Homework Assignment 1

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BMB: table of contents is a little bit of overkill?

1

dataset::longley is a macroeconomic data set with 7 economical variables, observed yearly from 1947 to 1962.

head(longley)

##		GNP.deflator	GNP	Unemployed	Armed.Forces	Population	Year	Employed
##	1947	83.0	234.289	235.6	159.0	107.608	1947	60.323
##	1948	88.5	259.426	232.5	145.6	108.632	1948	61.122
##	1949	88.2	258.054	368.2	161.6	109.773	1949	60.171
##	1950	89.5	284.599	335.1	165.0	110.929	1950	61.187
##	1951	96.2	328.975	209.9	309.9	112.075	1951	63.221
##	1952	98.1	346.999	193.2	359.4	113.270	1952	63.639

(a)

We first run a linear model, using the employment numbers as the response on all the rest of the variables. Then we show using VIF that there is multi-collineairty.

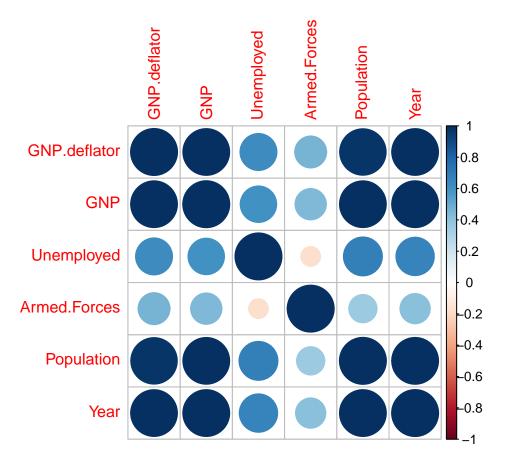
```
# Liner model for original data
m1 <- lm(Employed ~ . , data = longley)
summary(m1)
##
## Call:
## lm(formula = Employed ~ ., data = longley)
##
## Residuals:
##
       Min
                  1Q
                      Median
                                   3Q
                                           Max
## -0.41011 -0.15767 -0.02816 0.10155 0.45539
##
## Coefficients:
##
                 Estimate Std. Error t value Pr(>|t|)
## (Intercept) -3.482e+03 8.904e+02 -3.911 0.003560 **
## GNP.deflator 1.506e-02 8.492e-02
                                       0.177 0.863141
## GNP
               -3.582e-02 3.349e-02 -1.070 0.312681
## Unemployed
               -2.020e-02 4.884e-03 -4.136 0.002535 **
## Armed.Forces -1.033e-02 2.143e-03 -4.822 0.000944 ***
## Population
               -5.110e-02 2.261e-01 -0.226 0.826212
                1.829e+00 4.555e-01
                                       4.016 0.003037 **
## Year
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.3049 on 9 degrees of freedom
## Multiple R-squared: 0.9955, Adjusted R-squared: 0.9925
## F-statistic: 330.3 on 6 and 9 DF, p-value: 4.984e-10
```

It is suspicious that all of the coefficients are quite small. (**BMB**: why? doesn't that depend on the units of the predictors and response variable?) Furthermore the t values are quite small for some of the variables. If we look at the correlation plot for predictors we see

```
data <- longley %>%
    select(-Employed)
```

```
{\it \# Define \ function \ to \ convert \ covariance \ matrix \ to \ correlation \ matrix.}
cov_to_cor <- function(covmat) {</pre>
   # Extract std dev's and name
     SD <- diag(covmat) %>%
            sqrt()
     names(SD) <- colnames(covmat)</pre>
  # Extract dimensions of covariance matrix
  nrow <- dim(covmat)[1]</pre>
  ncol <- dim(covmat)[2]</pre>
  # Scale the covariance matrix
  for (i in 1: nrow ) {
   for (j in 1:ncol) {
      scale <- SD[i] * SD[j]</pre>
      covmat[i,j] <- covmat[i,j] / scale</pre>
  }
  }
  return(covmat)
```

corrplot::corrplot(cov_to_cor(cov(data)))



We observe that GNP.deflator, GNP and Population are very correlated in the correlation plot and also have small t-values for their corresponding coefficients in the linear model.

Now us compute the Variance Inflation Factor (VIF) for the predictors. It is a useful measure of multicollinearity in a data set. The VIF is the ratio of the variance of estimating some parameter in a model that includes multiple other predictors by the variance of a model constructed using only one term. It is defined as

$$VIF = \frac{1}{1 - R^2},$$

where R^2 is the coefficient of determination.

```
# Loop through each predictor to calculate its VIF
  for (i in seq_along(predictors)) {
    # Isolate the predictor of interest
    predictor_of_interest <- names(predictors)[i]</pre>
    # Create a formula to regress the predictor of interest on all other predictors
    formula <- as.formula(paste(predictor_of_interest, "~ ."))</pre>
    # Fit the linear model
    lm_model <- lm(formula, data = predictors)</pre>
    # Compute R-squared
    r_squared <- summary(lm_model)$r.squared</pre>
    # Compute VIF
    vif_values[i] <- 1 / (1 - r_squared)</pre>
  }
  # Add names
  names(vif_values) <- names(predictors)</pre>
  return(vif_values)
}
# Compute VIF values
vif_results <- compute_vif(longley, "Employed")</pre>
print(vif_results)
```

```
## GNP.deflator GNP Unemployed Armed.Forces Population Year
## 135.53244 1788.51348 33.61889 3.58893 399.15102 758.98060
```

The rule of thumb is that a VIF value above 5-10 indicates multicollinearity between that predictor and the other predictors. We observe that all variables other than Armed.Forces are **10.

Since the predictors are correlated and that makes our estimates of coefficients correlated, we can remove the multi-collinearity by using principal components as the predictors in our linear model. Therefore we will use all the predictors. After we compute the principal components we will interpret them in terms of the

Table 1: Proportions of Variance Explained by Components (Covariance Matrix)

Component	Proportion
Component 1	0.65
Component 2	0.30
Component 3	0.05
Component 4	0.00
Component 5	0.00
Component 6	0.00

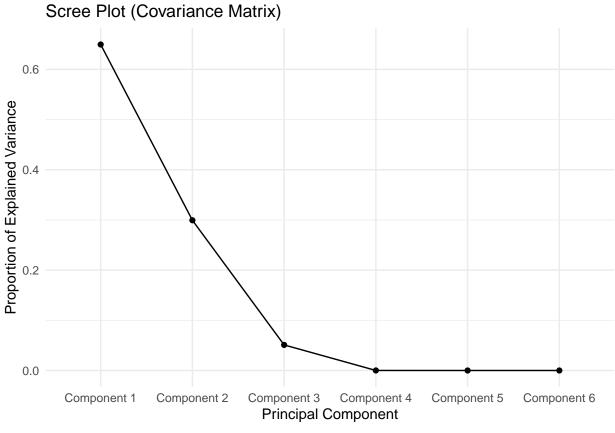
Table 2: Proportions of Variance Explained by Components (Correlation Matrix)

Component	Proportion
Component 1	0.51
Component 2	0.19
Component 3	0.14
Component 4	0.07
Component 5	0.06
Component 6	0.03

original predictors. We will do PCA using both the covariance and the correlation matrix, so they can be compared downstream when fitting the linear model.

```
# Define function to compute proportion of total population variation due to the
# k'th principal component.
prop_pc <- function(mat) {
    ei <- eigen(mat)
    evalues <- ei$values
    total <- sum(evalues)
    dim <- length(diag(mat))
    vec <- rep(NA, dim)

for (i in 1:dim) {
    vec[i] <- evalues[i]/total
    }
    return(vec)</pre>
```



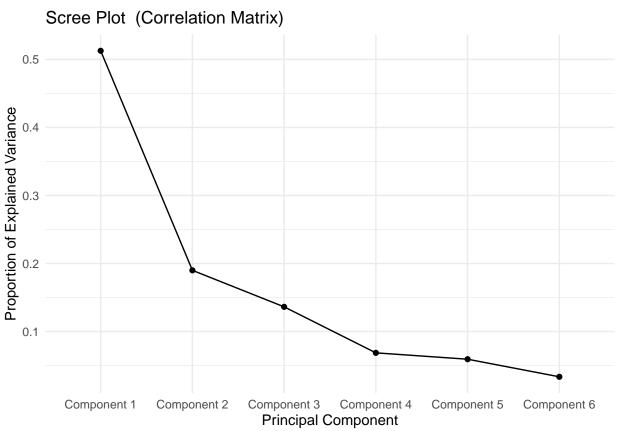


Table 3: Eigenvectors for the First Two Principal Components (Covariance)

	Component 1	Component 2
GNP.deflator	-0.082	0.034
GNP	-0.756	0.319
Unemployed	-0.626	-0.580
Armed.Forces	-0.158	0.748
Population	-0.054	0.013
Year	-0.037	0.012

Table 4: Eigenvectors for the First Four Principal Components (Correlation)

	Component 1	Component 2	Component 3
GNP.deflator	-0.471	0.003	0.073
GNP	-0.358	0.408	0.593
Unemployed	-0.434	0.404	0.064
Armed.Forces	-0.288	0.403	-0.794
Population	-0.440	-0.507	-0.014
Year	-0.430	-0.501	-0.090

Method	Component	Interpretation
Covariance-based	Component 1	Strong negative loadings on GNP and
Eigenvectors		Unemployed. This component might capture
		general economic factors related to both the
		gross national product and unemployment.
Covariance-based	Component 2	Strong positive loadings on
Eigenvectors		Armed.Forces, moderate positive loading on
		GNP and moderate negative loading on
		Unemployed. This component may capture
		factors related to the military and the
		economy.
Correlation-based	Component 1	Strong loadings on almost all variables except
Eigenvectors		for Armed. Forces. This could be a "general
		factor" capturing broad economic and
		demographic aspects.
Correlation-based	Component 2	Strong negative loadings on Population and
Eigenvectors		Year, and positive loadings on GNP,
		Unemployed, and Armed.Forces. This might
		capture more specific year-over-year changes
		or economic factors separate from
		demographic growth.

Method	Component	Interpretation
Correlation-based	Component 3	Strong negative loading on Armed.Forces and
Eigenvectors		strong positive loading on GNP. This could
		represent aspects of the economy not
		associated with military activity.

(b)

Table 6: Description of Variables in the Longley Dataset

Variable	Description	Unit	Interpretation	Threshold
GNP.deflator	Gross National Product implicit price deflator	Index (year $1954 = 100$)	Measure of level of prices	1 index point
GNP	Gross National Product	Billions of USD	Total value of goods and services	0.5 Billion USD
Unemployed	Number of unemployed	Thousands of people	People looking for a job	1K people
Armed.Forces	Number of people in the armed forces	Thousands of people	People in the military	1K people
Population	Non-institutionalized population	Thousands of people	People outside institutions	1K people
Year	Year of the observation	Years	Year recorded	1 Year
Employed	Number of people employed	Thousands of people	People who have jobs	1K people

(c)

```
# Center and Scale data
data_sc <- scale(data)</pre>
# Transform data with eigenvectors for covariation matrix
pc_scores_cov <- data_sc %*% eigenvectors_cov</pre>
# Transform data with eigenvectors for correlation matrix
pc_scores_cor <- data_sc %*% eigenvectors</pre>
# Fit new lm
response <- longley$Employed</pre>
mod_pc_cov <- lm(response ~ pc_scores_cov)</pre>
mod_pc_cor <- lm(response ~ pc_scores_cor)</pre>
summary(mod_pc_cov)
##
## Call:
## lm(formula = response ~ pc_scores_cov)
##
## Residuals:
##
        Min
                   1Q
                       Median
                                      3Q
                                               Max
## -2.44203 -0.67573 0.02242 0.86219 1.77752
```

```
##
## Coefficients:
##
                           Estimate Std. Error t value Pr(>|t|)
                                        0.3154 207.097 < 2e-16 ***
## (Intercept)
                            65.3170
## pc_scores_covComponent 1 -2.1029
                                        0.2272 -9.256 4.38e-07 ***
## pc_scores_covComponent 2
                                                 3.059 0.00915 **
                             0.9253
                                        0.3025
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 1.262 on 13 degrees of freedom
## Multiple R-squared: 0.8882, Adjusted R-squared: 0.871
## F-statistic: 51.62 on 2 and 13 DF, p-value: 6.542e-07
summary(mod_pc_cor)
##
## Call:
## lm(formula = response ~ pc_scores_cor)
##
## Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
## -0.99469 -0.19427 -0.04553 0.17479
                                      1.03934
##
## Coefficients:
##
                           Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                                       0.12297 531.180 < 2e-16 ***
                           65.31700
## pc_scores_corComponent 1 -1.43795
                                       0.06595 -21.805 5.07e-11 ***
## pc_scores_corComponent 2 -5.18685
                                       0.63432 -8.177 3.00e-06 ***
## pc_scores_corComponent 3 -1.63049
                                       0.25602 -6.369 3.56e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4919 on 12 degrees of freedom
## Multiple R-squared: 0.9843, Adjusted R-squared: 0.9804
## F-statistic: 250.9 on 3 and 12 DF, p-value: 4.349e-11
```

Both models seem to fit the data well. The R^2 value is high for both models but for the correlation model $R^2 = 99.85$ is very high. Both models show significant t-values for their coefficients, indicating that the

selected principal components are important predictors for the employment rate in both cases.

BMB: I would have said that with 15 observations and 6 potential predictors, you should have decided immediately that you needed to reduce the number of covariates ...

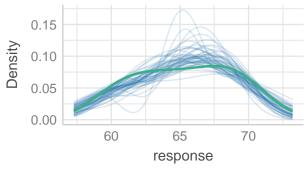
(d)

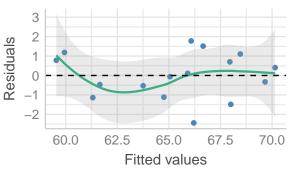
```
check_cov <- performance::check_model(mod_pc_cov)
par(mfrow= c(2,2))
plot(check_cov, title = "PCR (Covariance")</pre>
```

Posterior Predictive Check

Linearity

Model-predicted lines should resemble observed da Reference line should be flat and horizontal

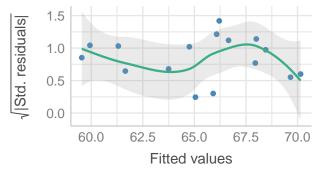




Observed data — Model-predicted data

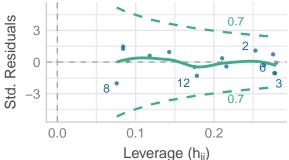
Homogeneity of Variance

Reference line should be flat and horizontal

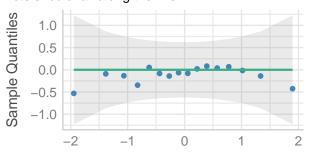


Influential Observations

Points should be inside the contour lines

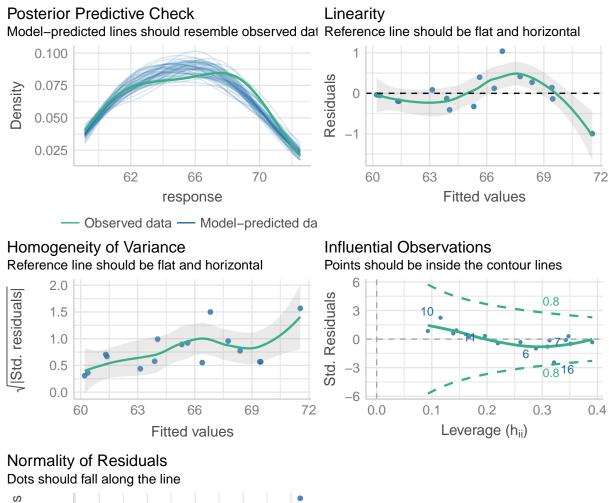


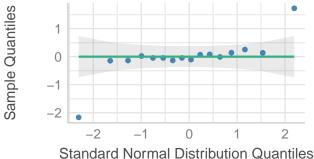
Normality of Residuals Dots should fall along the line



Standard Normal Distribution Quantiles

```
check_cor <- performance::check_model(mod_pc_cor)
par(mfrow= c(2,2))
plot(check_cor, title = "PCR (Correlation)")</pre>
```





The posterior check looks much better for the correlation PCR (PCRcor) then for the covariance PCR (PCRcov). The linearity for PCRcov has wider CI's but it is more rectangular then PCRcor which appears fairly non-linear after in the upper half. A similar argument holds for the homogeneity of variance. The PCRcor model appears to have a borderline influential observation for the 16th data point. The PCRcov model does not appear to any influential observation. The normality of residuals is very good for both model except that PCRcor has an extreme looking point on both side of the 2nd quantile.

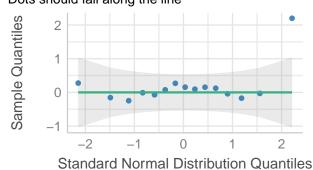
(e)

We will not drop any observations but for curiosity sake lets drop the possible influential observation for PCRcor and refit the model.

```
data2 <- data_sc[1:15,]</pre>
pc_scores_cor2 <- data2 %*% eigenvectors</pre>
response2<- response[1:15]</pre>
mod_pc_cor2 <- lm(response2 ~ pc_scores_cor2)</pre>
summary(mod_pc_cor)
##
## Call:
## lm(formula = response ~ pc_scores_cor)
##
## Residuals:
##
        Min
                  1Q
                       Median
                                     3Q
                                             Max
## -0.99469 -0.19427 -0.04553 0.17479 1.03934
## Coefficients:
##
                            Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                            65.31700
                                        0.12297 531.180 < 2e-16 ***
## pc_scores_corComponent 1 -1.43795
                                        0.06595 -21.805 5.07e-11 ***
## pc_scores_corComponent 2 -5.18685
                                        0.63432 -8.177 3.00e-06 ***
## pc_scores_corComponent 3 -1.63049
                                        0.25602 -6.369 3.56e-05 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4919 on 12 degrees of freedom
## Multiple R-squared: 0.9843, Adjusted R-squared: 0.9804
## F-statistic: 250.9 on 3 and 12 DF, p-value: 4.349e-11
check_cor2 <- performance::check_model(mod_pc_cor2)</pre>
par(mfrow= c(2,2))
plot(check_cor2, title = "PCR (Correlation)")
```

Linearity Posterior Predictive Check Model-predicted lines should resemble observed d Reference line should be flat and horizontal 0.5 0.08 Residuals Density 0.0 0.06 -0.50.04 60 64 68 60.0 62.5 65.0 67.5 70.0 response2 Fitted values Observed data — Model-predicted da Homogeneity of Variance Influential Observations Reference line should be flat and horizontal Points should be inside the contour lines 6 1.5 Std. residuals Std. Residuals 8.0 3 1.0 0 0.5 -3 0.0 -6 60.0 62.5 65.0 67.5 70.0 0.0 0.1 0.2 0.3 0.4 Leverage (hii) Fitted values

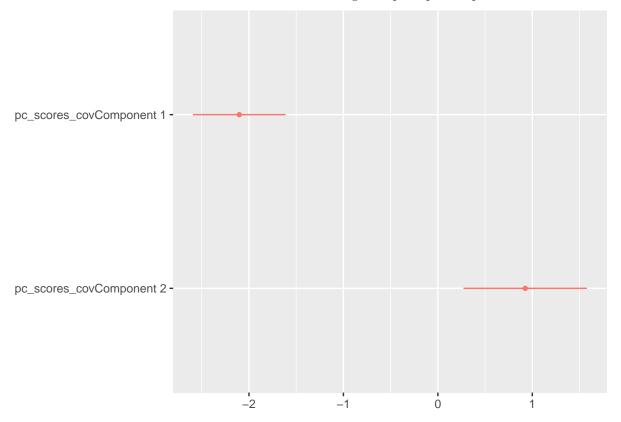
Normality of Residuals Dots should fall along the line

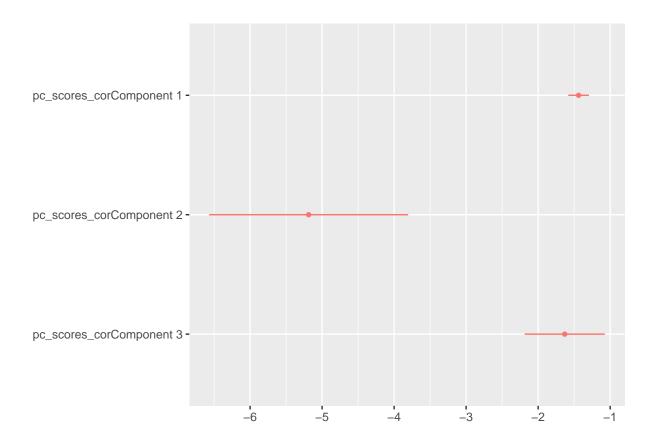


We see that the linearity and the homogeneity of variance both looks better now.

(f)

The data is scaled and centered as is custom when working with principal components.

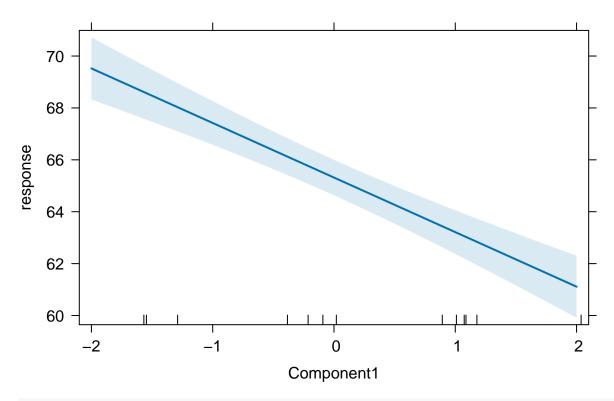




(g)

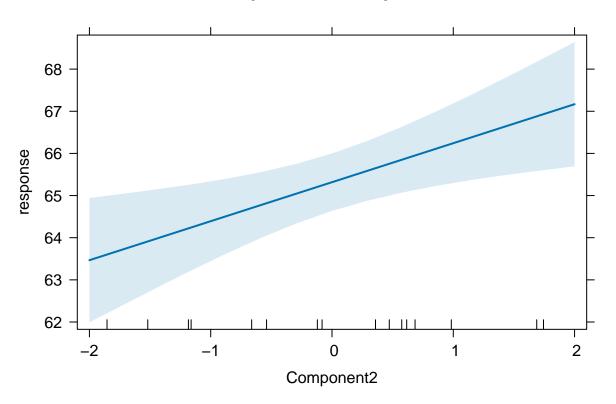
```
#effect() was very picky so had to make some changes below to make it happy.
# Rename columns to remove spaces and convert matrix to data frame
colnames(pc_scores_cov)<- c("Component1", "Component2")</pre>
pc_scores_cov_df <- as.data.frame(pc_scores_cov)</pre>
colnames(pc_scores_cor)<- c("Component1", "Component2", "Component3")</pre>
pc_scores_cor_df <- as.data.frame(pc_scores_cor)</pre>
# Fitting the model using the data frame
mod_pc_cov_updated <- lm(formula = response ~ Component1 + Component2,</pre>
                          data = pc_scores_cov_df)
mod_pc_cor_updated <- lm(formula = response ~ Component1 + Component2 +</pre>
                             Component3,
                           data = pc_scores_cor_df)
# Creating the effect objects for each component
eff1_cov <- effect("Component1", mod_pc_cov_updated)</pre>
eff2_cov <- effect("Component2", mod_pc_cov_updated)</pre>
eff1_cor <- effect("Component1", mod_pc_cor_updated)</pre>
eff2_cor <- effect("Component2", mod_pc_cor_updated)</pre>
eff3_cor <- effect("Component3", mod_pc_cor_updated)</pre>
# Plotting the effect for each component
par(mfrow = c(3,2))
plot(eff1_cov)
```

Component1 effect plot



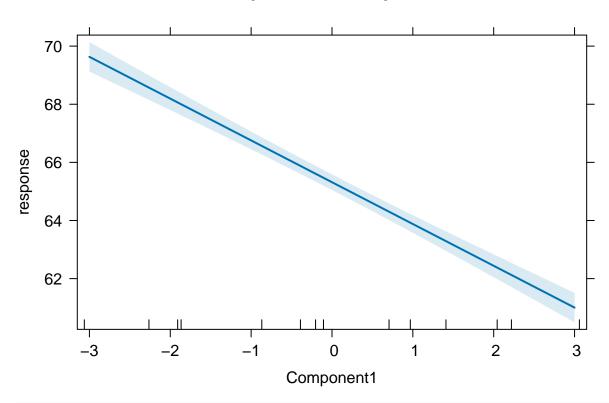
plot(eff2_cov)

Component2 effect plot



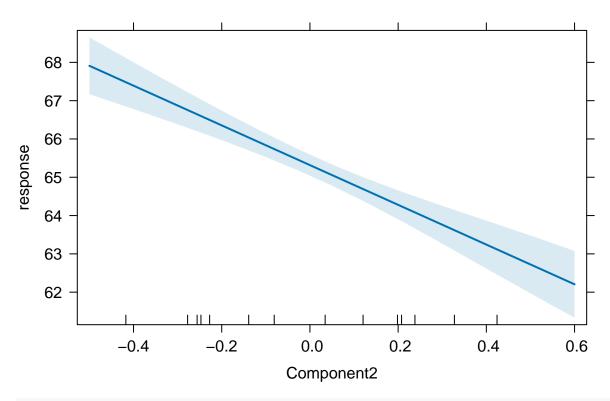
plot(eff1_cor)

Component1 effect plot



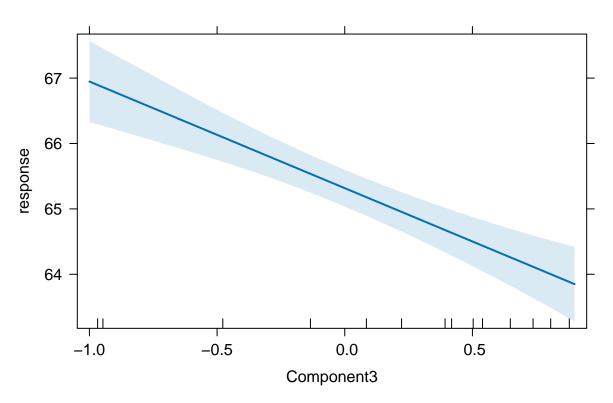
plot(eff2_cor)

Component2 effect plot



plot(eff3_cor)

Component3 effect plot



For PCRcov we see that the main effect for the 1st principal component has a large negative magnitude. This makes sense at the corresponding eigenvector had large negative loadings on Unemploymed and GNP. Therefore when Unemploymed and GNP are low, employment is high. in this component are associated with higher employment. The 2nd principal component has a large positive loading on Armed.Forces and a moderate negative loading on Unemployed. Therefore when unemployment is low and enrollment in armed forces is high, this component is associated with low employment.

A similar interpretaion can be done for the correlation plot (which I will not do).

Impact

```
# Create a data frame with the unique combinations of Period and Treatment
design <- expand.grid(</pre>
  Period = factor(c("Before", "After")),
  Treatment = factor(c("Control", "Impact"))
)
# Define the contrasts for Period and Treatment
contrasts(design$Period) <- matrix(c(-1, 1), 2)</pre>
contrasts(design$Treatment) <- matrix(c(-1, 1), 2)</pre>
# Construct the minimal model matrix
model_matrix_1 <- model.matrix(~ Period * Treatment, design)</pre>
print(model_matrix_1)
     (Intercept) Period1 Treatment1 Period1:Treatment1
##
## 1
                        1
                                   -1
                                                       -1
## 2
                                   -1
## 3
                1
                        1
                                    1
                                                        1
## 4
                1
                       -1
                                    1
                                                       -1
## attr(,"assign")
## [1] 0 1 2 3
## attr(,"contrasts")
## attr(,"contrasts")$Period
          [,1]
##
## After
            -1
## Before
             1
##
## attr(,"contrasts")$Treatment
##
            [,1]
## Control
             -1
```

First consider model_matrix_1 for the model ~ Period * Treatment. The first column of 1s represents the intercept. This is used to estimate the average value of Control and Impact during the Before period. The second column indicates the effect of moving from Control to Impact within the Before period. The

contrast matrix for Period is set up to estimate the difference $(\bar{\mu}A - \bar{\mu}B)$. The third column indicates the effect of moving from Before to After, averaged over the Control and Impact. The contrast matrix for Period is set up to estimate the difference, $(\delta(\mu_A) - \delta(\mu_B))$. The fourth column represents the interaction between Period and Treatment. It estimates the difference in the difference Control-Impact between the Before and After periods. Therefore model_matrix_1 allows for estimating the intercept, main effects and interaction.

```
# Create the model matrix for ~ 0 + Period:Treatment
model_matrix_2 <- model.matrix(~ 0 + Period:Treatment, design)
print(model_matrix_2)</pre>
```

```
PeriodAfter:TreatmentControl PeriodBefore:TreatmentControl
## 1
## 2
                                                                  0
                                 1
## 3
                                  0
                                                                  0
## 4
                                  0
                                                                  0
     PeriodAfter:TreatmentImpact PeriodBefore:TreatmentImpact
##
                                 0
## 1
                                                               0
## 2
                                 0
                                                               0
## 3
                                                               1
## 4
                                                               0
## attr(,"assign")
## [1] 1 1 1 1
## attr(,"contrasts")
## attr(,"contrasts")$Period
##
          [,1]
## After
            -1
## Before
             1
##
## attr(,"contrasts")$Treatment
           [,1]
##
## Control
             -1
## Impact
```

model_matrix_2 for the model ~ 0 + Period * Treatment does not have an intercept. Each column includes a single 1 for each of the 4 possible interactions. This means you can only estimate the interaction effect between Period and Treatment.

```
set.seed(123)
# Function to simulate data with different levels of violation
simulate_data <- function(n=100 , violation_level=1) {</pre>
  x \leftarrow rnorm(n)
  epsilon <- rnorm(n, mean = 0, sd = abs(violation_level * x))</pre>
 y \leftarrow 2 * x + 3 + epsilon
 data.frame(x, y)
}
# Simulate
simulate <- function(n_sims=1000, violation_level=1,alpha = 0.05) {</pre>
  true_slope <- 2</pre>
  slopes <- numeric(n_sims)</pre>
  coverages <- numeric(n_sims)</pre>
  reject_null <- numeric(n_sims)</pre>
  for (i in 1:n_sims) {
    data <- simulate_data(violation_level = violation_level)</pre>
    model \leftarrow lm(y \sim x, data = data)
    slopes[i] <- coef(model)[2]</pre>
    conf_int <- confint(model)[2, ]</pre>
    coverages[i] <- conf_int[1] <= true_slope && conf_int[2] >= true_slope
    test <- bptest(model)</pre>
    reject_null[i] <- test$p.value < alpha</pre>
  }
  bias <- mean(slopes - true_slope)</pre>
  rmse <- sqrt(mean((slopes - true_slope)^2))</pre>
  coverage <- mean(coverages)</pre>
  power <- mean(reject_null)</pre>
  list(bias = bias, rmse = rmse, coverage = coverage, power = power)
}
```

##		violation_level	bias	rmse	coverage	power
##	1	1	0.004054285	0.1721927	0.723	0.386
##	2	2	-0.016098659	0.3580873	0.722	0.359
##	3	3	0.009015272	0.5055129	0.757	0.336
##	4	4	0.033175944	0.6775691	0.747	0.364
##	5	5	0.035515563	0.8643396	0.744	0.351
##	6	6	-0.033579776	1.0542837	0.729	0.384
##	7	7	-0.014555320	1.2290865	0.722	0.368
##	8	8	-0.050130958	1.3233036	0.763	0.364
##	9	9	-0.023257116	1.5036724	0.752	0.379
##	10	10	0.016101423	1.7504208	0.729	0.366

Effect of Violating Heteroscedasticity on BIAS, Coverage, RMSE and Power by the function f(x) = violation level * sd

