

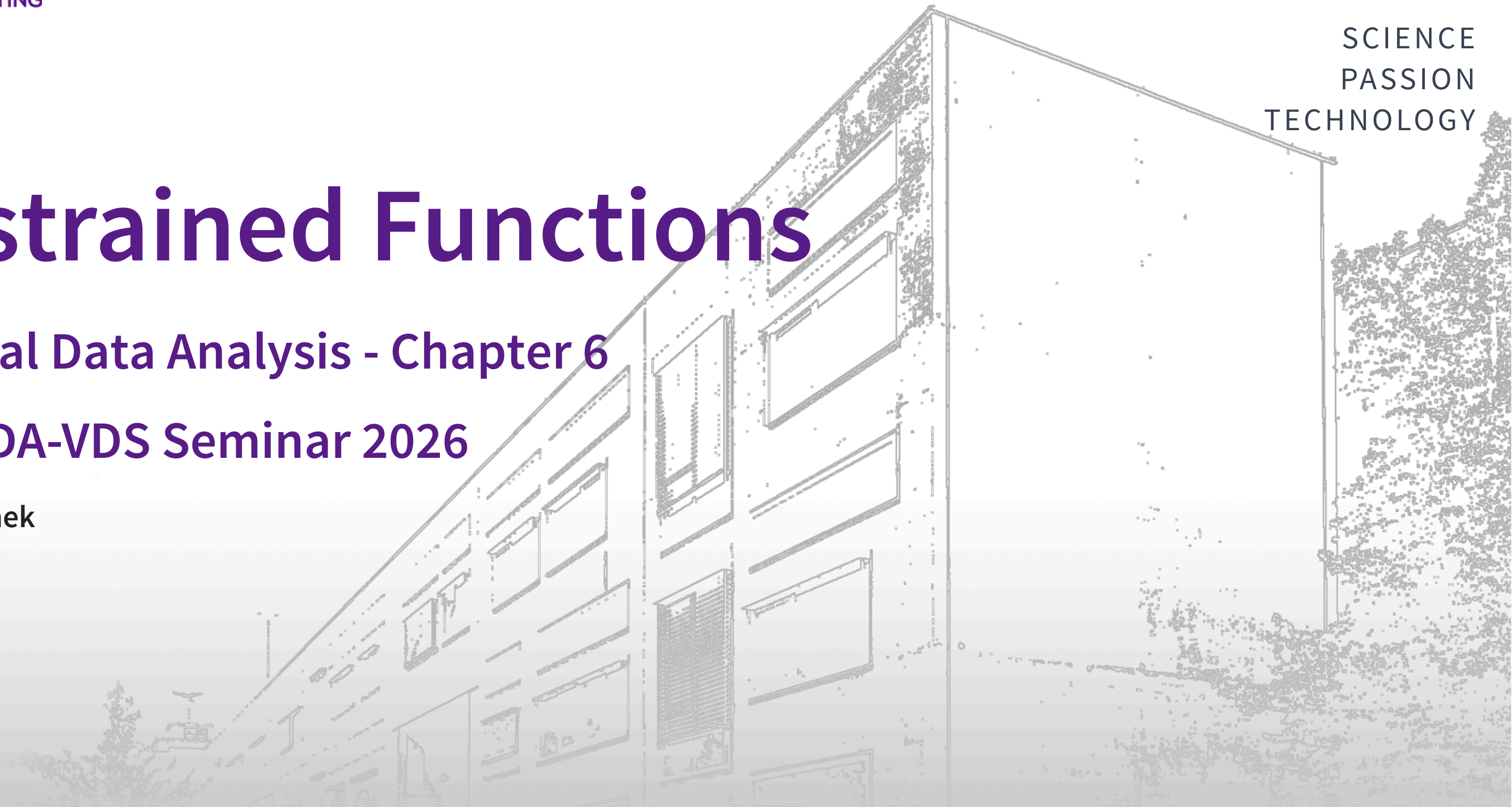
Constrained Functions

Functional Data Analysis - Chapter 6

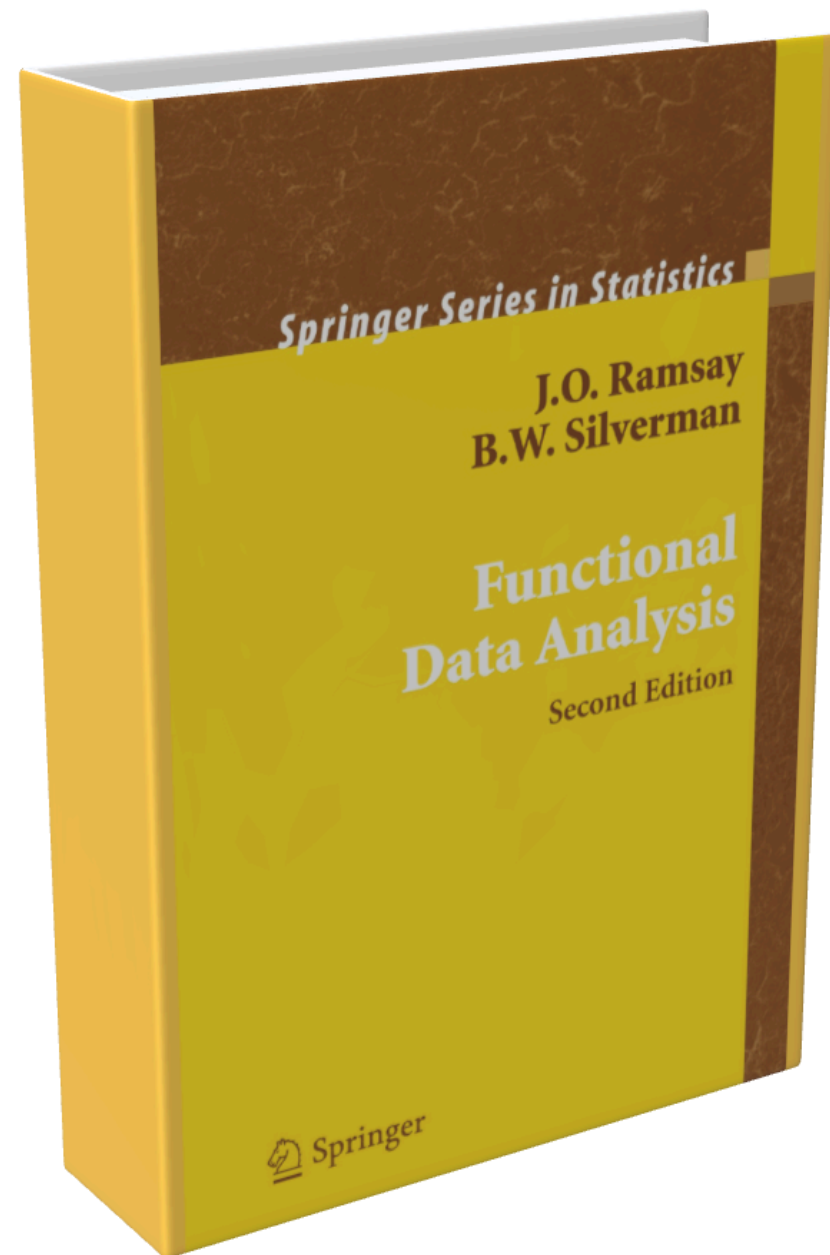
CEDAS FDA-VDS Seminar 2026

Julian Rakuschek

11.06.2026



In this lecture

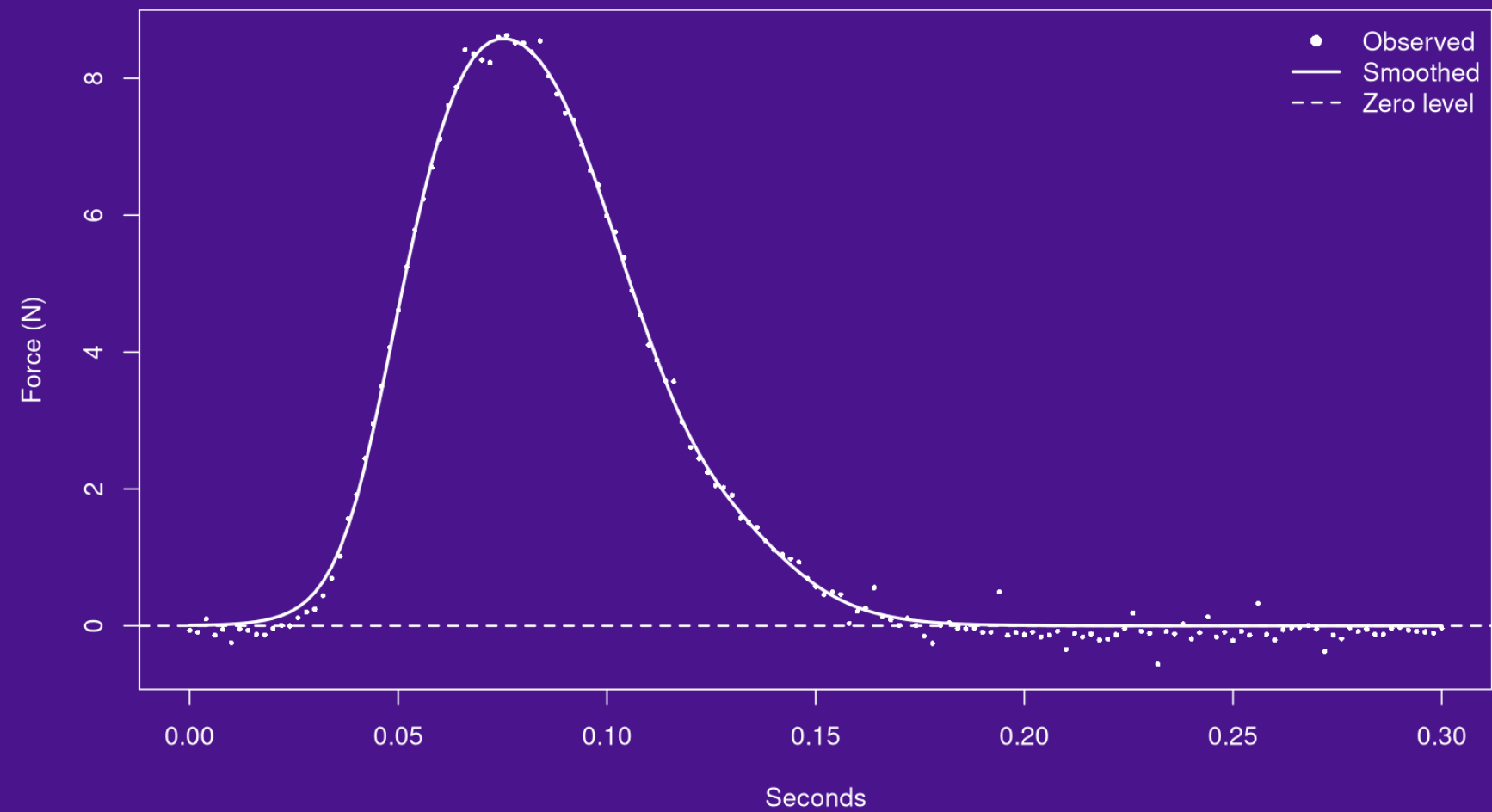


We are going to cover **chapter 6** in *Functional Data Analysis on Constrained Functions*

Focus:

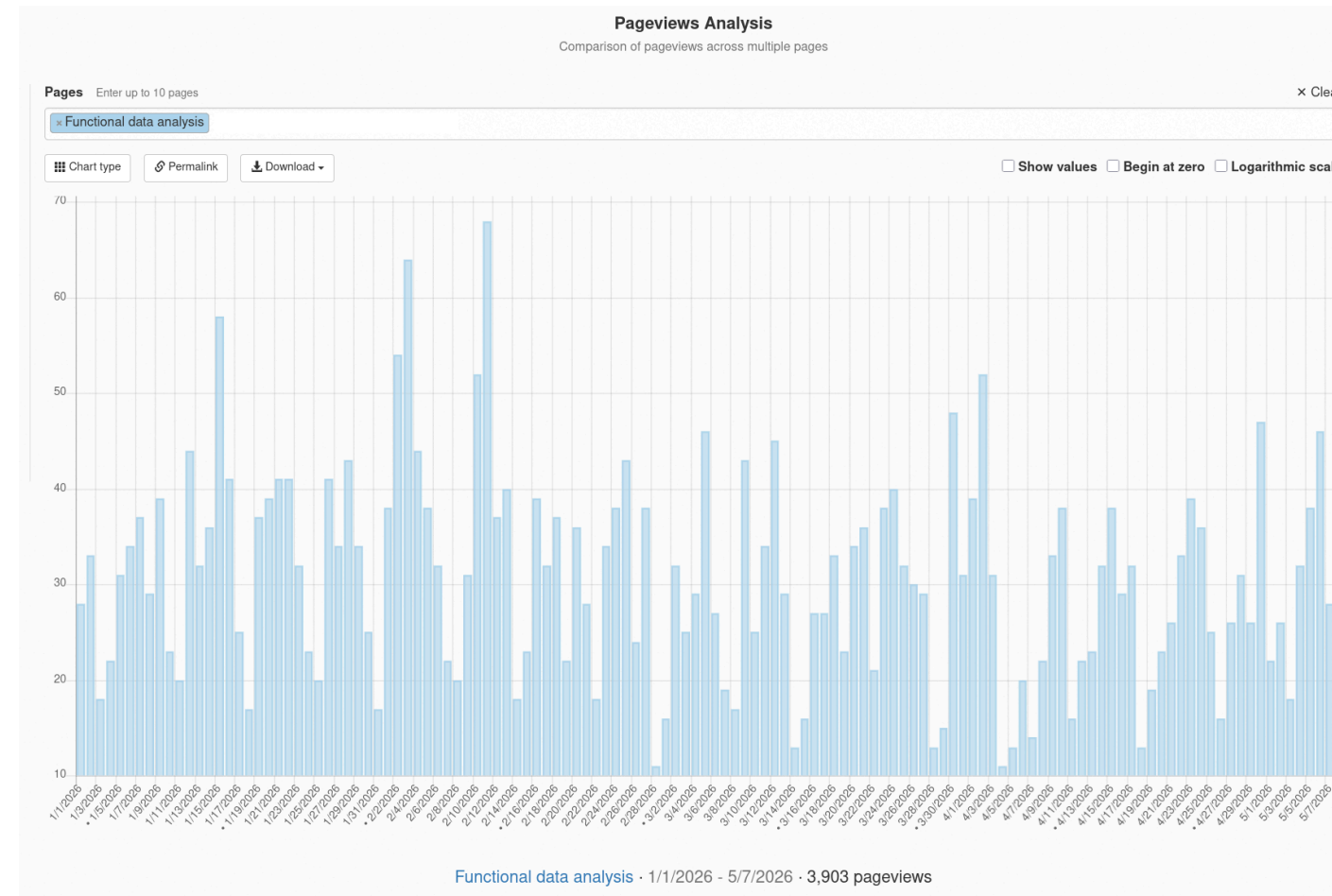
- Positive functions
- Monotonically increasing functions
- Probability density function estimation

Positive Functions



Which functions are positive? (1 / 3)

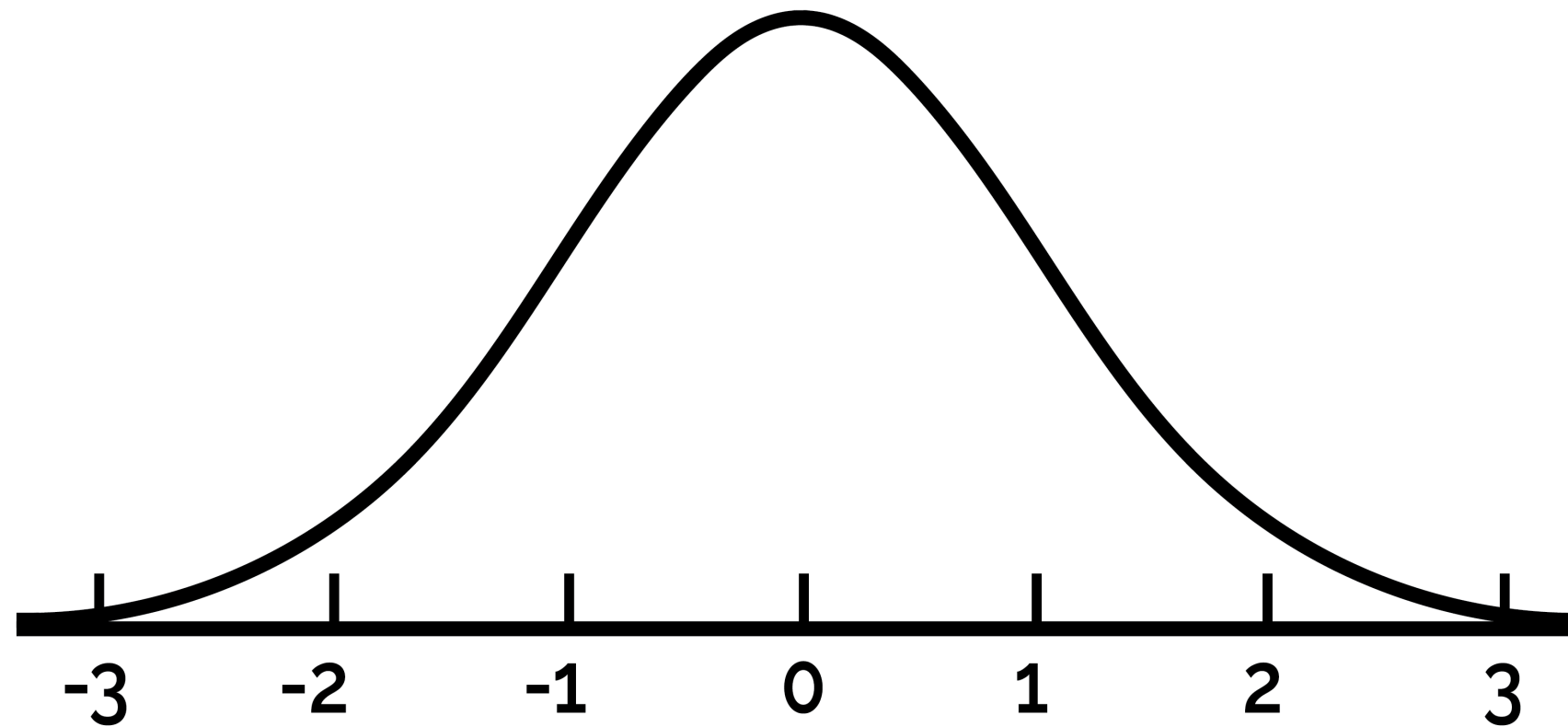
Counts, for example page views



Strictly speaking: This would be non-negative

Which functions are positive? (2 / 3)

Probability functions



Which functions are positive? (3 / 3)

Many quantities can only be positive

Energy

- Kinetic Energy
- Potential Energy

Measures

- Area
- Volume
- Length

Information Theoretic

- Entropy
- Mutual information
- Divergences

Pinch Force

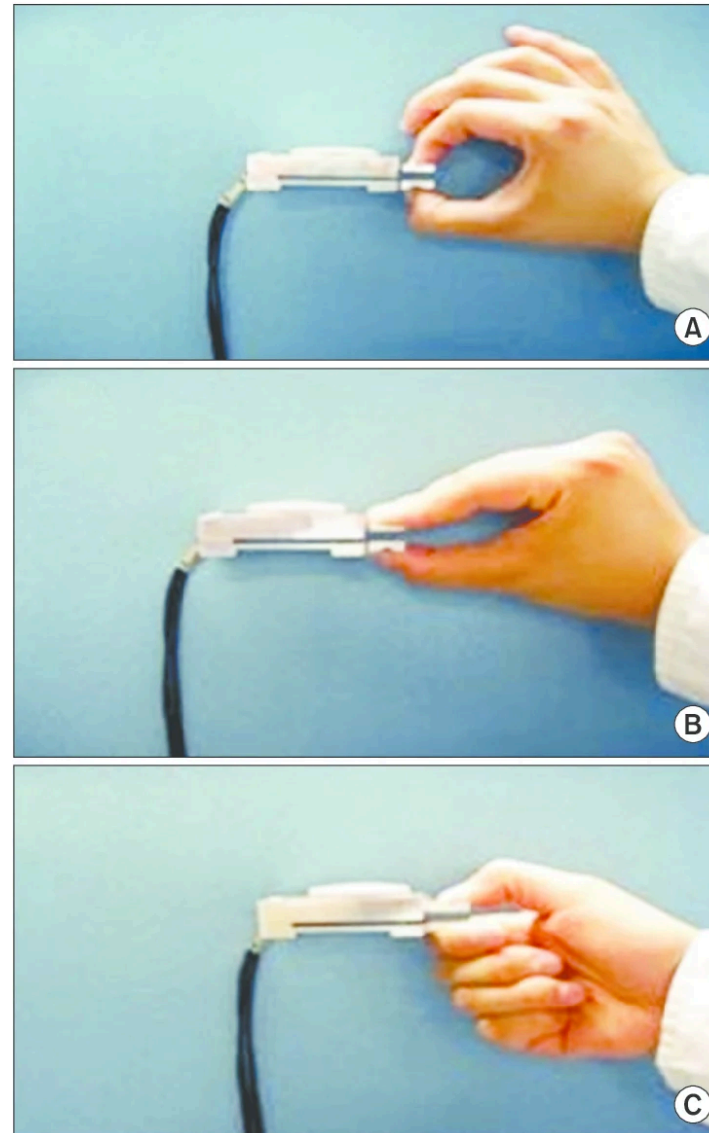
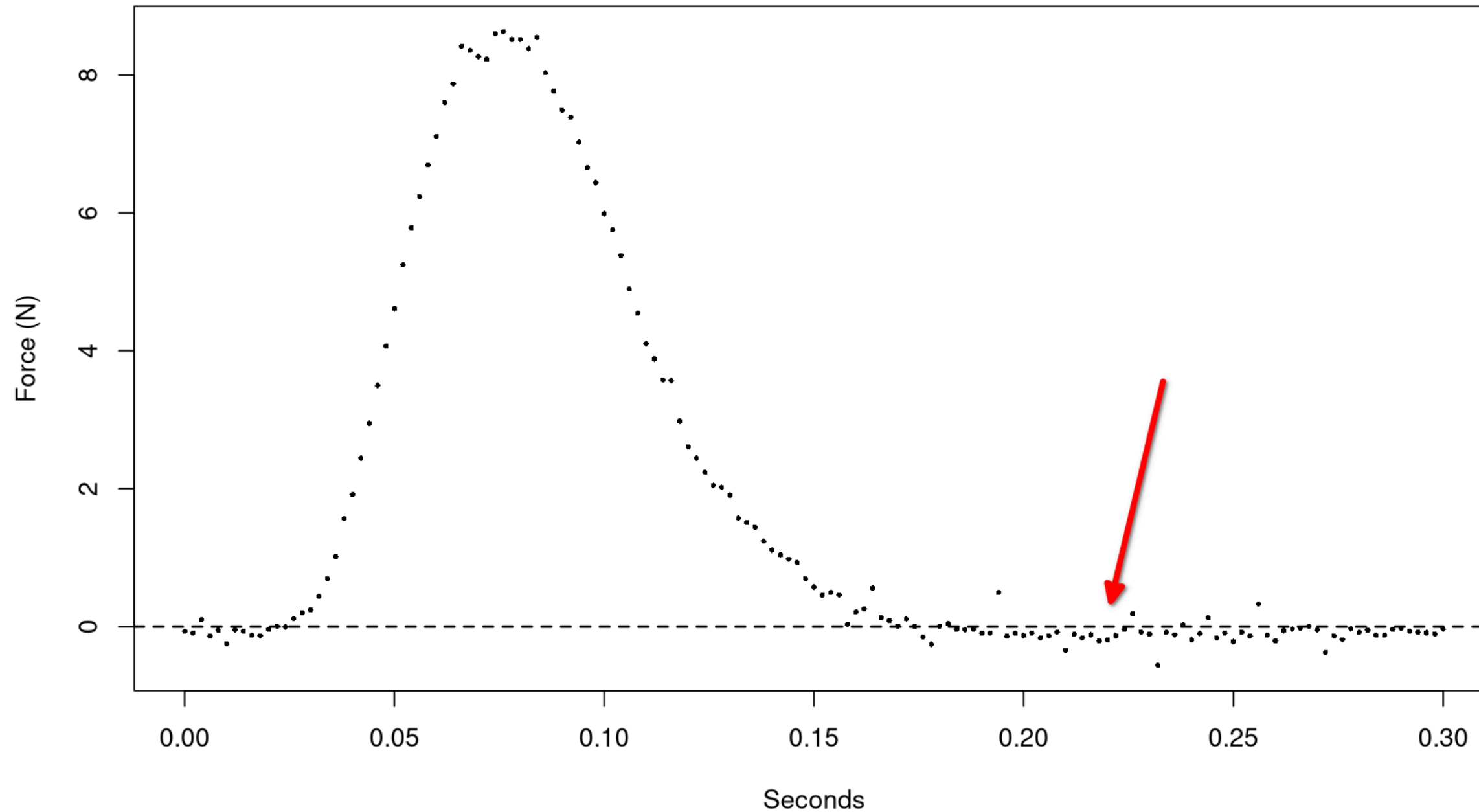


Image Source: Kong, Sangwon & Lee, Kyung & Kim, Junho & Jang, Seong. (2014). The Effect of Two Different Hand Exercises on Grip Strength, Forearm Circumference, and Vascular Maturation in Patients Who Underwent Arteriovenous Fistula Surgery. *Annals of Rehabilitation Medicine*. 38. 648. 10.5535/arm.2014.38.5.648.

Measurements can be noisy and drop below zero



The way we have fitted a function previously

Chapter 5 – smoothing with roughness penalty:

$$\text{PENSSE}_\lambda(x|\mathbf{y}) = \underbrace{[\mathbf{y} - x(\mathbf{t})]^T \mathbf{W} [\mathbf{y} - x(\mathbf{t})]^2}_{\text{Fit of the data}} + \lambda \underbrace{\int (D^2 x(s))^2 ds}_{\text{Function roughness based on curvature}}$$

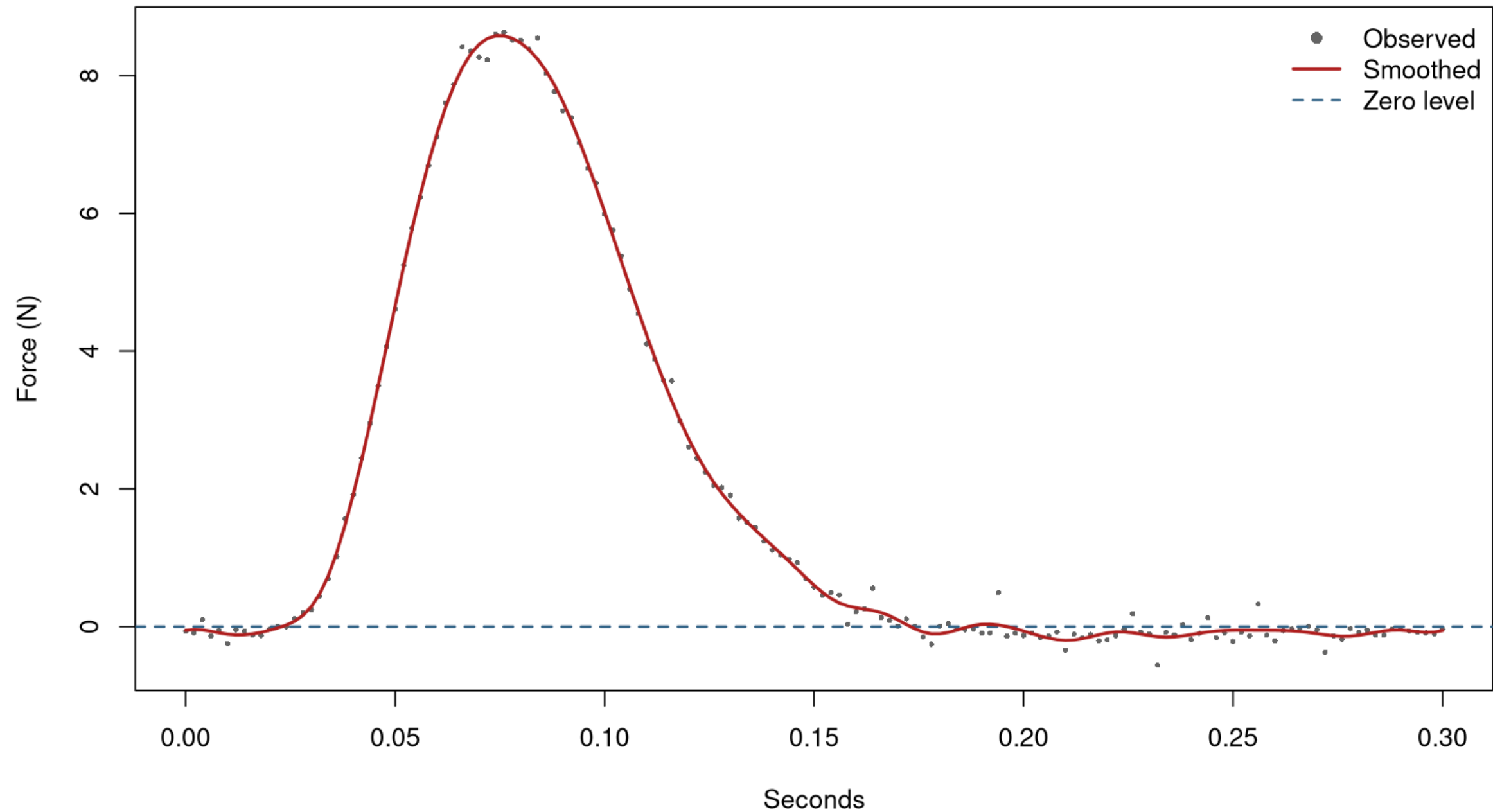
Remember, $x(t)$ is a linear expansion

$$x(t) = \sum_{k=1}^K c_k \phi_k(t)$$

In the FDA package

```
fdParobj <- fdPar(  
  fdobj = basis,  
  Lfdobj = 2,  
  lambda = 1e-8  
)  
  
fit <- smooth.basis(  
  argvals = x_values,  
  y = y_values,  
  fdParobj = fdParobj  
)
```

This creates some noisy tails



Defining positive smoothing functions

Instead of

$$x(t) = \sum_{k=1}^K c_k \phi_k(t)$$

we now define

$$x(t) = e^{\sum_{k=1}^K c_k \phi_k(t)} = e^{W(t)}$$

Positive smoothing function x can always be defined on the exponential of an unconstrained function W .

Remarks

$$x(t) = e^{W(t)}$$

Any other basis can be used as well!

Nota bene: This function can never be exactly zero. It can only get very close to zero.

The updated optimization criterion

$$\text{PENSSE}_\lambda(x|\mathbf{y}) = [\mathbf{y} - x(\mathbf{t})]^T \mathbf{W} [\mathbf{y} - x(\mathbf{t})]^2 + \lambda \int (D^2 x(s))^2 ds$$

now becomes

$$\text{PENSSE}_\lambda(\mathbf{W}|\mathbf{y}) = [\mathbf{y} - e^{\mathbf{W}(\mathbf{t})}]^T \mathbf{W} [\mathbf{y} - e^{\mathbf{W}(\mathbf{t})}]^2 + \lambda \int (D^2 \mathbf{W}(t))^2 dt$$

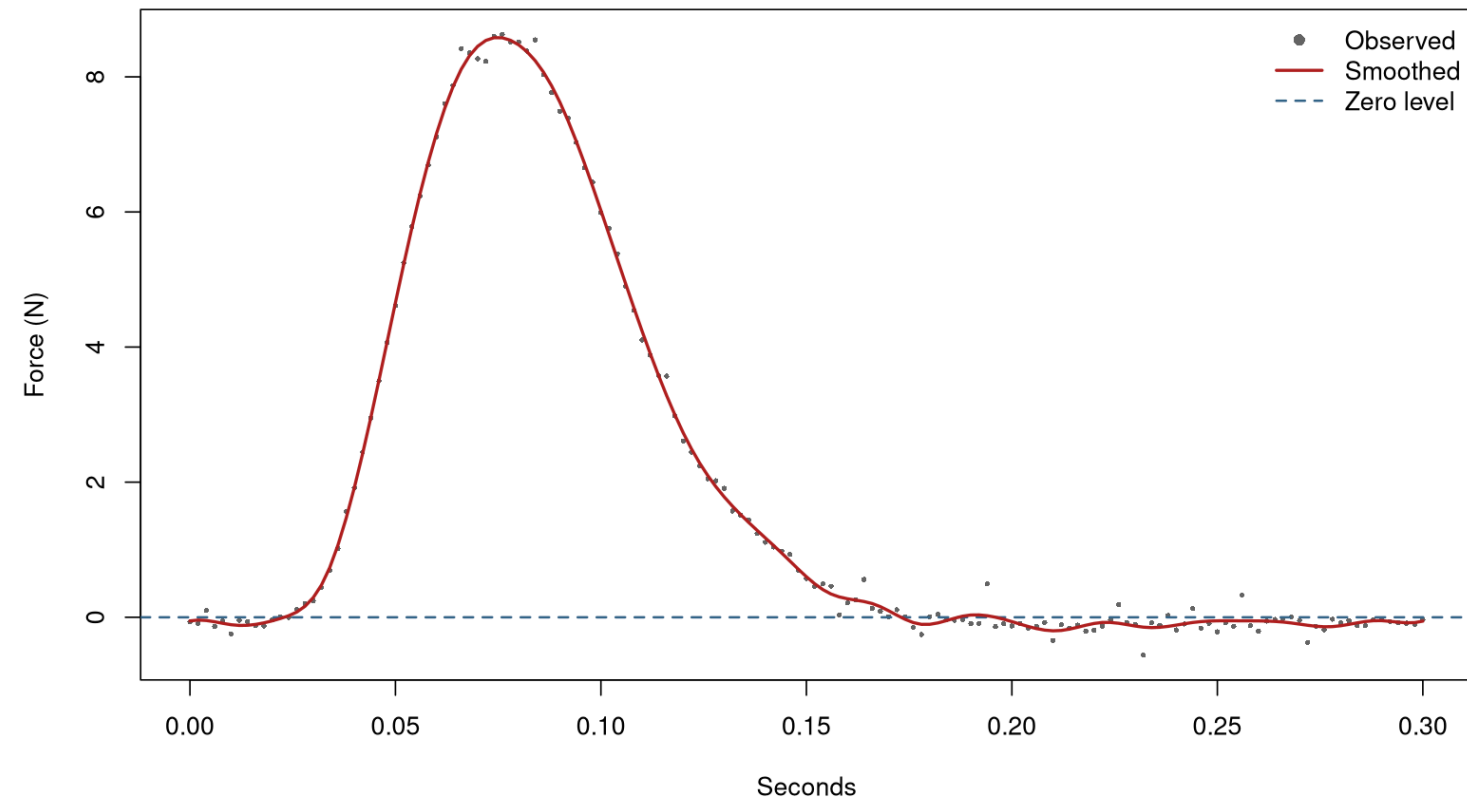
Needs to be solved through numerical methods. But these converge quickly because the expression is only mildly nonlinear.

smooth.basis \rightarrow smooth.pos

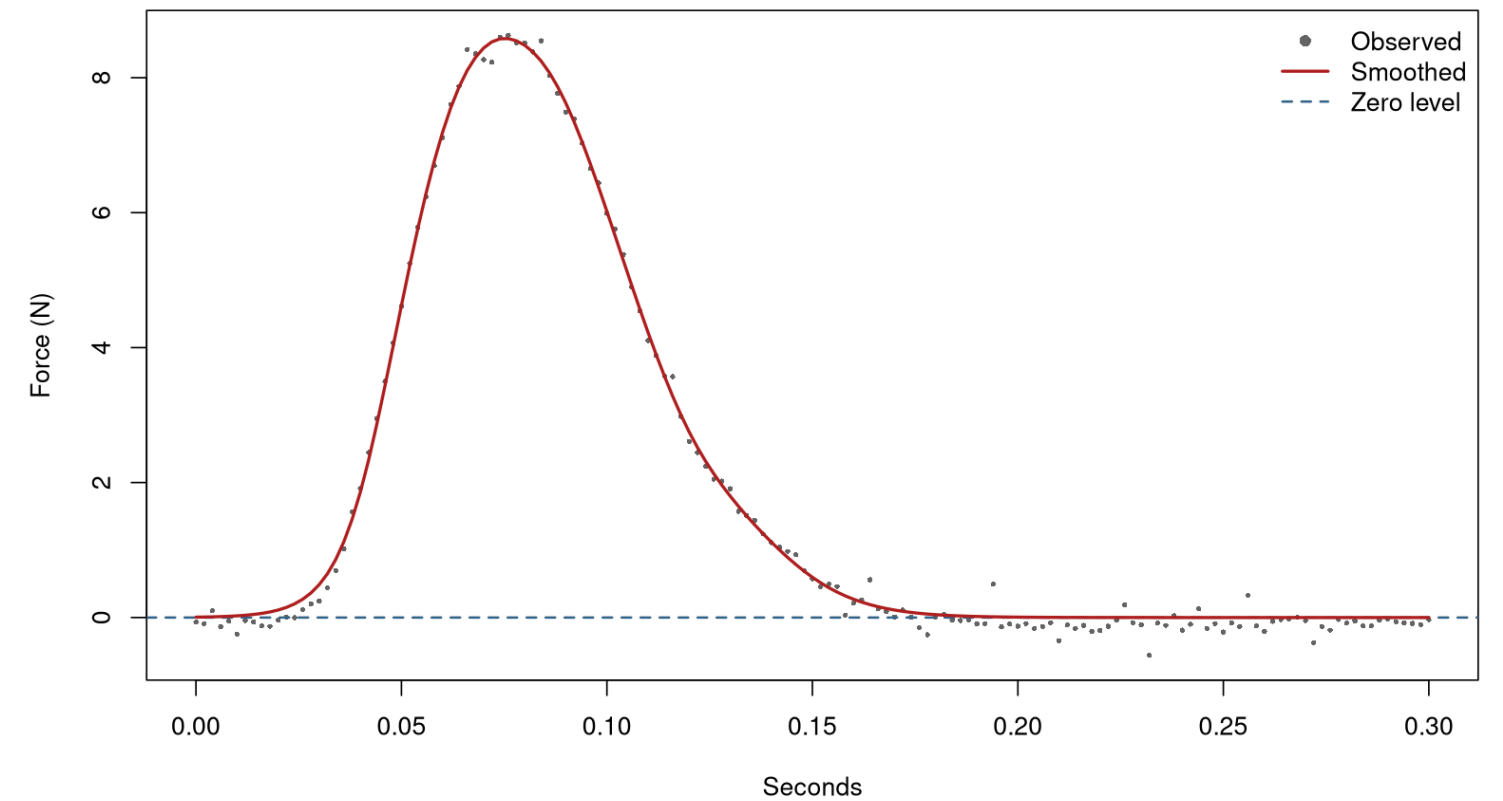
```
fdParobj <- fdPar(  
  fdobj = basis,  
  Lfdobj = 2,  
  lambda = 1e-8  
)  
  
fit <- smooth.pos(  
  argvals = x_values,  
  y = y_values,  
  WfdParobj = fdParobj,  
  dbglev = 0  
)
```

Comparison

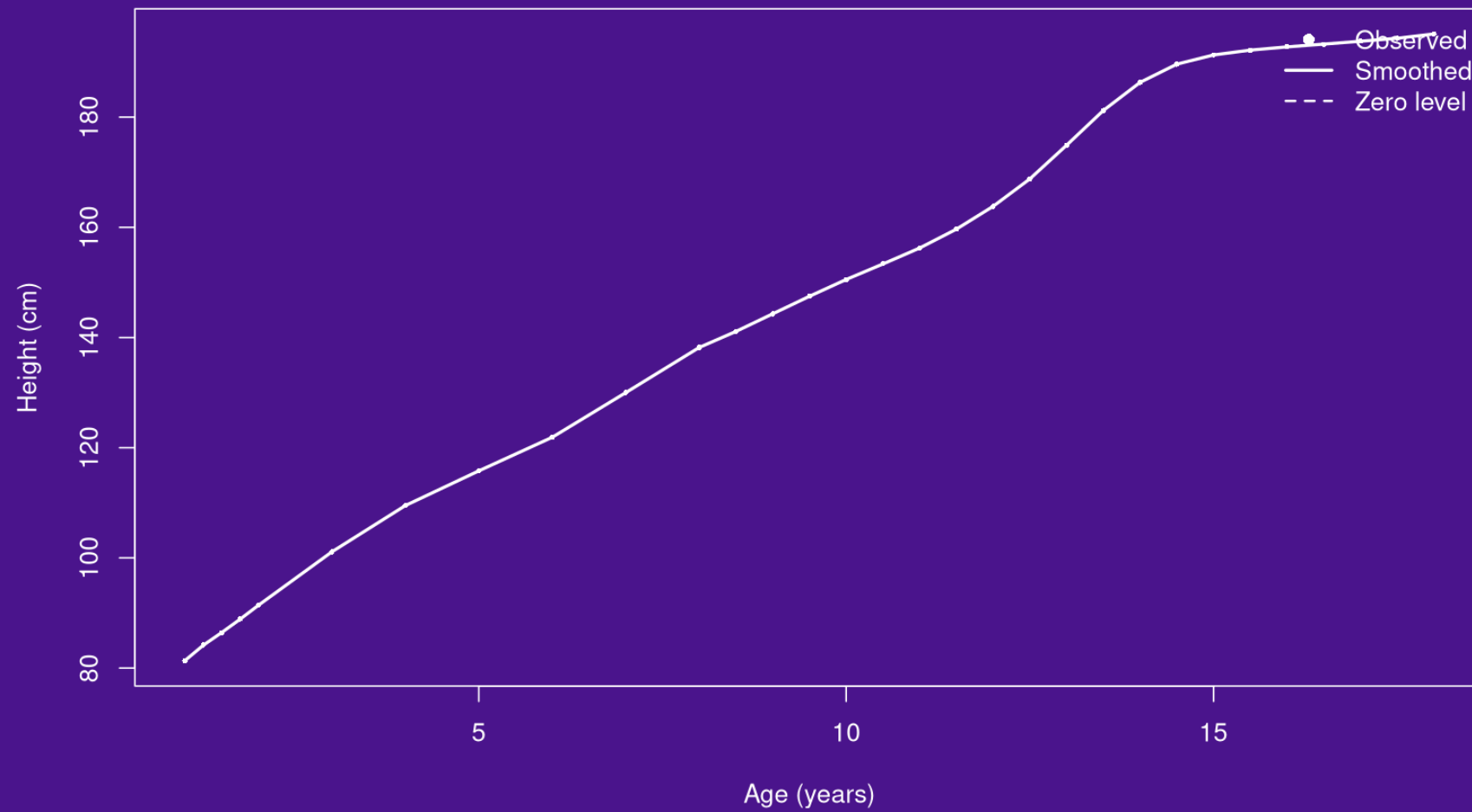
Unconstrained



Constrained

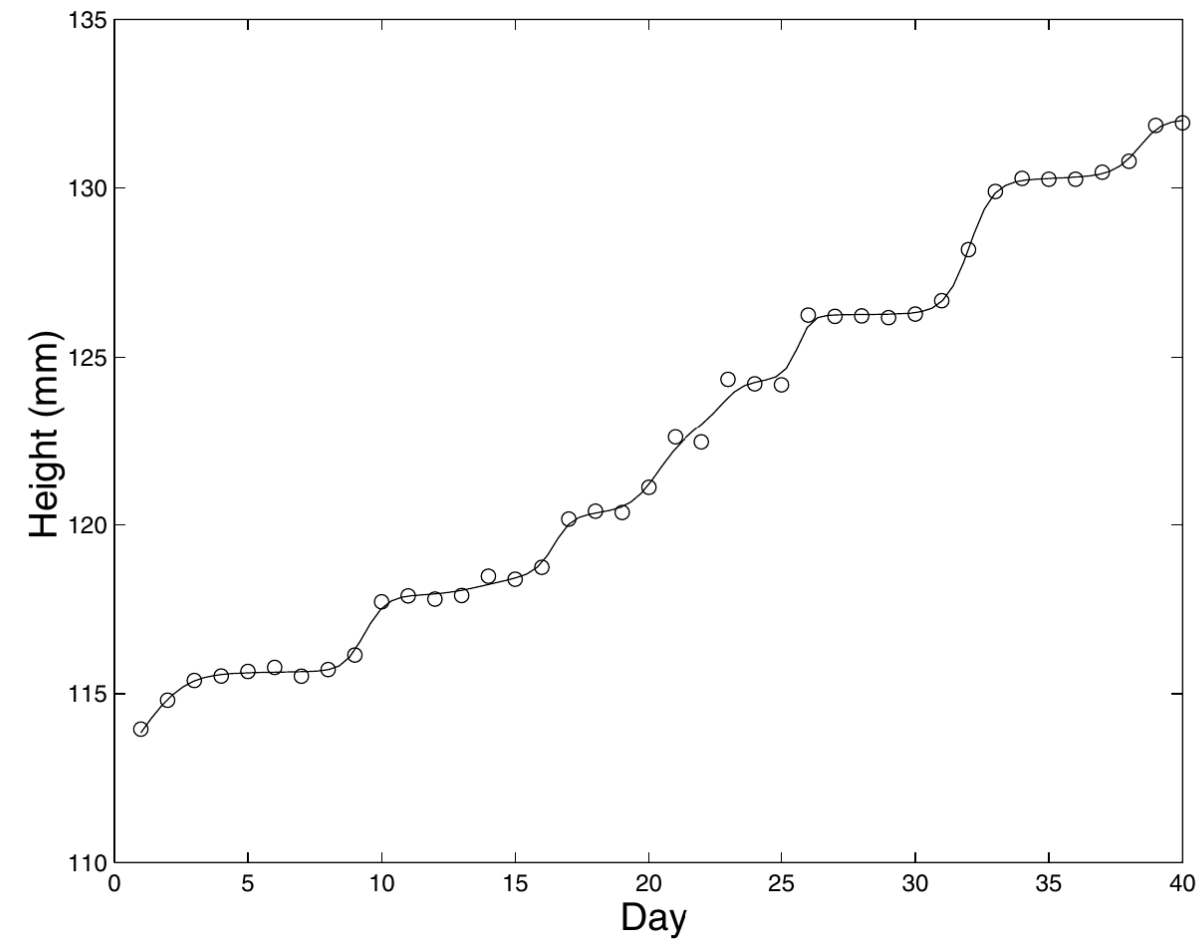


Monotone Functions



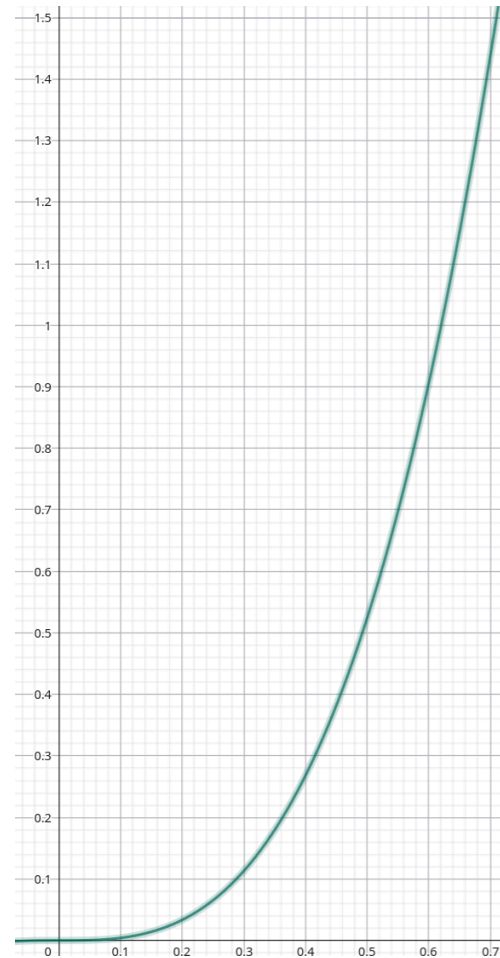
Which functions are monotone? (1 / 3)

Growth



Which functions are monotone? (2 / 3)

Growth

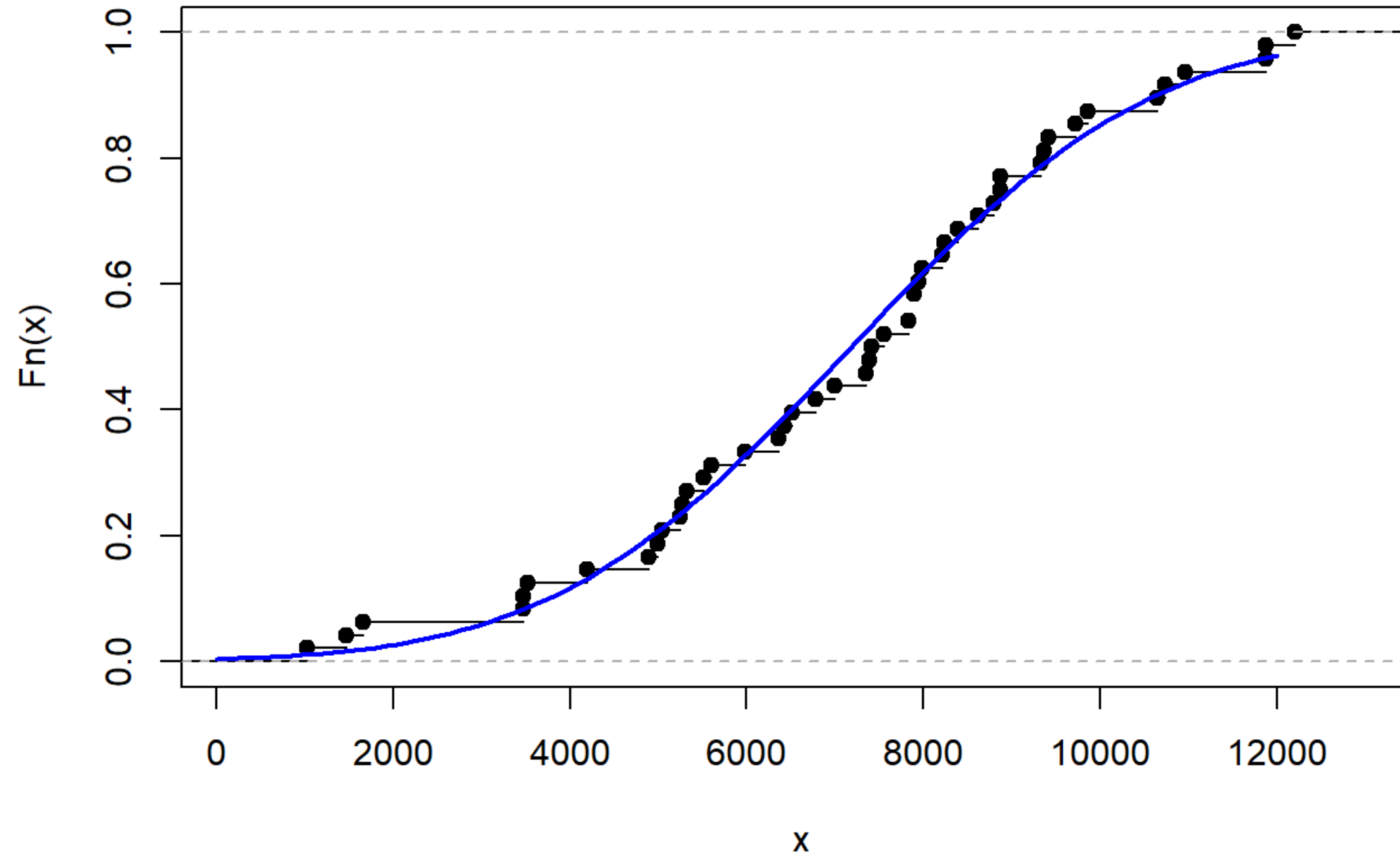


Volume is monotone increasing in the radius.

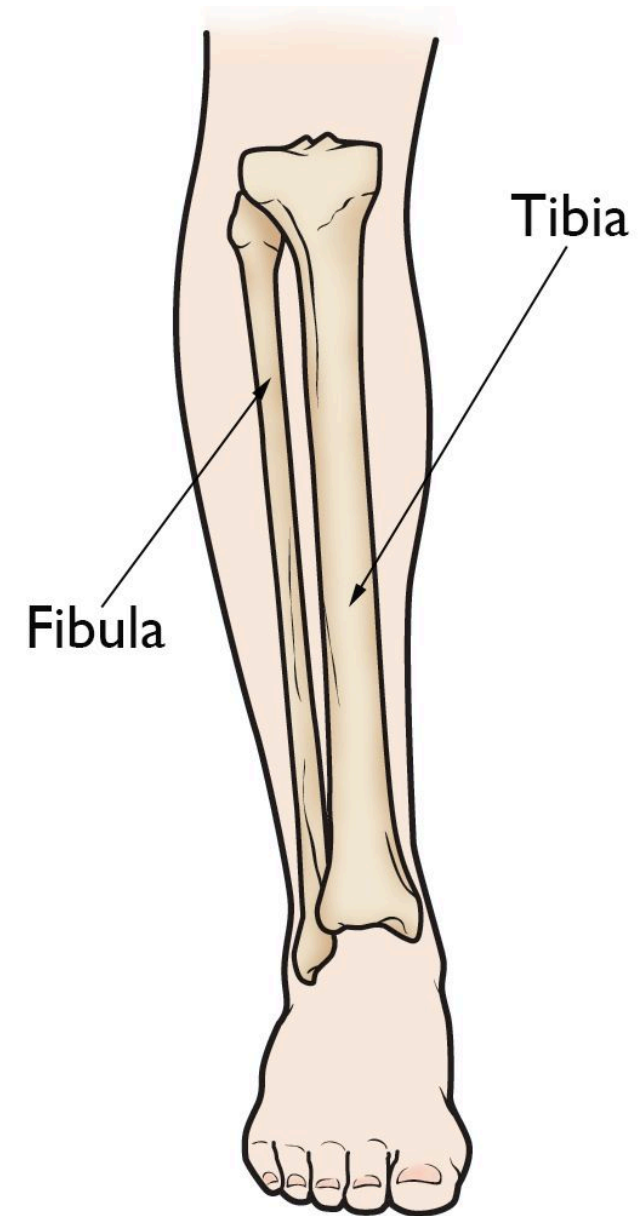
$$V(r) = \frac{4}{3}\pi r^3$$

Which functions are monotone? (3 / 3)

Cumulative Distribution Functions

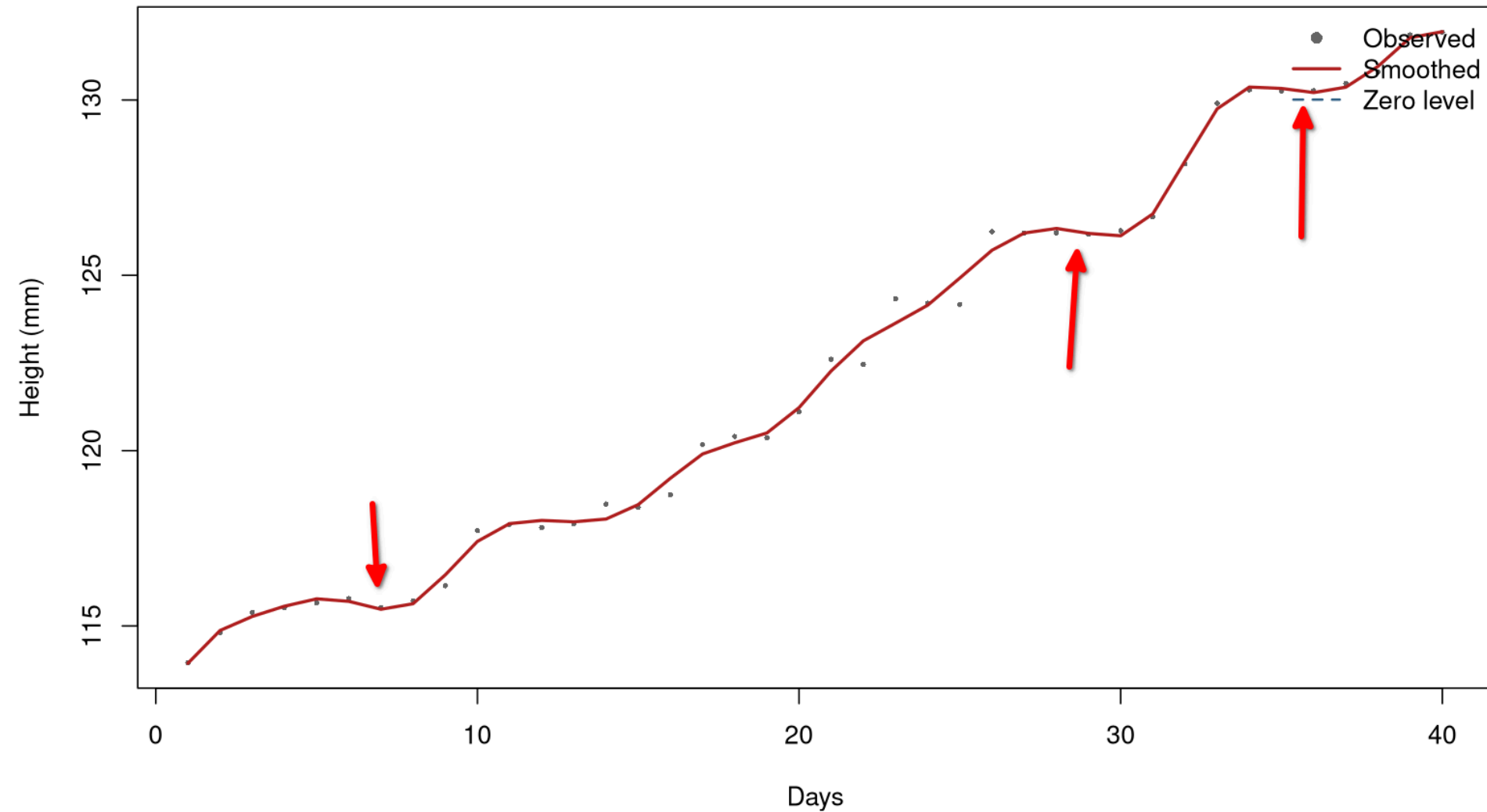


The Tibia Dataset



Fitting with the previous approach

Tibia growth for one infant



Ensuring monotone behavior

The first derivative must be positive everywhere.

$$Dx(t) = e^{W(t)}$$

Integrate both sides:

$$x(t) = c + \int_{t_0}^t e^{W(u)} du$$

c is a constant estimated from the data.

Depending on c , the function will also be positive everywhere once it crosses the 0-level.

But: A zero increase is not possible, the function is always **strictly** monotone.

The updated optimization criterion

$$\text{PENSSE}_\lambda(x|\mathbf{y}) = [\mathbf{y} - x(\mathbf{t})]^T \mathbf{W} [\mathbf{y} - x(\mathbf{t})]^2 + \lambda \int (D^2 x(s))^2 ds$$

now becomes

$$\begin{aligned} \text{PENSSE}_\lambda(\mathbf{W}|\mathbf{y}) = & \\ & \left[\mathbf{y} - \left(\mathbf{c} + \int_{t_0}^t e^{\mathbf{W}(u)} du \right) \right]^T \mathbf{W} \left[\mathbf{y} - \left(\mathbf{c} + \int_{t_0}^t e^{\mathbf{W}(u)} du \right) \right]^2 \\ & + \lambda \int (D^2 \mathbf{W}(t))^2 dt \end{aligned}$$

Needs to be solved through numerical methods.

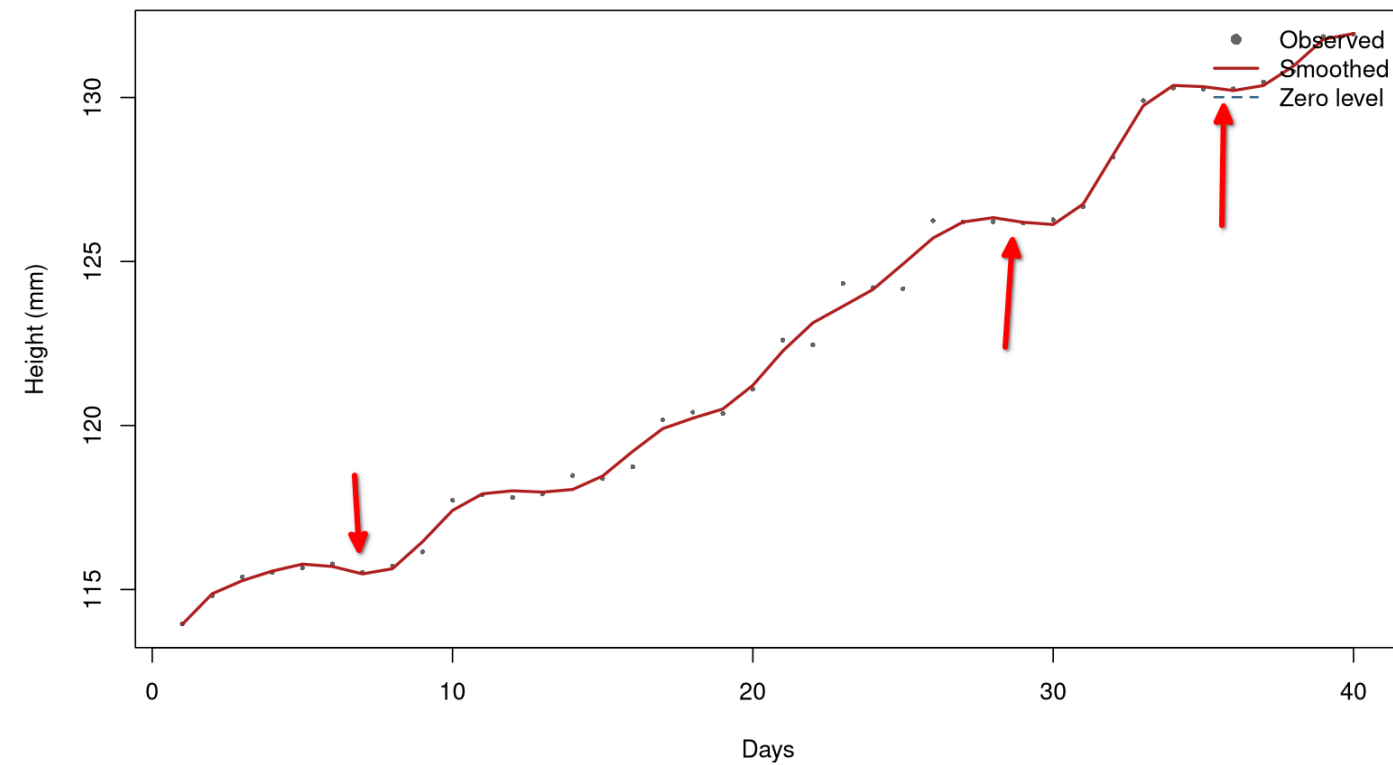
smooth.basis \rightarrow smooth.monotone

```
fdParobj <- fdPar(  
  fdobj = basis,  
  Lfdobj = 2,  
  lambda = 1e-8  
)  
  
fit <- smooth.monotone(  
  argvals = x_values,  
  y = y_values,  
  WfdParobj = fdParobj  
)
```

Comparison

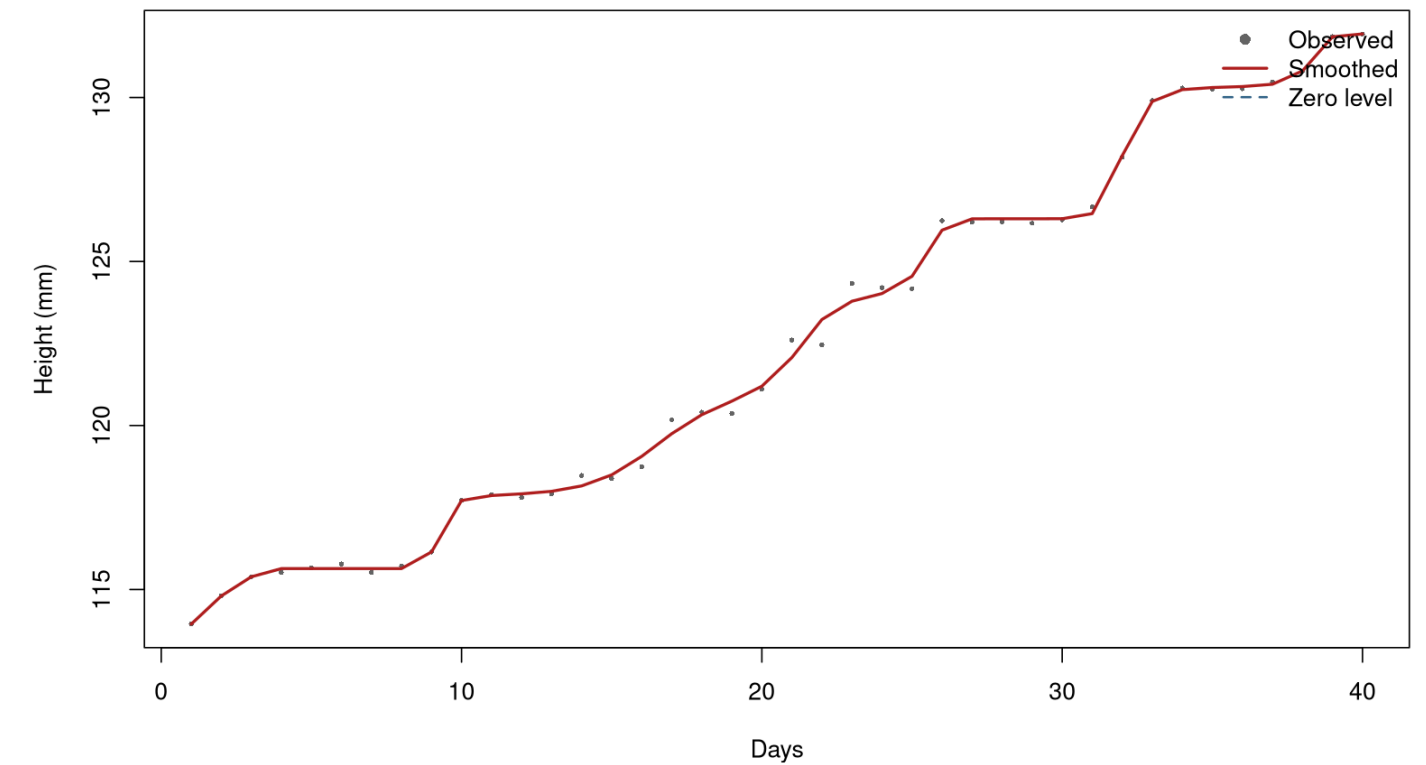
Unconstrained

Tibia growth for one infant

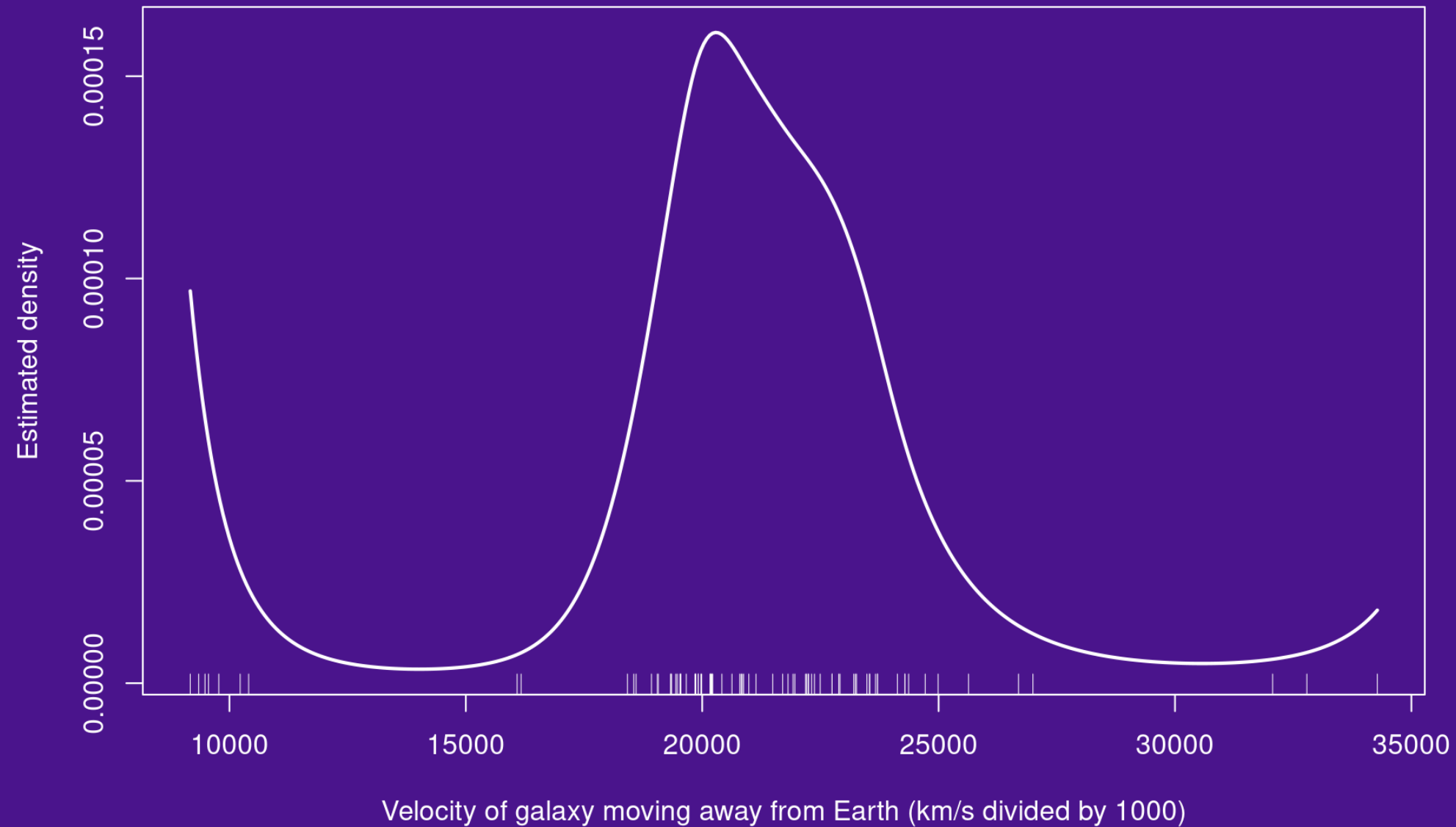


Constrained

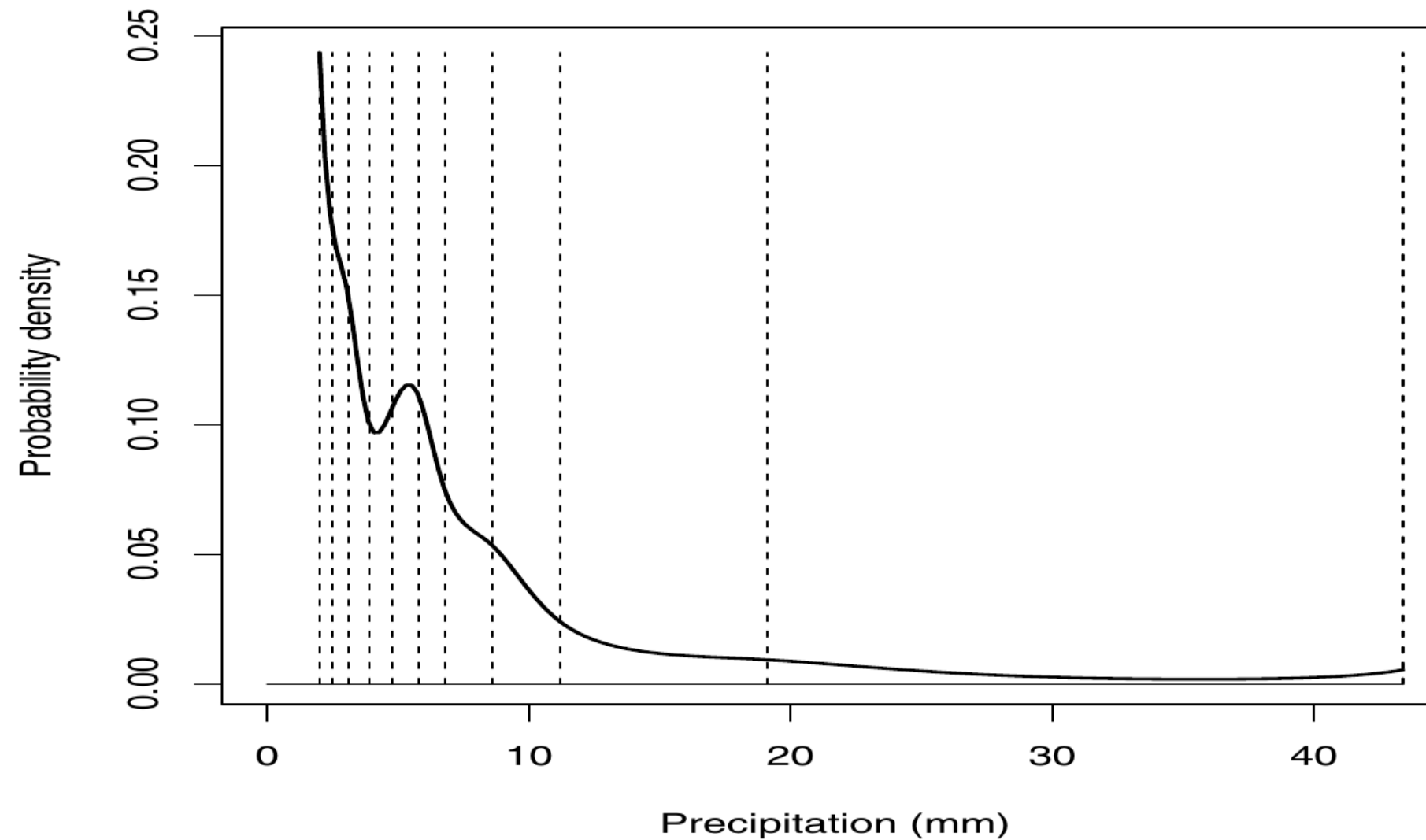
Tibia growth for one infant



Probability Density Functions



We now want to fit a probability density function



Each t has a probability of $p(t)$ to occur.

Two constraints for this function

$$p(t) > 0$$
$$\int p(t) dt = 1$$

$p(t)$ therefore looks like this:

$$p(t) = \frac{\overbrace{e^{W(t)}}^{\text{positive function, } p(t) > 0}}{\underbrace{\int e^{W(t)} dt}_{\text{normalizing constant, } \int p(t) dt = 1}}$$

This is our density model.

Remember our basis expansion

$$W(t) = \sum_{k=1}^K c_k \phi_k(t)$$

The goal is to find an optimal coefficient vector c for the model $p(t) = \frac{e^{W(t)}}{\int e^{W(t)} dt}$

Using **maximum likelihood estimation** is the way to go!

Likelihood Refresher

Likelihood for all N observations t_1, \dots, t_N :

$$L(W|c) = \prod_{i=1}^N p(t_i)$$

Maximum likelihood means finding a density that assigns high probability density to the observed samples.

Products are annoying

Therefore we apply the log to convert it to a sum:

$$\log L(W|c) = \sum_{i=1}^N \log p(t_i)$$

Insert our density model

$$\log L(W|c) = \sum_{i=1}^N \log p(t_i) = \sum_{i=1}^N \log \left(\frac{e^{W(t)}}{\int e^{W(t)} dt} \right)$$

$$\text{Apply } \log \left(\frac{A}{B} \right) = \log A - \log B$$

$$\begin{aligned} &= \sum_{i=1}^N \left[\log \left(e^{W(t)} \right) - \log \left(\int e^{W(t)} dt \right) \right] \\ &= \sum_{i=1}^N [W(t)] - N \log \left(\int e^{W(t)} dt \right) \end{aligned}$$

Insert basis expansion

$$\log L(W|c) = \sum_{i=1}^N [\mathbf{c}^T \phi(t_i)] - N \ln \left(\int e^{\mathbf{c}^T \phi(t)} dt \right)$$

So what do the terms mean?

$$\log L(W|\mathbf{c}) = \sum_{i=1}^N [\mathbf{c}^T \phi(t_i)] - N \ln \left(\int e^{\mathbf{c}^T \phi(t_i)} dt \right)$$

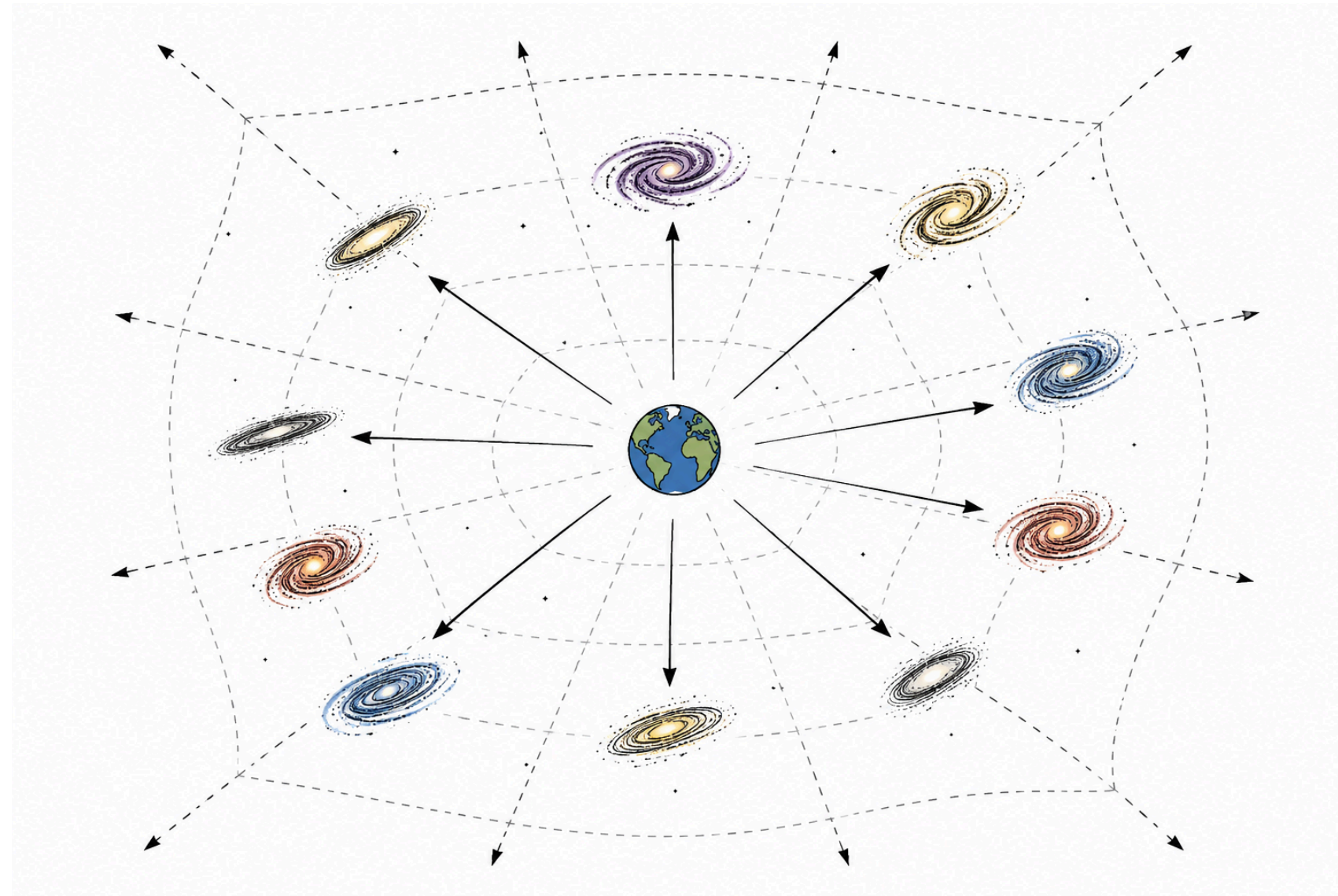
$$\sum_{i=1}^N [\mathbf{c}^T \phi(t_i)]$$

Rewards making W large where data points occur.

$$N \ln \left(\int e^{\mathbf{c}^T \phi(t_i)} dt \right)$$

Normalization penalty; without it, $W \rightarrow \infty$ everywhere, the term forces the density to integrate to 1.

Let's apply this to the R galaxies dataset!



82 observations of the velocities (in 1000 km/second) of distant galaxies diverging from our own

The dataset is a simple list of velocities



Velocity of galaxy moving away from Earth (km/s divided by 1000)

Constructing the R script (1)

Preliminaries

```
library(fda)  
library(MASS)  
data(galaxies)  
dataset <- galaxies  
N <- length(dataset)
```

Constructing the R script (2)

Construct a B-Spline basis

```
Wknots <- dataset[round(N * seq(1 / N, 1, length.out = 11), 0)]  
Wnbasis <- length(Wknots) + 2  
Wbasis <- create.bspline.basis(  
  rangeval = range(dataset),  
  nbasis   = Wnbasis,  
  norder   = 4,  
  breaks   = Wknots  
)
```

Constructing the R script (3)

Then we compute the fit using `density.fd`

```
Wlambda <- 0
Wfd0 <- fd(matrix(0, Wnbasis, 1), Wbasis)
WfdPar <- fdPar(Wfd0, Lfdobj = 2, lambda = Wlambda)

# Fit the density with density.fd.
# The result contains:
# - Wfdobj: fitted log-density component  $W(x)$ 
# - C: normalizing constant
densityList <- density.fd(dataset, WfdPar)
```

Attention: `density.fd` is only available up until version 6.2.0. It has been removed in version 6.3.0.

Constructing the R script (4)

Evaluate the fitted fda density on a fine grid.

```
Wfd <- densityList$Wfdobj
Cconst <- densityList$C
Zfine <- seq(min(dataset), max(dataset), length.out = 401)
Wfine <- eval.fd(Zfine, Wfd)
Pfine <- exp(Wfine) / Cconst
```

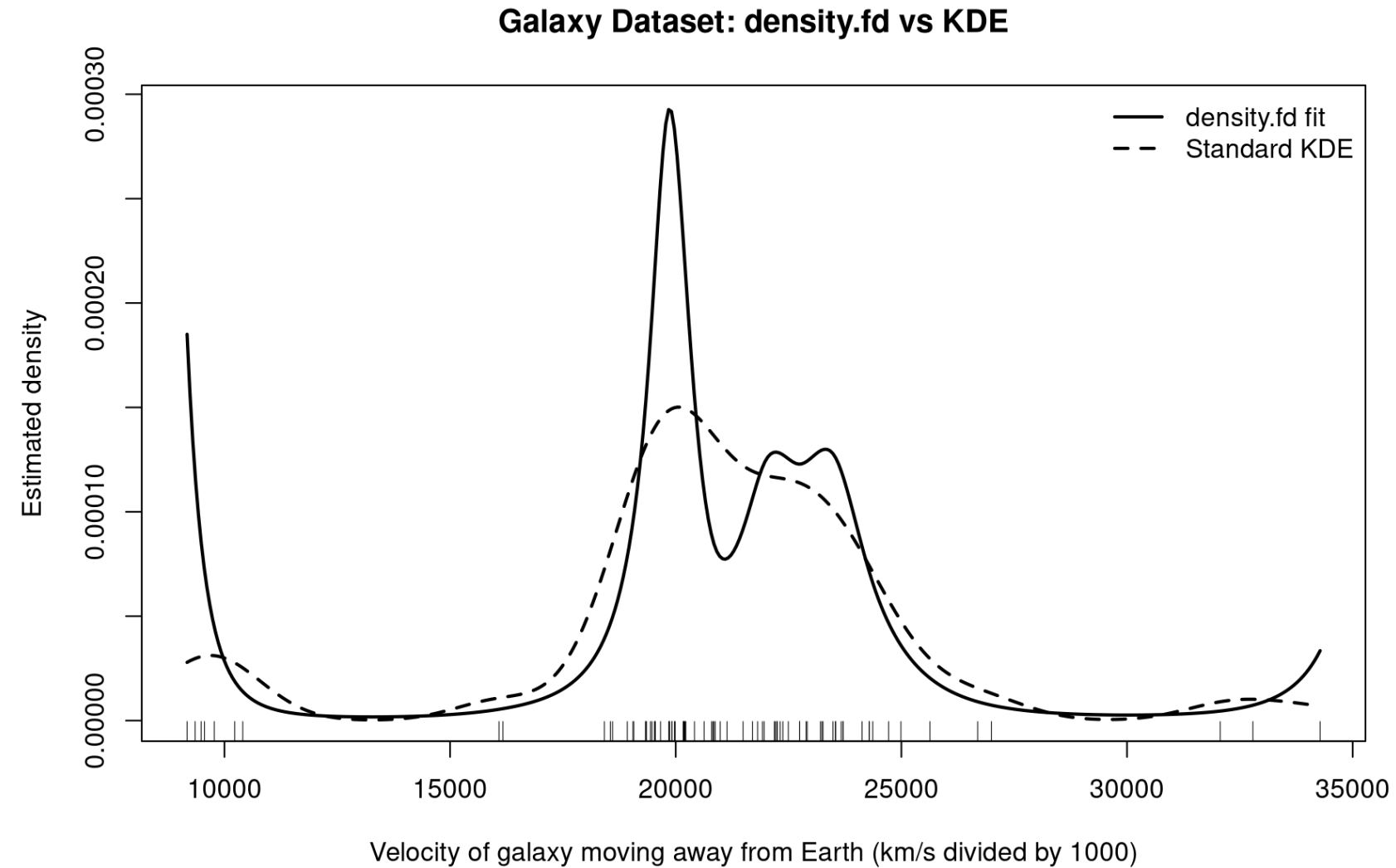
densityList only gives you $W(t)$ and the normalization constant, but not $p(t)$, so you need to stitch it together yourself.

Constructing the R script (5)

Finally plot it.

```
plot(  
  Zfine, Pfine,  
  type = "l",  
  lwd = 2,  
  xlab = "Velocity of galaxy moving away from Earth (km/s divided by 1000)"  
  ylab = "Estimated density",  
  main = "Galaxy Dataset: density.fd vs KDE"  
)  
  
lines(kde_fit$x, kde_fit$y, lwd = 2, lty = 2)
```

The initial approach overfits badly!



No smoothness control!

Let's extend it with a roughness penalty

$$\log L(W|\mathbf{c}) = \sum_{i=1}^N [\mathbf{c}^T \phi(t_i)] - N \ln \left(\int e^{\mathbf{c}^T \phi(t_i)} dt \right) - \underbrace{\lambda \int [D^3(\mathbf{c}^T \phi(t_i))]^2 dt}_{\text{PEN}_3(W)}$$

$\text{PEN}_3(W)$ penalizes changes in curvature:

- less wiggles
- less oscillations
- less rapid shape changes

λ is the smoothing parameter (similar to bandwidth in KDE).

Find coefficient vector \mathbf{c} via numerical optimization.

The `density.fd` function accepts λ as a parameter.

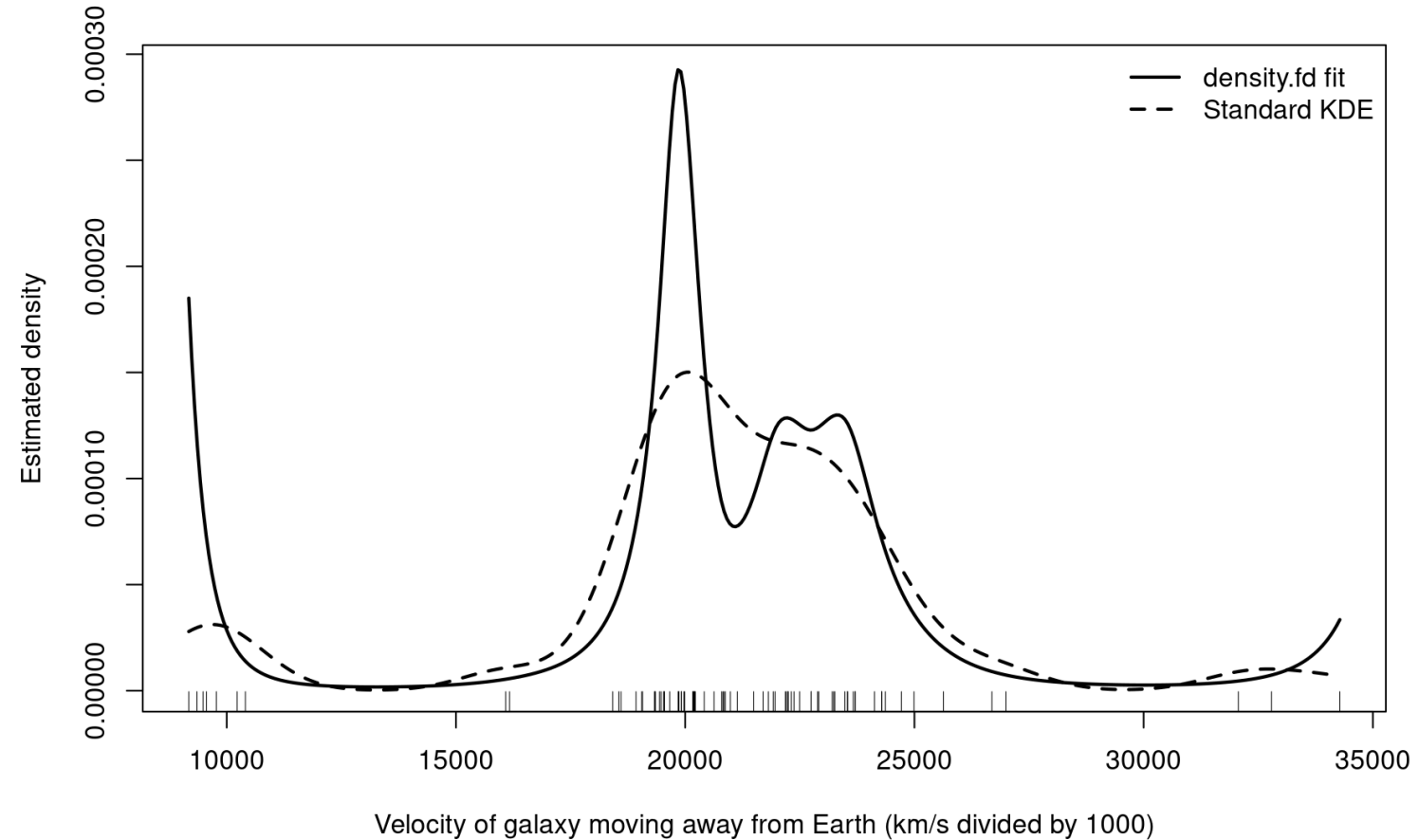
Adjust `Wlambda` accordingly

```
Wlambda <- 3e+9
Wfd0 <- fd(matrix(0, Wnbasis, 1), Wbasis)
WfdPar <- fdPar(Wfd0, Lfdobj = 2, lambda = Wlambda)

# Fit the density with density.fd.
# The result contains:
# - Wfdobj: fitted log-density component  $W(x)$ 
# - C: normalizing constant
densityList <- density.fd(dataset, WfdPar)
```

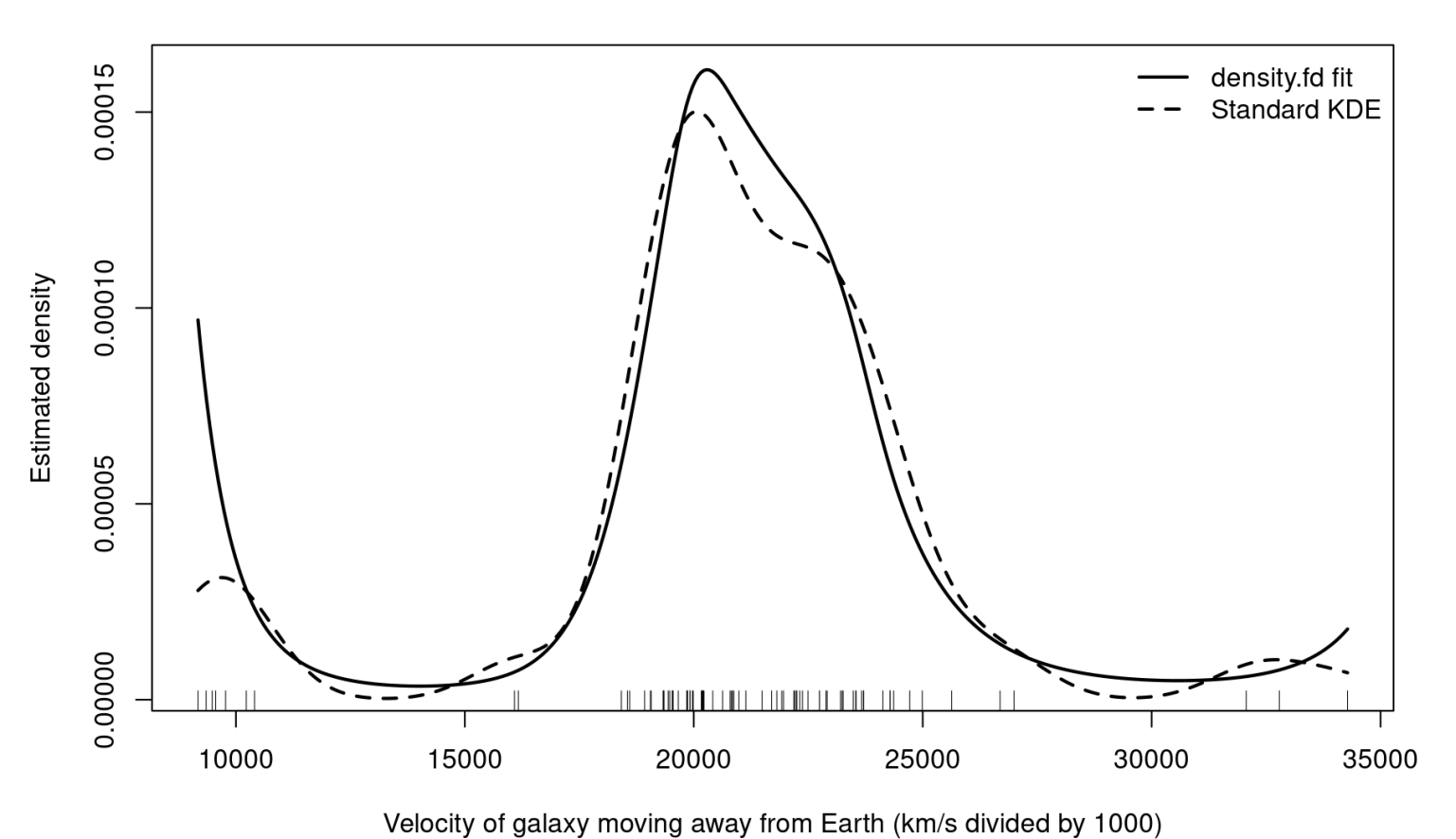
The effect of λ

Galaxy Dataset: density.fd vs KDE



$$\lambda = 0$$

Galaxy Dataset: density.fd vs KDE



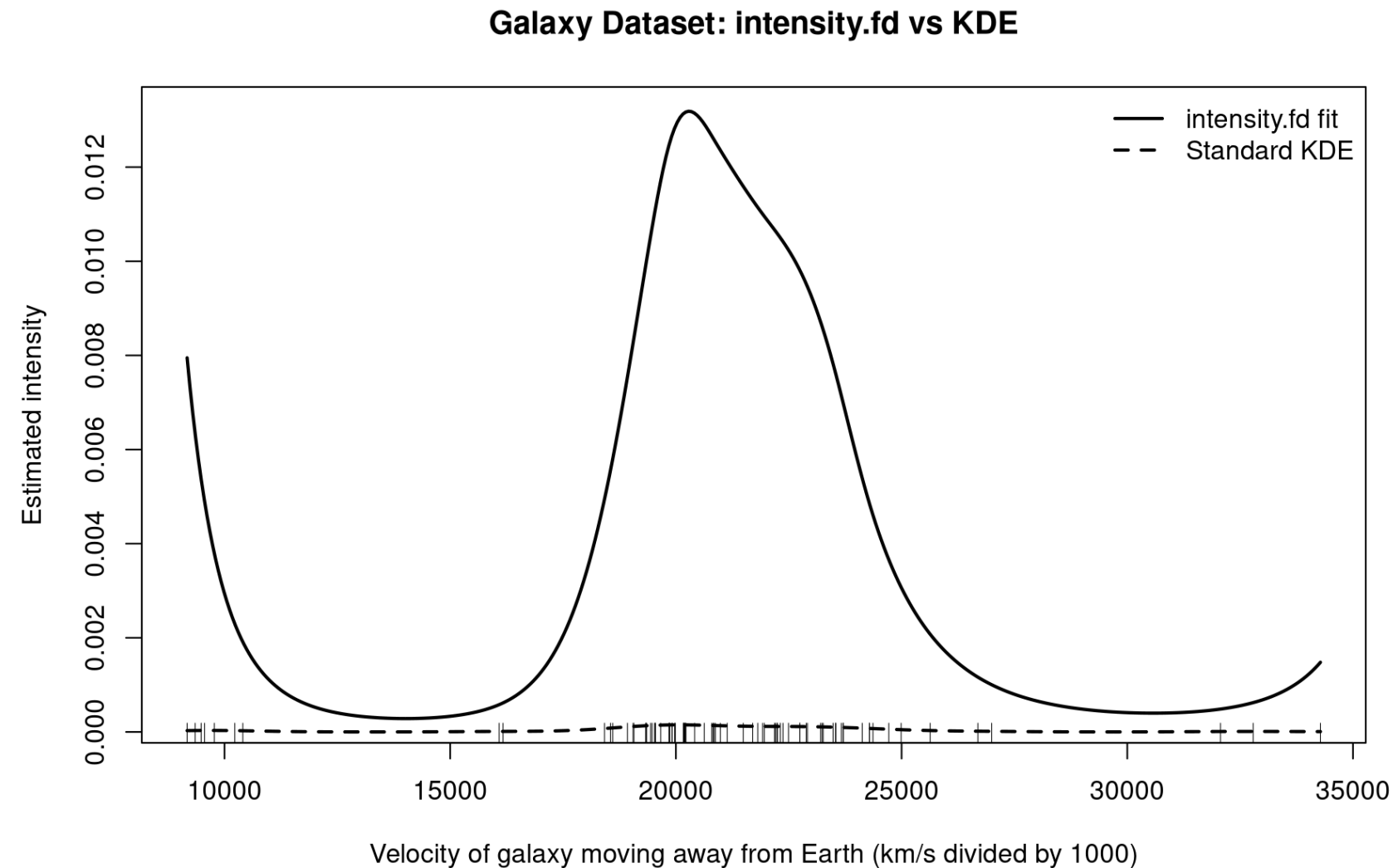
$$\lambda = 3 \times 10^9$$

For the fda package version 6.3.0, intensity.fd can be used

intensity.fd is essentially the same as density.fd, but does not normalize.

```
# Fit the intensity with intensity.fd.  
# The result contains:  
# - Wfdobj: fitted log-intensity component  $W(x)$   
densityList <- intensity.fd(dataset, WfdPar)  
  
Wfd <- densityList$Wfdobj  
  
# Evaluate the fitted intensity on a fine grid.  
Zfine <- seq(min(dataset), max(dataset), length.out = 401)  
Wfine <- eval.fd(Zfine, Wfd)  
Ifine <- as.numeric(exp(Wfine))
```

Same curve, but not normalized.



intensity.fd does not provide us a normalization constant.

Question to the experts here:

Why would we use this method instead of simply computing the usual KDE?

Exercise



Task 1

Task 2

Task 3

Questions?