

#### An introduction to Gaussian Processes

 $\label{lem:compared} And rew \ C \ Parnell \\ github.com/and rewcparnell/intro\_to\_gaussian\_processes$ 

# Why learn about Gaussian Processes?



- They are a great introduction to thinking about the multivariate normal distribution
- They are widely used in spatial statistics and machine learning
- They are very flexible and extendable
- Some of you will be using them for your PhD

To follow this presentation you will need to understand linear regression and a bit of matrix algebra

## Back to linear regression

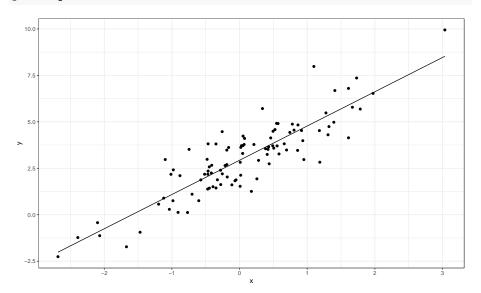


```
lr_dat = tibble(
  x = sort(rnorm(100)),
  y = rnorm(100, 3 + 2 * x, 1),
  fits = fitted(lm(y ~ x))
)
p = ggplot(lr_dat, aes(x, y)) +
  geom_point() +
  geom_line(aes(y = fits))
```

# Linear regression plot



#### print(p)



# Linear regression maths



We normally write this as:

$$y_i = \beta_0 + \beta_1 x_i + \epsilon_i$$
;  $i = 1, ..., n, \epsilon_i \sim N(0, \sigma^2)$ 

... so there are three parameters to estimate.

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... so there are three parameters to estimate.

Another way of writing this is:

$$y_i|x_i \sim N(\beta_0 + \beta_1 x_i, \sigma^2)$$

# Linear regression in matrices



#### Suppose we write:

$$y = \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{bmatrix}, X = \begin{bmatrix} 1 & x_1 \\ 1 & x_2 \\ \vdots \\ 1 & x_n \end{bmatrix}, \epsilon = \begin{bmatrix} \epsilon_1 \\ \epsilon_2 \\ \vdots \\ \epsilon_n \end{bmatrix}, \beta = \begin{bmatrix} \beta_0 \\ \beta_1 \end{bmatrix}$$

# Linear regression in matrices



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We can then write the model as:

$$\begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_n \end{bmatrix} = \begin{bmatrix} 1 & x_1 \\ 1 & x_2 \\ \vdots \\ 1 & x_n \end{bmatrix} \begin{bmatrix} \beta_0 \\ \beta_1 \end{bmatrix} + \begin{bmatrix} \epsilon_1 \\ \epsilon_2 \\ \vdots \\ \epsilon_n \end{bmatrix}$$

or

$$y = X\beta + \epsilon$$

# Fitting linear regression via maximum likelihood



One way to fit is to compute the (log) likelihood and maximise it:

$$\log L = \sum_{i=1}^{n} \log \left\{ \sqrt{\frac{1}{2\pi\sigma^2}} \exp \left[ -\frac{1}{2\sigma^2} (y_i - \beta_0 - \beta_1 x_i)^2 \right] \right\}$$

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```
## [1] -763.9474
```

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```
## [1] -140.7746
```

### LR via the multivariate normal distribution



If you simplify the calculation on the previous slide you get:

$$\log L = -\frac{n}{2}\log(\sigma^2) - \frac{1}{2\sigma^2}\sum_{i=1}^{n}(y_i - \beta_0 - \beta_1 x_i)^2$$

(I have dropped the the  $2\pi$  as it's just a constant)

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This can be re-written in matrix format

$$\log L = -\frac{1}{2} \log \left[ \det(\Sigma) \right] - \frac{1}{2} (y - X\beta)^{\mathsf{T}} \Sigma^{-1} (y - X\beta)$$

where  $\Sigma = \sigma^2 I$ 

#### LR via the multivariate normal distribution



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This is the likelihood for a multivariate normal distribution written  $y \sim MVN(X\beta, \Sigma)$ 

# Fitting linear regressions using the multivariate regressions using the multivariate



Two different ways to write the log likelihood:

```
beta 0 = 3; beta 1 = 2; sigma = 1
with(lr_dat, sum(dnorm(y, beta_0 + beta_1 * x, sd = sigma,
                       log = TRUE)))
```

```
## [1] -140.7746
```

# Fitting linear regressions using the multivariate r



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## [1] -140.7746
library(mvtnorm)
Sigma = diag(sigma, nrow(lr_dat))
X = with(lr dat, cbind(1,x))
beta = c(beta 0, beta 1)
with(lr_dat, dmvnorm(y, X%*%beta, Sigma,
                     log = TRUE))
```

```
## [1] -140.7746
```

# Changing the off-diagonals



The matrix Sigma currently looks like this:

$$\Sigma = \left[ \begin{array}{cccc} \sigma^2 & 0 & \dots & 0 \\ 0 & \sigma^2 & \dots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \dots & \sigma^2 \end{array} \right]$$

What would happen if we changed the off-diagonal terms? These represent the covariances (or scaled correlations) between the *y*-values

# Changing the off-diagonals



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What would happen if we changed the off-diagonal terms? These represent the covariances (or scaled correlations) between the y-values

#### Gaussian processes

The *key idea* in Gaussian Processes is to change the off-diagonals of  $\Sigma$  so that the you get higher correlations between  $y_i$  and  $y_j$  when  $x_i$  and  $x_j$  are close.

#### Autocovariance functions



The most common autocovariance function (ACF):

$$\Sigma_{ij} = au^2 \exp\left[-rac{(x_i - x_j)^2}{ heta}
ight]$$

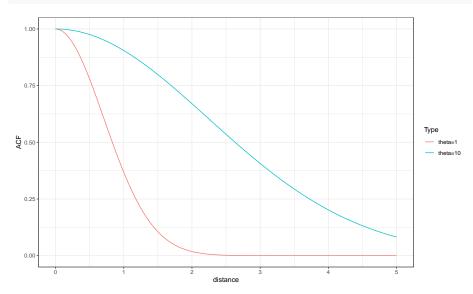
for  $i \neq j$  with  $\Sigma_{ii} = \tau^2 + \sigma^2$ 

This can be plotted:

```
sigma = 1; theta1 = 10; theta2 = 1
acv = tibble(
  distance = seq(0, 5, length = 100),
  `theta=10` = sigma^2 * exp(-(distance ^ 2) / theta1),
  `theta=1` = sigma^2 * exp(-(distance ^ 2) / theta2)
)
p = acv %>%
  gather(key = Type, value = ACF, -distance) %>%
  ggplot(aes(x = distance, y = ACF, colour = Type)) +
  geom_line()
```

# Pictures of squared exponential covariance funct

#### print(p)



## Fitting a model via the MVN



We can fit this model in exactly the same way via the log-likelihood:

```
suppressPackageStartupMessages(library(fields))
sigma = 1; tau = 1
Sigma = with(lr dat,
            diag(sigma^2, nrow(lr dat)) +
              tau^2 * Exp.cov(x, x,
                              theta = theta2, p = 2)
round(Sigma[1:3,1:3], 3)
##
       [,1] [,2] [,3]
## [1,] 2.000 0.918 0.710
## [2,] 0.918 2.000 0.918
## [3,] 0.710 0.918 2.000
with(lr dat, dmvnorm(y, X%*%beta, Sigma,
                    log = TRUE)
```

## The magic GP formula



Suppose you want to predict some new y values for some given new x values. Write these as  $y_{new}$  and  $x_{new}$ , and set up the problem as:

$$\left[\begin{array}{c} y \\ y_{new} \end{array}\right] \sim MVN\left(\left[\begin{array}{c} X\beta \\ X_{new}\beta \end{array}\right], \left[\begin{array}{cc} \Sigma & \Sigma_{new} \\ \Sigma_{new}^T & \Sigma_{new,new} \end{array}\right]\right)$$

where 
$$\Sigma_{new,i,j} = \tau^2 \exp\left[-\frac{(x_i - x_{new,j})^2}{\theta}\right]$$
,  $\Sigma_{new,new,i,j} = \tau^2 \exp\left[-\frac{(x_{new,i} - x_{new,j})^2}{\theta}\right]$ , and  $\Sigma_{new,new,i,i} = \sigma^2 + \tau^2$ 

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,  $\Sigma_{new,new,i,j} = \tau^2 \exp\left[-\frac{(x_{new,i} - x_{new,j})^2}{\theta}\right]$ , and  $\Sigma_{new,new,i,i} = \sigma^2 + \tau^2$ 

#### The magic formula gives you

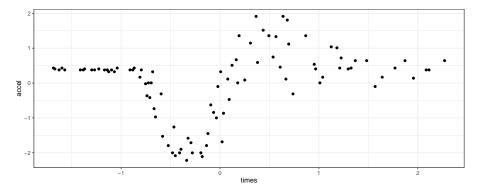
$$y_{new}|y \sim \textit{MVN}(X_{new}\beta + \Sigma_{new}^T \Sigma^{-1}(y - X\beta)), \Sigma_{new,new} - \Sigma_{new}^T \Sigma^{-1} \Sigma_{new})$$

This means that, once the parameters are estimated, we can produce instant predictions!

## A new regression data set



```
suppressPackageStartupMessages(library(boot))
scale2 = function(x) (x - mean(x))/sd(x)
motor2 = motor %>% mutate_all(scale2)
motor2 %>%
  ggplot(aes(x = times, y = accel)) +
  geom_point()
```



## Fitting a GP



```
nll = function(par) {
  tau = par[1]
  sigma = par[2]
  theta = par[3]
  Sigma = with(motor2,
             diag(sigma^2, nrow(motor2)) +
               tau^2 * Exp.cov(times, times,
                                theta = theta, p = 2)
  with(motor2, -dmvnorm(accel,
                         cbind(1. times)%*%beta.
                         sigma = Sigma, log = TRUE))
nll(c(1,1,1))
```

## [1] 138.4153

## Optimising the parameters



```
library(optimx)
answer = optimx(par = c(1,1,1),
                fn = nll,
                method = 'BFGS')
print(answer)
##
              p1
                         p2
                                   p3
                                         value fevals gevals r
  BFGS 3.441868 -0.4365699 0.7728277 81.24886
                                                    57
                                                            12
##
        kkt1 kkt2 xtime
## BFGS TRUE TRUE 0.102
```

# Making predictions using GPs



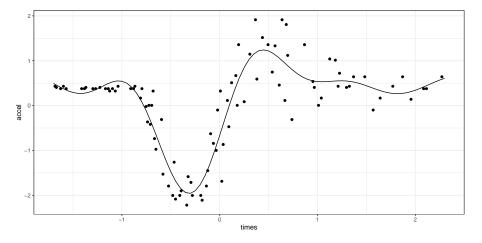
Now use the magic formula to predict for new  $\boldsymbol{x}$  values:

```
tau = answer$p1
sigma = answer$p2
theta = answer$p3
x_new = pretty(motor2\$times, n = 100)
Sigma new = with(motor2,
             tau^2 * Exp.cov(times, x_new,
                                 theta = theta, p = 2)
Sigma = with(motor2,
             diag(sigma^2,nrow(motor2)) + tau^2 * Exp.cov(time
                                 theta = theta, p = 2)
GP_pred = tibble(
  x_{new} = x_{new}
  mu pred = t(Sigma new) % * % solve(Sigma) % * % motor 2 $ accel
```

## Plot of predictions



```
ggplot(motor2, aes(x = times, y = accel)) +
geom_point() +
geom_line(data = GP_pred, aes(x = x_new, y = mu_pred))
```



# Shiny app



andrewcparnell.shinyapps.io/GP\_explore

or

 $is.gd/gp\_play$ 

# What else can you do with GPs?



- You can put the linear regression terms back in to the mean if you want
- If you drop the  $\sigma^2$  off the diagonal you can perform interpolation rather than smoothing
- The x values don't have to be one dimensional. Provided you can create a univariate distance between them you can run a GP. This is the foundation for spatial statistics and Kriging
- Lots of work on clever covariance functions
- Easy to make Bayesian by putting priors on parameters
- Big road block is calculation of  $\Sigma^{-1}$

# Summary



- You have now seen that a Gaussian Process is just a multivariate normal distribution
- The first clever trick is to make the observations correlated to each other by an autocovariance function
- The second clever trick is to use the magic multivariate normal formula to get instant access to predictions (and confidence intervals)

Thanks, and enjoy using Gaussian Processes!