Problems with the Error Solutions

- 1. Here we go...
- (a) With R,

```
> out <- Im(Lab ~ Field, pipeline)
> summary(out)
```

Call:

Im(formula = Lab ~ Field, data = pipeline)

Residuals:

Min 1Q Median 3Q Max -21.985 -4.072 -1.431 2.504 24.334

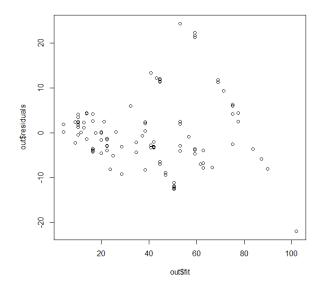
Coefficients:

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

Residual standard error: 7.865 on 105 degrees of freedom Multiple R-squared: 0.8941, Adjusted R-squared: 0.8931 F-statistic: 886.7 on 1 and 105 DF, p-value: < 2.2e-16

By non-constant variance, Faraway means non-constant variance of the residuals:

> plot(out\$residuals ~ out\$fit)



Holy \$#@! This certainly looks like a non-constant variance situation. Just for fun, I checked for serial correlation of the residuals and got nothing:

```
> cor(residuals(out)[-1], residuals(out)[-length(residuals(out))])
[1] 0.08754773
```

(b) Following Faraway's instructions...

This is certainly consistent with our residuals vs. fits plot from before; it seems the variance increases in the Field variable.

Next, we are to assume that $var(Lab) = a_0$ Field ^ a_1. Taking logs of both sides gives

$$log(var(Lab)) = log(a_0) + a_1 * log(Field).$$

Then we can regress like Faraway suggests using simple linear regression to estimate a_0 and a_1:

```
> logs.reg <- lm(log(varlab) ~ log(meanfield))
> summary(logs.reg)

Call:
lm(formula = log(varlab) ~ log(meanfield))

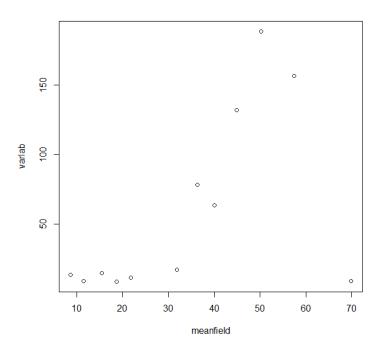
Residuals:
    Min    1Q Median    3Q Max
-2.2038 -0.6729    0.1656    0.7205    1.1891

Coefficients:
        Estimate Std. Error t value Pr(>|t|)
(Intercept)    -0.3538    1.5715    -0.225    0.8264
log(meanfield)    1.1244    0.4617    2.435    0.0351 *
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1

Residual standard error: 1.018 on 10 degrees of freedom
Multiple R-squared: 0.3723, Adjusted R-squared: 0.3095
```

F-statistic: 5.931 on 1 and 10 DF, p-value: 0.03513

The output suggests estimating a_0 with e ^ -.3538 = .7020 and estimating a_1 with 1.1244. I kept the last point. However, as Faraway suggests, you might probably want to exclude it because



So I'll throw out the 12th point and redo...

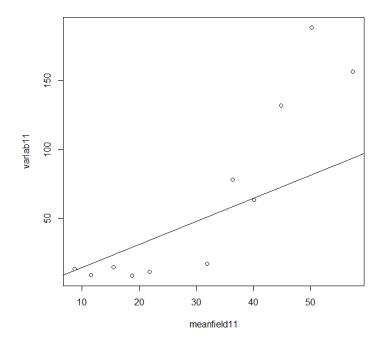
> varlab <- data.frame(varlab)

```
> meanfield <- data.frame(meanfield)
> varlab11 <- varlab[1:11,]
> meanfield11 <- meanfield[1:11,]
> logs11.reg <- lm(log(varlab11) ~ log(meanfield11))
> summary(logs11.reg)
Call:
lm(formula = log(varlab11) ~ log(meanfield11))
Residuals:
  Min
          1Q Median
                         3Q
                               Max
-1.00477 -0.42268 0.05989 0.37854 0.93815
Coefficients:
         Estimate Std. Error t value Pr(>|t|)
             -1.9352 1.0929 -1.771 0.110403
log(meanfield11) 1.6707 0.3296 5.070 0.000672 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
```

Residual standard error: 0.657 on 9 degrees of freedom Multiple R-squared: 0.7406, Adjusted R-squared: 0.7118

F-statistic: 25.7 on 1 and 9 DF, p-value: 0.0006723

```
> plot(varlab11 ~ meanfield11, xlab="meanfield11", ylab="varlab11")
> abline(logs11.reg)
>
```



Even though the trend does not appear linear, this clearly gives a better approximation for the a_0 and a_1: a_0 is approximately $e^{-1.9352} = 0.1444$ and a_1 is approximately 1.6707. Now it seems appropriate to set the WLS regression weights to $[a_0 * Field * a_1] * -1$:

```
> a0 = .1444
> a1 = 1.6707
> w <- 1/(a0 * pipeline$Field ^ a1)
> wlmod <- lm(Lab ~ Field, data = pipeline, weights=w)
> summary(wlmod)

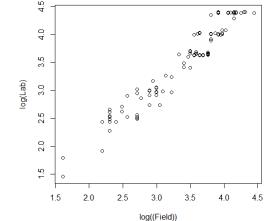
Call:
lm(formula = Lab ~ Field, data = pipeline, weights = w)

Weighted Residuals:
    Min    1Q Median    3Q Max
-1.7433 -0.6719 -0.2493    0.5967    2.7277
```

Coefficients:

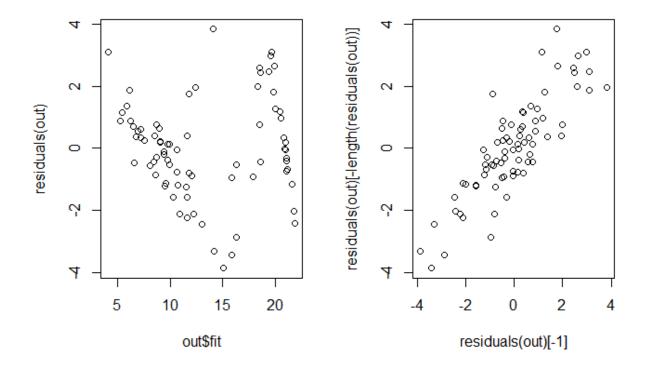
```
Estimate Std. Error t value Pr(>|t|)
(Intercept) -1.05531  0.69766 -1.513  0.133
Field
       Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 0.9846 on 105 degrees of freedom
Multiple R-squared: 0.921, Adjusted R-squared: 0.9202
F-statistic: 1224 on 1 and 105 DF, p-value: < 2.2e-16
      (c) Out of all of these, I think taking logs of both variables worked out nicely:
> outloglog <- Im(log(Lab) ~ log((Field)))
> summary(outloglog)
Call:
Im(formula = log(Lab) ~ log((Field)))
Residuals:
                          Max
  Min
        1Q Median
                      3Q
-0.40212 -0.11853 -0.03092 0.13424 0.40209
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 0.1837 on 105 degrees of freedom
Multiple R-squared: 0.9337, Adjusted R-squared: 0.9331
F-statistic: 1479 on 1 and 105 DF, p-value: < 2.2e-16
```

> plot(log(Lab) ~ log(Field))



2. Here we go!

```
> attach(divusa)
> out <- Im(divorce ~ unemployed + femlab + marriage + birth + military)
> summary(out)
Call:
Im(formula = divorce ~ unemployed + femlab + marriage + birth +
  military)
Residuals:
  Min 1Q Median 3Q Max
-3.8611 -0.8916 -0.0496 0.8650 3.8300
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
(Intercept) 2.48784 3.39378 0.733 0.4659
unemployed -0.11125  0.05592 -1.989  0.0505.
femlab 0.38365 0.03059 12.543 < 2e-16 ***
marriage 0.11867 0.02441 4.861 6.77e-06 ***
birth -0.12996 0.01560 -8.333 4.03e-12 ***
military -0.02673 0.01425 -1.876 0.0647.
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 1.65 on 71 degrees of freedom
Multiple R-squared: 0.9208, Adjusted R-squared: 0.9152
F-statistic: 165.1 on 5 and 71 DF, p-value: < 2.2e-16
Checking for serial correlation...
> cor(residuals(out)[-1], residuals(out)[-length(residuals(out))])
[1] 0.8469792
There certainly seems to be some serial correlation. Let's look at a couple of plots.
> par(mfrow=c(1,2))
> plot(residuals(out) ~ out$fit)
> plot(residuals(out)[-1], residuals(out)[-length(residuals(out))])
```



The first plot on the left indicates a sort-of down, up, down pattern of the residuals with the fits. The second plot indicates serial correlation in the residuals. In fact, I computed this correlation previously to be 0.8469792 (see above).

```
(b) > glmod <- gls(divorce ~ unemployed + femlab + marriage + birth + military, method= "ML",
correlation=corAR1(form= ~year), data = na.omit(divusa))
> summary(glmod)
Generalized least squares fit by maximum likelihood
Model: divorce ~ unemployed + femlab + marriage + birth + military
Data: na.omit(divusa)
          BIC logLik
   AIC
 179.9523 198.7027 -81.97613
Correlation Structure: AR(1)
Formula: ~year
Parameter estimate(s):
   Phi
0.9715486
Coefficients:
        Value Std.Error t-value p-value
```

(Intercept) -7.059682 5.547193 -1.272658 0.2073 unemployed 0.107643 0.045915 2.344395 0.0219

```
femlab 0.312085 0.095151 3.279878 0.0016
marriage 0.164326 0.022897 7.176766 0.0000
birth -0.049909 0.022012 -2.267345 0.0264
military 0.017946 0.014271 1.257544 0.2127
```

Correlation:

(Intr) unmply femlab marrig birth unemployed -0.420 femlab -0.802 0.240 marriage -0.516 0.607 0.307 birth -0.379 0.041 0.066 -0.094 military -0.036 0.436 -0.311 0.530 0.128

Standardized residuals:

Min Q1 Med Q3 Max -1.4509327 -0.9760939 -0.6164694 1.1375377 2.1593261

Residual standard error: 2.907665

Degrees of freedom: 77 total; 71 residual
> intervals(glmod, which="var-cov")

Approximate 95% confidence intervals

Correlation structure:

lower est. upper
Phi 0.6528097 0.9715486 0.9980192
attr(,"label")
[1] "Correlation structure:"

Residual standard error:

lower est. upper 0.7974404 2.9076645 10.6020628

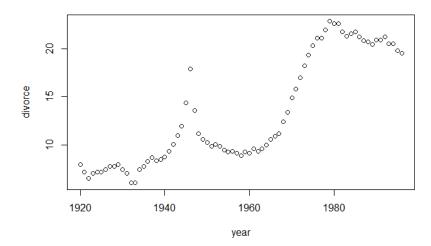
So it seems the phi value is very likely greater than 0, and probably around .97. That is, there is a high degree of serial correlation in the residuals.

The GLS model is pretty consistent with the lm() model. All the predictors are statistically significant with the possible exception of military. The intercept is likely not needed also.

(c) Why might there be correlation in the errors? Check this out:

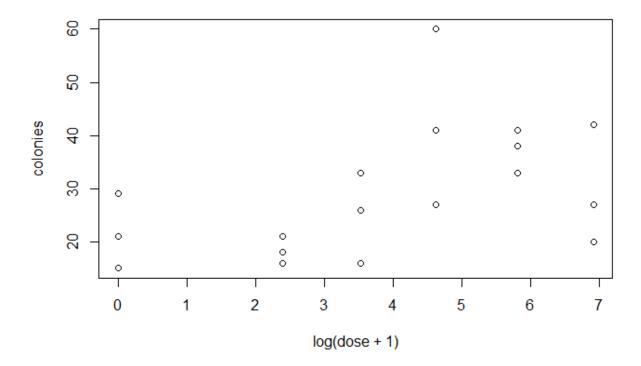
```
> plot(divorce ~ year)
```

Note the divorce rate shot up in the mid-1940s, perhaps having to do with husbands going away to war. It then fell back down through the 1950s, and began increasing again in the 1960s, when we know that divorce began to be more common.



3. Salmonella!

```
> attach(salmonella)
> out <- Im(colonies ~ log(dose+1))
> summary(out)
Call:
Im(formula = colonies ~ log(dose + 1))
Residuals:
  Min 1Q Median 3Q Max
-16.376 -6.882 -1.509 5.400 29.119
Coefficients:
       Estimate Std. Error t value Pr(>|t|)
(Intercept) 19.823 5.064 3.915 0.00123 **
log(dose + 1) 2.396 1.128 2.125 0.04955 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 10.84 on 16 degrees of freedom
Multiple R-squared: 0.2201, Adjusted R-squared: 0.1713
F-statistic: 4.514 on 1 and 16 DF, p-value: 0.04955
> plot(colonies ~ log(dose+1))
```



There could be a lack of linear fit here. Let's see:

> pureErrorAnova(out)

Analysis of Variance Table

```
Response: colonies
```

Df Sum Sq Mean Sq F value Pr(>F)

log(dose + 1) 1 530.71 530.71 5.8356 0.03257 *

Residuals 16 1881.06 117.57

Lack of fit 4 789.73 197.43 2.1709 0.13420

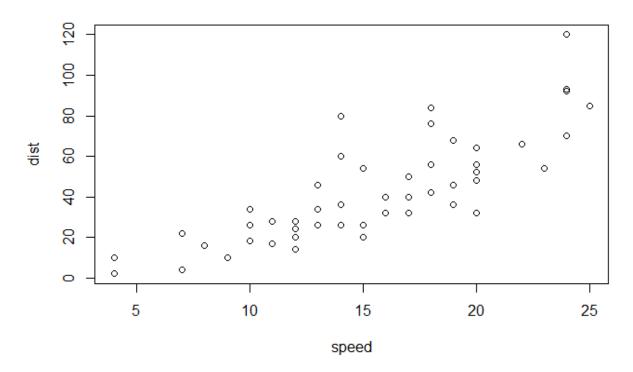
Pure Error 12 1091.33 90.94

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

The lack-of-linear fit p-value is .13420, and so I would conclude there is insufficient evidence to conclude there exists a lack of linear fit.

4. Cars!!!

```
> out <- Im(dist ~ speed)
> plot(dist ~ speed)
```



> summary(out)

Call:

lm(formula = dist ~ speed)

Residuals:

Min 1Q Median 3Q Max -29.069 -9.525 -2.272 9.215 43.201

Coefficients:

Estimate Std. Error t value Pr(>|t|)
(Intercept) -17.5791 6.7584 -2.601 0.0123 *
speed 3.9324 0.4155 9.464 1.49e-12 ***

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

Residual standard error: 15.38 on 48 degrees of freedom Multiple R-squared: 0.6511, Adjusted R-squared: 0.6438

```
F-statistic: 89.57 on 1 and 48 DF, p-value: 1.49e-12
```

> pureErrorAnova(out)

Analysis of Variance Table

Response: dist

Df Sum Sq Mean Sq F value Pr(>F)

speed 1 21185.5 21185.5 97.0836 4.558e-11 ***

Residuals 48 11353.5 236.5

Lack of fit 17 4588.7 269.9 1.2369 0.2948

Pure Error 31 6764.8 218.2

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

It doesn't appear that there is much statistical evidence pointing to lack of linear fit.

5. Here is the least-squares output:

```
> Isout <- Im(stack.loss ~ ., stackloss)
> summary(Isout)
Im(formula = stack.loss ~ ., data = stackloss)
Residuals:
 Min 1Q Median 3Q Max
-7.2377 -1.7117 -0.4551 2.3614 5.6978
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
Air.Flow 0.7156 0.1349 5.307 5.8e-05 ***
Water.Temp 1.2953 0.3680 3.520 0.00263 **
Acid.Conc. -0.1521  0.1563 -0.973  0.34405
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '' 1
Residual standard error: 3.243 on 17 degrees of freedom
Multiple R-squared: 0.9136, Adjusted R-squared: 0.8983
F-statistic: 59.9 on 3 and 17 DF, p-value: 3.016e-09
Next is the LAD output:
> require(quantreg)
> LADout <- rq(stack.loss ~., data = stackloss)
> summary(LADout)
Call: rq(formula = stack.loss ~ ., data = stackloss)
tau: [1] 0.5
Coefficients:
      coefficients lower bd upper bd
(Intercept) -39.68986 -41.61973 -29.67754
Air.Flow 0.83188 0.51278 1.14117
Water.Temp 0.57391 0.32182 1.41090
Acid.Conc. -0.06087 -0.21348 -0.02891
Next is the output from the Huber method:
> require(MASS)
> Huberout <- rlm(stack.loss ~ ., data = stackloss)
> summary(Huberout)
```

```
Call: rlm(formula = stack.loss ~ ., data = stackloss)
Residuals:
    Min 1Q Median 3Q Max
-8.91753 -1.73127 0.06187 1.54306 6.50163

Coefficients:require(MASS)

    Value Std. Error t value
(Intercept) -41.0265 9.8073 -4.1832
Air.Flow 0.8294 0.1112 7.4597
Water.Temp 0.9261 0.3034 3.0524
Acid.Conc. -0.1278 0.1289 -0.9922
```

Residual standard error: 2.441 on 17 degrees of freedom

Finally, below is the output from the least trimmed squares method. I did an exhaustive search since the data set is not that large:

```
> LTSout <- Itsreg(stack.loss ~ ., data = stackloss, nsamp="exact")
> summary(LTSout)
      Length Class Mode
       1 -none- numeric
crit
        1 -none- character
coefficients 4 -none- numeric
bestone 4 -none- numeric
fitted.values 21 -none- numeric
residuals 21 -none- numeric
scale
        2 -none- numeric
terms
        3 terms call
call
       5 -none- call
xlevels 0 -none- list
         4 data.frame list
model
> coef(LTSout)
(Intercept) Air.Flow Water.Temp Acid.Conc.
-3.580556e+01 7.500000e-01 3.333333e-01 3.489094e-17
```

The Acid.Conc variable doesn't seem very important.