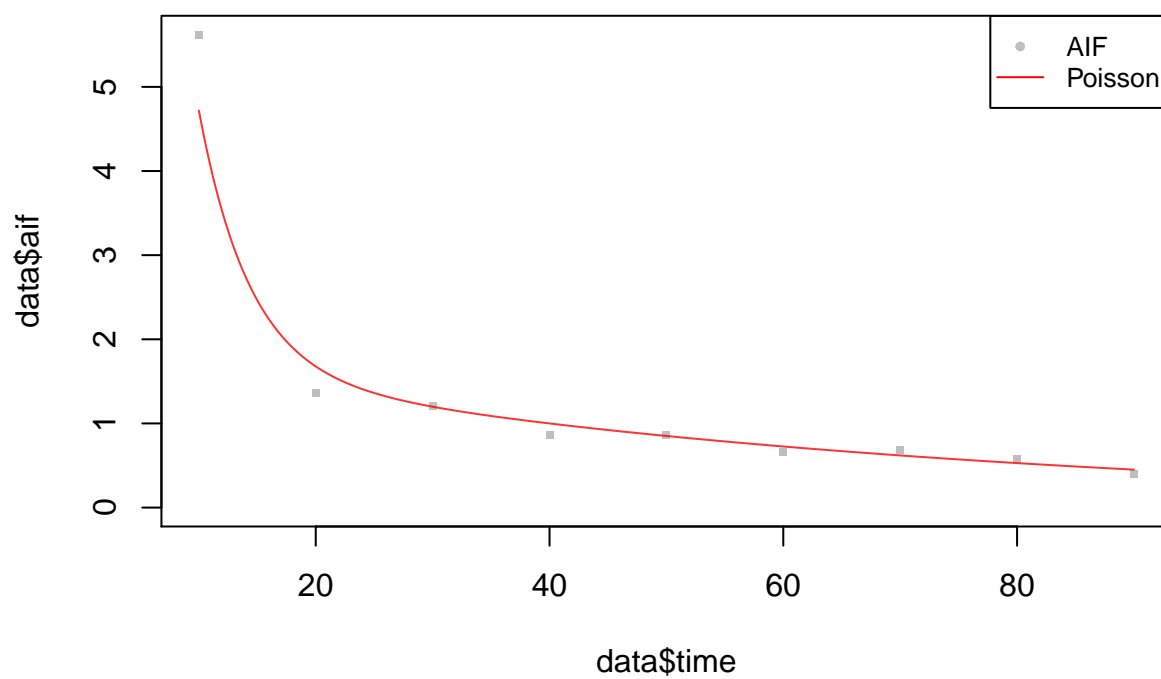
**patient:mhco1**



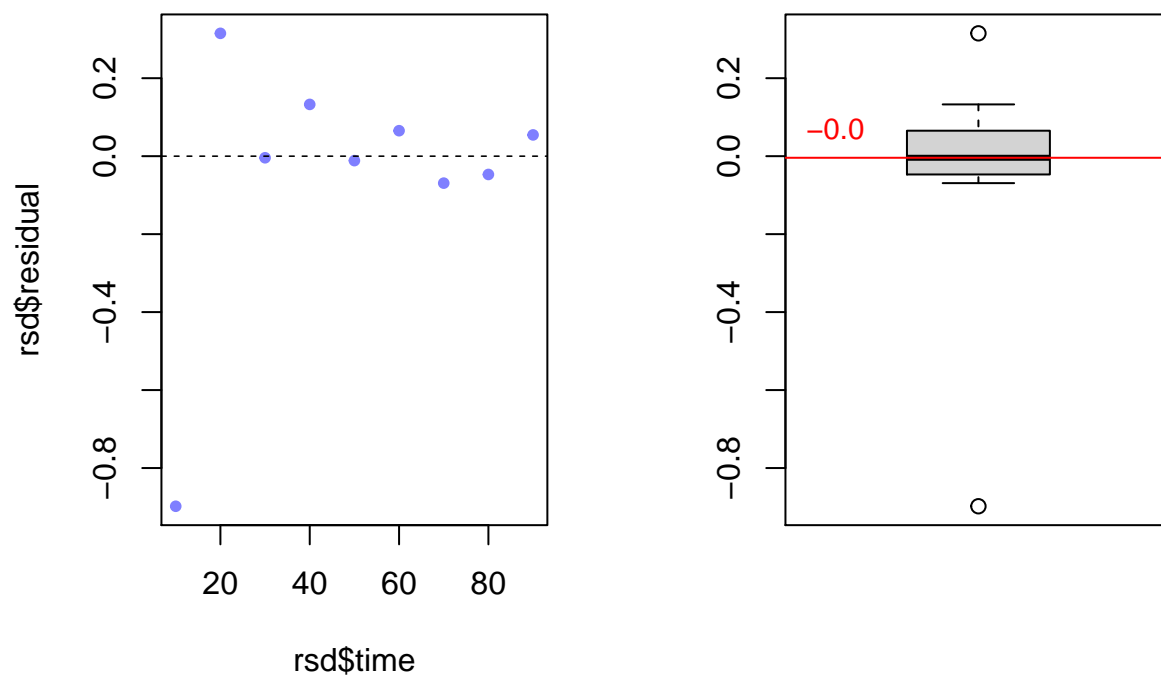
**patient:mhco1**



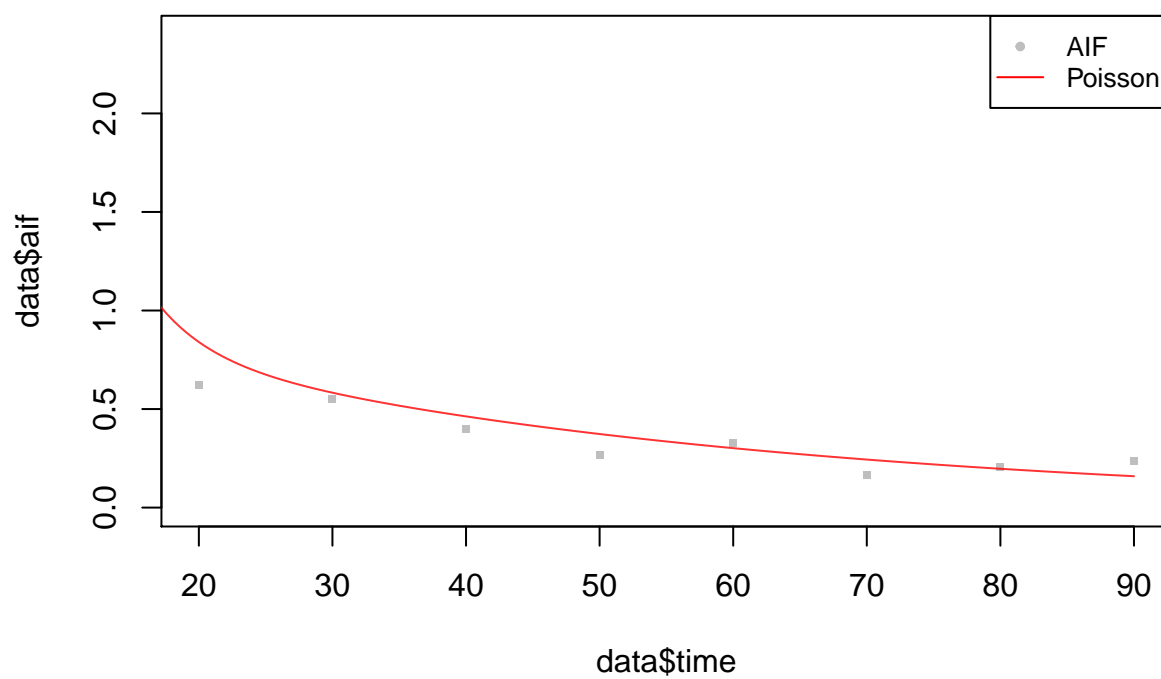
**patient:mhco2**



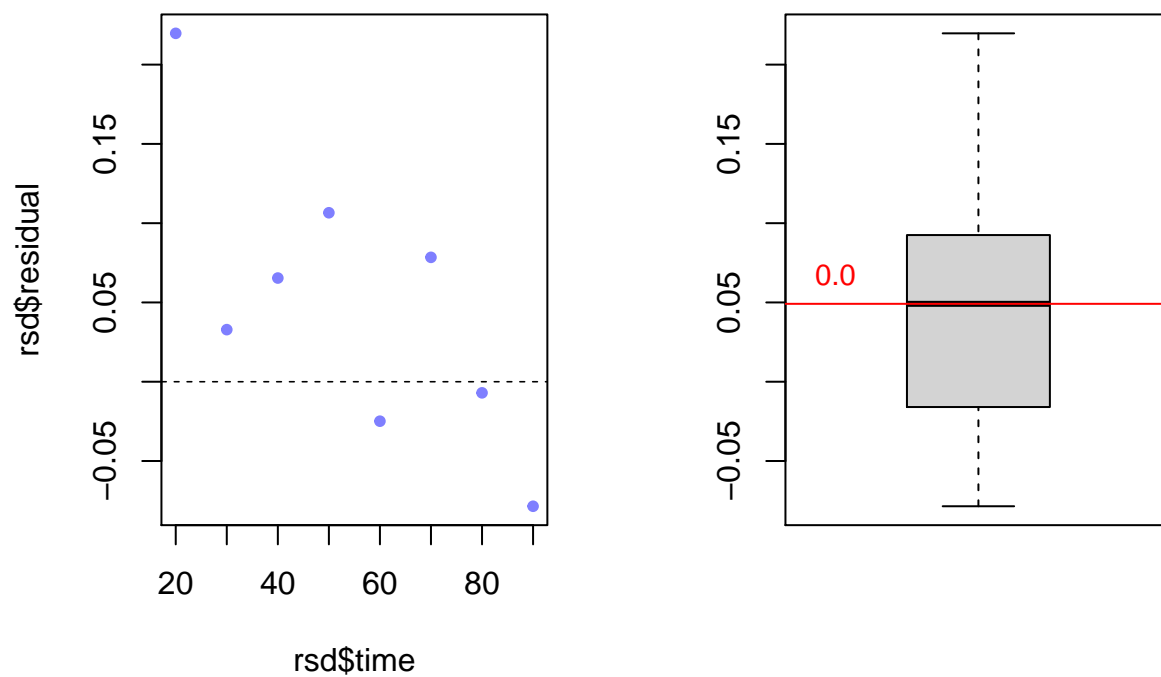
**patient:mhco2**



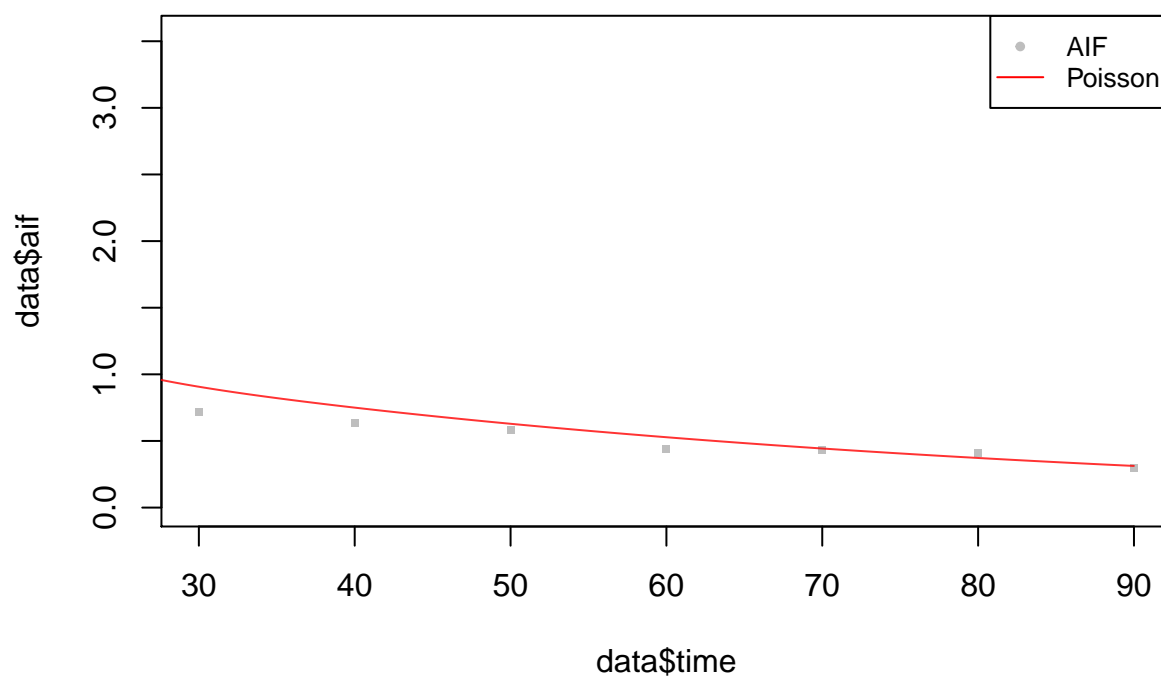
**patient:rwr1**



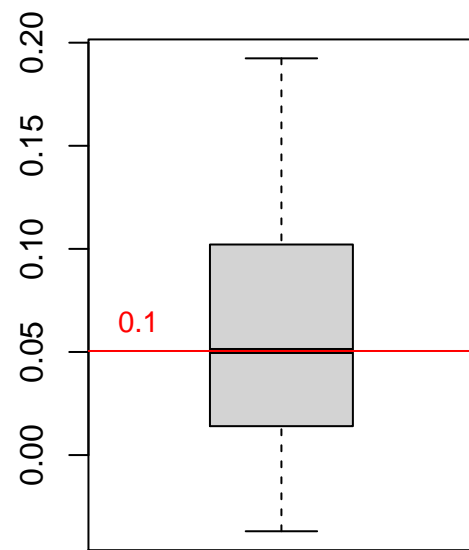
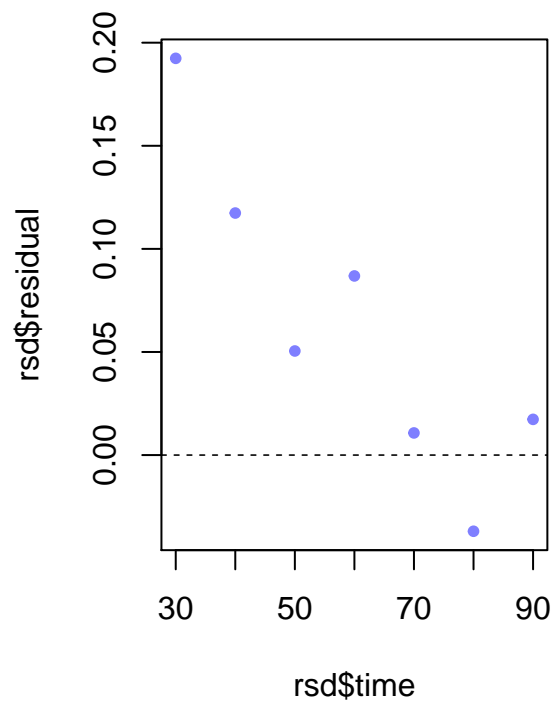
patient:rwr1



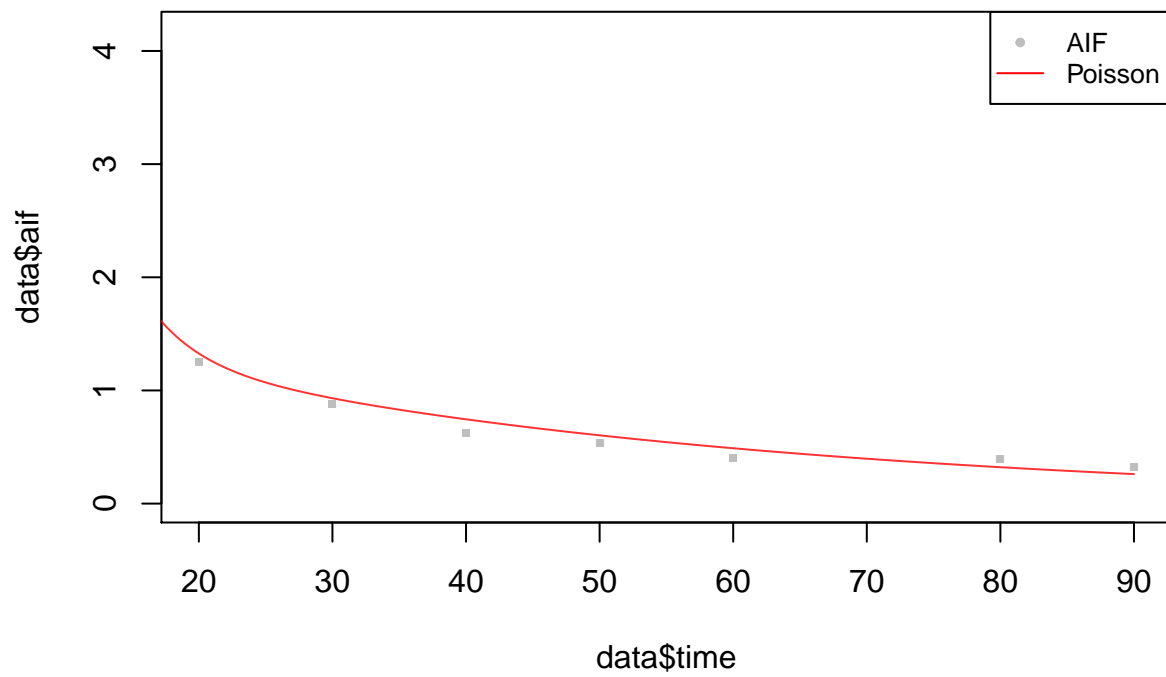
patient:rwr2



**patient:rwr2**

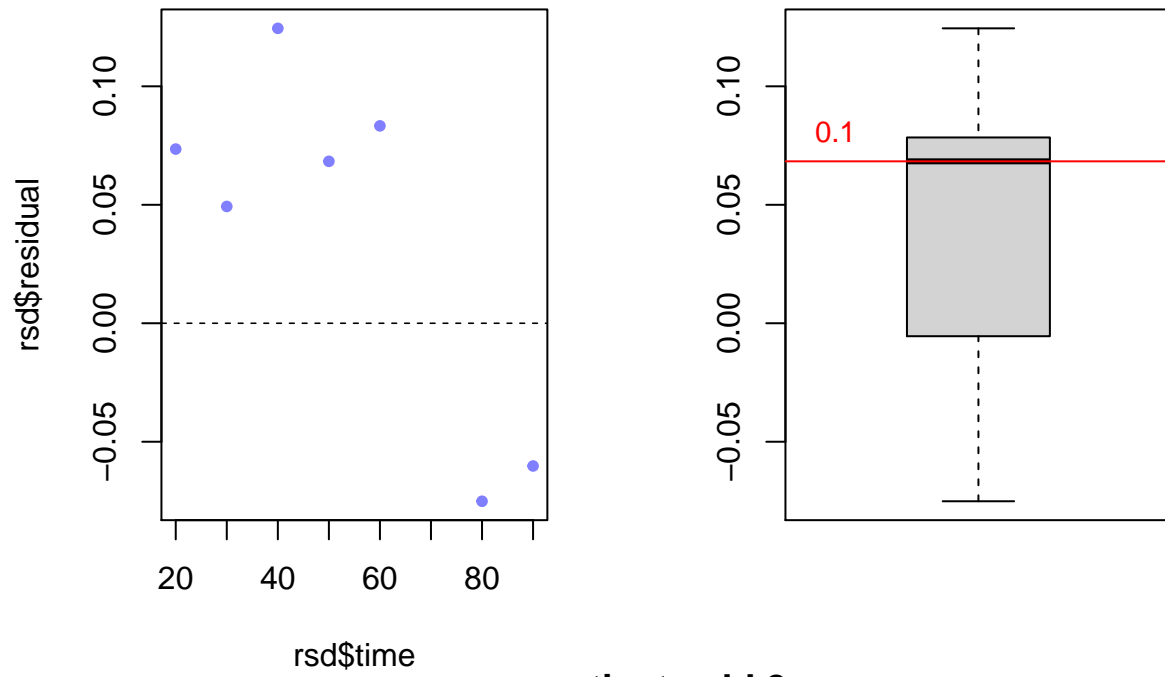


**patient:xehk1**

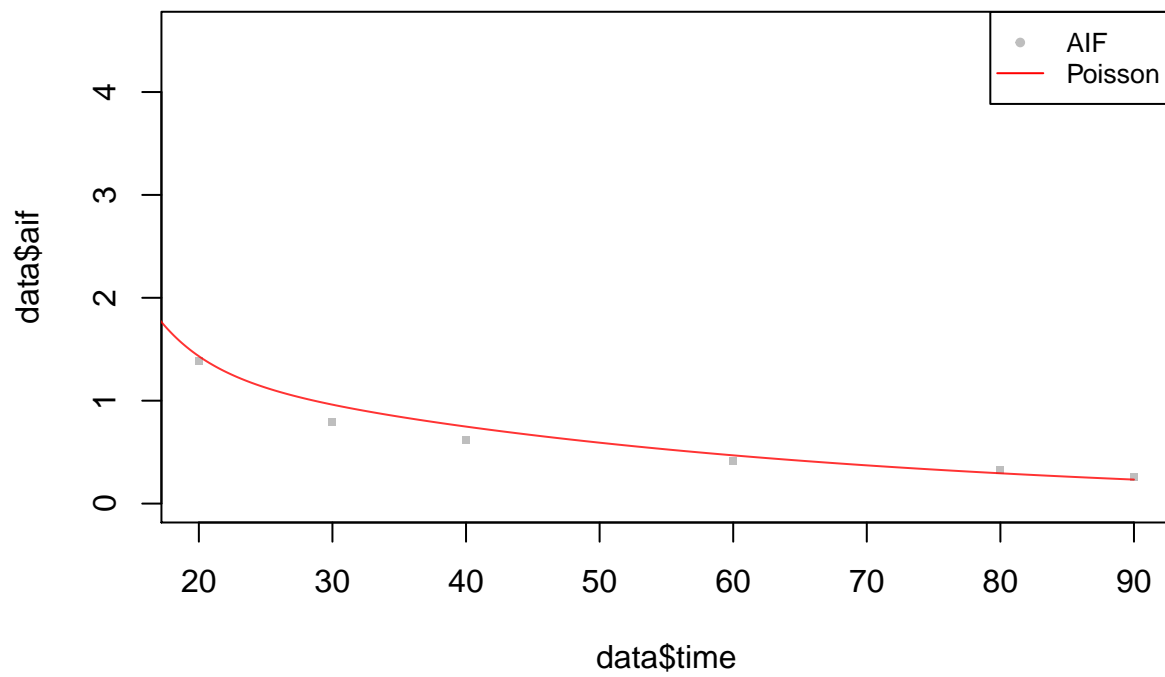




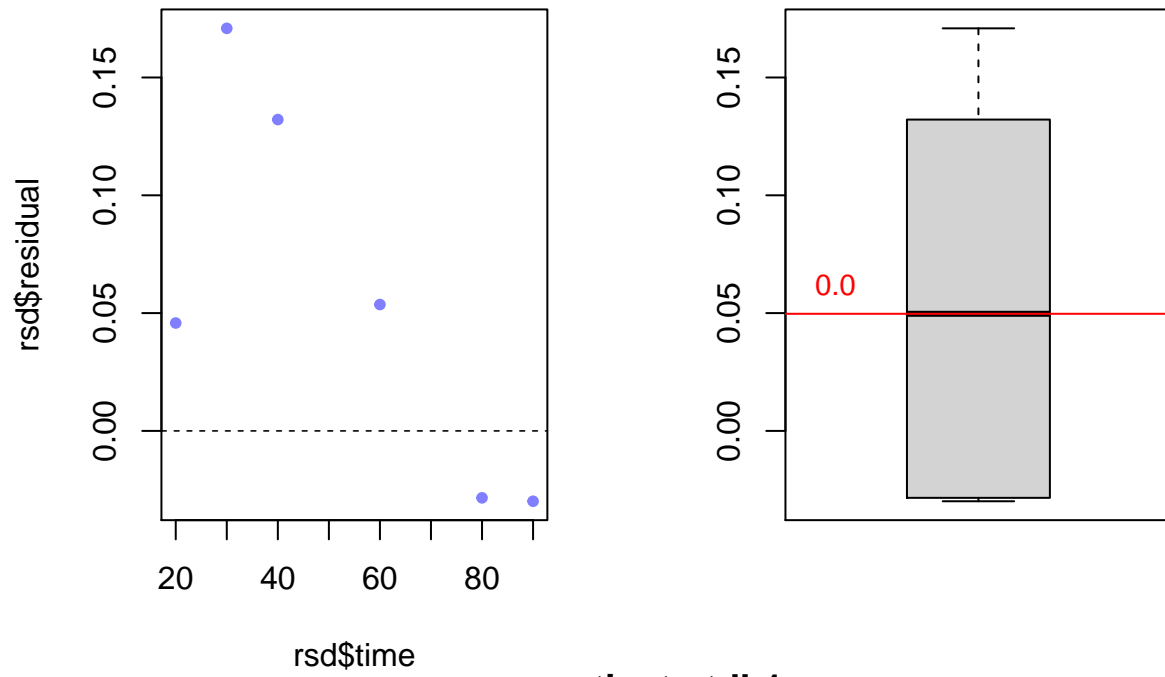
**patient:xehk1**



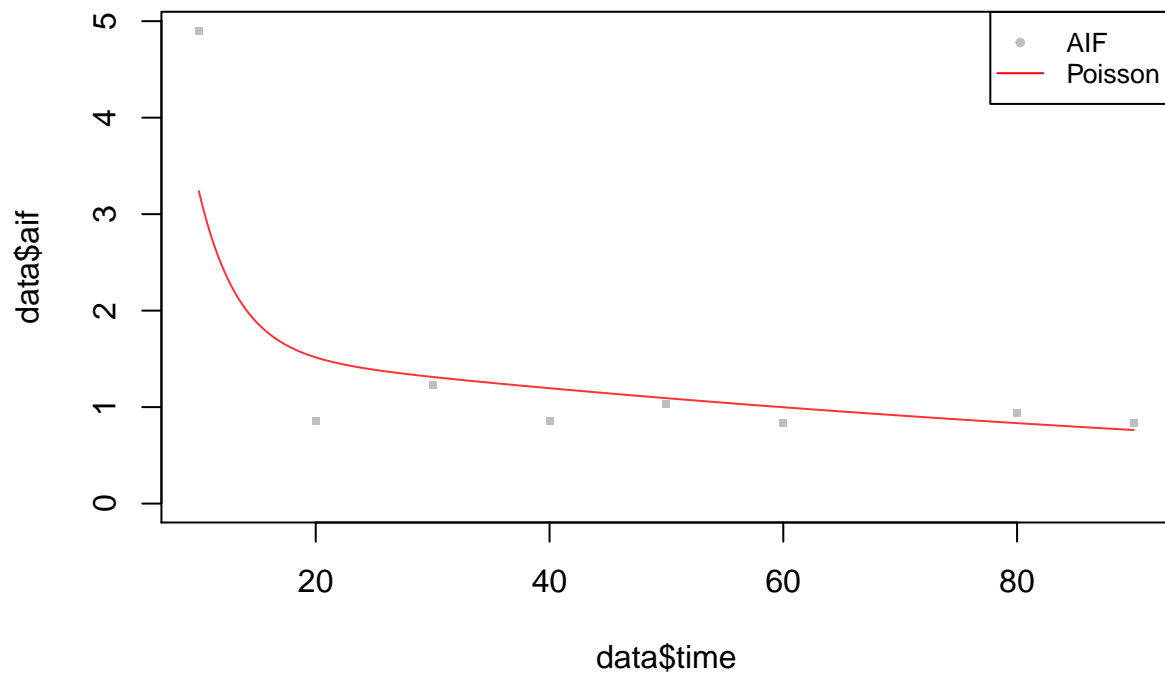
**patient:xehk2**



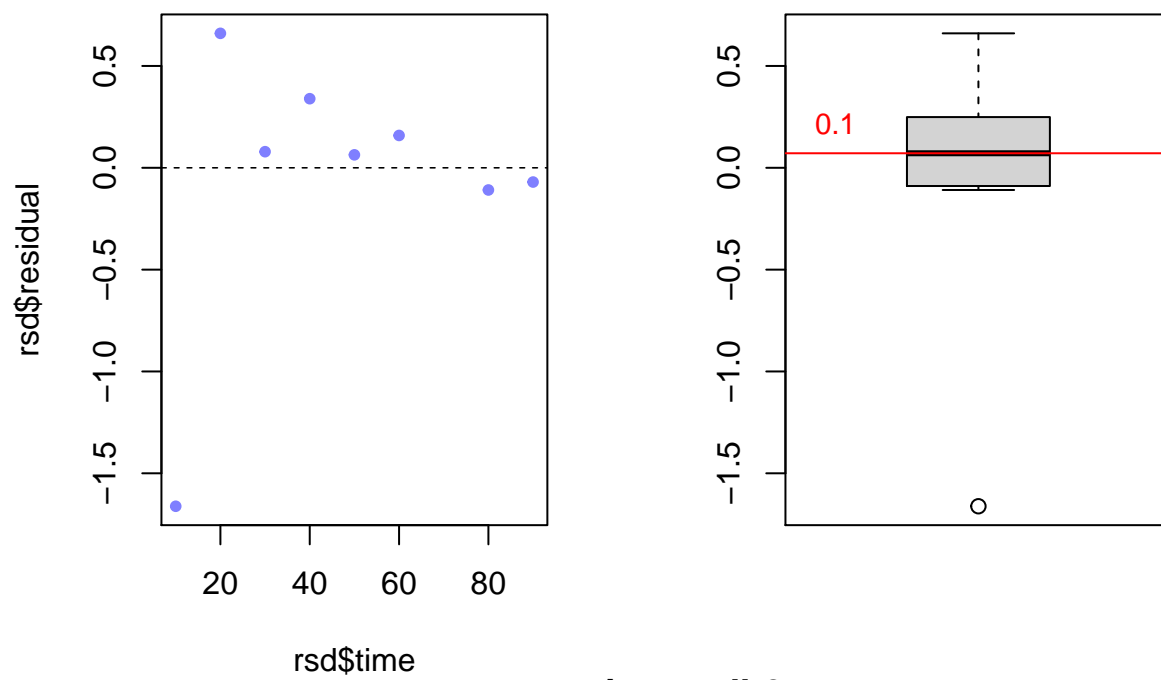
**patient:xehk2**



**patient:ytdh1**



**patient:ytdh1**



**patient:ytdh2**

