Not to turn in:

A. Practice to model splines. If you did it, please send in your graphs and equations and I can review it.

To turn in:

- (1) Patients with Chronic Beryllium Disease have ongoing visits at NJH to monitor their health. Once measure taken during their visits is AADO₂R (alveolar-arterial oxygen tension difference at rest; the lower the value, the better the health; see course notes for more detail). The goal is to estimate the effect of the disease on AADO₂R over time, after accounting for variables known to be related to it (age, gender, height). One modeling challenge is that subjects come in on different days, have different times between visits, and don't have the same number of visits
 - a. Consider the model for $AADO_2R$ as a function of time since first exposure (*ntep*, in years), age (*ageep*, in years), *height* (inches) and *gender*; the variables *ntep*, *ageep* and *height* are time-varying; a random intercept is included for subjects. Write out a statistical model.

Solution:

```
Y_{ij} = \beta_0 + \beta_1 ntep_{ij} + \beta_2 age_{ij} + \beta_3 height_{ij} + \beta_4 female_i + b_i + \varepsilon_{ij}  (full rank)
```

where *i* denotes subject, *j* denotes time; $Y=AADO_2R$; female = 1 for females, 0 for males; $b_i \sim N(0, \sigma_b^2)$, $\varepsilon_i \sim N(0, \sigma_\varepsilon^2 \mathbf{I})$. You can also use a variable such as θ_k , k=1,2 (for F, M) in place of β_4 female_i, inducing a less-than-full-rank model. This is the model SAS fits, which is

$$Y_{ijk} = \beta_0 + \beta_1 ntep_{ij} + \beta_2 age_{ij} + \beta_3 height_{ij} + \theta_k + b_i + \varepsilon_{kij}$$
 (less than full rank)

Be careful not to mix different types of notation. If you use the 'regression' approach (i.e., full-rank model), you need to use a dummy (0/1) variable; if you specify a variable that has values such as 1 or 2, M or F, you need to demonstrate it is a class variable with 2 levels.

b. Using software, fit the model. (Note: *ageep*, *height* and *gender* are 'a priori' covariates. They stay in the model regardless of their significance.)

c. Consider the same model as above, but include a spatial power covariance structure for repeated measures. Does the addition of the new \mathbf{R}_i structure improve the model fit?

Solution: Yes, the AIC drops a couple of points after adding the SP(EXP) structure.

```
proc mixed data=long.Be_study;
                                            Fit Statistics
class sex id;
model aado2r=ntep height sex ageep
                                            -2 Res Log Likelihood
                                                                        1542.8
                                            AIC (smaller is better)
                                                                        1548.8
   / solution;
random intercept / subject=id;
                                            AICC (smaller is better)
                                                                        1548.9
                                            BIC (smaller is better)
                                                                        1555.1
repeated / subject=id
type=sp(exp)(ntep); run;
```

d. Using relevant output and formulas (for the model in part c), estimate the correlation between $Y_{ij}|b_{0i}$ and $Y_{ij'}|b_{0i}$ for responses that are (i) 4 months apart, (ii) 1 year apart. Interpret the results for the study.

Covariance Parameter Estimates						
Cov Parm	Subject	Estimate				
Intercept	id	12.2551				
SP(POW)	id	0.04757				
Residual		28.3927				

Solution: here I am asking for elements of the R matrix (when conditioning on the random effects).

- (i) $0.04757^{(4/12)} = 0.3623$
- (ii) $0.04757^1 = 0.04757$

Interpretation for (i): For a given subject and their deviation from the population mean, two measures 3 months apart have a correlation of 0.36.

e. Repeat part d, but for the correlation between Y_{ij} and $Y_{ij'}$ (unconditional responses).

Solution: here I am asking for elements of the V matrix (not conditioning on the random effects)

- (i) $(28.3927*0.04757^{(4/12)} + 12.2551) / (28.3927+12.2551) = 0.5546$
- (ii) (28.3927*0.04757+12.2551) / (28.3927+12.2551) = 0.3347

Interpretation for (i): Two measures within a subject that are 3 months apart have a correlation of 0.55 (not conditioning on their random deviation from the mean).

f. Include a streamlined version of the output and write a few sentences of summary of the results. Make sure to include an estimate the progression of the illness (slope of *ntep* term), with a 95% confidence interval. For *AADO2R*, remember that the lower the value, the better the health. What can be concluded about the progression of the illness over time?

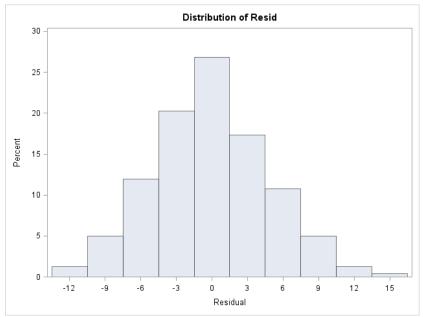
<u>Solution</u>: By adding CL as an option in the MODEL statement, we can get confidence limits. Below are parameter estimates in the model, for both fixed effects and covariance parameters. There is an average increase of 0.2117 units in AADO2R (units are mmHg) for each additional year with the disease, after removing effects of aging (95% CI: 0.055 to 0.368). SAS output follows on the next page.

Cov Parm	Subject	Estimate
Intercept	id	12.2621
SP(EXP)	id	0.3283
Residual		28.3902

Solution for Fixed Effects

			Standard						
Effect	sex	Estimate	Error	DF	t Value	Pr > t	Alpha	Lower	Upper
Intercept		-3.3030	14.5816	58	-0.23	0.8216	0.05	-32.4912	25.8852
ntep		0.2117	0.07919	179	2.67	0.0082	0.05	0.05544	0.3680
height		0.1326	0.2214	179	0.60	0.5501	0.05	-0.3043	0.5694
sex	1	-2.1333	1.8690	179	-1.14	0.2552	0.05	-5.8213	1.5547
sex	2	0							
ageep		0.007743	0.07610	179	0.10	0.9191	0.05	-0.1424	0.1579

g. Note that *AADO₂R* actually is a continuous variable with a clump at 0. The best model for these data might actually be a mixture model (to be discussed later, if time). However, examine a histogram of the residuals to determine whether the current model may be acceptable. Comment on what you see. [Aside: in this case, it is likely that the 0's should have been positive since a 0 value does not appear to be theoretically possible for this variable; from what I can see they can be attributed to measurement error, which may have affected other values as well.]



Solution: The residuals are pretty normally distributed! So, despite having a mixed distribution (continuous plus 0's), it seems that we've satisfied distributional assumptions for the model. Still, we can use mixture model techniques for these data, to be discussed soon. Also, a more thorough investigation would involve reviewing other model diagnostics, not just a histogram of the residuals...