# Manuscript progress report

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## 2024-02-06

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## Method

## Assumptions

• For each subject i in the population, a generalized outcome  $Y_i(t)$  is generated along a variable t (for example, time), where  $t \in (0, T)$ .

• The outcome, at any specific t, follows an exponential family distribution characterized by a (latent) continuous function  $\eta_i(t)$ :

$$g[E(Y_i(t))] = \eta_i(t) = \beta_0(t) + b_i(t)$$

$$p(Y_i(t)) = h(Y_i(t))exp\{\eta_i(t)T[Y_i(t)] - A(\eta_i(t))\}$$

• The continuous latent function consists of a population-level fixed process and an individual-level random process

$$\eta_i(t) = \beta_0(t) + b_i(t)$$

## Observed data

In practice we would observe the discrete realization of  $\{Y_i(t), t\}$  along a dense grid. For simplicity, we assume the observation grid is regular (same across sample). When we have J observations points in (0, T], then for the jth observation point, we denote the corresponding value of t as  $t_j$ , and the corresponding outcome at this point  $Y_i(t_j)$ .

### fGFPCA Algorithm

#### Bin data:

Choose a proper bin width w considering model complexity and identifiability. For now let's say the bins are equal-length and non-overlapping.

- Bin index s = 1...S
- Index of bin midpoints  $m_s$
- Value of t corresponding to bin midpoints  $t_{m_s}\,$
- Bin endpoints:  $(t_{m_s} \frac{w}{2}, t_{m_s} + \frac{w}{2}]$

#### Local GLMMs

At the every bin, we fit a local intercept-only model:

$$g[E(Y_i(t_i))] = \eta_i(t_{m_s}) = \beta_0(t_{m_s}) + b_i(t_{m_s})$$

where  $t_j \in (t_{m_s} - \frac{w}{2}, t_{m_s} + \frac{w}{2}].$ 

Here we are basically saying that the value of latent function is constant within the same bin, which clearly is a misspecification of the true latent process.

From the model above, we will be able to estimate a  $\hat{\eta}_i(t_{m_s})$  on the binned grid for every individual in the training sample.

#### **FPCA**

Here, we fit a FPCA model on the  $\hat{\eta}_i(t_{m_s})$  obtained from step 2:

$$\hat{\eta}_i(t_{m_s}) = f_0(t_{m_s}) + \sum_{k=1}^K \xi_{ik} \phi_k(t_{m_s}) + \epsilon_i(t_{m_s})$$

where  $\xi_{ik}$  independently follows normal distribution  $N(0, \lambda_k)$ , and  $\epsilon_i(t_{m_s})$  at each point follows  $N(0, \sigma_2)$ . From this model, we will be able to obtain the following estimates which are shared across population:

- Population mean  $\hat{f}_0(t_{m_s})$
- Basis functions  $\hat{\mathbf{\Phi}} = \{\hat{\phi}_1(t_{m_s}), ..., \hat{\phi}_K(t_{m_s})\}$
- Estimates of variance of scores  $\hat{\lambda}_1...\hat{\lambda}_K$

### **Projection and Debias**

The mean and basis functions are evaluated on the binned grid. To extend it to the original measurement grid data was collected on, we project the estimated eigenfunctions  $\hat{\Phi}$  back use spline basis. Now we have extend the  $\hat{\phi}_k(t_{m_s})$  to the original grid  $\hat{\phi}_k(t_j)$ 

Because of the misspecification of local GLMMs, the estimated eigenfunctions and eigenvalues are also biased by a constant multiplicative effect. Therefore, we use a GLMM to re-evaluate the mean function, eigenfunctions and eigenvalues.

## Out-of-sample prediction

Now, let's assume we have a new subject u with  $J_u$  observations  $(J_u < J)$ . Then the log-likelihood of this new subject would be:

$$l_{u} = \sum_{t_{j} < t_{J_{u}}} log(h(Y_{u}(t_{j}))) + \hat{\eta}_{u}(t_{j})T(Y_{u}(t_{j})) - log(A[\hat{\eta}_{u}(t_{j})])$$

where 
$$\hat{\eta}_u(t_j) = \hat{f}_0(t_j) + \sum_{k=1}^K \xi_{uk} \hat{\phi}(t_j)$$
.

With estimates for the population-level parameters from fGFPCA algorithms above, we can estimate  $\xi_{uk}$  by maximization of  $l_u$ . Direct maximization some times does not have closed form solution. Numeric maximization methods seem not very stable as well. So I have decided to used a Bayes approach (Laplace Approximation):

- Prior distribution:  $\xi_{uk} \sim N(0, \hat{\lambda}_k)$
- Posterior distribution: the likelihood of  $l_u = l(Y_u(t_j)|\xi_u)$

Laplace Approximation would get the posterior mode of  $\xi_{uk}$  through quadratic approximation.

## Reference method

## **GLMMadaptive**

• For large datasets, we can fit a model with random intercept and slope for time. It is doable on 500 datasets, but obviously too simple for the data generation scheme. We would expect it to perform terribly.

$$g(E(Y_i(t))) = \beta_0 + \beta_1 t + b_{i0} + b_{i1} t$$

• For small datasets, we would like to fit fGFPCA and GLMMadaptive on a dataset with smaller sample size and/or smaller measurement density. For the GLMMadaptive model, we would set it up with spline basis functions so that its flexibility is comparable with fGFPCA model, such as:

$$g(E(Y_i(t))) = \sum_{k=1}^{4} \zeta_k B_k(t) + \sum_{l=1}^{4} \xi_{il} \phi_l(t)$$

## Generalized functional-on-scalar regression

We would like to use a second reference method for predictive performance comparision. In this model, predictions on a specific interval are all made using the last few observations in the observed window. For example, if we are given observations on [0, 0.2] (j = 1...200), we may use L observations taken right before t=0.2 ( $t_j: j = 200,...200 - (L-1)$ ) as time-fixed covariate to predict any future time.

Let's write out the model expression (with questionable notation). If the observations if up to  $t_m$ , then:

$$g(E[Y_i(t)]) = \beta_0(t) + \sum_{l=1}^{L} \beta_l(t) Y_i(t_l)$$
$$l = m, \dots m - (j-1)$$
$$t > t_m$$

This is a simple function-on-scalar model with no random effect, meaning all subject with the same last observed outcome would have the same estimated/predicted latent track. This is clearly anti-intuitive. But adding random effects would make out-of-sample prediction impossible.

Under both frameworks, for each dataset we need to fit four models (similar to the GLMMadaptive method):

- Given 0-0.2, predict 0.2-1
- Given 0-0.4, predict 0.4-1
- Given 0-0.6, predict 0.6-1
- Given 0-0.8, predict 0.8-1

But prediction performance will be reported by equal-length time window.

## Larger-scale simulation

### Simulation set up

Here we simulate binary data from cyclic latent process:

```
Y_{i}(t) \sim Bernoulli(\frac{exp(\eta_{i}(t))}{1 + exp(\eta_{i}(t))})
\eta_{i}(t) = f_{0}(t) + \xi_{i1}\sqrt{2}sin(2\pi t) + \xi_{i2}\sqrt{2}cos(2\pi t) + \xi_{i3}\sqrt{2}sin(4\pi t) + \xi_{i4}\sqrt{2}cos(4\pi t)
```

where:

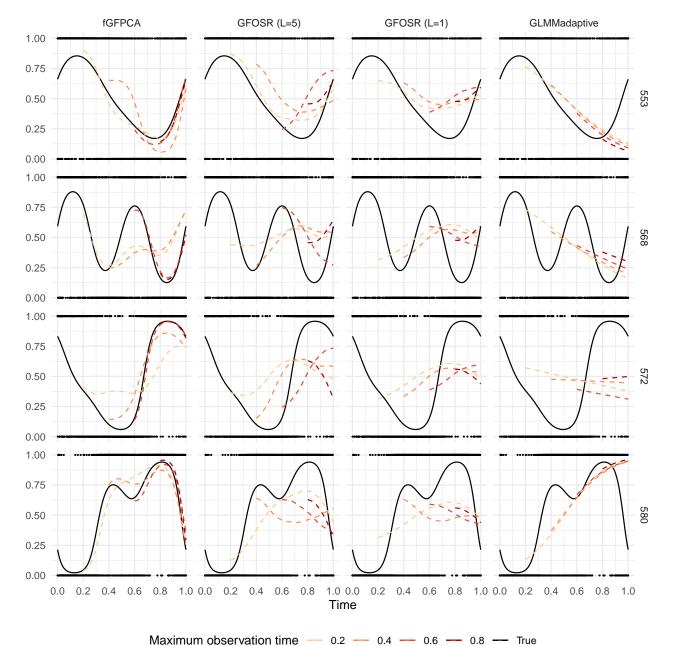
- t is 1000 equal-spaced observations points on [0, 1] (J = 1000).
- $f_0(t) = 0$
- $\xi_k \sim N(0, \lambda_k)$ , and  $\lambda_k = 1, 0.5, 0.25, 0.125$  for k = 1, 2, 3, 4 respectively.
- Sample size N = 500
- In the binning step, we bin every 10 observations
- 500 simulations were implemented

## **Figures**

```
load(here("Data/SimOutput_fGFPCA.RData"))
load(here("Data/SimOutput_GLMMadaptive.RData"))
load(here("Data/SimOutput_GFOSR_L1.RData"))
pred_list_gfosr_l1 <- pred_list_gfofr
t_vec_gfosr_l1 <- t_vec_gfofr
rm(pred_list_gfofr, t_vec_gfofr)
load(here("Data/SimOutput_GFOSR_L5.RData"))
pred_list_gfosr_l5 <- pred_list_gfofr
t_vec_gfosr_l5 <- t_vec_gfofr
rm(pred_list_gfofr, t_vec_gfofr)
rm(pred_list_gfofr, t_vec_gfofr)
rand_id <- sample(501:600, size = 4)</pre>
```

```
bind rows(
  pred_list_all[[1]] %>% filter(id %in% rand_id) %>%
   mutate(method="fGFPCA"),
  pred_list_ref[[1]] %>% filter(id %in% rand_id) %>%
   mutate(method = "GLMMadaptive"),
  pred_list_gfosr_l1[[1]] %>% filter(id %in% rand_id) %>%
   mutate(method = "GFOSR (L=1)") %>%
   rename(pred0.2=pred_w1, pred0.4=pred_w2, pred0.6=pred_w3, pred0.8=pred_w4),
  pred_list_gfosr_l5[[1]] %>% filter(id %in% rand_id) %>%
   mutate(method = "GFOSR (L=5)") %>%
    rename(pred0.2=pred_w1, pred0.4=pred_w2, pred0.6=pred_w3, pred0.8=pred_w4)
) %>% mutate at(vars(eta i, pred0.2, pred0.4, pred0.6, pred0.8),
                .funs = function(x)\{\exp(x)/(1+\exp(x))\}) %>%
  mutate(method=factor(method,
         levels = c("fGFPCA", "GFOSR (L=5)", "GFOSR (L=1)", "GLMMadaptive"))) %>%
  ggplot()+
```

```
geom_point(aes(x=t, y=Y), size = 0.2)+
geom_line(aes(x=t, y=eta_i, col = "True"))+
geom_line(aes(x=t, y=pred0.2, col = "0.2"), linetype="dashed", na.rm=T)+
geom_line(aes(x=t, y=pred0.4, col = "0.4"), linetype="dashed", na.rm=T)+
geom_line(aes(x=t, y=pred0.6, col = "0.6"), linetype="dashed", na.rm=T)+
geom_line(aes(x=t, y=pred0.8, col = "0.8"), linetype="dashed", na.rm=T)+
facet_grid(rows = vars(id), cols = vars(method))+
labs(col = "Maximum observation time", x = "Time", y="")+
scale_x_continuous(breaks = seq(0, 1, by = 0.2))+
theme(legend.position = "bottom")+
scale_color_manual(values = cols)
```



### ISE and AUC

```
## ISE container
ise fgfpca <- ise adglmm <- ise gfosr 11 <- ise gfosr 15 <-
  array(NA, dim = c(length(window)-2, length(window)-2, M))
# dims: prediction window, max obs time, simulation iter
## calculation
for(m in 1:M){
  # this_df <- pred_list_all[[m]]</pre>
  ise_tb <- pred_list_all[[m]] %>%
    mutate(err1 = (pred0.2-eta i)^2,
           err2 = (pred0.4-eta_i)^2,
           err3 = (pred0.6-eta i)^2,
           err4 = (pred0.8-eta_i)^2) %>%
    select(id, t, starts_with("err")) %>%
    mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group_by(window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != "[0,0.2]") %>%
    select(starts_with("err"))
  ise_fgfpca[, ,m] <- as.matrix(ise_tb)</pre>
}
mean_ise_fgfpca <- apply(ise_fgfpca, c(1, 2), mean)</pre>
# mean ise <- data.frame(mean ise) %>%
  mutate(Window = c("(0.2, 0.4]", "(0.4, 0.6]", "(0.6, 0.8]", "(0.8, 1.0]"),
           .before = 1)
colnames(mean_ise_fgfpca) <- c("0.2", "0.4", "0.6", "0.8")</pre>
## calculation
for(m in 1:M){
  # this_df <- pred_list_gfosr_l1[[m]]</pre>
  ise_tb <- pred_list_ref[[m]] %>%
    mutate(err1 = (pred0.2-eta_i)^2,
           err2 = (pred0.4-eta_i)^2,
           err3 = (pred0.6-eta_i)^2,
           err4 = (pred0.8-eta_i)^2) %>%
    select(id, t, starts_with("err")) %>%
    mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group_by(window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != "[0,0.2]") %>%
    select(starts with("err"))
  ise_adglmm[, ,m] <- as.matrix(ise_tb)</pre>
```

```
mean_ise_adglmm <- apply(ise_adglmm, c(1, 2), mean)</pre>
colnames(mean_ise_adglmm) <- c("0.2", "0.4", "0.6", "0.8")</pre>
## calculation
for(m in 1:M){
  # this_df <- pred_list_ref[[m]]</pre>
  ise_tb <- pred_list_gfosr_l1[[m]] %>%
    mutate(err1 = (pred_w1-eta_i)^2,
           err2 = (pred_w2-eta_i)^2,
           err3 = (pred_w3-eta_i)^2,
           err4 = (pred_w4-eta_i)^2) %>%
    select(id, t, starts_with("err"), window) %>%
    # mutate(window = factor(window, levels = 1:5,
                              labels = c("[0,0.2]", "(0.2,0.4]", "(0.4,0.6]",))) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group by (window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != 1) %>%
    select(starts_with("err"))
  ise_gfosr_l1[, ,m] <- as.matrix(ise_tb)</pre>
mean_ise_gfosr_l1 <- apply(ise_gfosr_l1, c(1, 2), mean)</pre>
colnames(mean_ise_gfosr_l1) <- c("0.2", "0.4", "0.6", "0.8")</pre>
## calculation
for(m in 1:M){
  # this_df <- pred_list_ref[[m]]</pre>
  ise_tb <- pred_list_gfosr_15[[m]] %>%
    mutate(err1 = (pred_w1-eta_i)^2,
           err2 = (pred_w2-eta_i)^2,
           err3 = (pred_w3-eta_i)^2,
           err4 = (pred_w4-eta_i)^2) %>%
    select(id, t, starts_with("err"), window) %>%
    # mutate(window = factor(window, levels = 1:5,
                              labels = c("[0,0.2]", "(0.2,0.4]", "(0.4,0.6]",))) %>%
    group by (window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group by (window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != 1) %>%
    select(starts_with("err"))
  ise_gfosr_15[, ,m] <- as.matrix(ise_tb)</pre>
mean_ise_gfosr_15 <- apply(ise_gfosr_15, c(1, 2), mean)</pre>
colnames(mean_ise_gfosr_15) <- c("0.2", "0.4", "0.6", "0.8")</pre>
## a function to calculate AUC
get_auc <- function(y, pred){</pre>
```

if(sum(is.na(y))>0 | sum(is.na(pred))>0){

```
auc <- NA
  }
  else{
    this_perf <- performance(prediction(pred, y), measure = "auc")</pre>
    auc <- this_perf@y.values[[1]]</pre>
 return(auc)
}
## auc container
auc_fgfpca <- auc_adglmm <- auc_gfosr_11 <- auc_gfosr_15 <-</pre>
  array(NA, dim = c(length(window)-2, length(window)-2, M))
for(m in 1:M){
  this_df <- pred_list_all[[m]]</pre>
  auc_tb <- this_df %>%
    mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
    select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred0.2),
              auc2 = get_auc(Y, pred0.4),
              auc3 = get auc(Y, pred0.6),
              auc4 = get_auc(Y, pred0.8)) %>%
    filter(window != "[0,0.2]") %>%
    select(starts_with("auc"))
  auc_fgfpca[, ,m] <- as.matrix(auc_tb)</pre>
mean_auc_fgfpca <- apply(auc_fgfpca, c(1, 2), mean)</pre>
# mean_auc <- data.frame(mean_auc) %>%
\# mutate(Window = c("(0.2, 0.4]", "(0.4, 0.6]", "(0.6, 0.8]", "(0.8, 1.0]"),
#
           .before = 1)
colnames(mean_auc_fgfpca) <- c("0.2", "0.4", "0.6", "0.8")</pre>
for(m in 1:M){
  this_df <- pred_list_ref[[m]]</pre>
  auc_tb <- this_df %>%
    mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
    select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred0.2),
              auc2 = get_auc(Y, pred0.4),
              auc3 = get_auc(Y, pred0.6),
              auc4 = get_auc(Y, pred0.8)) %>%
    filter(window != "[0,0.2]") %>%
    select(starts_with("auc"))
  auc_adglmm[, ,m] <- as.matrix(auc_tb)</pre>
}
mean_auc_adglmm <- apply(auc_adglmm, c(1, 2), mean)</pre>
```

```
colnames(mean_auc_adglmm) <- c("0.2", "0.4", "0.6", "0.8")</pre>
for(m in 1:M){
  this_df <- pred_list_gfosr_l1[[m]]</pre>
  auc tb <- this df %>%
    select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred_w1),
              auc2 = get_auc(Y, pred_w2),
              auc3 = get_auc(Y, pred_w3),
              auc4 = get_auc(Y, pred_w4)) %>%
    filter(window != 1) %>%
    select(starts_with("auc"))
  auc_gfosr_l1[, ,m] <- as.matrix(auc_tb)</pre>
}
mean_auc_gfosr_l1 <- apply(auc_gfosr_l1, c(1, 2), mean)</pre>
colnames(mean_auc_gfosr_l1) <- c("0.2", "0.4", "0.6", "0.8")</pre>
for(m in 1:M){
  this_df <- pred_list_gfosr_15[[m]]</pre>
  auc_tb <- this_df %>%
    select(Y, starts with("pred"), window) %>%
    group by (window) %>%
    summarise(auc1 = get_auc(Y, pred_w1),
              auc2 = get_auc(Y, pred_w2),
              auc3 = get_auc(Y, pred_w3),
              auc4 = get_auc(Y, pred_w4)) %>%
    filter(window != 1) %>%
    select(starts_with("auc"))
  auc_gfosr_15[, ,m] <- as.matrix(auc_tb)</pre>
}
mean_auc_gfosr_15 <- apply(auc_gfosr_15, c(1, 2), mean)</pre>
colnames(mean_auc_gfosr_15) <- c("0.2", "0.4", "0.6", "0.8")</pre>
fgfpca1 <- rbind(mean_ise_fgfpca ,mean_auc_fgfpca) %>% as_tibble() %>%
  mutate(Window = rep(c("(0.2, 0.4]", "(0.4, 0.6]", "(0.6, 0.8]", "(0.8, 1.0]"), 2), .before=1)
gfosr_15_1 <- rbind(mean_ise_gfosr_15 ,mean_auc_gfosr_15) %>% as_tibble()
gfosr_l1_1 <- rbind(mean_ise_gfosr_l1 ,mean_auc_gfosr_l1) %>% as_tibble()
adglmm1 <- rbind(mean_ise_adglmm ,mean_auc_adglmm) %>% as_tibble()
bind_cols(fgfpca1, gfosr_15_1, gfosr_11_1, adglmm1, .name_repair = "minimal") %>%
  kable(digits = 3, booktabs=T,
        table.attr="style=\"color:black;\"") %>%
 kable_styling(full_width = F) %>%
  add_header_above(c(" " = 1, "fGFPCA" = 4, "GFOSR (L=5)" = 4,
                      "GFOSR (L=1)" = 4, "GLMMadaptive" = 4)) %>%
  add_header_above(c(" " = 1, "Maximum observation time" = 16)) %>%
```

```
group_rows(index = c("ISE"= 4, "AUC" = 4)) %>%
landscape()
```

-
_
10

	Maximum observation time															
		fGFF	CA			GFOSE	R (L=5)			GFOSE	R (L=1)		GLMMadaptive			
Window	0.2	0.4	0.6	0.8	0.2	0.4	0.6	0.8	0.2	0.4	0.6	0.8	0.2	0.4	0.6	
ISE																
(0.2, 0.4]	146.407				274.947				362.476				387.708			
(0.4, 0.6]	183.967	74.977			277.438	220.209			286.614	262.552			291.579	269.799		
(0.6, 0.8]	218.265	49.275	15.776		322.338	373.108	325.017		385.701	410.508	389.305		315.778	282.736	278.242	
(0.8, 1.0]	108.918	77.981	17.747	12.005	290.982	318.079	350.850	333.580	328.482	341.274	354.211	347.067	563.011	477.485	597.746	
$\mathbf{AUC}$																
(0.2, 0.4]	0.748				0.686				0.624				0.591			
(0.4, 0.6]	0.664	0.734			0.563	0.630			0.543	0.590			0.524	0.596		
(0.6, 0.8]	0.715	0.790	0.803		0.669	0.628	0.676		0.604	0.577	0.615		0.669	0.694	0.687	
(0.8, 1.0]	0.740	0.755	0.781	0.784	0.626	0.606	0.552	0.584	0.588	0.564	0.537	0.551	0.514	0.556	0.526	

Method	Time
fGFPCA	2.317
GLMMadaptive	2.304
GFOSR (L=1)	0.025
GFOSR $(L=5)$	0.144

## Time (minutes)

I chose not to report fitting and prediction time separately because it is too difficult to tell them apart for the GFOSR models.

## Small-scale simulation

I have used 100 subjects for training and testing, and repeated 500 times. When fitting GLMMadpative, I reduce the number of measurements in the training dataset to 1/10 by taking one every 10 observations. The prediction is on the original grid.

```
load(here("Data/SubSimOutput_GEPCA.RData"))

load(here("Data/SubSimOutput_fGFPCA.RData"))

pred_subset_adglmm <- pred_subset_adglmm[!num_probs]

fit_time_subset_adglmm <- fit_time_subset_adglmm[!num_probs]

pred_time_subset_adglmm <- pred_time_subset_adglmm[!num_probs]

load(here("Data/SubSimOutput_GFOSR_L1.RData"))

pred_subset_gfosr_l1 <- pred_list_gfofr_subset

t_subset_gfosr_l1 <- t_vec_gfofr_subset

rm(pred_list_gfofr_subset, t_vec_gfofr_subset)

load(here("Data/SubSimOutput_GFOSR_L5.RData"))

pred_subset_gfosr_l5 <- pred_list_gfofr_subset

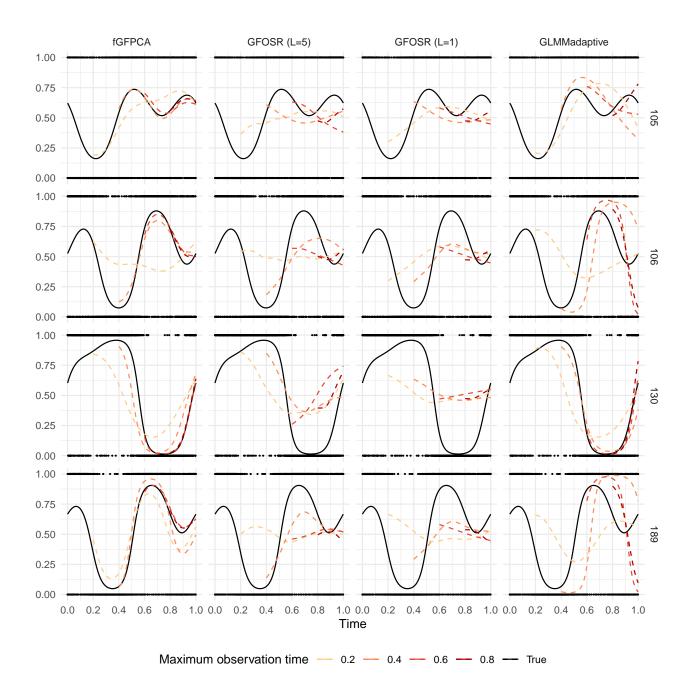
t_subset_gfosr_l5 <- t_vec_gfofr_subset

rm(pred_list_gfofr_subset, t_vec_gfofr_subset)

rand_id <- sample(101:200, size = 4)</pre>
```

## **Figure**

```
bind_rows(
  pred_subset_fGFPCA[[1]] %>% filter(id %in% rand_id) %>% mutate(method="fGFPCA"),
  pred_subset_gfosr_15[[1]] %>% filter(id %in% rand_id) %>% mutate(method = "GFOSR (L=5)") %>%
   rename(pred0.2=pred_w1, pred0.4=pred_w2, pred0.6=pred_w3, pred0.8=pred_w4),
  pred_subset_gfosr_l1[[1]] %>% filter(id %in% rand_id) %>% mutate(method = "GFOSR (L=1)") %>%
   rename(pred0.2=pred_w1, pred0.4=pred_w2, pred0.6=pred_w3, pred0.8=pred_w4),
  pred_subset_adglmm[[1]] %>% filter(id %in% rand_id) %>% mutate(method = "GLMMadaptive")) %>%
  mutate_at(vars(eta_i, pred0.2, pred0.4, pred0.6, pred0.8),
                .funs = function(x){exp(x)/(1+exp(x))}) \%%
  mutate(method=factor(method,
        levels = c("fGFPCA", "GFOSR (L=5)", "GFOSR (L=1)", "GLMMadaptive"))) %%
  ggplot()+
  geom point(aes(x=t, y=Y), size = 0.2)+
  geom_line(aes(x=t, y=eta_i, col = "True"))+
  geom_line(aes(x=t, y=pred0.2, col = "0.2"), linetype="dashed", na.rm=T)+
  geom_line(aes(x=t, y=pred0.4, col = "0.4"), linetype="dashed", na.rm=T)+
  geom_line(aes(x=t, y=pred0.6, col = "0.6"), linetype="dashed", na.rm=T)+
  geom_line(aes(x=t, y=pred0.8, col = "0.8"), linetype="dashed", na.rm=T)+
  facet_grid(rows = vars(id), cols = vars(method))+
  labs(col = "Maximum observation time", x = "Time", y="")+
  scale_x_continuous(breaks = seq(0, 1, by = 0.2))+
  theme(legend.position = "bottom")+
  scale_color_manual(values = cols)
```



## ISE and AUC

```
## ISE container
ise_fgfpca2 <- ise_adglmm2 <- ise_gfosr_l1_2 <- ise_gfosr_l5_2 <-
array(NA, dim = c(length(window)-2, length(window)-2, M))
# dims: prediction window, max obs time, simulation iter

for(m in 1:length(pred_subset_fGFPCA)){
    this_df <- pred_subset_fGFPCA[[m]]
    ise_tb_m <- this_df %>%
        mutate(err1 = (pred0.2-eta_i)^2,
```

```
err2 = (pred0.4-eta_i)^2,
           err3 = (pred0.6-eta_i)^2,
           err4 = (pred0.8-eta_i)^2) %>%
    select(id, t, starts_with("err")) %>%
    mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group by (window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != "[0,0.2]") %>%
    select(starts_with("err")) %>% as.matrix()
  ise_fgfpca2[,,m] <- ise_tb_m</pre>
mean_ise_fgfpca2 <- apply(ise_fgfpca2, c(1, 2), mean)</pre>
colnames(mean_ise_fgfpca2) <- c("0.2", "0.4", "0.6", "0.8")</pre>
for(m in 1:length(pred_subset_adglmm)){
  this_df <- pred_subset_adglmm[[m]]</pre>
  ise tb m <- this df %>%
    mutate(err1 = (pred0.2-eta_i)^2,
           err2 = (pred0.4-eta_i)^2,
           err3 = (pred0.6-eta_i)^2,
           err4 = (pred0.8-eta_i)^2) %>%
    select(id, t, starts_with("err")) %>%
    mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group_by(window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != "[0,0.2]") %>%
    select(starts_with("err")) %>% as.matrix()
  ise_adglmm2[,,m] <- ise_tb_m</pre>
mean_ise_adglmm2<- apply(ise_adglmm2, c(1, 2), mean, na.rm = T)</pre>
colnames(mean ise adglmm2) \leftarrow c("0.2", "0.4", "0.6", "0.8")
for(m in 1:length(pred_subset_gfosr_l1)){
  this_df <- pred_subset_gfosr_l1[[m]]</pre>
  ise_tb_m <- this_df %>%
    mutate(err1 = (pred_w1-eta_i)^2,
           err2 = (pred_w2-eta_i)^2,
           err3 = (pred_w3-eta_i)^2,
           err4 = (pred_w4-eta_i)^2) %>%
    select(id, t, starts_with("err"), window) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group by (window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != 1) %>%
```

```
select(starts_with("err")) %>% as.matrix()
 ise_gfosr_l1_2[,,m] <- ise_tb_m</pre>
mean_ise_gfosr_l1_2<- apply(ise_gfosr_l1_2, c(1, 2), mean, na.rm = T)</pre>
colnames(mean_ise_gfosr_11_2) <- c("0.2", "0.4", "0.6", "0.8")</pre>
for(m in 1:length(pred_subset_gfosr_15)){
  this_df <- pred_subset_gfosr_15[[m]]</pre>
  ise_tb_m <- this_df %>%
    mutate(err1 = (pred_w1-eta_i)^2,
           err2 = (pred_w2-eta_i)^2,
           err3 = (pred_w3-eta_i)^2,
           err4 = (pred_w4-eta_i)^2) %>%
    select(id, t, starts_with("err"), window) %>%
    group_by(window, id) %>%
    summarise_at(vars(err1, err2, err3, err4), sum) %>%
    group_by(window) %>%
    summarize_at(vars(err1, err2, err3, err4), mean) %>%
    filter(window != 1) %>%
    select(starts_with("err")) %>% as.matrix()
  ise_gfosr_15_2[,,m] <- ise_tb_m</pre>
mean_ise_gfosr_15_2 <- apply(ise_gfosr_15_2, c(1, 2), mean, na.rm = T)</pre>
colnames(mean_ise_gfosr_15_2) <- c("0.2", "0.4", "0.6", "0.8")
## auc container
auc_fgfpca2 <- auc_adglmm2 <- auc_gfosr_11_2 <- auc_gfosr_15_2 <-</pre>
  array(NA, dim = c(length(window)-2, length(window)-2, M))
for(m in 1:length(pred_subset_fGFPCA)){
  this_df <- pred_subset_fGFPCA[[m]]</pre>
  auc_tb_m <- this_df %>%
  mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
  select(Y, starts_with("pred"), window) %>%
  group_by(window) %>%
  summarise(auc1 = get_auc(Y, pred0.2),
            auc2 = get_auc(Y, pred0.4),
            auc3 = get_auc(Y, pred0.6),
            auc4 = get_auc(Y, pred0.8)) %>%
  filter(window != "[0,0.2]") %>%
  select(starts_with("auc")) %>% as.matrix()
 auc_fgfpca2[, , m] <- auc_tb_m</pre>
}
mean_auc_fgfpca2 <- apply(auc_fgfpca2, c(1,2), mean)</pre>
colnames(mean_auc_fgfpca2) <- c("0.2", "0.4", "0.6", "0.8")</pre>
```

```
for(m in 1:length(pred_subset_adglmm)){
  this_df <- pred_subset_adglmm[[m]]</pre>
  auc_tb_m <- this_df %>%
  mutate(window = cut(t, breaks = window, include.lowest = T)) %>%
  select(Y, starts_with("pred"), window) %>%
  group_by(window) %>%
  summarise(auc1 = get_auc(Y, pred0.2),
            auc2 = get auc(Y, pred0.4),
            auc3 = get_auc(Y, pred0.6),
            auc4 = get_auc(Y, pred0.8)) %>%
  filter(window != "[0,0.2]") %>%
  select(starts_with("auc")) %>% as.matrix()
 auc_adglmm2[, , m] <- auc_tb_m</pre>
}
mean_auc_adglmm2 <- apply(auc_adglmm2, c(1,2), mean, na.rm = T)</pre>
colnames(mean_auc_adglmm2) <- c("0.2", "0.4", "0.6", "0.8")</pre>
for(m in 1:length(pred_subset_gfosr_l1)){
  this_df <- pred_subset_gfosr_l1[[m]]</pre>
  auc_tb_m <- this_df %>%
  select(Y, starts_with("pred"), window) %>%
  group_by(window) %>%
  summarise(auc1 = get_auc(Y, pred_w1),
            auc2 = get_auc(Y, pred_w2),
            auc3 = get_auc(Y, pred_w3),
            auc4 = get_auc(Y, pred_w4)) %>%
  filter(window != 1) %>%
  select(starts_with("auc")) %>% as.matrix()
 auc_gfosr_l1_2[, , m] <- auc_tb_m</pre>
}
mean_auc_gfosr_l1_2 <- apply(auc_gfosr_l1_2, c(1,2), mean)</pre>
colnames(mean_auc_gfosr_11_2) <- c("0.2", "0.4", "0.6", "0.8")
for(m in 1:length(pred_subset_gfosr_15)){
  this_df <- pred_subset_gfosr_15[[m]]</pre>
  auc_tb_m <- this_df %>%
  select(Y, starts_with("pred"), window) %>%
  group_by(window) %>%
  summarise(auc1 = get_auc(Y, pred_w1),
            auc2 = get_auc(Y, pred_w2),
            auc3 = get_auc(Y, pred_w3),
            auc4 = get_auc(Y, pred_w4)) %>%
  filter(window != 1) %>%
  select(starts with("auc")) %>% as.matrix()
  auc_gfosr_15_2[, , m] <- auc_tb_m</pre>
```

```
}
mean_auc_gfosr_15_2 <- apply(auc_gfosr_15_2, c(1,2), mean)</pre>
colnames(mean_auc_gfosr_15_2) <- c("0.2", "0.4", "0.6", "0.8")</pre>
fgfpca2 <- rbind(mean_ise_fgfpca2 ,mean_auc_fgfpca2) %>% as_tibble() %>%
  mutate(Window = rep(c("(0.2, 0.4]", "(0.4, 0.6]", "(0.6, 0.8]", "(0.8, 1.0]"), 2), .before=1)
gfosr 15 2 <- rbind(mean ise gfosr 15 2 ,mean auc gfosr 15 2) %>% as tibble()
gfosr_11_2 <- rbind(mean_ise_gfosr_11_2 ,mean_auc_gfosr_11_2) %>% as_tibble()
adglmm2 <- rbind(mean_ise_adglmm2 ,mean_auc_adglmm2) %>% as_tibble()
bind_cols(fgfpca2, gfosr_15_2, gfosr_11_2, adglmm2, .name_repair = "minimal") %>%
  kable(digits = 3, booktabs=T,
        table.attr="style=\"color:black;\"") %>%
  kable_styling(full_width = F) %>%
  add_header_above(c(" " = 1, "fGFPCA" = 4, "GFOSR (L=5)" = 4,
                     "GFOSR (L=1)" = 4, "GLMMadaptive" = 4)) %>%
  add_header_above(c(" " = 1, "Maximum observation time" = 16)) %>%
  group_rows(index = c("ISE"= 4, "AUC" = 4)) %>%
  landscape()
```

2
0

	Maximum observation time															
		fGFF	PCA			GFOSF	R (L=5)			GFOSE	t (L=1)		GLMMadaptive			
Window	0.2	0.4	0.6	0.8	0.2	0.4	0.6	0.8	0.2	0.4	0.6	0.8	0.2	0.4	0.6	
ISE																
(0.2, 0.4]	150.641				286.853				367.282				374.836		J	
(0.4, 0.6]	188.065	77.380			286.628	227.225			290.781	265.809			268.390	463.337		
(0.6, 0.8]	224.001	51.533	16.879		328.612	374.572	326.522		386.705	408.427	388.427		319.731	287.001	233.510	
(0.8, 1.0]	112.163	81.368	19.592	13.399	301.165	324.817	353.888	335.139	331.488	342.935	354.800	347.627	228.335	469.769	331.509	
$\mathbf{AUC}$															ļ	
(0.2, 0.4]	0.746				0.679				0.620				0.672		1	
(0.4, 0.6]	0.662	0.734			0.549	0.627			0.536	0.590			0.622	0.671	J	
(0.6, 0.8]	0.710	0.787	0.801		0.656	0.617	0.668		0.598	0.574	0.613		0.689	0.698	0.731	
(0.8, 1.0]	0.740	0.754	0.781	0.784	0.611	0.588	0.551	0.578	0.580	0.556	0.536	0.551	0.681	0.629	0.679	

Method	Time
fGFPCA	1.629
GLMMadaptive	5.479
GFOSR (L=1)	0.010
GFOSR $(L=5)$	0.054

Among 500 hundred iterations, 6 did not converge for the GLMMadaptive model. No numeric issue for fGFPCA.

### Time

## NHANES data application

We take 60% subjects (N = 5257) for training, 40% (n = 3506) for out-of-sample prediction. When using the full sample The debiase step (mgcv::bam) in fGFPCA took a really a long time, and GLMM adaptive only works for intercept-only model. In the debias step, re-evaluated eigenvalues also lost their decreasing nature.

```
# df_nhanes <- read_rds(here("Data/nhanes_bi.rds"))
load(here("Data/ApplOutput_GEPCA.RData"))
load(here("Data/ApplOutput_GLMMadaptive.RData"))
load(here("Data/Appl_GFOSR_L5.RData"))
load(here("Data/Appl_GFOSR_L1.RData"))

pred_appl_gfosr_l1 <- pred_appl_gfosr_l1 %>%
    rename(pred360=pred_w1, pred720=pred_w2, pred1080=pred_w3)
pred_appl_gfosr_l5 <- pred_appl_gfosr_l5 %>%
    rename(pred360=pred_w1, pred720=pred_w2, pred1080=pred_w3)
```

```
df_nhanes %>% select(SEQN, Z, sind) %>%
  filter(SEQN %in% unique(nhanes_pred_adglmm$id) | SEQN %in% unique(nhanes_pred_fgfpca$id)) %>%
  mutate(Z = factor(Z, levels = 0:1, labels = c("Inactive", "Active"))) %>%
  mutate(SEQN = factor(SEQN)) %>%
  ggplot()+
  geom_tile(aes(x=sind, y = SEQN, fill = Z))+
  labs(x="Time", y="Subject", title = "Overview of NHANES binary activity indicator")+
  theme(axis.text.y = element_blank())+
```

### **AUC**

```
# window
window <- seq(0, 1440, by = 360)</pre>
```

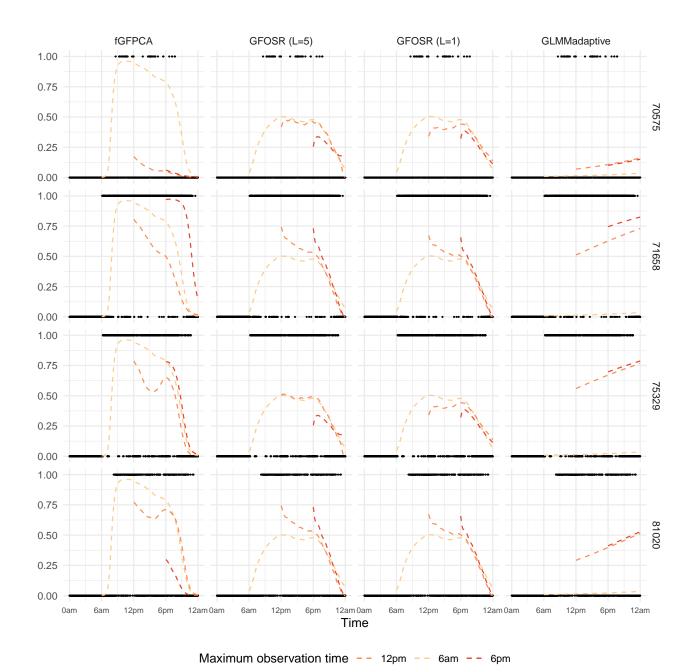
```
auc_appl_fgfpca <- pred_nhanes_fgfpca %>%
    mutate(window = cut(sind, breaks = window, include.lowest = T)) %>%
    select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred360),
              auc2 = get_auc(Y, pred720),
              auc3 = get_auc(Y, pred1080)) %>%
   filter(window != "[0,360]") %>%
    select(starts_with("auc"))
auc_appl_gfosr_15 <- pred_appl_gfosr_15 %>%
  select(-window) %>%
   mutate(window = cut(sind, breaks = window, include.lowest = T)) %>%
   select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred360),
              auc2 = get_auc(Y, pred720),
              auc3 = get_auc(Y, pred1080)) %>%
   filter(window != "[0,360]") %>%
    select(starts_with("auc"))
auc_appl_gfosr_l1 <- pred_appl_gfosr_l1 %>%
  select(-window) %>%
   mutate(window = cut(sind, breaks = window, include.lowest = T)) %>%
    select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred360),
              auc2 = get_auc(Y, pred720),
              auc3 = get_auc(Y, pred1080)) %>%
   filter(window != "[0,360]") %>%
    select(starts_with("auc"))
auc_appl_adglmm <- pred_nhanes_adglmm %>%
    mutate(window = cut(sind, breaks = window, include.lowest = T)) %>%
    select(Y, starts_with("pred"), window) %>%
    group_by(window) %>%
    summarise(auc1 = get_auc(Y, pred360),
              auc2 = get_auc(Y, pred720),
              auc3 = get_auc(Y, pred1080)) %>%
   filter(window != "[0,360]") %>%
    select(starts_with("auc"))
colnames(auc_appl_fgfpca) <- colnames(auc_appl_gfosr_15) <- colnames(auc_appl_gfosr_11) <- colnames(auc_appl_gfosr_15)
```

$\sim$
4

	Maximum observation time													
		fGFPCA	1	GF	OSR (L	GF	OSR (L	=1)	GLMMadaptive					
Window	6am	12pm	6pm	6am	12pm	6pm	6am	12pm	6pm	6am	12pm	6pm		
6am-12pm	0.703			0.689			0.684			0.581				
$12 \mathrm{pm}\text{-}6 \mathrm{pm}$	0.536	0.709		0.519	0.699		0.520	0.654		0.532	0.701			
6 pm- 12 am	0.716	0.679	0.773	0.677	0.678	0.723	0.677	0.676	0.708	0.514	0.565	0.626		

## **Figure**

```
rand_id <- sample(unique(pred_nhanes_adglmm$id), 4)</pre>
cols <- cols[1:3]</pre>
names(cols) <- c("6am", "12pm", "6pm")</pre>
bind_rows(
  pred_nhanes_fgfpca %>% filter(id %in% rand_id) %>% mutate(method="fGFPCA"),
  pred_appl_gfosr_15 %>% filter(id %in% rand_id) %>% mutate(method = "GFOSR (L=5)"),
  pred_appl_gfosr_l1 %>% filter(id %in% rand_id) %>% mutate(method = "GFOSR (L=1)"),
 pred_nhanes_adglmm %>% filter(id %in% rand_id) %>% mutate(method = "GLMMadaptive")) %>%
  mutate at(vars(pred360, pred720, pred1080),
                .funs = function(x)\{\exp(x)/(1+\exp(x))\}) %>%
  mutate(method=factor(method,
         levels = c("fGFPCA", "GFOSR (L=5)", "GFOSR (L=1)", "GLMMadaptive"))) %>%
  ggplot()+
  geom_point(aes(x=sind, y=Y), size = 0.2)+
  geom_line(aes(x=sind, y=pred360, col = "6am"), linetype="dashed", na.rm=T)+
  geom_line(aes(x=sind, y=pred720, col = "12pm"), linetype="dashed", na.rm=T)+
  geom_line(aes(x=sind, y=pred1080, col = "6pm"), linetype="dashed", na.rm=T)+
  facet_grid(rows = vars(id), cols = vars(method))+
  labs(col = "Maximum observation time", x = "Time", y="")+
  scale_x_continuous(breaks = seq(0, 1440, by = 360),
                     labels = c("0am", "6am", "12pm", "6pm", "12am"))+
  theme(legend.position = "bottom", axis.text.x = element_text(size = 7))+
  scale_color_manual(values = cols)
```



Additional figures