

# MTH3045: Statistical Computing

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## Week 6 lecture 1

# Challenges I

- Go to Challenges I of the week 6 lecture 1 challenges at <https://byoungman.github.io/MTH3045/challenges>

## Sherman-Morrison formula / Woodbury matrix identity

- In general, unless we actually need the inverse of a matrix (such as for standard errors of regression coefficients in a linear model), we should solve systems of linear equations
- Sometimes, though, we do need – or might just have – an inverse, and want to calculate something related to it
- The following are a set of formulae, which go by various names, that can be useful in this situation

## Woodbury's formula

- Consider an  $m \times m$  matrix  $\mathbf{A}$ , with  $m$  large, and for which we have the inverse,  $\mathbf{A}^{-1}$
- Suppose  $\mathbf{A}$  receives a small update of the form  $\mathbf{A} + \mathbf{U}\mathbf{V}^T$ , for  $m \times n$  matrices  $\mathbf{U}$  and  $\mathbf{V}$  where  $n \ll m$
- Then, by **Woodbury's formula**,

$$(\mathbf{A} + \mathbf{U}\mathbf{V}^T)^{-1} = \mathbf{A}^{-1} - \mathbf{A}^{-1}\mathbf{U}(\mathbf{I}_n + \mathbf{V}^T\mathbf{A}^{-1}\mathbf{U})^{-1}\mathbf{V}^T\mathbf{A}^{-1}$$

- *Remark:* What's important to note here is that we're looking to calculate an  $m \times m$  inverse with  $m$  large, and so in general this will be an  $O(m^3)$  calculation based on the LHS
- However, in the above, the RHS only requires that we to invert an  $n \times n$  matrix, at cost  $O(n^3)$ , which is much less than that of the LHS, if we already have  $\mathbf{A}^{-1}$ .

# Sherman-Morrison-Woodbury formula

- Woodbury's formula above generalises to the so-called **Sherman-Morrison-Woodbury formula** by introducing the  $n \times n$  matrix **C**, so that

$$(\mathbf{A} + \mathbf{UCV}^T)^{-1} = \mathbf{A}^{-1} - \mathbf{A}^{-1}\mathbf{U}(\mathbf{C}^{-1} + \mathbf{V}^T\mathbf{A}^{-1}\mathbf{U})^{-1}\mathbf{V}^T\mathbf{A}^{-1}$$

# Challenges II

- Go to Challenges II of the week 6 lecture 1 challenges at <https://byoungman.github.io/MTH3045/challenges>



# Sherman-Morrison formula

- The **Sherman-Morrison formula** is the special case of Woodbury's formula (and hence the Woodbury-Sherman-Morrison formula) in which the update to  $\mathbf{A}$  can be considered in terms of  $m$ -vectors  $\mathbf{u}$  and  $\mathbf{v}$ , so that

$$(\mathbf{A} + \mathbf{u}\mathbf{v}^T)^{-1} = \mathbf{A}^{-1} - \frac{\mathbf{A}^{-1}\mathbf{u}\mathbf{v}^T\mathbf{A}^{-1}}{1 + \mathbf{v}^T\mathbf{A}^{-1}\mathbf{u}}$$

- *Remark:* The Sherman-Morrison formula is particularly useful because it requires no matrix inversion

## Example: Bayesian linear regression I

- Recall from MTH2006 the normal linear model where

$$Y_i \sim N(\mathbf{x}_i^T \boldsymbol{\beta}, \sigma^2)$$

with independent errors  $\varepsilon_i = Y_i - \mathbf{x}_i^T \boldsymbol{\beta}$ , for  $i = 1, \dots, n$ , where  $\mathbf{x}_i^T = (1, x_{i1}, \dots, x_{ip})$  is the  $i$ th row of design matrix  $\mathbf{X}$  and where  $\boldsymbol{\beta} = (\beta_0, \beta_1, \dots, \beta_p)^T$

- Hence

$$\mathbf{Y} \mid \mathbf{X}\boldsymbol{\beta} \sim MVN_n(\mathbf{X}\boldsymbol{\beta}, \sigma^2 \mathbf{I}_n)$$

- In Bayesian linear regression, the elements of  $\boldsymbol{\beta}$  and  $\sigma^2$  are not fixed, unknown parameters: they are random variables, and we must declare *a priori* our beliefs about their distribution
- The conjugate prior is that

$$\boldsymbol{\beta} \sim MVN_{p+1}(\boldsymbol{\mu}_\beta, \boldsymbol{\Sigma}_\beta^{-1})$$

- Integrating out  $\boldsymbol{\beta}$  gives

$$\mathbf{Y} \sim MVN_n(\mathbf{X}\boldsymbol{\mu}_\beta, \sigma^2 \mathbf{I}_n + \mathbf{X}\boldsymbol{\Sigma}_\beta^{-1}\mathbf{X}^T)$$

## Example: Bayesian linear regression II

- Now suppose that we want to evaluate the marginal likelihood for an observation,  $\mathbf{y}$ , say. Recall the MVN pdf
- The Mahalanobis distance then involves the term

$$(\mathbf{y} - \mathbf{X}\boldsymbol{\mu}_{\beta})^T (\sigma^2 \mathbf{I}_n + \mathbf{X}\boldsymbol{\Sigma}_{\beta}^{-1} \mathbf{X}^T)^{-1} (\mathbf{y} - \mathbf{X}\boldsymbol{\mu}_{\beta})$$

- The covariance of the marginal distribution,  $\sigma^2 \mathbf{I}_n + \mathbf{X}\boldsymbol{\Sigma}_{\beta}^{-1} \mathbf{X}^T$ , is typically dense, expensive to form, and leads to expensive solutions to systems of linear equations
- Its inverse, however, can be computed through the Sherman-Morrison-Woodbury formula with  $\mathbf{A}^{-1} = \sigma^{-2} \mathbf{I}_n$ ,  $\mathbf{U} = \mathbf{V} = \mathbf{X}$ , and  $\mathbf{C} = \boldsymbol{\Sigma}_{\beta}^{-1}$

## Bibliographic notes

- For more details on matrix decompositions, consider Wood (2015, Appendix B) for a concise overview
- For fuller details consider Monahan (2011, chaps. 3, 4 and 6) or Press et al. (2007, chap. 2 and 11)

## Chapter 4: Numerical Calculus

# Numerical Calculus: Motivation

- In statistics, we often rely on the Normal distribution with pdf

$$\phi(x; \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp \left[ -\frac{(x - \mu)^2}{2\sigma^2} \right]$$

and cdf

$$\Phi(x; \mu, \sigma^2) = \int_{-\infty}^x \phi(x; \mu, \sigma^2) dx.$$

- Unfortunately no closed form exists for  $\Phi(x; \mu, \sigma^2)$
- However, if  $x$ ,  $\mu$  and  $\sigma$  are stored in R as `x`, `mu` and `sigma`, we can still evaluate  $\Phi(x; \mu, \sigma^2)$  with `pnorm(x, mu, sigma)`
- This is one example of a frequently-occurring situation in which we somehow want to evaluate an intractable integral
- This raises the question: are there generic methods that let us evaluate intractable integrals?
- The answer is often **numerical integration**

# Numerical Calculus: Motivation

- *Remark:* In this chapter, we'll consider generic methods for integration, i.e. that work in many scenarios
- Sometimes, such as evaluating  $\Phi(x; \mu, \sigma^2)$ , specific algorithms will give more accurate results
- This is what R does for `pnorm()`

# Numerical Differentiation

- In Chapter 5 we will cover optimisation of functions, such as numerically finding maximum likelihood estimates when analytical solutions aren't available
- We'll see that supplying derivatives can considerably improve estimation, typically in terms of reducing computation time
- Here we'll cover some useful results in terms of differentiation of matrices, which will later prove useful
- No knowledge of analytical matrix calculus beyond these results will be needed for MTH3045
- The matrix cookbook (Petersen and Pedersen 2012), however, can provide you with a much more thorough set of differentiation rules, should you ever need them



# Differentiation definitions I

- **Definition:** Consider  $f : \mathbb{R}^n \rightarrow \mathbb{R}$ , which we'll consider a function of vector  $\mathbf{x} = (x_1, \dots, x_n)^T$
- The **gradient operator**,  $\nabla$ , is defined as

$$\nabla f(\mathbf{x}) = \begin{pmatrix} \frac{\partial f(\mathbf{x})}{\partial x_1} \\ \frac{\partial f(\mathbf{x})}{\partial x_2} \\ \vdots \\ \frac{\partial f(\mathbf{x})}{\partial x_n} \end{pmatrix}$$

- Note that in MTH3045 we will have no cause to consider multivariate functions, i.e. to consider  $\mathbf{f} : \mathbb{R}^n \rightarrow \mathbb{R}^m$ , for  $m > 1$

## Differentiation definitions II

- **Definition:** Consider again  $f : \mathbb{R}^n \rightarrow \mathbb{R}$
- The **Hessian matrix** is the matrix of second derivatives of  $f$ , whereas the gradient operator considered first derivatives, and is given by

$$\nabla^2 f(\mathbf{x}) = \begin{pmatrix} \frac{\partial^2 f}{\partial x_1^2} & \frac{\partial^2 f}{\partial x_1 \partial x_2} & \cdots & \frac{\partial^2 f}{\partial x_1 \partial x_n} \\ \frac{\partial^2 f}{\partial x_2 \partial x_1} & \frac{\partial^2 f}{\partial x_2^2} & \cdots & \frac{\partial^2 f}{\partial x_2 \partial x_n} \\ \vdots & \vdots & \ddots & \vdots \\ \frac{\partial^2 f}{\partial x_n \partial x_1} & \frac{\partial^2 f}{\partial x_n \partial x_2} & \cdots & \frac{\partial^2 f}{\partial x_n^2} \end{pmatrix}$$

- The Hessian matrix plays an important role in statistics, in particular for estimating parameter uncertainty via the Fisher information, which is covered in MTH3028
- In the next chapter, we'll also see that it's important for optimisation

## Differentiation rules I

- Now consider  $g : \mathbb{R}^n \rightarrow \mathbb{R}$  a function of  $\mathbf{x}$  that, for fixed  $\mathbf{A}$ , takes the quadratic form  $g(\mathbf{x}) = \mathbf{x}^\top \mathbf{A} \mathbf{x}$  for  $n \times n$  matrix  $\mathbf{A}$  and  $n$ -vector  $\mathbf{x}$
- Then

$$\nabla g(\mathbf{x}) = (\mathbf{A} + \mathbf{A}^\top) \mathbf{x} \quad \text{and} \quad \nabla^2 g(\mathbf{x}) = \mathbf{A} + \mathbf{A}^\top$$

- Note that in the case of symmetric  $\mathbf{A}$ ,  $\nabla g(\mathbf{x}) = 2\mathbf{A}\mathbf{x}$  and  $\nabla^2 g(\mathbf{x}) = 2\mathbf{A}$

## Differentiation rules II

- Next consider  $h : \mathbb{R}^n \rightarrow \mathbb{R}$  a function of  $n$ -vector  $\mathbf{x}$  and  $p$ -vector  $\mathbf{y}$  that, for fixed  $n \times n$  matrix  $\mathbf{A}$  and  $n \times p$  matrix  $\mathbf{B}$ , takes the quadratic form  $h(\mathbf{x}, \mathbf{y}) = (\mathbf{x} + \mathbf{B}\mathbf{y})^T \mathbf{A}(\mathbf{x} + \mathbf{B}\mathbf{y})$
- Then

$$\frac{\partial h(\mathbf{x}, \mathbf{y})}{\partial \mathbf{x}} = (\mathbf{A} + \mathbf{A}^T)(\mathbf{x} + \mathbf{B}\mathbf{y}) \quad \text{and} \quad \frac{\partial h(\mathbf{x}, \mathbf{y})}{\partial \mathbf{y}} = \mathbf{B}^T(\mathbf{A} + \mathbf{A}^T)(\mathbf{x} + \mathbf{B}\mathbf{y}),$$

and also

$$\frac{\partial^2 h(\mathbf{x}, \mathbf{y})}{\partial \mathbf{x} \partial \mathbf{x}^T} = \mathbf{A} + \mathbf{A}^T \quad \text{and} \quad \frac{\partial^2 h(\mathbf{x}, \mathbf{y})}{\partial \mathbf{y} \partial \mathbf{y}^T} = \mathbf{B}^T(\mathbf{A} + \mathbf{A}^T)\mathbf{B}$$

- Note that above we use partial derivative notation, i.e.  $\partial$ , as opposed to gradient operator notation, i.e.  $\nabla$ , as the derivatives are not w.r.t. all variables.

# Challenges III

- Go to Challenges III of the week 6 lecture 1 challenges at <https://byoungman.github.io/MTH3045/challenges>

## Week 6 lecture 2

## Example: Maximum likelihood estimates of regression coefficients in the normal linear model via matrix calculus I

- Consider the normal linear model  $\mathbf{Y} = \mathbf{X}\beta + \varepsilon$ 
  - $\mathbf{Y} = (Y_1, \dots, Y_n)^\top$
  - $\mathbf{X}$  is an  $n \times (p+1)$  design matrix
  - $\beta$  is a  $(p+1)$ -vector of regression coefficients
  - $\varepsilon = (\varepsilon_1, \dots, \varepsilon_n)^\top$  with independent  $\varepsilon_i \sim N(0, \sigma^2)$
- The maximum likelihood estimate of  $\beta$ , denoted  $\hat{\beta}$ , minimises the RSS, i.e. minimises

$$(\mathbf{y} - \mathbf{X}\beta)^\top (\mathbf{y} - \mathbf{X}\beta)$$

if we observe  $\mathbf{y} = (y_1, \dots, y_n)^\top$

- Differentiating w.r.t.  $\beta$  gives

$$-\mathbf{X}^\top (\mathbf{y} - \mathbf{X}\beta) - (\mathbf{y} - \mathbf{X}\beta)^\top \mathbf{X}$$

## Example: Maximum likelihood estimates of regression coefficients in the normal linear model via matrix calculus II

- Then

$$-\mathbf{X}^T(\mathbf{y} - \mathbf{X}\beta) - (\mathbf{y} - \mathbf{X}\beta)^T\mathbf{X}$$

simplifies to

$$-2\mathbf{X}^T(\mathbf{y} - \mathbf{X}\beta)$$

as  $\mathbf{X}^T(\mathbf{y} - \mathbf{X}\beta)$  and  $(\mathbf{y} - \mathbf{X}\beta)^T\mathbf{X}$  are both  $n$ -vectors

- The derivative of the RSS is zero at  $\hat{\beta}$  and so

$$-2\mathbf{X}^T(\mathbf{y} - \mathbf{X}\hat{\beta}) = 0$$

- Therefore  $\hat{\beta}$  is the solution of

$$\mathbf{X}^T\mathbf{X}\hat{\beta} = \mathbf{X}^T\mathbf{y}$$

- Alternatively

$$\hat{\beta} = (\mathbf{X}^T\mathbf{X})^{-1}\mathbf{X}^T\mathbf{y}$$

as we were given in MTH2006



# Finite-differencing

- Consider again  $f(\mathbf{x})$ , a function of vector  $\mathbf{x}$
- **Definition:** Consider  $f : \mathbb{R}^n \rightarrow \mathbb{R}$  for  $n$ -vector  $\mathbf{x}$
- Let,  $\mathbf{e}_i$  be the  $n$ -vector comprising entirely zeros, except for its  $i$ th element, which is one
- Then the **partial derivative** of  $f(\mathbf{x})$  w.r.t.  $x_i$ , the  $i$ th element of  $\mathbf{x}$ , is

$$\frac{\partial f(\mathbf{x})}{\partial x_i} = \lim_{h \rightarrow 0} \frac{f(\mathbf{x} + h\mathbf{e}_i) - f(\mathbf{x})}{h}$$

- The preceding definition leads us to the **finite-difference** partial derivative approximation

$$\frac{\partial f(\mathbf{x})}{\partial x_i} \simeq \frac{f(\mathbf{x} + \delta\mathbf{e}_i) - f(\mathbf{x})}{\delta},$$

where  $\delta$  is small

## Example: Finite-differencing of $\sin(x)$ I

- Use finite-differencing to approximate the derivative of  $f(x) = \sin(x)$  for  $x \in [0, 2\pi]$ , and compare its accuracy to the true derivative
- First, let's calculate  $f$  and its *true* derivative, i.e.  $f'(x) = \cos(x)$ , and store this as `partial0` for  $\{x_i\}$ ,  $i = 1, \dots, 100$ , a set of equally-spaced points on  $[0, 2\pi]$

```
n <- 100
x <- seq(0, 2 * pi, l = 100)
f <- sin(x)
partial0 <- cos(x)
```

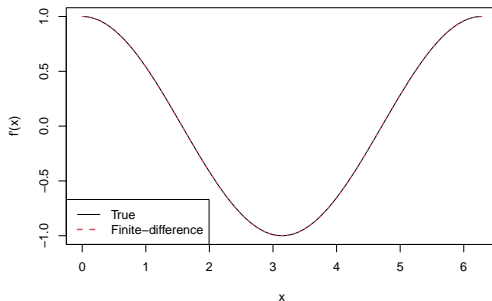
- Now we'll set  $\delta = 10^{-6}$  and calculate  $x_i + \delta$ , for each  $x_i$ , i.e. each element of `x`, so that the finite-difference approximation to the derivative is given by  $[\sin(x_i + 10^{-6}) - \sin(x_i)]/10^{-6}$ , which is calculated below and stored as `partial1`

```
delta1 <- 1e-6
x1 <- x + delta1
f1 <- sin(x1)
partial1 <- (f1 - f) / delta1
```

## Example: Finite-differencing of $\sin(x)$ II

- We'll then plot the  $f'(x)$  against its finite-difference approximation

```
matplot(x, cbind(partial0, partial1), type = 'l', xlab = 'x', ylab = "f'(x)")  
legend('bottomleft', lty = 1:2, col = 1:2, lwd = 1:2,  
       legend = c('True', 'Finite-difference'))
```



- In fact, the true derivative and its finite-difference approximation are so similar that it's difficult to distinguish the two, but they're both there in the plot!

## Example: Finite-differencing of $\sin(x)$ III

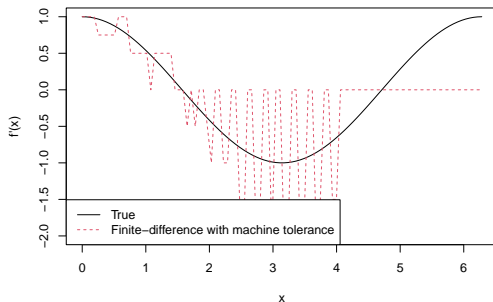
- We might be tempted to choose  $\delta$  as small as possible
- Suppose we were to repeat Example 4.3 with  $\delta = \epsilon_m$ , i.e. R's machine tolerance
- The following calculates the finite-difference approximation as `partial2`

```
delta2 <- .Machine$double.eps  
x2 <- x + delta2  
f2 <- sin(x2)  
partial2 <- (f2 - f) / delta2
```

## Example: Finite-differencing of $\sin(x)$ III

- The the following plots this against the true value of  $f'(x)$

```
matplot(x, cbind(partial0, partial2), type = 'l', xlab = 'x', ylab = "f'(x)")  
legend('bottomleft', lty = 1:2, col = 1:2, bg = 'white',  
       legend = c('True', 'Finite-difference with machine tolerance'))
```



- Using  $\delta = \epsilon_m$  gives a terrible approximation to  $f'(x)$ , which gets worse as  $x$  increases
- We've actually encountered an example *calculation error*, which was introduced in Chapter 2

# Challenges I

- Go to Challenges I of the week 6 lecture 2 challenges at <https://byoungman.github.io/MTH3045/challenges>

## Example: Finite differencing of the multivariate Normal log-likelihood I

- Find  $\partial \log f(\mathbf{y} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) / \partial y_i$  analytically, for  $i = 1, \dots, p$ , where  $f(\mathbf{y} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma})$  is the  $MVN_p(\boldsymbol{\mu}, \boldsymbol{\Sigma})$  pdf, for arbitrary  $\mathbf{y}$ ,  $\boldsymbol{\mu}$  and  $\boldsymbol{\Sigma}$
- Evaluate this for  $\mathbf{y}$ ,  $\boldsymbol{\mu}$  and  $\boldsymbol{\Sigma}$  as in Example 3.2
- Then approximate the same derivative using finite-differencing
- The logarithm of the  $MVN_p(\boldsymbol{\mu}, \boldsymbol{\Sigma})$  pdf is given by

$$\log f(\mathbf{y} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma}) = -\frac{1}{2} [p \log(2\pi) + \log(|\boldsymbol{\Sigma}|) + (\mathbf{y} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1} (\mathbf{y} - \boldsymbol{\mu})]$$

which we know, from Example 3.13, we can evaluate with `dmvn3()`

- Using the above properties

$$\frac{\partial \log f(\mathbf{y} \mid \boldsymbol{\mu}, \boldsymbol{\Sigma})}{\partial \mathbf{y}} = -\boldsymbol{\Sigma}^{-1}(\mathbf{y} - \boldsymbol{\mu})$$

since  $\boldsymbol{\Sigma}$  is symmetric

## Example: Finite differencing of the multivariate Normal log-likelihood II

- We can evaluate this for  $\mathbf{y}$ ,  $\boldsymbol{\mu}$  and  $\boldsymbol{\Sigma}$  as in Example 3.2 with the following

```
y <- c(.7, 1.3, 2.6)
mu <- 1:3
Sigma <- matrix(c(4, 2, 1, 2, 3, 2, 1, 2, 2), 3, 3)
deriv1 <- -solve(Sigma, y - mu)
```

which gives

```
as.vector(deriv1)
```

```
## [1] -0.08  0.38 -0.14
```



## Example: Finite differencing of the multivariate Normal log-likelihood III

- To approximate the derivative by finite-differencing, it makes sense to write a multi-purpose function for finite differencing, which we'll call `fd()`

```
fd <- function(x, f, delta = 1e-6, ...) {  
  # Function to evaluate derivative w.r.t. vector by finite-differencing  
  # x is a p-vector  
  # fn is the function for which the derivative is being calculated  
  # delta is the finite-differencing step, which defaults to 10^{-6}  
  # returns a vector of length x  
  f0 <- f(x, ...)  
  p <- length(x)  
  f1 <- numeric(p)  
  for (i in 1:p) {  
    ei <- replace(numeric(p), i, 1)  
    f1[i] <- f(x + delta * ei, ...)  
  }  
  (f1 - f0) / delta  
}
```

## Example: Finite differencing of the multivariate Normal log-likelihood IV

- Then we can use this with `dmvn3()` with the following

```
deriv2 <- fd(y, dmvn3, mu = mu, Sigma = Sigma)
```

which gives

```
deriv2
```

```
## [1] -0.0800002  0.3799993 -0.1400008
```

and is the same as the analytical result

```
all.equal(deriv1, deriv2)
```

```
## [1] "Mean relative difference: 2.832689e-06"
```

once we allow for error in the finite-difference approximation

# Challenges II

- Go to Challenges II of the week 6 lecture 2 challenges at <https://byoungman.github.io/MTH3045/challenges>

## Week 6 lecture 3

# Quadrature

- Another common requirement in statistics is that some integral needs to be evaluated
- To start, let's consider a simple integral of the form

$$I = \int_a^b f(x)dx$$

- We'll first take a look at some *deterministic* approaches to numerically evaluating integrals
- In fact, these all boil down to assuming that

$$I \simeq \sum_{i=1}^N w_i f(x_i^*)$$

for some weights  $w_i$  and nodes  $x_i^*$ ,  $i = 1, \dots, N$

- Note that here we're considering the so-called *composite* approach to approximating an interval, i.e. in which a rule is applied over a collection of sub-intervals

# Midpoint rule

- Perhaps the first numerical integration scheme we come across is the midpoint rule
- Put simply, we divide  $[a, b]$  into  $N$  equally-sized intervals, and use the midpoints of these as the nodes,  $x_i^*$
- This gives

$$\int_a^b f(x)dx \simeq h \sum_{i=1}^N f(x_i^*),$$

where  $x_i^* = a + (i - 0.5)(b - a)/N$  and  $h = (b - a)/N$

- The error in the approximation is  $O(h^2)$
- Thus more intervals reduces  $h$  and gives a more accurate approximation
- We can measure accuracy through **relative absolute error**
- **Definition:** The **relative absolute error**, or sometimes just *relative error*, of an estimate of some true value is given by

$$\left| \frac{\text{true value} - \text{estimate}}{\text{true value}} \right|$$

## Example: Midpoint rule I

- Consider the integral  $\int_0^1 \exp(x)dx = \exp(1) - 1 \simeq 1.7182818$
- Use R and the midpoint rule to estimate the integral with  $N = 10, 100$  and  $1000$
- Then compare the relative absolute error of each
- We'll start by calculating the true value of the integral, which we'll store as `true`

```
true <- exp(1) - 1
```

- Then we'll store the values of  $N$  that we're testing as `N_vals`

```
N_vals <- 10^c(1:3)
```

## Example: Midpoint rule II

- The following then creates a vector, `midpoint`, in which to store the integral approximations, and calculates the approximations with a `for` loop
- Inside the loop the integration nodes (i.e. the midpoints) and  $h$  are calculated

```
midpoint <- numeric(length(N_vals))
for (i in 1:length(N_vals)) {
  N <- N_vals[i]
  nodes <- (1:N - .5) / N
  h <- 1 / N
  midpoint[i] <- h * sum(exp(nodes))
}
midpoint
```

```
## [1] 1.717566 1.718275 1.718282
```



## Example: Midpoint rule III

- The relative absolute error for each is then given in the vector `rel_err_mp` below

```
rel_err_mp <- abs((true - midpoint) / true)
rel_err_mp
```

```
## [1] 4.165452e-04 4.166655e-06 4.166667e-08
```

- We clearly see that the absolute error reduces by two factors of ten for each factor of ten increase in  $N$ , which is consistent with the above comment of  $O(h^2)$  error, where here  $h = 1/N$

# Challenges III

- Go to Challenges III of the week 6 lecture 2 challenges at <https://byoungman.github.io/MTH3045/challenges>

# Simpson's rule

- The midpoint rule works simply by approximating  $f(x)$  over a sub-interval of  $[a, b]$  by a horizontal line
- The trapezium rule (which we'll overlook) assumes a straight line
- Simpson's rule is derived from a quadratic approximation and given by

$$\int_a^b f(x)dx \simeq \frac{h}{6} \left( f(a) + 4 \sum_{i=1}^N f(x_{1i}^*) + 2 \sum_{i=1}^{N-1} f(x_{2i}^*) + f(b) \right),$$

where  $x_{1i}^* = a + h(2i - 1)/2$ ,  $x_{2i}^* = a + ih$  and  $h = (b - a)/N$

- Note that Simpson's rule requires  $N + 1$  more evaluations of  $f$  than the midpoint rule
  - however, a benefit of those extra evaluations is that its error reduces to  $O(h^4)$

## Example: Simpson's rule I

- Now use R and Simpson's rule to approximate the integral  $\int_0^1 \exp(x) dx = \exp(1) - 1$  with  $N = 10, 100$  and  $1000$ , compare the relative absolute error for each, and against those of the midpoint rule in Example 4.5
- We already have `true` and `N_vals` from Example 4.5, and we can use a similar `for` loop to approximate the integral using Simpson's rule
- The main difference is that we create two sets of nodes, `nodes1` and `nodes2`, which correspond to the  $x_{1i}$ s and  $x_{2i}$ s in Equation (4.1), respectively

## Example: Simpson's rule II

- The integral approximations are stored as `simpson`

```
simpson <- numeric(length(N_vals))
N_vals <- 10^c(1:3)
for (i in 1:length(N_vals)) {
  N <- N_vals[i]
  h <- 1 / N
  simpson[i] <- 1 + exp(1)
  nodes1 <- h * (2*c(1:N) - 1) / 2
  simpson[i] <- simpson[i] + 4 * sum(exp(nodes1))
  nodes2 <- h * c(1:(N - 1))
  simpson[i] <- simpson[i] + 2 * sum(exp(nodes2))
  simpson[i] <- h * simpson[i] / 6
}
print(simpson, digits = 12)
```

```
## [1] 1.71828188810 1.71828182847 1.71828182846
```

- We print this to 11 decimal places so we can see where the approximations changes with  $N$

## Example: Simpson's rule III

- Finally we calculate the relative absolute errors, `rel_err_simp`,

```
rel_err_simp <- abs((true - simpson) / true)
rel_err_simp
```

```
## [1] 3.471189e-08 3.472270e-12 6.461239e-16
```

- We see a dramatic improvement in the accuracy of approximation that Simpson's rule brings, with relative absolute errors of the same order of magnitude as those from the midpoint rule using  $N = 1000$  achieved with  $N = 10$  for Simpson's rule
- Note, though, that for given  $N$ , Simpson's rule requires  $N + 1$  more evaluations of  $f()$

# Challenges I

- Go to Challenges I of the week 6 lecture 3 challenges at <https://byoungman.github.io/MTH3045/challenges>

# Gaussian quadrature I

- We've seen that Simpson's rule can considerably improve on the midpoint rule for approximating integrals
- However, we might still consider both restrictive in that they consider an equally-spaced set of nodes
- **Definition:** Consider  $g(x)$ , a polynomial of degree  $2N - 1$ , and a fixed weight function  $w(x)$
- Then, the **Gauss-Legendre quadrature rule** states that

$$\int_a^b w(x)g(x)dx = \sum_{i=1}^N w_i g(x_i),$$

where, for  $i = 1, \dots, N$ ,  $w_i$  and  $x_i$  depend on  $w(x)$ ,  $a$  and  $b$ , but not  $g(x)$



## Gaussian quadrature II

- The Gauss-Legendre quadrature rule is the motivation for **Gaussian quadrature**, whereby we assume that the integral we're interested in can be well-approximated by a polynomial
- This results in the approximation

$$\int_a^b f(x)dx \simeq \sum_{i=1}^N w_i f(x_i)$$

for a fixed set of  $x$  values,  $x_i$  with corresponding weights  $w_i$ , for  $i = 1, \dots, N$

- There are many rules for choosing the weights,  $w_i$ , but (perhaps fortunately) we won't go into them in detail in MTH3045
- Instead, we'll just consider the function `pracma::gaussLegendre()` (for which you'll need to install the `pracma` package), where `pracma::gaussLegendre(N, a, b)` produces  $N$  nodes and corresponding weights on the interval  $[a,b]$ , with  $N = N$ ,  $a = a$  and  $b = b$

# Gaussian quadrature III

- The following produces nodes and weights for  $N = 10$  on  $[0, 1]$

```
gq <- pracma::gaussLegendre(10, 0, 1)
```

```
gq
```

```
## $x
```

```
## [1] 0.01304674 0.06746832 0.16029522 0.28330230 0.42556283 0.57443717
```

```
## [7] 0.71669770 0.83970478 0.93253168 0.98695326
```

```
##
```

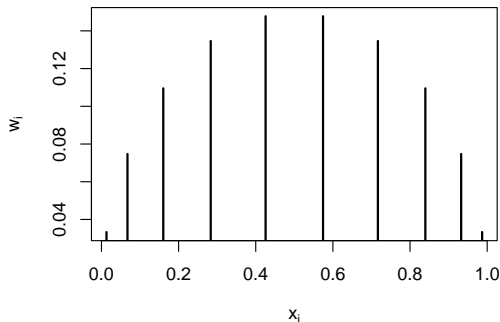
```
## $w
```

```
## [1] 0.03333567 0.07472567 0.10954318 0.13463336 0.14776211 0.14776211
```

```
## [7] 0.13463336 0.10954318 0.07472567 0.03333567
```

## Gaussian quadrature IV

- The plots below shows the nodes and their weights



- The nodes are spread further apart towards the middle of the  $[0, 1]$  range, but given more weight
- Note that `pracma::gaussLegendre()` is named so because it implements Gauss-Legendre quadrature, i.e. Gaussian quadrature with Legendre polynomials<sup>1</sup>

---

<sup>1</sup>Wikipedia has a useful pages [on Gaussian quadrature](#) and [on Legendre polynomials](#), should you want to read more on them

## Example: Gaussian quadrature I

- Now use R and Gauss-Legendre quadrature to approximate the integral  $\int_0^1 \exp(x) dx$  with  $N = 10$
- Explore what value of  $N$  gives a comparable estimate to that of the midpoint rule with  $N = 100$  based on relative absolute error
- We can re-use `true` from Example 4.5 and then we'll consider  $N = 10, 4$  and 3, which we'll call `N_vals`, and store the resulting integral approximations in `gauss`.

```
N_vals <- c(10, 4, 3)
gauss <- numeric(length(N_vals))
for (i in 1:length(N_vals)) {
  N <- N_vals[i]
  xw <- pracma::gaussLegendre(N, 0, 1)
  gauss[i] <- sum(xw$w * exp(xw$x))
}
gauss
```

```
## [1] 1.718282 1.718282 1.718281
```

## Example: Gaussian quadrature II

- The relative absolute errors, `rel_err_gauss`,

```
rel_err_gauss <- abs((true - gauss) / true)
rel_err_gauss
```

```
## [1] 2.584496e-16 5.429651e-10 4.795992e-07
```

show that, having considered  $N = 3, 4, 10$ , choosing  $N = 3$  for Gaussian quadrature gives closest relative absolute error to that of the midpoint rule with  $N = 100$ , which really is quite impressive

- Note, though, that  $f(x) = \exp(x)$  is a very smooth function
  - for wiggler functions, larger  $N$  is likely to be needed, and improvements in performance, such as Gaussian quadrature over the midpoint rule, might be significantly less

# Challenges II

- Go to Challenges II of the week 6 lecture 3 challenges at <https://byoungman.github.io/MTH3045/challenges>

## Example: Poisson marginal approximation using Gaussian quadrature I

- Consider a single random variable  $Y \mid \lambda \sim \text{Poisson}(\lambda)$ , where we can characterise our prior beliefs about  $\lambda$  as  $\lambda \sim \text{N}(\mu, \sigma^2)$
- Use Gaussian quadrature with  $N = 7$  to estimate the marginal pdf of  $Y$  if  $\mu = 10$  and  $\sigma = 3$
- The marginal pdf of  $Y$  is given by

$$f(y) = \int_{-\infty}^{\infty} f(y \mid \lambda) f(\lambda) d\lambda$$

- *Remark:* The *three sigma rule* is a heuristic rule of thumb that 99.7% of values lie within three standard deviations of the mean

## Example: Poisson marginal approximation using Gaussian quadrature II

- Hence for the  $N(10, 3^2)$  distribution we should expect 99.7% of values to lie within  $10 \pm 3 \times 3$
- Hence we'll take this as our range for the Gaussian quadrature nodes.

```
mu <- 10
sigma <- 3
N <- 7 # no. of nodes
xw <- pracma::gaussLegendre(
  N,
  mu - 3 * sigma, # left-hand end
  mu + 3 * sigma # right-hand end
)
xw
```

```
## $x
## [1] 1.458029 3.326219 6.347394 10.000000 13.652606 16.673781 18.541971
##
## $w
## [1] 1.165365 2.517349 3.436470 3.761633 3.436470 2.517349 1.165365
```

which are stored as `xw$x` with corresponding weights `xw$w`,  $w_1, \dots, w_N$



## Example: Poisson marginal approximation using Gaussian quadrature III

- Next we want a set of values at which to evaluate  $f(y)$ , and for this we'll choose  $0, 1, \dots, 30$ , which we can create in R with

```
y_vals <- 0:30
```

- Then we can estimate  $f(y)$  as

$$\hat{f}(y) \simeq \sum_{i=1}^N w_i f(y \mid \lambda_i^*) f(\lambda_i^*)$$

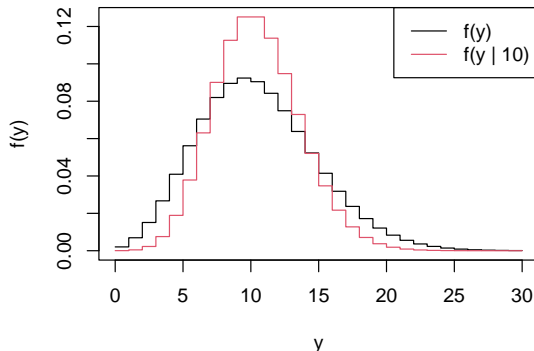
- The following code gives  $\hat{f}(y)$  as fhat for y in the set of values y\_vals

```
m <- length(y_vals)
fhat <- numeric(m)
for (i in 1:m) {
  fhat[i] <- sum(xw$w * dpois(y_vals[i], xw$x) *
                 dnorm(xw$x, mu, sigma))
}
```

## Example: Poisson marginal approximation using Gaussian quadrature IV

- Finally, we'll plot  $\hat{f}(y)$  against the pdf of the Poisson(10) distribution

```
matplot(y_vals, cbind(fhat, dpois(y_vals, mu)), lwd = 1,  
        col = 1:2, lty = 1, type = 'l', xlab = 'y', ylab = 'f(y)')  
legend('topright', c("f(y)", "f(y | 10)"),  
      lty = 1, col = 1:2)
```



- $f(y)$  is broader than  $f(y | 10)$ , which is to be expected given that  $f(y)$  integrates out the variability in  $\lambda$  given by the  $N(10, 3^2)$  distribution

# One-dimensional numerical integration in R

- Unsurprisingly, R has a function for one-dimensional numerical integration
- It's called `integrate()`
- It uses a method that builds on Gaussian quadrature, but we won't go into its details
- Use of `integrate()`, however, is fairly straightforward

## Example: Integration with `integrate()`

- Evaluate the integral  $\int_0^1 \exp(x)dx = \exp(1) - 1$  using R's `integrate()` function with  $N = 10$  and report its relative absolute error
- We can use the following code, where the first argument to `integrate()` is the function we're integrating, the second and third are the lower and upper ranges of the definite integral, and `subdivisions` is the maximum number of nodes to use in the approximation, which defaults to 100.

```
true <- exp(1) - 1
estimate <- integrate(function(x) exp(x), 0, 1, subdivisions = 10)
estimate
```

```
## 1.718282 with absolute error < 1.9e-14
```

```
rel_err <- abs((true - estimate$value) / true)
rel_err
```

```
## [1] 1.292248e-16
```

- Note above that the absolute error is similarly tiny to that of Gaussian quadrature above
- The values themselves, being so close to the machine tolerance, are incomparable
  - but we can be sure that the approximation is incredibly accurate

# References

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