Case study: asphalt

The asphalt data

- 31 asphalt pavements prepared under different conditions. How does quality of pavement depend on these?
- Variables:
 - pct.a.surf Percentage of asphalt in surface layer
 - pct.a.base Percentage of asphalt in base layer
 - fines Percentage of fines in surface layer
 - voids Percentage of voids in surface layer
 - ▶ rut.depth Change in rut depth per million vehicle passes
 - viscosity Viscosity of asphalt
 - run 2 data collection periods: 1 for run 1, 0 for run 2.
- rut.depth response. Depends on other variables, how?

Packages for this section

```
library(MASS, exclude = "select")
library(tidyverse)
library(broom)
library(leaps)
```

Make sure to load MASS before tidyverse (for annoying technical reasons), or to load MASS excluding its select (as above).

Getting set up

```
my_url <- "http://ritsokiguess.site/datafiles/asphalt.txt"
asphalt <- read_delim(my_url, " ")</pre>
```

- Quantitative variables with one response: multiple regression.
- Some issues here that don't come up in "simple" regression; handle as we go. (STAB27/STAC67 ideas.)

The data (some)

asphalt

```
A tibble: 31 \times 7
  pct.a.surf pct.a.base fines voids rut.depth viscosity
                                                           run
        <dbl>
                   <dbl> <dbl> <dbl>
                                         <dbl>
                                                   <dbl> <dbl>
        4.68
                    4.87
                           8.4
                               4.92
                                          6.75
                                                    2.8
         5.19
                    4.5
                           6.5 4.56
                                         13
                                                    1.4
3
        4.82
                    4.73 7.9 5.32
                                         14.8
                                                    1.4
        4.85
                    4.76 8.3 4.86
                                         12.6
                                                    3.3
5
        4.86
                    4.95 8.4 3.78
                                          8.25
                                                    1.7
6
         5.16
                    4.45 7.4 4.40
                                         10.7
                                                    2.9
        4.82
                    5.05
                           6.8
                               4.87
                                          7.28
                                                    3.7
8
        4.86
                    4.7
                           8.6
                               4.83
                                         12.7
                                                    1.7
9
        4.78
                    4.84
                           6.7 4.86
                                         12.6
                                                    0.92
10
                    4.76
         5.16
                           7.7
                                4.03
                                         20.6
                                                    0.68
   21 more rows
```

Plotting response "rut depth" against everything else

Same idea as for plotting separate predictions on one plot:

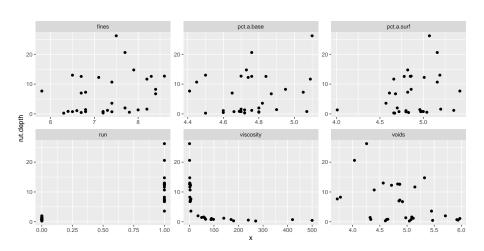
```
asphalt %>%
  pivot_longer(
    -rut.depth,
    names_to="xname", values_to="x"
) %>%
  ggplot(aes(x = x, y = rut.depth)) + geom_point() +
  facet_wrap(~xname, scales = "free") -> g
```

"collect all the x-variables together into one column called x, with another column xname saying which x they were, then plot these x's against rut.depth, a separate facet for each x-variable."

I saved this graph to plot later (on the next page).

The plot

g



Interpreting the plots

- One plot of rut depth against each of the six other variables.
- Get rough idea of what's going on.
- Trends mostly weak.
- viscosity has strong but non-linear trend.
- run has effect but variability bigger when run is 1.
- Weak but downward trend for voids.
- Non-linearity of rut.depth-viscosity relationship should concern us.

Log of viscosity: more nearly linear?

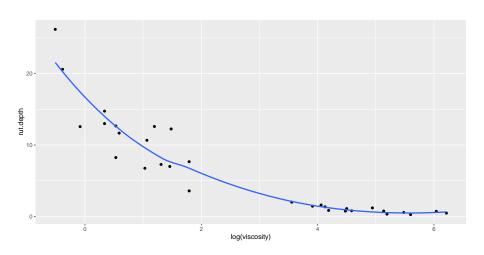
• Take this back to asphalt engineer: suggests log of viscosity:

```
ggplot(asphalt, aes(y = rut.depth, x = log(viscosity))) +
  geom_point() + geom_smooth(se = FALSE) -> g
```

(plot overleaf)

Rut depth against log-viscosity

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Comments and next steps

- Not very linear, but better than before.
- In multiple regression, hard to guess which x's affect response. So typically start by predicting from everything else.
- Model formula has response on left, squiggle, explanatories on right joined by plusses:

```
rut.1 <- lm(rut.depth ~ pct.a.surf + pct.a.base + fines +
  voids + log(viscosity) + run, data = asphalt)</pre>
```

Regression output:

summary(rut.1)

```
Call:
lm(formula = rut.depth ~ pct.a.surf + pct.a.base + fines + voids +
   log(viscosity) + run, data = asphalt)
Residuals:
            10 Median
   Min
                          30
                                 Max
-4.1211 -1.9075 -0.7175 1.6382 9.5947
Coefficients:
             Estimate Std. Error t value Pr(>|t|)
                         26.2188 -0.496 0.6247
(Intercept)
             -12.9937
pct.a.surf
               3.9706
                         2.4966 1.590 0.1248
pct.a.base
               1.2631 3.9703 0.318 0.7531
fines
               0.1164 1.0124 0.115 0.9094
voids
               0.5893 1.3244 0.445 0.6604
              -3.1515 0.9194 -3.428 0.0022 **
log(viscosity)
run
              -1.9655
                         3.6472 -0.539
                                         0.5949
---
Signif. codes:
              0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Residual standard error: 3.324 on 24 degrees of freedom

Comments

- R-squared 81%, not so bad.
- P-value in glance asserts that something helping to predict rut.depth.
- Table of coefficients says log(viscosity).
- But confused by clearly non-significant variables: remove those to get clearer picture of what is helpful.

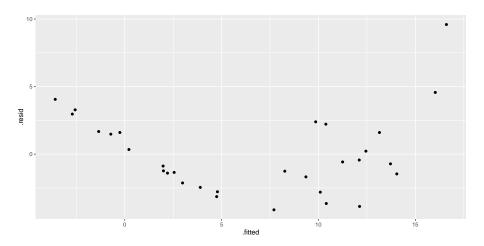
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Before we do anything, look at residual plots:

- • of residuals against each explanatory.
- Problem fixes:
 - with (a): fix response variable;
 - with some plots in (b): fix those explanatory variables.

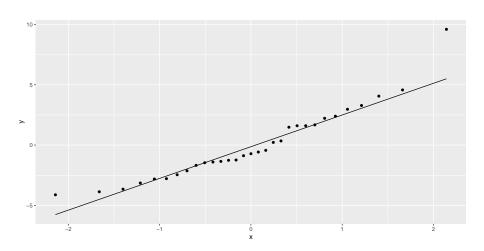
Plot fitted values against residuals

 $ggplot(rut.1, aes(x = .fitted, y = .resid)) + geom_point()$



Normal quantile plot of residuals

```
ggplot(rut.1, aes(sample = .resid)) + stat_qq() +
   stat_qq_line()
```



Plotting residuals against x variables

- Problem here is that residuals are in the fitted model, and the observed x-values are in the original data frame asphalt.
- Package broom contains a function augment that combines these two together so that they can later be plotted: start with a model first, and then augment with a data frame:

```
rut.1 %>% augment(asphalt) -> rut.1a
rut.1a
```

```
# A tibble: 31 x 13
  pct.a.surf pct.a.base fines voids rut.depth viscosity
                                                        run .fitted .resid
       dbl>
                  <dbl> <dbl> <dbl>
                                       <dbl>
                                                 <dbl> <dbl>
                                                              <dbl> <dbl>
                                        6.75
                                                 2.8
                                                               10.4 - 3.65
1
        4.68
                  4.87
                         8.4 4.92
2
        5.19
                  4.5 6.5 4.56
                                       13
                                                  1.4
                                                               13.7 - 0.718
3
        4.82
                  4.73
                         7.9 5.32
                                       14.8
                                                 1.4
                                                               13.1 1.60
        4.85
                  4.76 8.3 4.86
                                       12.6
                                                 3.3
                                                               10.4 2.22
 5
        4.86
                4.95 8.4 3.78
                                        8.25
                                                 1.7
                                                               12.1 - 3.87
6
        5.16
                  4.45 7.4 4.40
                                       10.7
                                                 2.9
                                                               11.2 - 0.577
        4.82
                   5.05
                         6.8 4.87
                                        7.28
                                                 3.7
                                                               10.1 - 2.81
        4.86
                  4.7
                         8.6 4.83
                                       12.7
                                                  1.7
                                                               12.4 0.221
        4.78
                  4.84 6.7 4.86
                                       12.6
                                                 0.92
                                                               14.0 -1.46
10
        5.16
                  4.76
                          7.7 4.03
                                       20.6
                                                  0.68
                                                               16.0 4.57
   21 more rows
```

What does rut.1a contain?

names(rut.1a)

```
[1] "pct.a.surf" "pct.a.base" "fines" "voids" "rut.depth
[6] "viscosity" "run" ".fitted" ".resid" ".hat"
[11] ".sigma" ".cooksd" ".std.resid"
```

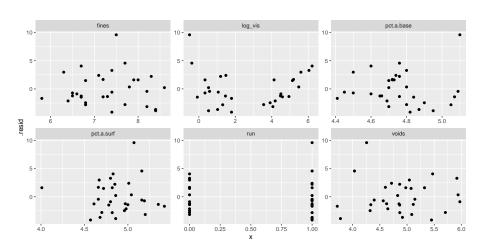
- all the stuff in original data frame, plus:
- quantities from regression (starting with a dot)

Plotting residuals against x-variables

```
rut.1a %>%
  mutate(log_vis=log(viscosity)) %>%
  pivot_longer(
    c(pct.a.surf:voids, run, log_vis),
    names_to="xname", values_to="x"
) %>%
  ggplot(aes(x = x, y = .resid)) +
  geom_point() + facet_wrap(~xname, scales = "free") -> g
```

The plot

g



Comments

- There is serious curve in plot of residuals vs. fitted values. Suggests a transformation of y.
- The residuals-vs-x's plots don't show any serious trends. Worst probably that potential curve against log-viscosity.
- Also, large positive residual, 10, that shows up on all plots. Perhaps transformation of y will help with this too.
- If residual-fitted plot OK, but some residual-x plots not, try transforming those x's, eg. by adding x^2 to help with curve.

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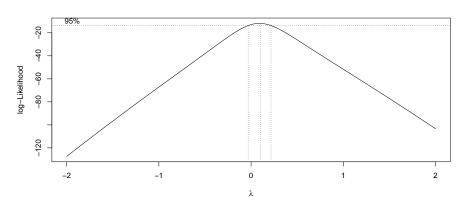
Which transformation?

- Best way: consult with person who brought you the data.
- Can't do that here!
- No idea what transformation would be good.
- Let data choose: "Box-Cox transformation".
- Scale is that of "ladder of powers": power transformation, but 0 is log.

Running Box-Cox

From package MASS:

```
boxcox(rut.depth ~ pct.a.surf + pct.a.base + fines + voids +
  log(viscosity) + run, data = asphalt)
```



Comments on Box-Cox plot

- λ represents power to transform y with.
- \bullet Best single choice of transformation parameter λ is peak of curve, close to 0.
- Vertical dotted lines give CI for λ , about (-0.05, 0.2).
- $\lambda = 0$ means "log".
- Narrowness of confidence interval mean that these not supported by data:
 - ▶ No transformation ($\lambda = 1$)
 - Square root $(\lambda = 0.5)$
 - ▶ Reciprocal $(\lambda = -1)$.

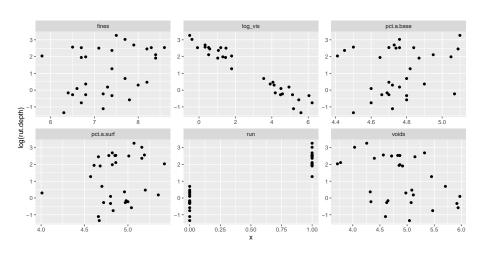
Relationships with explanatories

 As before: plot response (now log(rut.depth)) against other explanatory variables, all in one shot:

```
asphalt %>%
  mutate(log_vis=log(viscosity)) %>%
  pivot_longer(
    c(pct.a.surf:voids, run, log_vis),
    names_to="xname", values_to="x"
) %>%
  ggplot(aes(y = log(rut.depth), x = x)) + geom_point() +
  facet_wrap(~xname, scales = "free") -> g3
```

The new plots

g3



Modelling with transformed response

- These trends look pretty straight, especially with log.viscosity.
- Values of log.rut.depth for each run have same spread.
- Other trends weak, but are straight if they exist.
- Start modelling from the beginning again.
- Model log.rut.depth in terms of everything else, see what can be removed:

```
rut.2 <- lm(log(rut.depth) ~ pct.a.surf + pct.a.base +
fines + voids + log(viscosity) + run, data = asphalt)</pre>
```

• use tidy from broom to display just the coefficients.

Output

tidy(rut.2)

```
# A tibble: 7 \times 5
                estimate std.error statistic
                                                 p.value
 term
 <chr>>
                   <dbl>
                                                   <dbl>
                             <dbl>
                                       <dbl>
1 (Intercept)
                 -1.57
                            2.44
                                      -0.646 \ 0.525
2 pct.a.surf
                  0.584
                            0.232
                                       2.52
                                             0.0190
3 pct.a.base
                 -0.103
                            0.369
                                      -0.2800.782
                  0.0978
                            0.0941
4 fines
                                       1.04 0.309
5 voids
                  0.199
                            0.123
                                       1.62 0.119
                                      -6.53 0.000000945
6 log(viscosity)
                 -0.558
                            0.0854
7 run
                  0.340
                            0.339
                                       1.00 0.326
```

Taking out everything non-significant

• Try: remove everything but pct.a.surf and log.viscosity:

```
rut.3 <- lm(log(rut.depth) ~ pct.a.surf + log(viscosity), data = asphalt)
tidy(rut.3)</pre>
```

Check that removing all those variables wasn't too much

```
anova(rut.3, rut.2)
```

Analysis of Variance Table

- 2 24 2.2888 4 0.59216 1.5523 0.2191
 - H_0 : two models equally good; H_a : bigger model better.
 - Null not rejected here; small model as good as the big one, so prefer simpler smaller model rut.3.

Find the largest P-value by eye:

tidy(rut.2)

```
# A tibble: 7 \times 5
               estimate std.error statistic
                                            p.value
 term
 <chr>>
                 <dbl>
                          <dbl>
                                   <dbl>
                                              <dbl>
1 (Intercept)
               -1.57
                         2.44
                                  -0.646 0.525
                0.584
                         0.232
                                   2.52
2 pct.a.surf
                                         0.0190
                         0.369
                                  -0.2800.782
3 pct.a.base
              -0.103
                         0.0941
4 fines
                0.0978
                                   1.04 0.309
5 voids
                0.199
                         0.123 1.62 0.119
                                  -6.53 0.000000945
6 log(viscosity)
                -0.558
                         0.0854
7 run
                0.340
                         0.339
                                   1.00 0.326
```

- Largest P-value is 0.78 for pct.a.base, not significant.
- So remove this first, re-fit and re-assess.
- Or, as over.

Get the computer to find the largest P-value for you

• Output from tidy is itself a data frame, thus:

```
tidy(rut.2) %>% arrange(p.value)
```

```
# A tibble: 7 x 5
 term
              estimate std.error statistic p.value
 <chr>>
                 <dbl>
                         <dbl>
                                  <dbl>
                                            <dbl>
1 log(viscosity)
               -0.558
                        0.0854
                                 -6.53 0.000000945
2 pct.a.surf
                0.584
                        0.232 2.52 0.0190
3 voids
                0.199
                        0.123 1.62 0.119
4 fines
                0.0978
                        0.0941 1.04 0.309
5 run
                0.340
                        0.339 1.00 0.326
             -1.57
                        2.44
                                 -0.6460.525
 (Intercept)
                        0.369
                                 -0.2800.782
7 pct.a.base
              -0.103
```

Largest P-value at the bottom.

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Take out pct.a.base

A tibble: 6×2

Copy and paste the lm code and remove what you're removing:

```
term p.value <chr> <chr> <chr> 1 log(viscosity) 0.000000448
2 pct.a.surf 0.0143
0.109
4 (Intercept) 0.208
5 run 0.279
6 fines 0.316
```

• fines is next to go, P-value 0.32.

"Update"

Another way to do the same thing:

```
rut.4 <- update(rut.2, . ~ . - pct.a.base)
tidy(rut.4) %>% arrange(p.value)
```

```
# A tibble: 6 x 5
 term
             estimate std.error statistic p.value
 <chr>>
                <dbl>
                      <dbl>
                                <dbl> <dbl>
1 log(viscosity) -0.552 0.0818
                               -6.75 0.000000448
            0.593 0.225 2.63 0.0143
2 pct.a.surf
3 voids
            0.200 0.121 1.66 0.109
4 (Intercept) -2.08 1.61 -1.29 0.208
               0.360 0.325
                             1.11 0.279
5 run
               0.0889
                       0.0870 1.02 0.316
6 fines
```

Again, fines is the one to go. (Output identical as it should be.)

Take out fines:

```
rut.5 <- update(rut.4, . ~ . - fines)
tidy(rut.5) %>% arrange(p.value) %>% select(term, p.value)
```

Can't take out intercept, so run, with P-value 0.36, goes next.

Take out run:

```
rut.6 <- update(rut.5, . ~ . - run)
tidy(rut.6) %>% arrange(p.value) %>% select(term, p.value)
```

Again, can't take out intercept, so largest P-value is for voids, 0.044. But this is significant, so we shouldn't remove voids.

Comments

0.9001389

- Here we stop: pct.a.surf, voids and log.viscosity would all make fit significantly worse if removed. So they stay.
- Different final result from taking things out one at a time (top), than by taking out 4 at once (bottom):

• Point: Can make difference which way we go.

0.3911481 -0.6185628

Comments on variable selection

- Best way to decide which x's belong: expert knowledge: which of them should be important.
- Best automatic method: what we did, "backward selection".
- Do not learn about "stepwise regression"! eg. here
- R has function step that does backward selection, like this:

```
step(rut.2, direction = "backward", test = "F")
```

Gets same answer as we did (by removing least significant x).

- Removing non-significant x's may remove interesting ones whose P-values happened not to reach 0.05. Consider using less stringent cutoff like 0.20 or even bigger.
- Can also fit all possible regressions, as over (may need to do install.packages("leaps") first).

All possible regressions (output over)

Uses package leaps:

The output

```
d %>% rownames_to_column("model") %>% arrange(desc(rsq))
```

```
        model
        rsq pct.a.surf
        pct.a.base fines
        voids
        log.viscosity.
        run

        1
        6
        (1) 0.9609642
        *
        *
        *
        *
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        *
        *
        *</
```

Comments

- Problem: even adding a worthless x increases R-squared. So try for line where R-squared stops increasing "too much", eg. top line (just log.viscosity), first 3-variable line (backwards-elimination model). Hard to judge.
- One solution (STAC67): adjusted R-squared, where adding worthless variable makes it go down.
- data.frame rather than tibble because there are several columns in outmat.

All possible regressions, adjusted R-squared

```
with(s, data.frame(adjr2, outmat)) %>%
  rownames_to_column("model") %>%
  arrange(desc(adjr2))
```

```
        model
        adjr2 pct.a.surf pct.a.base fines voids log.viscosity. run

        1 3 (1) 0.9531812
        *
        *
        *

        2 5 (1) 0.9530038
        *
        *
        *
        *

        3 4 (1) 0.9529226
        *
        *
        *
        *

        4 4 (2) 0.9526007
        *
        *
        *
        *

        5 6 (1) 0.9512052
        *
        *
        *
        *

        6 5 (2) 0.9511918
        *
        *
        *
        *

        7 3 (2) 0.9482845
        *
        *
        *
        *

        8 2 (1) 0.9473550
        *
        *
        *

        9 2 (2) 0.9442365
        *
        *
        *

        10 1 (1) 0.9433685
        *
        *
        *

        11 1 (2) 0.8576662
        *
        *
        *
```

Revisiting the best model

Best model was our rut.6:

```
tidy(rut.6)
```

```
# A tibble: 4 x 5
 term
               estimate std.error statistic p.value
 <chr>>
                 <dbl>
                          <dbl>
                                   <dbl>
                                          <dbl>
1 (Intercept)
                -1.02 1.36
                                  -0.748 4.61e- 1
2 pct.a.surf
                 0.555 0.220 2.52 1.80e- 2
                 0.245 0.116 2.12 4.36e- 2
3 voids
4 log(viscosity)
              -0.646
                         0.0288 -22.5 5.29e-19
```

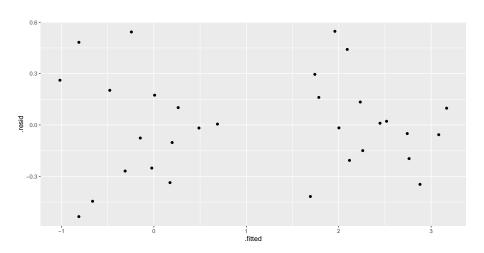
Revisiting (2)

- Regression slopes say that rut depth increases as log-viscosity decreases, pct.a.surf increases and voids increases. This more or less checks out with out scatterplots against log.viscosity.
- We should check residual plots again, though previous scatterplots say it's unlikely that there will be a problem:

```
g <- ggplot(rut.6, aes(y = .resid, x = .fitted)) +
geom_point()</pre>
```

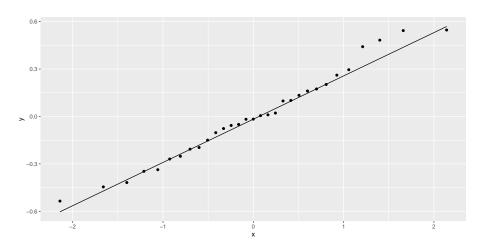
Residuals against fitted values

g



Normal quantile plot of residuals

```
ggplot(rut.6, aes(sample = .resid)) + stat_qq() + stat_qq_line
```



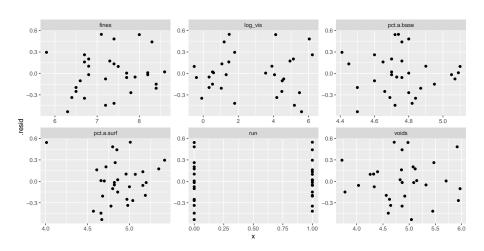
Plotting residuals against x's

• Do our trick again to put them all on one plot:

```
augment(rut.6, asphalt) %>%
  mutate(log_vis=log(viscosity)) %>%
  pivot_longer(
    c(pct.a.surf:voids, run, log_vis),
    names_to="xname", values_to="x",
) %>%
  ggplot(aes(y = .resid, x = x)) + geom_point() +
  facet_wrap(~xname, scales = "free") -> g2
```

Residuals against the x's

g2



Comments

- None of the plots show any sort of pattern. The points all look random on each plot.
- On the plot of fitted values (and on the one of log.viscosity), the points seem to form a "left half" and a "right half" with a gap in the middle. This is not a concern.
- One of the pct.a.surf values is low outlier (4), shows up top left of that plot.
- Only two possible values of run; the points in each group look randomly scattered around 0, with equal spreads.
- Residuals seem to go above zero further than below, suggesting a mild non-normality, but not enough to be a problem.

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Variable-selection strategies

- Expert knowledge.
- Backward elimination.
- All possible regressions.
- Taking a variety of models to experts and asking their opinion.
- Use a looser cutoff to eliminate variables in backward elimination (eg. only if P-value greater than 0.20).
- If goal is prediction, eliminating worthless variables less important.
- If goal is understanding, want to eliminate worthless variables where possible.
- Results of variable selection not always reproducible, so caution advised.