

Exercise

Parameter estimation with R

A regression model for disease forecasting

The exercise

Parameter estimation with R

- Two functions for non linear models
 - `nls()` is specifically adapted to minimization of sums of squares and regression analysis.
 - `optim()` is a general purpose minimizer (minimize sum of squared errors)
 - If `nls()` works, that gives more information. If not use `optim()`
 - They require different functions to evaluate model.
Sorry.

Example

Grain filling of wheat

- The model for grain filling

$$y^{\text{mod}} = \frac{W}{1 + \exp(B - (c)(DD))}$$

- y is grain weight (mg)
- DD is degree days from anthesis (the explanatory variable)
- W , B , c are parameters, to be estimated

R

```
rm(list=ls())  
seedwt<-function(DD,W,B,c)  
{  
  weight<-W/(1+exp(B-c*DD))  
  return(weight)  
}
```

My function to calculate seed weight as a function of **explanatory variables** and **parameters**

```
wt<-c(8,13,17,21,26,27,28,27,27,29,28)  
DD<-c(195,240,280,330,380,410,480,530,580,620,680)
```

The data. Observed values and explanatory variables

```
ols<-nls(y~seedwt(time,W,B,c),  
  data=list(y=wt,time=DD),  
  start=list(W=30,B=4,c=0.019))  
summary(ols)
```

Call to nls. data has observed values and **explanatory variables**. start has **parameters**.

Result of nls

Some rules

- The function must return a vector of simulated values, in the same order as the observed values
- The arguments in the call must match the arguments in the function
 - First argument in call becomes first argument in function etc.
 - (Other possibility – use names)
- The start argument of nls defines parameters and gives starting values
- The data argument of nls gives values of everything else (observed data and explanatory variables)

Displaying and analyzing the results

summary(ols)

Formula: $y \sim \text{seedwt}(\text{time}, W, B, c)$

Parameters:

	Estimate	Std. Error	t value	Pr(> t)	
W	28.171255	0.408284	69.00	2.17e-12	***
B	4.160223	0.360084	11.55	2.86e-06	***
c	0.016492	0.001415	11.65	2.68e-06	***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.8364 on 8 degrees of freedom

Number of iterations to convergence: 5

Achieved convergence tolerance: 1.961e-06

to retrieve coefficients for further calculation

coef(ols)

W	B	c
---	---	---

28.17125533	4.16022349	0.01649195
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```
# fitted() and residuals() use OLS parameters  
cbind(wt,fitted(ols),residuals(ols))
```

```
wt
```

```
[1,] 8 7.888909 0.1110908  
[2,] 13 12.666715 0.3332847  
[3,] 17 17.252820 -0.2528196  
[4,] 21 22.052755 -1.0527547  
[5,] 26 25.116207 0.8837928  
[6,] 27 26.226215 0.7737854  
[7,] 28 27.527683 0.4723169  
[8,] 27 27.885439 -0.8854391  
[9,] 27 28.045232 -1.0452322  
[10,] 29 28.105958 0.8940420  
[11,] 28 28.146945 -0.1469453
```

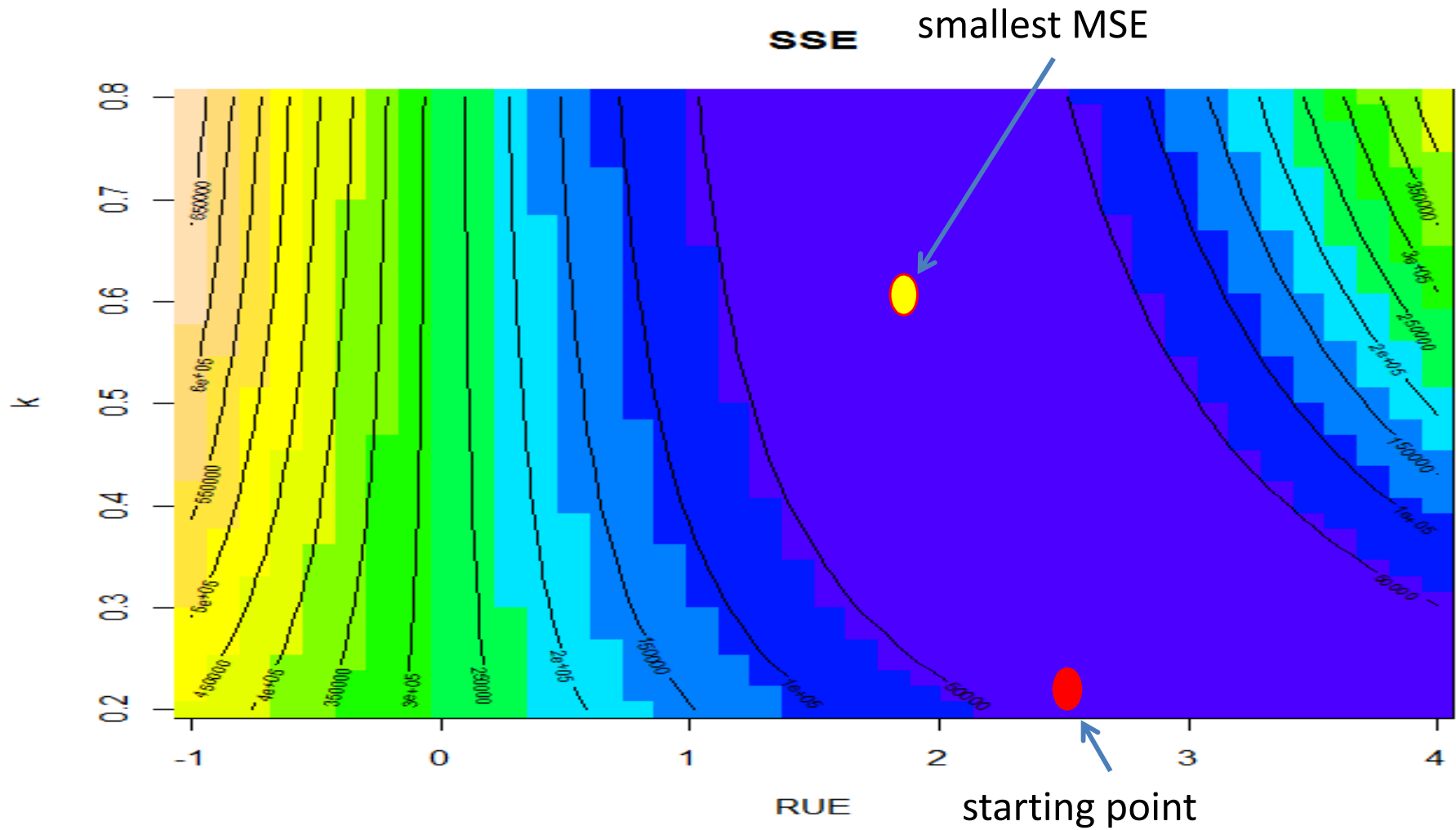
```
# predict() gives new predictions using OLS parameters
predict(ols,newdata=list(time=195:240))
[1] 7.888909 7.982918 8.077601 8.172953 8.268967
8.365637 8.462956 8.560919 8.659517
[10] 8.758743 8.858590 8.959050 9.060115 9.161777
9.264026 9.366854 9.470251 9.574209
[19] 9.678718 9.783767 9.889346 9.995446 10.102055
10.209162 10.316756 10.424826 10.533361
[28] 10.642348 10.751775 10.861630 10.971900
11.082573 11.193636 11.305076 11.416879 11.529032
[37] 11.641521 11.754333 11.867452 11.980866
12.094560 12.208519 12.322729 12.437175 12.551842
[46] 12.666715
```

OLS in pictures

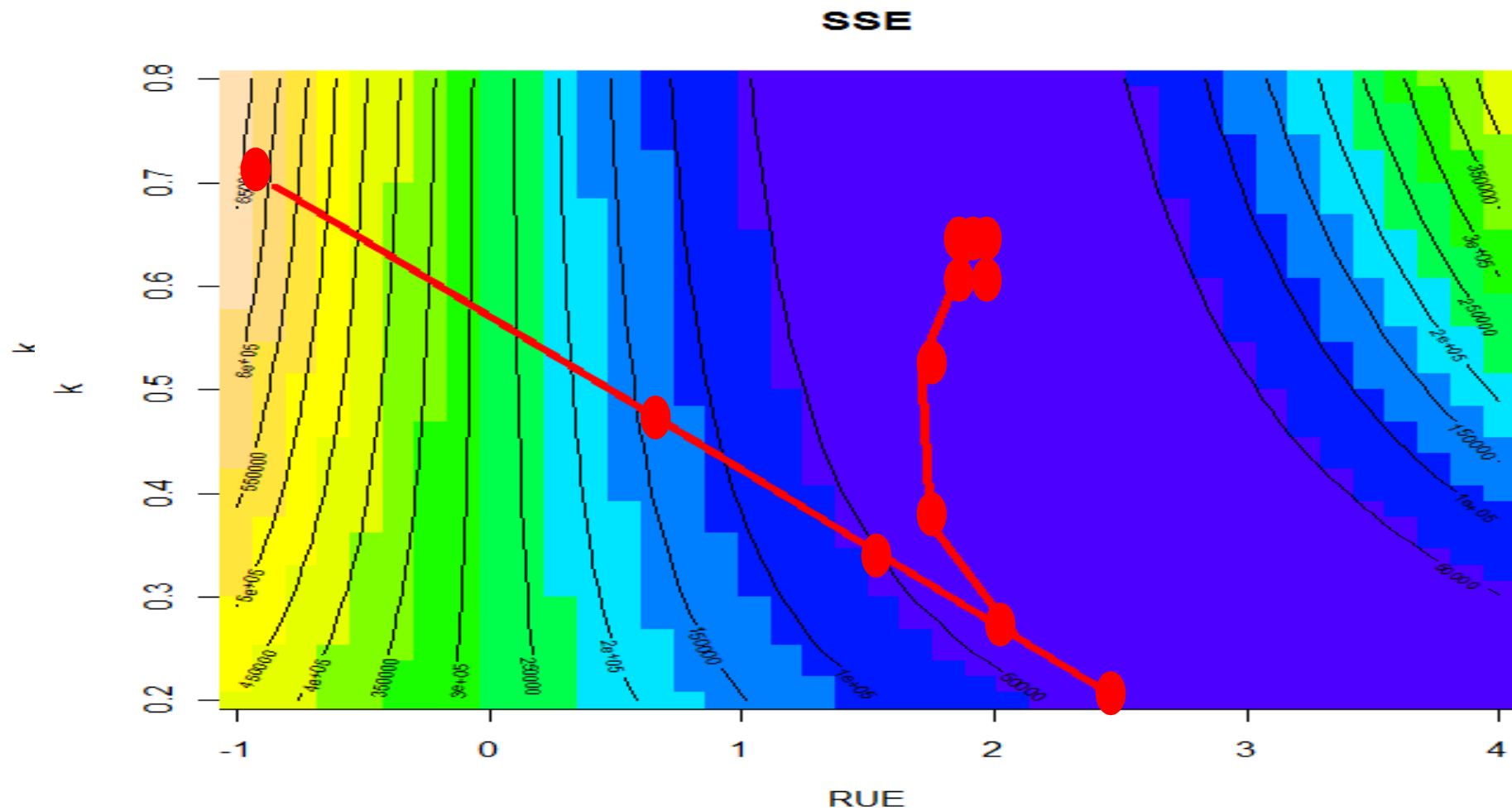
Gradient method

- Gauss-Newton. Default for `nls()`
 - Linearize model. (Need to calculate derivatives)
 - Calculate parameters that minimize MSE of linear problem (analytical)
 - Iterate (necessary because of non-linearity)

Path of Gauss Newton?

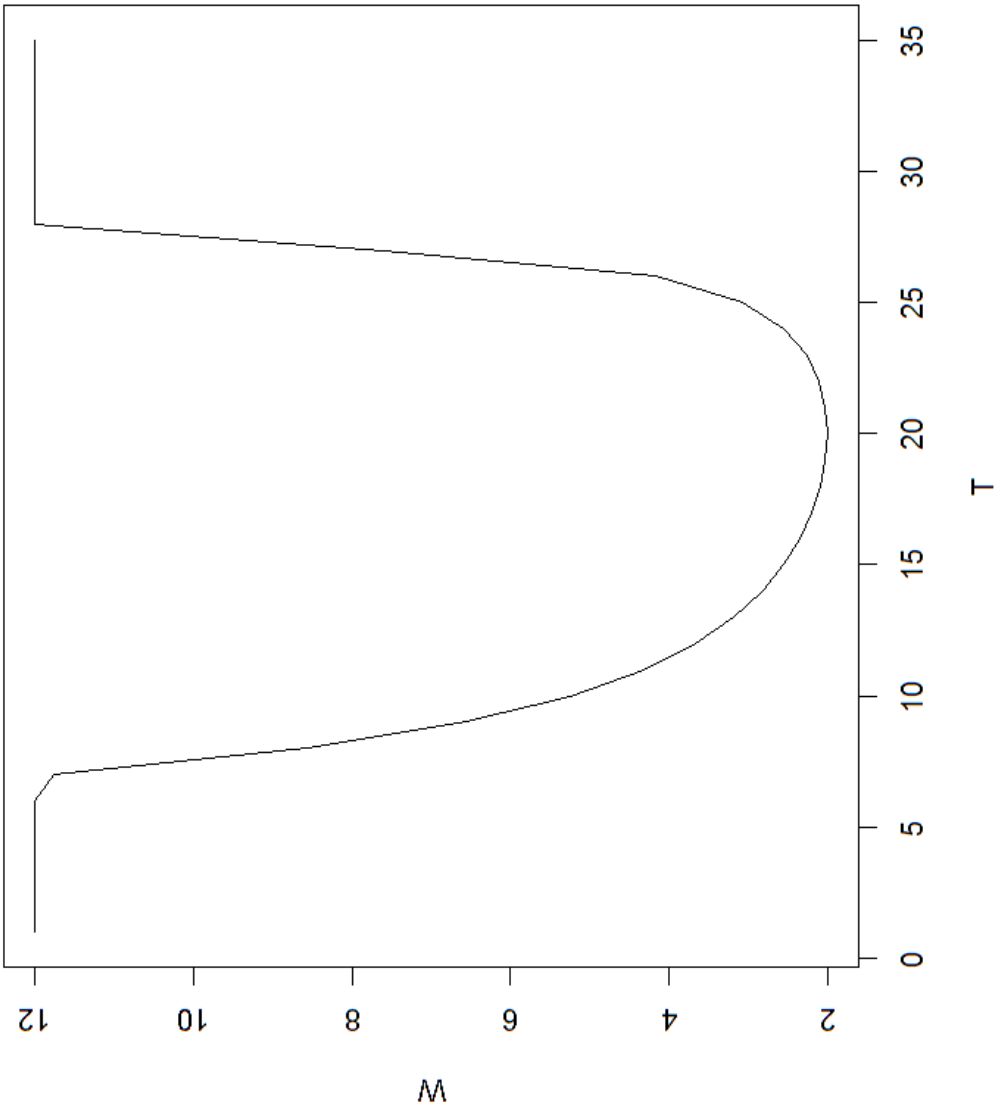


Path of Gauss Newton



Model of leaf wetness requirement

- Infection models of fungal foliar pathogens: often success of infection depends on temperature T and duration of leaf wetness W .
- $W=f(T;\theta)$ Each pathogen has requirement of W for infection to occur, depends on T
- Get parameters θ by fitting past data
 - Experiments: multiple T , W combinations
 - Each T , identify minimum W such that there is 20% disease incidence or 5% disease severity on an infected plant part at nonlimiting inoculum concentration.



Exercise

- Calibrate a model of leaf wetness requirement for fungal foliar pathogens

- Model is

$$W = W_{min} / fT$$

$$fT = \left[\frac{T_{max} - T}{T_{max} - T_{opt}} \right] \left[\frac{T - T_{min}}{T_{opt} - T_{min}} \right]^{(T_{opt} - T_{min}) / (T_{max} - T_{opt})}$$

$$W = W_{max} \text{ if } W > W_{max}$$

$$W = W_{max} \text{ if } T < T_{min} \text{ or } T > T_{max}$$

- 5 parameters: T_{min} , T_{opt} , T_{max} , W_{min} , W_{max}

- Data for *Pseudoperonospora cubensis* (downy mildew on cucumber)

T (°C)	W (hours)
4	12.0
8	8.6
12	3.7
16	2.3
20	2.0
24	2.5
28	12.0
32	12.0

- The disease might have parameters similar to *Bremia lactucae* on lettuce
 - $T_{min}=1$, $T_{opt}=15$, $T_{max}=25$, $W_{min}=4$, $W_{max}=10$

To do

1. Write a function that calculates W as a function of T given T_{min} , T_{opt} , T_{max} , W_{min} , W_{max}
2. Use the parameters of *Bremia lactucae* to calculate W as a function of temperature. Draw the graph.
3. Add the points for *Pseudoperonospora cubensis* to the graph
4. Use the data for *Pseudoperonospora cubensis* to calibrate the model for that disease. Start from *Bremia lactucae* parameters
5. Examine the output of `nls`. Are all the parameters well estimated?
6. Use the calibrated model to predict W for $T <- 0:40$
7. Graph the model results for T from 0 to 40°C (`type="l"`)
8. Add the data points. How well does the model fit the data?
9. Plot residuals (observed- simulated) versus simulated values.
10. Is this a "correct model"
11. Are the residuals "homoscedastic"?