Centene Daily Closing Price

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Introduction

Our goal is to identify the most effective multiple variable model for predicting the daily closing price of Centene (CNC). To achieve this objective, we will leverage the predictive power of four key stocks - Anthem (ANTM), Cigna (CI), United Health Group (UNH), and Humana (HUM) - and examine their relationship with CNC.

Once we have established the most effective predictive model, we will be able to use it to inform our investment decisions and optimize our portfolio. By leveraging data-driven insights, we can make informed investment choices that lead to greater returns and success in the marketplace.

Overall, our analysis will provide valuable insights into the relationship between these stocks and CNC, and help us make more informed decisions when it comes to investing in the healthcare sector.

The Data

```
importdata <- read_excel("ClosingPrices.xlsx")
mydata <- importdata[,c(19,9,22,34,57)]
sum(is.na(mydata))</pre>
```

[1] 0

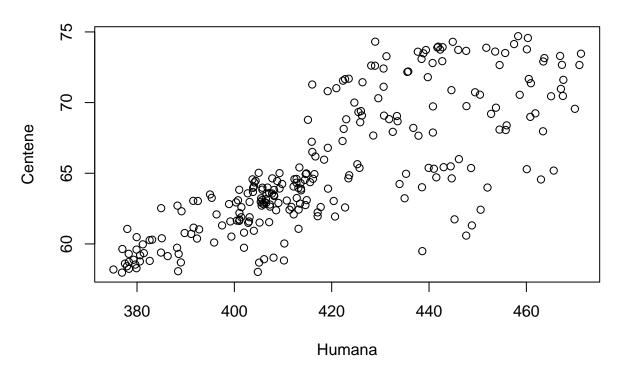
I loaded in the data into R from the Excel file. The file contains 58 different healthcare stocks daily closing price, but I chose only the 5 stocks that are in the same GICS sub-industry category as Centene, Managed Healthcare. I checked to see any missing values or error in the data set, but there is none. This data set was collected by the NASDAQ historical quotes from November 2, 2020 to October 28, 2021.

Analyzing the Variables

I made a scatter plot with each predictor variable to visually analyze the graph and identify any relationship with Centene's daily closing price. This will give us an idea of which predictor variable will be most significant towards our final model.

Scatter Plots of CNC VS HUM

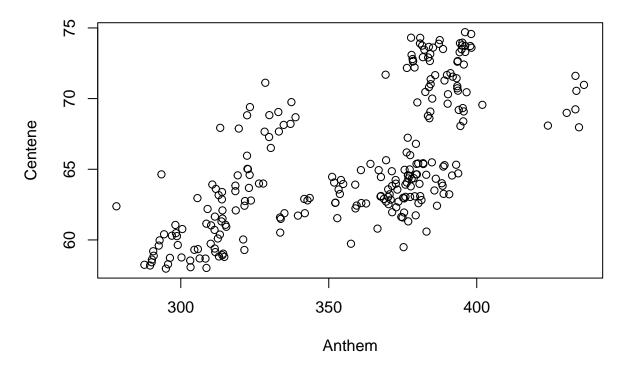
Scatterplot of CNC vs HUM



We see a moderate linear relationship between Centene and Humana with moderate dispersion in the upper right region.

Scatter Plots of CNC VS ANTM

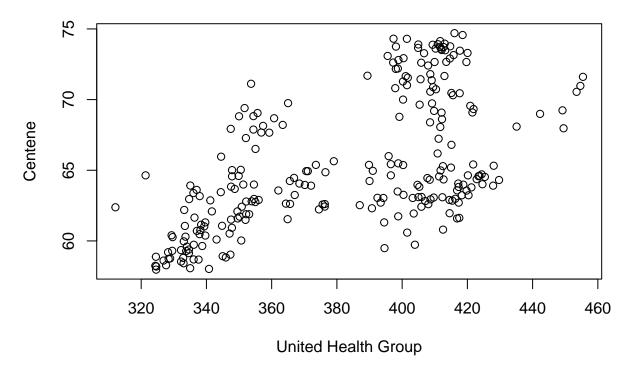
Scatterplot of CNC vs ANTM



We see a moderate linear relationship between Centene and Anthem with moderate dispersion in the upper right region.

Scatter Plots of CNC VS UNH

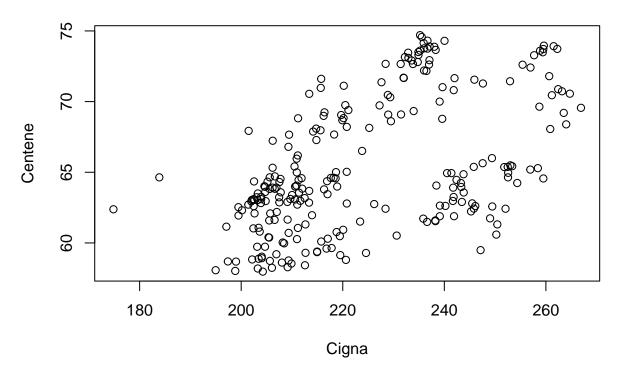
Scatterplot of CNC vs UNH



We see a moderate linear relationship between Centene and United Health Group with moderate dispersion in the upper region.

Scatter Plots of CNC VS CI

Scatterplot of CNC vs CI



We see a moderate linear relationship between Centene and Cigna with moderate dispersion in the upper region.

Correlation Matrix

```
mydata.rcorr = rcorr(as.matrix(mydata))
mydata.rcorr
```

```
##
         CNC ANTM
                     CI
                        HUM
        1.00 0.66 0.52 0.78 0.59
   ANTM 0.66 1.00 0.47 0.72 0.96
        0.52 0.47 1.00 0.57 0.31
##
   CI
        0.78 0.72 0.57 1.00 0.65
       0.59 0.96 0.31 0.65 1.00
## UNH
##
## n= 250
##
##
## P
        CNC ANTM CI HUM UNH
##
## CNC
                          0
##
   ANTM
         0
                   0
                      0
                          0
## CI
         0
             0
                      0
                          0
                   0
                          0
## HUM
```

```
## UNH O O O O
```

We want to determine numerically which predictor variables are liearly correlated with Centene. Scatter plots alone is not enough to determine which variables to use. Looking at the correlation matrix, each variable had a linear coefficient value greater than 0.50 and a p-value less than 0.05 making each predictor variable significant. We can see that Humana has the highest linear coefficient value.

Variable Selection Process

Based off the scatter plots and correlation matrix, each predictor variable seems significant enough to be included towards the model. However, to make sure we have selected the right variables to create our model, we will run a Stepwise Akaike Information Criterion (AIC) regression. It involves a step-by-step process of adding or removing variables from a model based on their statistical significance and their contribution to the overall goodness of fit of the model.

The AIC is a measure of the relative quality of a statistical model, and it is based on the likelihood function of the model and the number of parameters included in the model. The lower the AIC value, the better the model fits the data.

Stepwise AIC Variable Selection

##

- mydata\$HUM added

```
model <- lm(mydata$CNC ~ mydata$ANTM + mydata$CI + mydata$HUM + mydata$UNH, data = mydata)
ols_step_both_aic(model, details = TRUE)
## Stepwise Selection Method
##
##
## Candidate Terms:
##
## 1 . mydata$ANTM
## 2 . mydata$CI
## 3 . mydata$HUM
## 4 . mydata$UNH
##
##
   Step 0: AIC = 1482.59
##
   mydata$CNC ~ 1
##
##
## Variables Entered/Removed:
##
##
                             Enter New Variables
##
                DF
                        AIC
                                              RSS
## Variable
                                  Sum Sq
                                                        R-Sq
                                                                Adj. R-Sq
  ______
## mydata$HUM
                 1
                      1248.159
                                 3315.184
                                            2105.311
                                                        0.612
                                                                    0.610
                                                                    0.438
## mydata$ANTM
                 1
                      1339.435
                                 2387.440
                                            3033.054
                                                        0.440
## mydata$UNH
                 1
                      1375.523
                                 1916.437
                                            3504.058
                                                        0.354
                                                                    0.351
## mvdata$CI
                 1
                      1404.172
                                 1490.972
                                            3929.522
                                                        0.275
                                                                    0.272
```

```
##
##
## Step 1 : AIC = 1248.159
## mydata$CNC ~ mydata$HUM
##
                  Enter New Variables
       DF AIC Sum Sq RSS R-Sq Adj. R-Sq
## -----
## mydata$ANTM 1 1235.461
                     3435.386 1985.108 0.634
                                           0.631
## mydata$UNH
          1 1242.199 3381.156 2039.338 0.624
                                           0.621
## mydata$CI 1 1243.631 3369.443 2051.051 0.622 0.619
##
## - mydata$ANTM added
##
##
## Step 2 : AIC = 1235.461
##
 mydata$CNC ~ mydata$HUM + mydata$ANTM
##
##
               Remove Existing Variables
## ------
## Variable DF AIC Sum Sq RSS R-Sq Adj. R-Sq
## -----
## mydata$ANTM 1 1248.159 3315.184 2105.311 0.612
                                           0.610
## mydata$HUM 1 1339.435 2387.440 3033.054 0.440
##
                 Enter New Variables
## -----
## Variable DF AIC Sum Sq RSS R-Sq
                                        Adj. R-Sq
## ------
## mydata$CI 1 1232.748 3472.460 1948.034 0.641
                                           0.636
## mydata$UNH 1 1233.387 3467.478 1953.016 0.640
                                           0.635
##
## - mydata$CI added
##
##
## Step 3 : AIC = 1232.748
## mydata$CNC ~ mydata$HUM + mydata$ANTM + mydata$CI
##
               Remove Existing Variables
## -----
## Variable DF AIC Sum Sq RSS R-Sq Adj. R-Sq
## mydata$CI 1 1235.461 3435.386 1985.108 0.634 0.631
## mydata$ANTM 1 1243.631 3369.443 2051.051 0.622
## mydata$HUM 1 1314.644 2695.663 2724.831 0.497
                                           0.619
                                           0.493
##
              Enter New Variables
## -----
## Variable DF AIC Sum Sq RSS R-Sq Adj. R-Sq
```

##									
##	mydata\$UNH	1 1233	3.872 3	479.275	1941.	219 0.64	12	0.636	
##									
##									
##	No more varia	ables to be	added or	removed.					
##									
	Final Model C								
##									
##			Model Sum						
##			0.800			2.814			
	R-Squared		0.641						
	Adj. R-Square Pred R-Square					7.919 2.133			
##	RMSE: Root M	Mean Square	Error						
	MSE: Mean So								
	MAE: Mean Ab	solute Erro	or						
##			Δ	NOVA					
	ANOVA								
##		Sum of							
##		Squares	DF	Mean	Square	F	Sig.		
##	Regression Residual	1948.034	246		7.919				
##	Total	5420.494	249						
##								_	
##	Parameter Estimates								
##	model	Beta 	Std. Erro	r Std 	. Beta	t 	Sig	lower	upper
##	(Intercept) mydata\$HUM	0.111	0.01	1	0.583	1.376 9.904	0.000	0.089	0.133
	mydata\$ANTM	0.026	0.00	7	0.199	3.607	0.000	0.012	0.040
##	mydata\$CI	0.024	0.01	1	0.101	2.164	0.031	0.002	0.047
##									
##									
##									
##	Stepwise Summary								
##		·							
	Variable	Method 	AIC		RSS 	Sum Sq 	R-Sq	Ac 	lj. R-Sq
	mydata\$HUM	addition	1248.1	59 21	05.311	3315.184	0.6116	0	0.61004
	mydata\$ANTM	addition	1235.4		85.108	3435.386	0.6337		0.63081
	mydata\$CI	addition	1232.7		48.034	3472.460	0.6406		0.63623
##									

From the four variables, the selection process did not include United Health Group into the final model as it determines it was not significant towards the model. We will use Anthem (ANTM), Cigna (CI), and

Humana (HUM) in our model. Currently, the model has an Adjusted R-Squared of 0.636 and RMSE of 2.814. Hopefully adding more complex terms will improve the overall model.

Interaction Terms

Since all the stocks are in the same sub-industry of healthcare, it is logical to think that each stock may be related to one another. We will add the interaction terms into the model and analyze each interaction variables p-values in hopes of improvement.

```
##
## Call:
## lm(formula = mydata$CNC ~ mydata$ANTM + mydata$CI + mydata$HUM +
##
       mydata$ANTM * mydata$CI + mydata$ANTM * mydata$HUM + mydata$CI *
##
       mydata$HUM, data = mydata)
##
## Residuals:
##
      Min
                10 Median
                               30
                                      Max
  -8.7221 -1.4866 0.2449
                           1.6766
##
## Coefficients:
##
                           Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                          -1.221e+02 4.428e+01 -2.759 0.00625 **
                          -3.171e-01 1.209e-01
## mydata$ANTM
                                                -2.623 0.00928 **
## mydata$CI
                          3.942e-01 1.963e-01
                                                 2.008 0.04572 *
## mydata$HUM
                                                 5.909 1.16e-08 ***
                          8.114e-01 1.373e-01
## mydata$ANTM:mydata$CI
                          2.179e-03 4.493e-04
                                                 4.850 2.20e-06 ***
## mydata$ANTM:mydata$HUM -2.892e-04 1.991e-04
                                                -1.452 0.14766
## mydata$CI:mydata$HUM
                          -2.754e-03 6.019e-04
                                               -4.575 7.60e-06 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 2.669 on 243 degrees of freedom
## Multiple R-squared: 0.6806, Adjusted R-squared: 0.6727
## F-statistic: 86.28 on 6 and 243 DF, p-value: < 2.2e-16
```

From the summary, we can see that the interaction term ANTM*HUM is not significant and will be removed from the model. We also see that adding the interaction terms have increased the Adjusted R-Squared value.

Quadratic Terms

We will next add quadratic terms to the model and determine if they are significant to the model.

```
sqANTM <- mydata$ANTM^2
sqCI <- mydata$CI^2
sqHUM <- mydata$HUM^2
sqUNH <- mydata$UNH^2</pre>
```

```
QuadraticModel <- lm(mydata$CNC ~ mydata$ANTM + mydata$CI + mydata$HUM +
                         mydata$ANTM*mydata$CI + mydata$CI*mydata$HUM + sqANTM + sqCI + sqHUM,
                          data = mydata)
summary(QuadraticModel)
##
## Call:
  lm(formula = mydata$CNC ~ mydata$ANTM + mydata$CI + mydata$HUM +
       mydata$ANTM * mydata$CI + mydata$CI * mydata$HUM + sqANTM +
       sqCI + sqHUM, data = mydata)
##
##
## Residuals:
##
      Min
               1Q Median
                                30
                                       Max
## -8.6761 -1.3554 0.0267 1.5825 5.9083
##
## Coefficients:
##
                          Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                         -1.293e+02 5.072e+01
                                               -2.550
                                                         0.0114 *
## mydata$ANTM
                         -8.417e-01 1.316e-01 -6.394 8.32e-10 ***
## mydata$CI
                         1.558e+00 2.263e-01
                                                6.883 5.04e-11 ***
## mydata$HUM
                         6.716e-01 2.690e-01
                                                2.496
                                                         0.0132 *
## sqANTM
                         -4.097e-04 2.047e-04
                                               -2.002
                                                         0.0464 *
## sqCI
                        -5.967e-03 7.227e-04
                                               -8.255 1.00e-14 ***
## sqHUM
                         -1.955e-04 4.410e-04
                                               -0.443
                                                         0.6580
## mydata$ANTM:mydata$CI 5.481e-03
                                    6.107e-04
                                                8.975
                                                       < 2e-16 ***
## mydata$CI:mydata$HUM -1.982e-03 7.957e-04
                                               -2.490
                                                         0.0134 *
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
##
```

From the summary, we can see that the quadratic term sqHUM² is not significant and will be removed from the model. We also see that adding the quadratic terms have increased the R-Squared value.

Residual standard error: 2.346 on 241 degrees of freedom
Multiple R-squared: 0.7553, Adjusted R-squared: 0.7472
F-statistic: 93.01 on 8 and 241 DF, p-value: < 2.2e-16</pre>

Multicollinearity

2.079838

##

1.490767

Since we are using multiple predictor variables in this model, it is important to confirm none of the variables are correlated to one another.

Based on the Variance Inflation factor, we see each predictor variable is less than 10. There is no multi-collinearity.

2.369435

Final Model

```
##
## Call:
  lm(formula = mydata$CNC ~ mydata$ANTM + mydata$CI + mydata$HUM +
      mydata$ANTM * mydata$CI + mydata$CI * mydata$HUM + sqANTM +
##
##
      sqCI, data = mydata)
##
## Residuals:
##
      Min
               1Q Median
                               3Q
                                      Max
## -8.6222 -1.3570 0.0363 1.5715 5.9970
##
## Coefficients:
                          Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                        -1.141e+02 3.721e+01 -3.066 0.002414 **
## mydata$ANTM
                        -8.162e-01 1.182e-01 -6.907 4.35e-11 ***
## mydata$CI
                         1.586e+00 2.170e-01
                                                7.308 3.94e-12 ***
## mydata$HUM
                                                4.835 2.37e-06 ***
                         5.642e-01 1.167e-01
## sqANTM
                        -4.764e-04 1.387e-04 -3.435 0.000698 ***
## sqCI
                        -5.867e-03 6.857e-04 -8.557 1.35e-15 ***
## mydata$ANTM:mydata$CI 5.584e-03 5.639e-04
                                                9.903 < 2e-16 ***
## mydata$CI:mydata$HUM -2.245e-03 5.281e-04 -4.251 3.04e-05 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 2.342 on 242 degrees of freedom
## Multiple R-squared: 0.7551, Adjusted R-squared: 0.7481
## F-statistic: 106.6 on 7 and 242 DF, p-value: < 2.2e-16
```

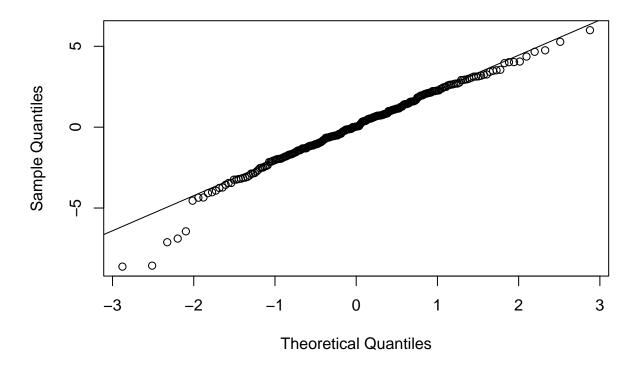
Our final model is:

```
Y = -114.1 - 0.8162(X_1, ANTM) + 1.586(X_2, CI) + 0.5642(X_3, HUM) - 0.0004764(X_4, ANTM^2) - 0.005867(X_5, CI^2) + 0.005584(X_6, ANTM * CI) - 0.002245(X_7, CI * HUM)
```

The final model has an R-Square Adjusted = 0.7481 which means that 74.81% of the variation in Centene's daily closing price are explained by the predictor variables. The R-Squared Adjusted is greater than the acceptable value for R-Squared Adjusted of 50%. The F-Statistic = 106.6 and P-Value = <0.0001. I also noticed when we removed the non-significant interaction and quadratic term, our overall model and the independent variables t-test statistics have improved, so it is safe to say we have found our best prediction model.

```
resids <- FinalModel$residuals
qqnorm(resids, main="Normal Q-Q Plot of Residuals from FinalModel")
qqline(resids)</pre>
```

Normal Q-Q Plot of Residuals from FinalModel



As we can see, the Normal Q-Q plot shows the residuals are normally distibuted with majority of data points on the line with a few outliers on both ends of the line.

Conclusion

I can confidently conclude that we have developed the best model for predicting Cenetene's daily closing price using predictor variables: ANTM, CI, HUM, ANTM^2, CI^2, ANTM CI, CIHUM.

Dataset Citation

"Market Activity Market Activity -> Stocks Options Etfs Mutual Funds Indexes Commodities Cryptocurrency Currencies Futures Fixed Income Global Markets Quick Links Real-Time Quotes after-Hours Quotes Pre-Market Quotes NASDAQ-100 Symbol Screener Online Brokers Glossary Sustainable Bond Network Symbol Change History IPO Performance Ownership Search Dividend History Investing Lists Rulebooks & Regulations Fundinsight Market Events Economic Calendar Earnings IPO Calendar Dividend Calendar Spo Calendar Holiday Schedule Analyst Activity Analyst Recommendations Daily Earnings Surprise Forecast Changes Commodities -> Gold Copper Crude Oil Natural Gas Nasdaq Data Statistical Milestones Total Returns Daily Market Statistics Most Active See All Market Activity ->." Nasdaq, www.nasdaq.com/market-activity/stocks/cnc/historical.