

Covariance models for hierarchical data

560 Hierarchical modeling

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Fixed occasion longitudinal data

Data measured at regular time intervals on each subject.

- $y_{t,j}$ = observation on subject j at time t .
- $j = 1, \dots, m$;
- $t = 1, \dots, T$.

$$\begin{pmatrix} y_{1,1} & y_{2,1} & \cdots & y_{T,1} \\ y_{1,2} & y_{2,2} & \cdots & y_{T,2} \\ \vdots & & & \vdots \\ y_{1,m} & y_{2,m} & \cdots & y_{T,m} \end{pmatrix}$$

Fixed occasion longitudinal data

Structurally, this can be thought of as two-level hierarchical data:

- A subject is a *group*, as we have multiple observations within a subject.
- $t = 1, \dots, T$ indexes the multiple observations within a group (subject).

One goal of HM is to *account for correlation of observations within a group*.

For longitudinal data, this translates to the goal of

accounting for temporal correlation within a subject.

The difference: There is some structure to the “units” within a group:

- time is an ordered quantity;
- there might be some similarity of “units” *across* groups, as they correspond to common times.

Longitudinal data - an alternative viewpoint

Alternatively, you could structure the data as

- $y_{i,t}$ = observation on subject i at time t .
- $i = 1, \dots, n$
- $j = 1, \dots, T$.

You could think of each time point as being a group, within which there are multiple observations (subjects).

Problem with this perspective: In most applications, we are

- concerned with temporal dependence *within a subject*,
- not dependence of subjects *within a timepoint*.

However, there are some situations when this perspective is reasonable:

- if time can be viewed as a “sampled” quantity without an ordered effect.
- if the similarity due to common time session dominates similarity due to other factors.

Example: Sleep deprivation study

$y_{t,j}$ = reaction time of subject j , t days after beginning of study.

$t = 0, \dots, 9$.

$j = 1, \dots, 18$.

```
sleep[1:15,]
```

##	Reaction	Days	Subject
## 1	249.5600	0	308
## 2	258.7047	1	308
## 3	250.8006	2	308
## 4	321.4398	3	308
## 5	356.8519	4	308
## 6	414.6901	5	308
## 7	382.2038	6	308
## 8	290.1486	7	308
## 9	430.5853	8	308
## 10	466.3535	9	308
## 11	222.7339	0	309
## 12	205.2658	1	309
## 13	202.9778	2	309
## 14	204.7070	3	309
## 15	207.7161	4	309

Example: Life satisfaction

$y_{t,j}$ = life satisfaction of j at age t .

$t = 55, 56, \dots, 59, 60$.

$j = 1, \dots, 1237$.

```
happy[1:10,]
```

##	id	j	gender	age	married	nchildren	yearsEd	emplStat	totIncBeforeG
## 1	101	1985	1	55	1	0	15.0	1	23219.03
## 2	101	1986	1	56	1	0	15.0	1	23263.78
## 3	101	1987	1	57	1	0	15.0	1	23273.50
## 4	101	1988	1	58	1	0	15.0	1	23559.82
## 5	101	1989	1	59	1	0	15.0	1	23110.63
## 6	901	2006	2	55	2	0	10.5	1	22501.25
## 7	1701	2003	1	55	1	0	15.0	1	68116.78
## 8	1701	2004	1	56	1	0	15.0	1	78862.02
## 9	1701	2005	1	57	1	0	15.0	1	77910.88
## 10	1701	2006	1	58	1	0	15.0	1	79684.43
##	totIncAfterG	laborInc	state	healthSat	lifeSat	emplStat55			
## 1	16164.22	23060.29	0	8	8	1			
## 2	16072.46	23263.78	0	7	10	1			
## 3	16108.25	23273.50	0	5	8	1			
## 4	16268.29	23559.82	0	8	8	1			
## 5	16110.02	22857.30	0	7	8	1			
## 6	14876.25	22396.00	0	7	6	1			
## 7	45437.78	67975.36	0	9	8	1			
## 8	52091.36	78720.60	0	9	10	1			
## 9	50958.88	77769.45	0	9	10	1			
## 10	52345.42	79543.00	0	10	9	1			

Example: Life satisfaction

Notice:

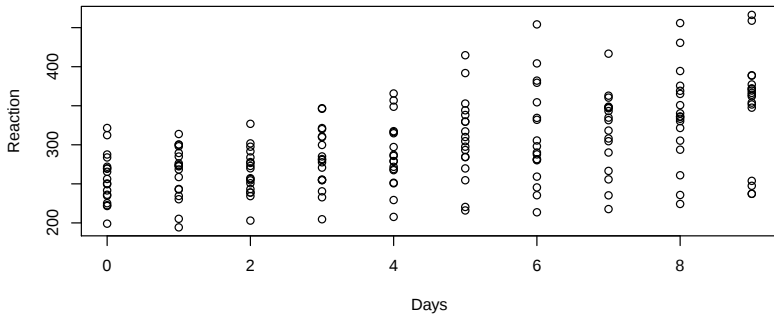
- The “times” are not actually at the same time.
- Subjects have varying amounts of data in the study.
- All data corresponds to one of the six ages (fixed occasion data).

Model fitting, diagnostics and model building

We will first develop some methods by analyzing the sleep data.

Questions:

- What are the effects of sleep deprivation on reaction time, on average?
- How do these effects vary across people?

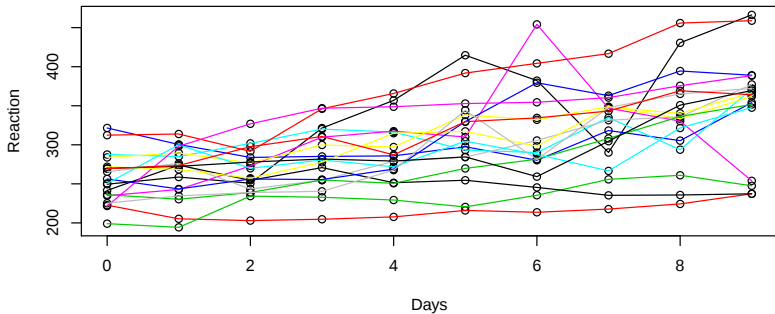


Model fitting, diagnostics and model building

We will first develop some methods by analyzing the sleep data.

Questions:

- What are the effects of sleep deprivation on reaction time, on average?
- How do these effects vary across people?



Start simple, then criticize

Time as a categorical predictor:

```
fit_fact<-lm( Reaction~as.factor(Days), data=sleep)

BIC(fit_fact)

## [1] 1955.84
```

Time as a continuous predictor:

```
fit<-lm( Reaction~Days, data=sleep)

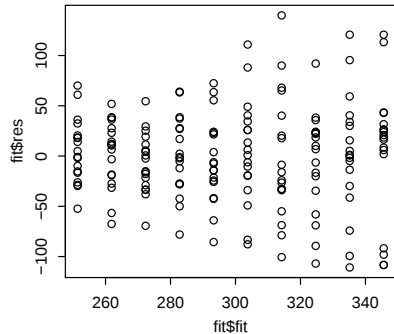
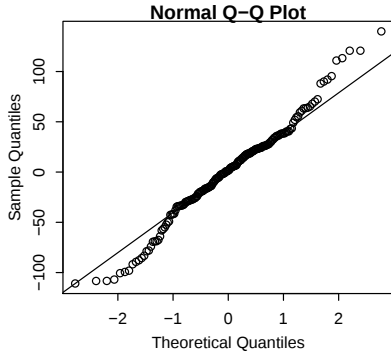
BIC(fit)

## [1] 1915.872
```

```
summary(fit)$coef
```

	Estimate	Std. Error	t value	Pr(> t)
## (Intercept)	251.40510	6.610154	38.033169	2.156888e-87
## Days	10.46729	1.238195	8.453663	9.894096e-15

Start simple, then criticize

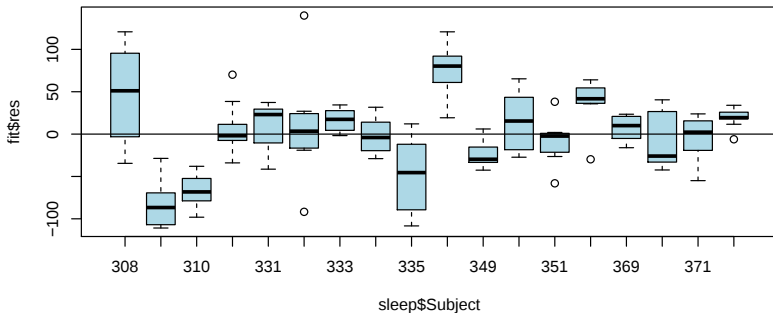


Recall: General order of importance of modeling assumptions:

1. independence;
2. constant variance;
3. normality.

Checking independence

```
plot(fit$res~sleep$Subject,col="lightblue") ; abline(h=0)
```



There is clear evidence that the observations are not independent.

- many subjects have either all residuals above zero, or all residuals below.

Compound symmetry/exchangeable covariance model

Q: How can we model dependence within a subject?

A1: You already have one tool at your disposal, the HNM:

Random effects representation:

$$\begin{aligned}y_{t,j} &= \beta_0 + \beta_1 \times t + b_j + \epsilon_{t,j} \\ \{b_j\} &\sim iid N(0, \tau^2) \\ \{\epsilon_{t,j}\} &\sim iid N(0, \sigma^2)\end{aligned}$$

Correlated data representation:

$$y_{t,j} = \beta_0 + \beta_1 \times t + e_{t,j}$$

$$\text{Cov} \begin{pmatrix} e_{1,j} \\ e_{2,j} \\ \vdots \\ e_{T,j} \end{pmatrix} = \begin{pmatrix} \tau^2 + \sigma^2 & \tau^2 & \dots & \tau^2 \\ \tau^2 & \tau^2 + \sigma^2 & \dots & \tau^2 \\ \vdots & \vdots & \ddots & \vdots \\ \tau^2 & \dots & \tau^2 & \tau^2 + \sigma^2 \end{pmatrix}$$

Recall, under this model $\text{Cor}[y_{t_1,j}, y_{t_2,j}] = \frac{\tau^2}{\tau^2 + \sigma^2}$.

Compound symmetry/exchangeable covariance model

```
fit.lm<-lm(Reaction~Days , data=sleep)  
fit.lme<-lmer(Reaction~Days+(1|Subject) , data=sleep)
```

```
BIC(fit.lm)
```

```
## [1] 1915.872
```

```
BIC(fit.lme)
```

```
## [1] 1807.237
```

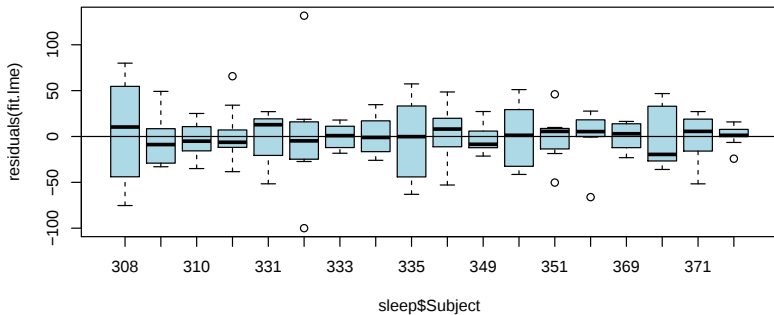
```
summary(fit.lme)$coef
```

```
##           Estimate Std. Error t value  
## (Intercept) 251.40510   9.7467163 25.79383  
## Days        10.46729   0.8042214 13.01543
```

Checking residuals

$$\hat{\epsilon}_{t,j} = y_{t,j} - (\hat{\beta}_0 + \hat{\beta}_1 \times t + \hat{b}_j)$$

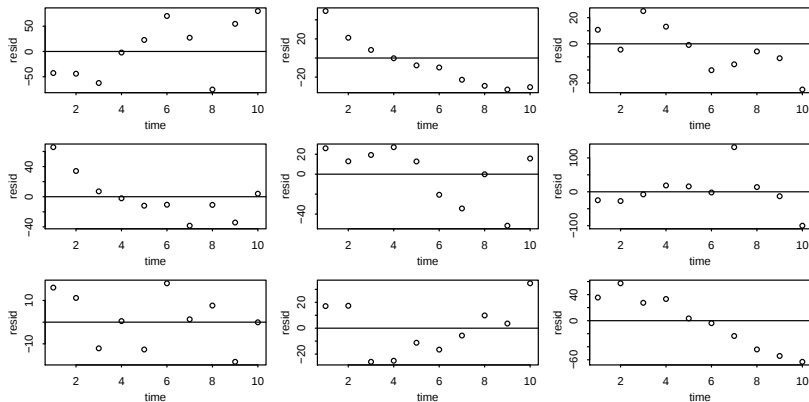
```
plot( residuals(fit.lme)~ sleep$Subject,col="lightblue") ; abline(h=0)
```



Looks much better!

Closer inspection of residuals

$$\hat{\epsilon}_{t,j} = y_{t,j} - (\hat{\beta}_0 + \hat{\beta}_1 \times t + \hat{b}_j)$$



What kind of model might fit better?

Group-specific time trends

```
fit.lme2<-lmer(Reaction~Days+(Days|Subject) , data=sleep)
```

```
BIC(fit.lme)
```

```
## [1] 1807.237
```

```
BIC(fit.lme2)
```

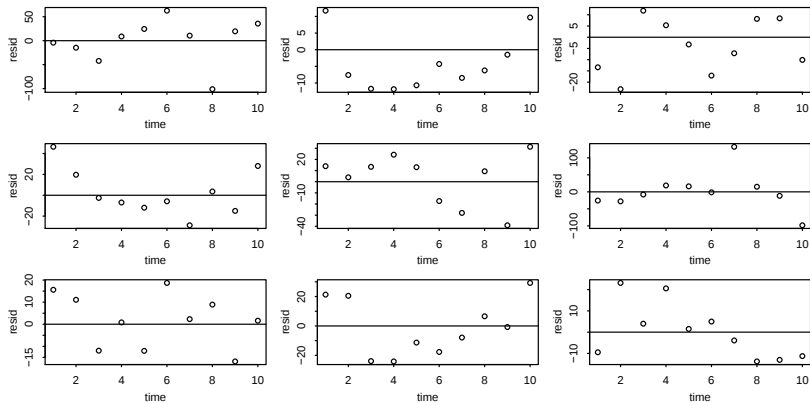
```
## [1] 1774.786
```

```
summary(fit.lme2)$coef
```

##	Estimate	Std. Error	t value
## (Intercept)	251.40510	6.824557	36.838306
## Days	10.46729	1.545789	6.771485

Residual analysis

$$\hat{\epsilon}_{t,j} = y_{t,j} - (\hat{\beta}_0 + \hat{\beta}_1 \times t + \hat{b}_{0j} + \hat{b}_{1j} \times t)$$



Better still, but there still seems to be residual dependence.

Residual correlation

```
RES<- tapply(residuals(fit.lme2),list(sleep$Subject,sleep$Days),"c")
round(RES,2)
```

##	0	1	2	3	4	5	6	7	8	9
## 308	-4.10	-14.63	-42.20	8.78	24.52	62.70	10.54	-101.18	19.59	35.69
## 309	11.73	-7.59	-11.72	-11.84	-10.68	-4.28	-8.46	-6.21	-1.49	9.68
## 310	-13.39	-23.13	11.84	5.34	-3.21	-17.08	-7.13	8.18	8.42	-10.10
## 330	46.45	19.65	-2.55	-6.92	-11.91	-5.77	-28.77	3.60	-14.97	28.08
## 331	13.94	3.94	13.36	24.26	13.02	-17.33	-27.97	9.37	-39.10	31.34
## 332	-25.58	-27.83	-7.87	18.74	16.24	-1.42	132.55	15.02	-11.71	-98.34
## 333	15.60	11.07	-11.96	0.83	-12.05	18.70	2.32	8.89	-16.83	1.60
## 334	21.30	20.49	-23.89	-24.13	-11.32	-17.69	-7.90	6.56	-0.76	29.25
## 335	-9.46	23.16	3.99	20.59	1.52	4.99	-3.91	-13.77	-13.04	-11.26
## 337	26.07	8.41	-32.88	2.54	3.05	10.07	3.39	-3.27	16.80	0.76
## 349	9.91	-7.52	-10.55	-6.20	-22.05	-14.62	-14.47	0.42	16.96	20.68
## 350	17.96	-11.96	-16.29	-34.05	-37.74	5.98	38.62	5.01	19.50	-3.02
## 351	-5.46	36.62	-0.99	2.25	-13.96	11.39	-12.95	-41.55	5.94	24.51
## 352	-50.59	11.92	26.60	32.58	20.46	10.54	-1.86	-9.86	-8.65	-9.76
## 369	17.24	2.42	-20.12	-11.04	14.78	5.84	-24.58	14.07	-5.12	9.78
## 370	-0.53	-6.56	-17.47	-31.19	-19.41	41.95	-36.38	14.77	17.05	8.83
## 371	17.67	10.75	6.73	1.14	-10.96	-15.10	-49.82	-13.94	22.74	31.94
## 372	5.69	-2.00	10.37	11.66	-23.55	7.13	0.25	-2.76	11.41	-5.36

Residual covariance

```
round(cov(RES),1)
```

##	0	1	2	3	4	5	6	7	8	9
## 0	468.4	85.5	-157.8	-222.2	-165.6	-97.3	-326.3	107.2	16.7	343.9
## 1	85.5	285.9	50.0	21.9	-28.3	-39.3	-329.3	-42.1	-69.1	243.9
## 2	-157.8	50.0	305.2	176.1	-0.1	-180.4	-100.6	147.7	-115.1	-81.9
## 3	-222.2	21.9	176.1	337.8	209.2	-23.2	136.7	-115.4	-141.8	-163.6
## 4	-165.6	-28.3	-0.1	209.2	293.8	76.0	134.5	-147.3	-116.4	-104.8
## 5	-97.3	-39.3	-180.4	-23.2	76.0	442.2	59.6	-344.9	106.8	34.1
## 6	-326.3	-329.3	-100.6	136.7	134.5	59.6	1512.7	53.1	-53.9	-1012.4
## 7	107.2	-42.1	147.7	-115.4	-147.3	-344.9	53.1	754.5	-152.8	-294.7
## 8	16.7	-69.1	-115.1	-141.8	-116.4	106.8	-53.9	-152.8	281.5	80.2
## 9	343.9	243.9	-81.9	-163.6	-104.8	34.1	-1012.4	-294.7	80.2	926.9

Residual correlation

```
round(cor(RES),2)
```

##	0	1	2	3	4	5	6	7	8	9
## 0	1.00	0.23	-0.42	-0.56	-0.45	-0.21	-0.39	0.18	0.05	0.52
## 1	0.23	1.00	0.17	0.07	-0.10	-0.11	-0.50	-0.09	-0.24	0.47
## 2	-0.42	0.17	1.00	0.55	0.00	-0.49	-0.15	0.31	-0.39	-0.15
## 3	-0.56	0.07	0.55	1.00	0.66	-0.06	0.19	-0.23	-0.46	-0.29
## 4	-0.45	-0.10	0.00	0.66	1.00	0.21	0.20	-0.31	-0.40	-0.20
## 5	-0.21	-0.11	-0.49	-0.06	0.21	1.00	0.07	-0.60	0.30	0.05
## 6	-0.39	-0.50	-0.15	0.19	0.20	0.07	1.00	0.05	-0.08	-0.86
## 7	0.18	-0.09	0.31	-0.23	-0.31	-0.60	0.05	1.00	-0.33	-0.35
## 8	0.05	-0.24	-0.39	-0.46	-0.40	0.30	-0.08	-0.33	1.00	0.16
## 9	0.52	0.47	-0.15	-0.29	-0.20	0.05	-0.86	-0.35	0.16	1.00

Covariance models

Each model we've considered so far corresponds to a different covariance model:

Linear model: $y_{t,j} = \beta_1 + \beta_2 \times t + \epsilon_{t,j}$

$$\text{Cov}[y_{s,j}, y_{t,j}] = 0$$

HLM with random intercepts: $y_{t,j} = \beta_1 + \beta_2 \times t + b_j + \epsilon_{t,j}$

$$\text{Cov}[y_{s,j}, y_{t,j}] = \tau^2, \text{ where } \text{Var}[b_j] = \tau^2$$

HLM with random intercepts and time trends:

$$\text{Cov}[y_{s,j}, y_{t,j}] = \tau_1^2 + \tau_{12} \times (t + s + t \times s \times \tau_1^2), \text{ where } \text{Cov} \begin{pmatrix} b_{0,j} \\ b_{1,j} \end{pmatrix} = \begin{pmatrix} \tau_1^2 & \tau_{12} \\ \tau_{12} & \tau_2^2 \end{pmatrix}$$

Residual covariance models

In all the models considered so far, we've assumed

$$\text{Cov} \begin{pmatrix} \epsilon_{1,j} \\ \vdots \\ \epsilon_{n,j} \end{pmatrix} = \sigma^2 \mathbf{I} = \begin{pmatrix} \sigma^2 & 0 & \cdots & 0 \\ 0 & \sigma^2 & \cdots & 0 \\ \vdots & & & \vdots \\ 0 & \cdots & 0 & \sigma^2 \end{pmatrix}$$

- “micro” level errors are independent, or equivalently
- all within-group dependence is explained by random effects.

In cases where it appears random effects are not accounting for the dependence, we may want to allow for non-independence among the $\epsilon_{i,j}$'s.

$$\text{Cov} \begin{pmatrix} \epsilon_{1,j} \\ \vdots \\ \epsilon_{n,j} \end{pmatrix} = \Sigma_{\epsilon} = \begin{pmatrix} \sigma_1^2 & \sigma_{1,2} & \cdots & \sigma_{1,T} \\ \sigma_{1,2} & \sigma_2^2 & \cdots & \sigma_{2,T} \\ \vdots & & & \vdots \\ \sigma_{1,T} & \cdots & \sigma_{T-1,T} & \sigma_T^2 \end{pmatrix}$$

Model fitting with nlme

```
fit.lme4<-lmer(Reaction~Days+(Days|Subject), data=sleep,REML=FALSE)
logLik(fit.lme4)
## 'log Lik.' -875.9697 (df=6)
```

```
library(nlme)
fit.nlme<-lme(Reaction~Days, random=~Days|Subject, data=sleep,method="ML")
logLik(fit.nlme)
## 'log Lik.' -875.9697 (df=6)
```

nlme output

```
summary(fit.nlme)

## Linear mixed-effects model fit by maximum likelihood
## Data: sleep
##      AIC      BIC    logLik
## 1763.939 1783.097 -875.9697
##
## Random effects:
## Formula: ~Days | Subject
## Structure: General positive-definite, Log-Cholesky parametrization
##           StdDev    Corr
## (Intercept) 23.780376 (Intr)
## Days        5.716807 0.081
## Residual    25.591842
##
## Fixed effects: Reaction ~ Days
##           Value Std.Error DF t-value p-value
## (Intercept) 251.40510  6.669396 161 37.69533      0
## Days        10.46729  1.510647 161  6.92901      0
## Correlation:
##      (Intr)
## Days -0.138
##
## Standardized Within-Group Residuals:
##           Min           Q1           Med           Q3           Max
## -3.94156355 -0.46559311  0.02894656  0.46361051  5.17933587
##
## Number of Observations: 180
## Number of Groups: 18
```

Correlation models

?corClasses

Correlation Structure Classes

Description:

Standard classes of correlation structures (`corStruct`) available in the nlme package.

Value:

Available standard classes:

`corAR1`: autoregressive process of order 1.

`corARMA`: autoregressive moving average process, with arbitrary orders for the autoregressive and moving average components.

`corCAR1`: continuous autoregressive process (AR(1) process for a continuous time covariate).

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`corRatio`: Rational quadratics spatial correlation.

`corSpher`: spherical spatial correlation.

`corSymm`: general correlation matrix, with no additional structure.

Correlation models for longitudinal data

Several of the correlation models are designed specifically for longitudinal data.

- **corAR1**: First-order discrete-time (fixed occasion) autocorrelation model.
- **corARMA**: A general class of discrete-time models.
- **corCAR1**: A first order continuous time model.

In particular, AR1 model is a two-parameter covariance model for which

$$\text{Cov} \begin{pmatrix} \epsilon_{1,j} \\ \vdots \\ \epsilon_{T,j} \end{pmatrix} = \sigma^2 \begin{pmatrix} 1 & \rho & \rho^2 & \cdots & \rho^{T-1} \\ \rho & 1 & \rho & \cdots & \rho^{T-2} \\ \vdots & & & & \vdots \\ \rho^{T-1} & \rho^{T-2} & \cdots & \rho & 1 \end{pmatrix},$$

where $\sigma^2 > 0$ and $\rho \in (-1, 1)$.

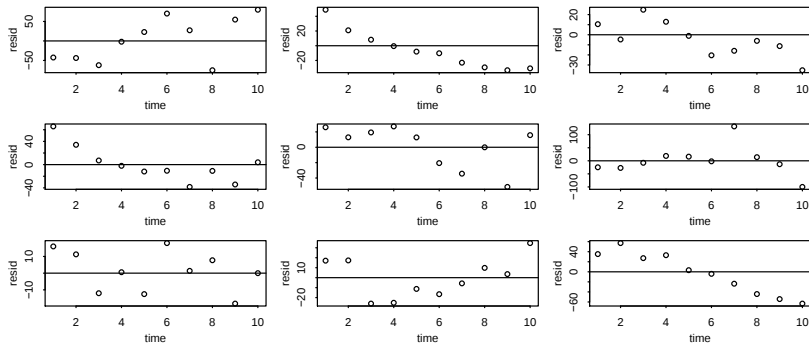
Exercise: Describe the correlation between two observations as a function of their time difference.

Model comparison and selection

Random intercept:

```
fit1<-lme(Reaction~Days, random=~1|Subject, data=sleep,method="ML")
```

$$\hat{\epsilon}_{t,j} = y_{t,j} - (\hat{\beta}_0 + \hat{\beta}_1 \times t + \hat{b}_j)$$



What kind of model might fit better? Random time trend or AR1?

Model comparison

Random intercept and time trend:

```
fit2a<-lme(Reaction~Days, random=~Days|Subject, data=sleep,method="ML")  
BIC(fit2a)  
## [1] 1783.097
```

Random intercept and autocorrelated errors:

```
fit2b<-lme(Reaction~Days, random=~1|Subject,correlation=corAR1(), data=sleep,method="ML")  
BIC(fit2b)  
## [1] 1770.795
```

Limitations

```
fit2b<-lme(Reaction~Days, random=~Days|Subject,correlation=corAR1(), data=sleep,method="ML")  
## Error in lme.formula(Reaction ~ Days, random = ~Days | Subject, correlation =  
corAR1(), : nlminb problem, convergence error code = 1  
## message = iteration limit reached without convergence (10)
```

Longitudinal data with covariates

Male subset of SOEP data

```
mappy[1:10, c(1, 14, 4, 16, 17, 7) ]
```

##	id	lifeSat	age	birthyear	logincome	yearsEd
## 1	101	8	55	1930	10.04591	15
## 2	101	10	56	1930	10.05470	15
## 3	101	8	57	1930	10.05511	15
## 4	101	8	58	1930	10.06734	15
## 5	101	8	59	1930	10.03707	15
## 7	1701	8	55	1948	11.12692	15
## 8	1701	10	56	1948	11.27367	15
## 9	1701	10	57	1948	11.26152	15
## 10	1701	9	58	1948	11.28407	15
## 13	2301	7	55	1946	11.01356	18

We'll take `lifeSat` as the response. Covariates include

- `age`: age at the time of the survey;
- `birthyear`: year of birth;
- `logincome`: log pre-tax income;
- `yearsEd`: years of formal education.

Types of variables

Macro variables: Vary across individuals but not time - `birthyear`, `yearsEd`

Micro variables: Vary within individuals/ across time - `age`, `logincome`.

The types effects we may want to include are

- fixed effects for macro and micro variables, and their interactions.
- random effects for micro variables, and their interactions.

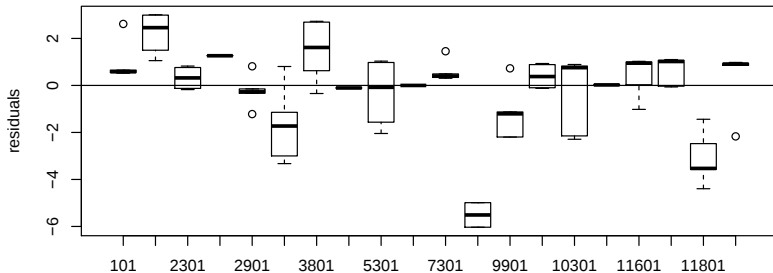
Random effects model with independent errors and main effects

```
fit0<-lm(lifeSat ~ age + birthyear + logincome + yearsEd, data=mappy )
```

```
summary(fit0)$coef
```

##		Estimate	Std. Error	t value	Pr(> t)
##	(Intercept)	58.97223503	8.772117828	6.722691	1.989586e-11
##	age	0.03301564	0.015410178	2.142456	3.220633e-02
##	birthyear	-0.02845157	0.004463747	-6.373920	2.013280e-10
##	logincome	0.09934413	0.009347222	10.628198	4.255265e-26
##	yearsEd	0.03166438	0.010570551	2.995527	2.753540e-03

Residual plots:

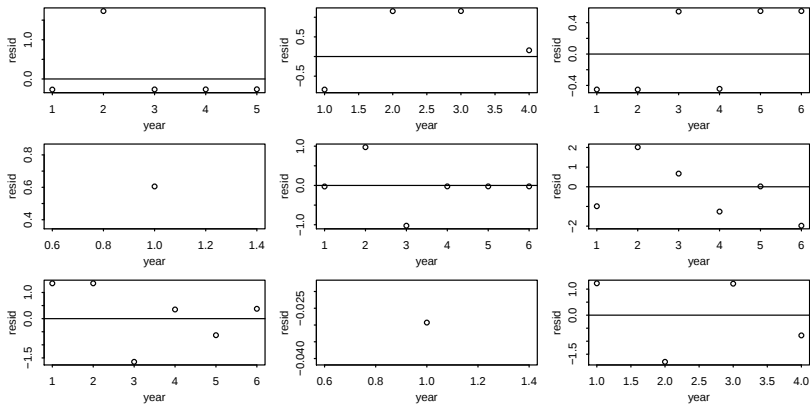


Random effects model with independent errors and main effects

```
fit1<-lme(lifeSat ~ age + birthyear + logincome + yearsEd,  
          random= ~1|id,data=mappy , method="ML")  
  
fit2<-lme(lifeSat ~ age + birthyear + logincome + yearsEd,  
          random= ~1+age|id,data=mappy , method="ML")  
  
fit3<-lme(lifeSat ~ age + birthyear + logincome + yearsEd,  
          random= ~1+age+logincome|id,data=mappy , method="ML")
```

```
BIC(fit0)  
## [1] 19471.85  
  
BIC(fit1)  
## [1] 17602.67  
  
BIC(fit2)  
## [1] 17619.65  
  
BIC(fit3)  
## [1] 17625.13
```

Temporal dependence of residuals



AR1 covariance model

```
fit4<-lme(lifeSat ~ age + birthyear + logincome + yearsEd,  
          random= ~1|id,correlation=corAR1(),data=mappy,method="ML")  
  
BIC(fit4)  
  
## [1] 17590.9
```

Looking for interactions

```
add1(fit4,"age*birthyear",data=mappy,test="Chisq")

## Single term additions
##
## Model:
## lifeSat ~ age + birthyear + logincome + yearsEd
##           Df    AIC    LRT Pr(>Chi)
## <none>           17539
## age*birthyear  1 17528 12.859 0.0003359 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
add1(fit4,"age*logincome",data=mappy,test="Chisq")

## Single term additions
##
## Model:
## lifeSat ~ age + birthyear + logincome + yearsEd
##           Df    AIC    LRT Pr(>Chi)
## <none>           17539
## age*logincome  1 17538 2.544  0.1107
```

```
add1(fit4,"age*yearsEd",data=mappy,test="Chisq")
```

```
## Single term additions
##
## Model:
## lifeSat ~ age + birthyear + logincome + yearsEd
##           Df    AIC    LRT Pr(>Chi)
```

More interactions

```
add1(fit4,"birthyear*logincome",data=mappy,test="Chisq")
```

```
## Single term additions
##
## Model:
## lifeSat ~ age + birthyear + logincome + yearsEd
##           Df    AIC    LRT Pr(>Chi)
## <none>           17539
## birthyear*logincome  1 17540 1.0472  0.3061
```

```
add1(fit4,"birthyear*yearsEd",data=mappy,test="Chisq")
```

```
## Single term additions
##
## Model:
## lifeSat ~ age + birthyear + logincome + yearsEd
##           Df    AIC    LRT Pr(>Chi)
## <none>           17539
## birthyear*yearsEd  1 17541 0.3618  0.5475
```

```
add1(fit4,"logincome*yearsEd",data=mappy,test="Chisq")
```

```
## Single term additions
##
## Model:
## lifeSat ~ age + birthyear + logincome + yearsEd
##           Df    AIC    LRT Pr(>Chi)
## <none>           17539
## logincome*yearsEd  1 17540 0.51157  0.4745
```

“Final” model

```
fit5<-lme(lifeSat ~ age + birthyear + logincome + yearsEd + age*birthyear,
          random= ~1|id,correlation=corAR1(),data=mappy,method="ML")

summary(fit5)

## Linear mixed-effects model fit by maximum likelihood
## Data: mappy
##      AIC      BIC    logLik
## 17528.13 17586.53 -8755.065
##
## Random effects:
## Formula: ~1 | id
##      (Intercept) Residual
## StdDev:      1.297903 1.238987
##
## Correlation Structure: AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##      Phi
## 0.09557115
## Fixed effects: lifeSat ~ age + birthyear + logincome + yearsEd + age * birthyear
##              Value Std.Error   DF   t-value p-value
## (Intercept)  907.3259 232.77705 3868   3.897832  0.0001
## age          -14.7224   4.10112 3868  -3.589834  0.0003
## birthyear    -0.4652   0.12016  988  -3.871912  0.0001
## logincome     0.0426   0.00983 3868   4.336077  0.0000
## yearsEd       0.0482   0.01833 3868   2.628823  0.0086
## age:birthyear  0.0076   0.00212 3868   3.590019  0.0003
## Correlation:
##      (Intr) age    brthyr logncm versEd
```