

Load Relevant R Libraries

```
library(rmarkdown); library(knitr); library(moments);  
library(scatterplot3d); library(corrplot); library(pso)  
library(psych); library(GPArotation); library(lavaan)
```

```
##SEM
```

```
data_matrix_csv <- read.csv("C:/Users/User/OneDrive - University of St. Thomas/Classes/STAT360/STAT360 I  
life_expectancy <- as.matrix(data_matrix_csv)
```

```
col_names_unedited <- c(  
  "Life expectancy at birth, total (years)",  
  "CO2 emissions (metric tons per capita)",  
  "Access to electricity (% of population)",  
  "Current health expenditure (% of GDP)",  
  "Out-of-pocket expenditure (% of current health expenditure)",  
  "Domestic private health expenditure per capita, PPP (current international $)",  
  "Domestic general government health expenditure per capita, PPP (current international $)",  
  "Renewable internal freshwater resources per capita (cubic meters)",  
  "Prevalence of HIV, total (% of population ages 15-49)",  
  "Unemployment, total (% of total labor force) (national estimate)",  
  "Government Effectiveness: Estimate",  
  "Income share held by highest 10%",  
  "Prevalence of current tobacco use (% of adults)",  
  "Total alcohol consumption per capita (liters of pure alcohol, projected estimates, 15+ years of age)",  
  "Political Stability and Absence of Violence/Terrorism: Estimate",  
  "Population density (people per sq. km of land area)"  
)
```

```
# Better for displaying
```

```
col_names <- c(  
  "Life_expectancy",  
  "CO2_emissions",  
  "Electricity",  
  "Health_expenditure",  
  "Out_of_pocket",  
  "Private_health_expenditure",  
  "Govt_health_expenditure",  
  "Freshwater_resources",  
  "HIV_prevalence",  
  "Unemployment",  
  "Govt_effectiveness",  
  "Income_share",  
  "Tobacco_use",  
  "Alcohol_consumption",  
  "Political_stability",  
  "Population_density"  
)
```

```
colnames(life_expectancy) <- col_names
```

```
SIG <- cov(life_expectancy, use = "pairwise.complete.obs")
```

```
EQN <- '
```

```
  # Measurement Model (Factor Definition)
```

```
  Healthcare_Spending =~ Private_health_expenditure + Health_expenditure
```

```
  SocioEconomicHealth =~ HIV_prevalence + Unemployment
```

```
  #structural model
```

```
  Life_expectancy ~ SocioEconomicHealth + Healthcare_Spending + CO2_emissions + Electricity + Govt_effect
```

```
  Political_stability + Political_stability + Freshwater_resources + Out_of_pocket + Govt_effect
```

```
'
```

```
MOD <- sem(EQN, data = scale(life_expectancy), sample.nobs = 217)
```

```
## Warning in lav_object_post_check(object): lavaan WARNING: some estimated ov  
## variances are negative
```

```
MOD
```

```
## lavaan 0.6.16 ended normally after 109 iterations
```

```
##
```

```
## Estimator ML
```

```
## Optimization method NLMINB
```

```
## Number of model parameters 21
```

```
##
```

```
## Used Total
```

```
## Number of observations 81 217
```

```
##
```

```
## Model Test User Model:
```

```
##
```

```
## Test statistic 172.425
```

```
## Degrees of freedom 39
```

```
## P-value (Chi-square) 0.000
```

```
parameterEstimates(MOD)[1:4,]
```

PRACTICAL SIGNIFICANCE

```
##           lhs op           rhs est   se    z pvalue  
## 1 Healthcare_Spending =~ Private_health_expenditure 1.00 0.000   NA    NA  
## 2 Healthcare_Spending =~ Health_expenditure 0.14 0.321 0.436 0.663  
## 3 SocioEconomicHealth =~ HIV_prevalence 1.00 0.000   NA    NA  
## 4 SocioEconomicHealth =~ Unemployment 0.26 0.143 1.815 0.070  
## ci.lower ci.upper  
## 1 1.000 1.00  
## 2 -0.490 0.77  
## 3 1.000 1.00  
## 4 -0.021 0.54
```

The Healthcare_Spending factor had a stronger effect on Private_health_expenditure than Health_expenditure because of its regression coefficient of 1.00 compared to 0.14.

The SocioEconomicHealth factor had a stronger effect on Hiv_prevalence than Unemployment because of its regression coefficient of 1.00 compared to 0.26.

```
parameterEstimates(MOD)[5:15,c("lhs","op", "rhs", "est", "z")]
```

STATISTICAL SIGNIFICANCE

##	lhs	op	rhs	est	z
## 5	Life_expectancy	~	SocioEconomicHealth	-0.138	-1.896
## 6	Life_expectancy	~	Healthcare_Spending	0.012	0.375
## 7	Life_expectancy	~	CO2_emissions	-0.021	-0.436
## 8	Life_expectancy	~	Electricity	0.609	8.383
## 9	Life_expectancy	~	Govt_health_expenditure	0.169	2.630
## 10	Life_expectancy	~	Tobacco_use	0.087	1.967
## 11	Life_expectancy	~	Alcohol_consumption	-0.107	-2.084
## 12	Life_expectancy	~	Political_stability	0.232	3.063
## 13	Life_expectancy	~	Freshwater_resources	-0.015	-0.542
## 14	Life_expectancy	~	Out_of_pocket	-0.203	-3.672
## 15	Life_expectancy	~	Govt_effectiveness	0.104	1.084

Electricity is the most significant predictor of Life_expectancy as it has the z-score with the highest magnitude (8.383). It is statistically significant because the p-value (0.000) is below the common alpha level of 0.05. There is positive relationship, such that for a 1-unit increase in a country's Electricity we would expect a 0.609 unit increase in Life_expectancy according to our model.

Out_of_pocket is the second-most significant predictor of Life_expectancy as it has the z-score with the second highest magnitude (3.672). It is statistically significant because the p-value (0.000) is below the common alpha level of 0.05. There is a negative relationship, such that for a 1-unit increase in in a country's Out_of_pocket we would expect a -0.203 unit decrease in Life_expectancy according to our model.

Political_stability is the third-most significant predictor of Life_expectancy as it has the z-score with the third highest magnitude (3.063). It is statistically significant because the p-value (0.002) is below the common alpha level of 0.05. There is a positive relationship, such that for a 1-unit increase in in a country's Political_stability we would expect a 0.232 unit positive in Life_expectancy according to our model.

Govt_health_expenditure is the fourth-most significant predictor of Life_expectancy as it has the z-score with the fourth highest magnitude (2.630). It is statistically significant because the p-value (0.009) is below the common alpha level of 0.05. There is a positive relationship, such that for a 1-unit increase in in a country's Govt_health_expenditure we would expect a 0.169 unit increase in Life_expectancy according to our model.

Alcohol_consumption is the fifth-most significant predictor of Life_expectancy as it has the z-score with the fifth highest magnitude (2.084). It is statistically significant because the p-value (0.037) is below the common alpha level of 0.05. There is a negative relationship, such that for a 1-unit increase in in a country's Alcohol_consumption we would expect a 0.107 unit decrease in Life_expectancy according to our model.

Tabacoo_use is the sixth-most significant predictor of Life_expectancy as it has the z-score with the sixth highest magnitude (1.967). It is statistically significant because the p-value (0.049) is below the common alpha level of 0.05. There is a positive relationship, such that for a 1-unit increase in in a country's Tabacoo_use we would expect a 0.087 unit increase in Life_expectancy according to our model.

```
indicies <- fitMeasures(MOD)
indicies
```

Model Fit

```
##          npar          fmin          chisq
##          21.000          1.064          172.425
##          df          pvalue    baseline.chisq
##          39.000          0.000          412.206
##    baseline.df    baseline.pvalue          cfi
##          55.000          0.000          0.626
##          tli          nnfi          rfi
##          0.473          0.473          0.410
##          nfi          pnfi          ifi
##          0.582          0.412          0.642
##          rni          logl    unrestricted.logl
##          0.626          -441.813          -355.601
##          aic          bic          ntotal
##          925.626          975.910          81.000
##          bic2          rmsea    rmsea.ci.lower
##          909.683          0.206          0.175
##    rmsea.ci.upper    rmsea.ci.level    rmsea.pvalue
##          0.237          0.900          0.000
##    rmsea.close.h0    rmsea.notclose.pvalue    rmsea.notclose.h0
##          0.050          1.000          0.080
##          rmr          rmr_nomean          srmr
##          0.191          0.191          0.198
##    srmr_bentler    srmr_bentler_nomean          crmr
##          0.198          0.198          0.211
##    crmr_nomean    srmr_mplus    srmr_mplus_nomean
##          0.211          0.197          0.197
##          cn_05          cn_01          gfi
##          26.636          30.327          0.899
##          agfi          pgfi          mfi
##          0.729          0.334          0.439
##          ecvi
##          2.647
```

Comparative Fit Indices

```
indicies['nfi']
```

```
##          nfi
## 0.5817025
```

```
indicies['nnfi']
```

```
##          nnfi
## 0.4732364
```

```
indicies['ifi']
```

```
##      ifi  
## 0.6424904
```

```
indicies['cfi']
```

```
##      cfi  
## 0.6264768
```

Our Normed Fit Index lies at 0.582, our Non-normed Fit Index lies at 0.473, our Incremental Fit Index lies at 0.642, and our Comparative Fit Index lies at 0.626. These are below the common threshold (0.90–0.95) that is considered a good fit, so our model does not provide an adequate fit when compared to a fully independent model according to these comparative fit indices.

```
indicies['rmsea']
```

```
##      rmsea  
## 0.2055149
```

Our Root Mean Square Error of Approximation lies at 0.206 which is above the common threshold of 0.08. This means that our model does not provide an adequate fit to the data when compared to a fully saturated model according to this index of fit.

Absolute Fit Indices

```
indicies['mfi']
```

```
##      mfi  
## 0.438845
```

Our McDonald & Marsh Fit Index lies at 0.439 which is far below the common threshold of 0.90–0.95. This means our model does not provide an adequate fit to the original covariance structure.

```
indicies['gfi']
```

```
##      gfi  
## 0.8994171
```

```
indicies['agfi']
```

```
##      agfi  
## 0.7291999
```

Our goodness-of-fit index lies at 0.899 and our adjusted goodness-of-fit lies at 0.729. This means that about 72.9% of the covariability in the original data matrix can be explained by the structural equation model.