Attitudes Towards Mental Health in the Technology Workplace: Structural Equation Modeling

EPPS 7318: Structural Equation and Multilevel (Hierarchical) Modeling

Why focus on mental health in the tech workplace?

- Mental health needs to be addressed by employers in order to support the work environment (Joyce at al. 2016)
- Difficult to address mental health due to stigma in the workplace (Krupa et al. 2009)
- Study determined that employees with serious mental illness are aware when they face discrimination (Baldwin and Marcus 2006)
- Knowledge, attitudes, and behavior are three problems that cause mental health stigma (Brohan and Thornicroft 2010)

Data: Open Sourcing Mental Illness (OSMI)

- Non-profit conducting annual Mental Health in Tech Survey
 - Understand and examine mental health concerns in the tech community
- Raw data used from 2019 version of survey
 - Missing values
- Sample size of 352 participants
- Includes 82 different survey questions
 - Likert Scale survey questions

Variables

. su

| Variable | 0bs | Mean | Std. Dev. | Min | Max |
|------------|-----|----------|-----------|-----|-----|
| emp_phi | 304 | 6.328947 | 2.298734 | 0 | 10 |
| emp_mhi | 304 | 4.878289 | 2.609007 | 0 | 10 |
| pemp_phi | 296 | 5.587838 | 2.549653 | 0 | 10 |
| pemp_mhi | 296 | 3.527027 | 2.418701 | 0 | 10 |
| share_ff | 352 | 6.272727 | 2.65908 | 0 | 10 |
| a_career | 18 | 3.833333 | 2.202939 | 0 | 8 |
| work_react | 352 | 5.275568 | 2.208679 | 0 | 10 |
| support | 352 | 2.602273 | .9641999 | 1 | 5 |

Spearman's Correlation

Ordinal variables with a monotonic relationship

- Spearman's correlation coefficient indicates a mediocre monotonic relationship among variables
 - Results are statistically significant

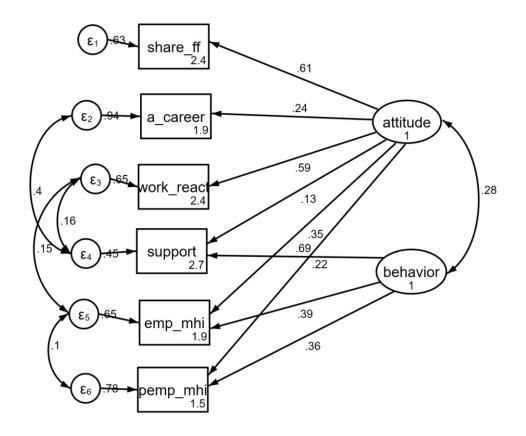
Hypothesis and Variables

- 2 models with 2 exogenous latent constructs which covary
 - Hypothesis: survey questions accurately measure attitude and behavior towards mental health
- Error covariances only observed between:
 - support and work_react
- Error covariances observed between:
 - support and work_react
 - support and a_career
 - emp_mhi and work_react
 - emp_mhi and pemp_mhi

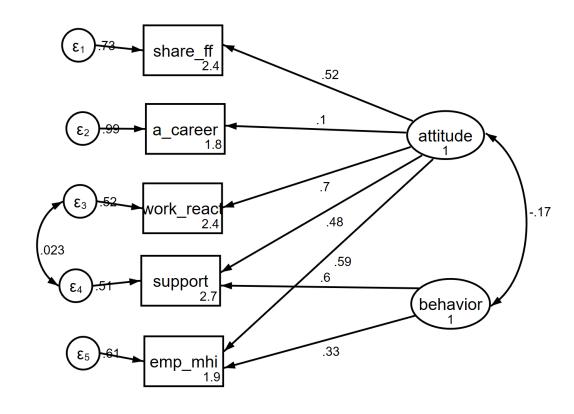
- Behavior concerning mental health
 - emp_mhi
 - support
 - pemp_mhi
- Overall attitude towards mental health
 - support
 - work_react
 - a_career
 - share ff
 - emp_mhi
 - pemp_mhi

Confirmatory Factor Analysis

Model 1



Model 2



Goodness of Fit

Model 1

. estat gof, stats(all)

| Fit statistic | Value | Description |
|----------------------|----------|--|
| Likelihood ratio | | |
| chi2_ms(1) | 6.194 | model vs. saturated |
| p > chi2 | 0.013 | |
| chi2_bs(15) | 226.947 | baseline vs. saturated |
| p > chi2 | 0.000 | |
| Population error | | |
| RMSEA | 0.122 | Root mean squared error of approximation |
| 90% CI, lower bound | 0.045 | |
| upper bound | 0.221 | |
| pclose | 0.061 | Probability RMSEA <= 0.05 |
| Information criteria | | |
| AIC | 6930.854 | Akaike's information criterion |
| BIC | 7031.308 | Bayesian information criterion |
| Baseline comparison | | |
| · CFI | 0.975 | Comparative fit index |
| TLI | 0.632 | Tucker-Lewis index |
| Size of residuals | | |
| CD | 0.839 | Coefficient of determination |
| , | | |

Note: SRMR is not reported because of missing values.

Model 2

. estat gof, stats(all)

| Fit statistic | Value | Description |
|----------------------|----------|--|
| Likelihood ratio | | |
| chi2 ms(1) | 4.798 | model vs. saturated |
| p > chi2 | 0.028 | |
| chi2 bs(10) | 170.824 | baseline vs. saturated |
| p > chi2 | 0.000 | |
| Population error | | |
| RMSEA | 0.104 | Root mean squared error of approximation |
| 90% CI, lower bound | 0.027 | |
| upper bound | 0.205 | |
| pclose | 0.106 | Probability RMSEA <= 0.05 |
| Information criteria | | |
| AIC | 5609.699 | Akaike's information criterion |
| BIC | 5683.108 | Bayesian information criterion |
| Baseline comparison | | |
| CFI | 0.976 | Comparative fit index |
| TLI | 0.764 | Tucker-Lewis index |
| Size of residuals | | |
| CD | 0.807 | Coefficient of determination |

Method of Analysis

- Confirmatory Factor Analysis demonstrates that the second model is better based on both a lower AIC and BIC and a higher CFI and TLI, however, the RMSEA is a poor fit for both models
- Reject the null
 - Therefore the hypothesized structure does not fit the data well enough
- Exploratory Factor Analysis is executed in order to gain a better understanding about the structure of the variables and how many dimensions are in a set of variables (Byrne 2013)

EFA 1

. factor work_react support emp_mhi pemp_mhi share_ff a_career
(obs=14)

| Factor analysis/correlation | Number of obs = | 14 |
|-----------------------------|--------------------|----|
| Method: principal factors | Retained factors = | 4 |
| Rotation: (unrotated) | Number of params = | 15 |

| Factor | Eigenvalue | Difference | Proportion | Cumulative |
|---------|------------|------------|------------|------------|
| Factor1 | 1.88958 | 1.17384 | 0.6931 | 0.6931 |
| Factor2 | 0.71575 | 0.42992 | 0.2625 | 0.9556 |
| Factor3 | 0.28583 | 0.11452 | 0.1048 | 1.0604 |
| Factor4 | 0.17131 | 0.24494 | 0.0628 | 1.1233 |
| Factor5 | -0.07363 | 0.18879 | -0.0270 | 1.0962 |
| Factor6 | -0.26241 | | -0.0962 | 1.0000 |

LR test: independent vs. saturated: chi2(15) = 21.69 Prob>chi2 = 0.1162

Factor loadings (pattern matrix) and unique variances

| Variable | Factor1 | Factor2 | Factor3 | Factor4 | Uniqueness |
|------------|---------|---------|---------|---------|------------|
| work_react | 0.7865 | -0.4300 | 0.0861 | 0.1145 | 0.1759 |
| support | 0.4293 | 0.4034 | 0.1688 | -0.2336 | 0.5698 |
| emp_mhi | 0.8988 | 0.0595 | -0.0656 | -0.1113 | 0.1720 |
| pemp_mhi | 0.1916 | 0.1762 | 0.3768 | 0.2111 | 0.7457 |
| share_ff | 0.4797 | 0.0301 | -0.2863 | 0.1019 | 0.6767 |
| a_career | 0.1098 | 0.5768 | -0.1473 | 0.1905 | 0.5973 |
| | | | | | |

EFA 2

. factor work_react support emp_mhi pemp_mhi share_ff
(obs=253)

| Factor analysis/correlation | Number of obs = | 253 |
|-----------------------------|----------------------|-----|
| Method: principal factors | Retained factors = | 2 |
| Rotation: (unrotated) | Number of params $=$ | 9 |

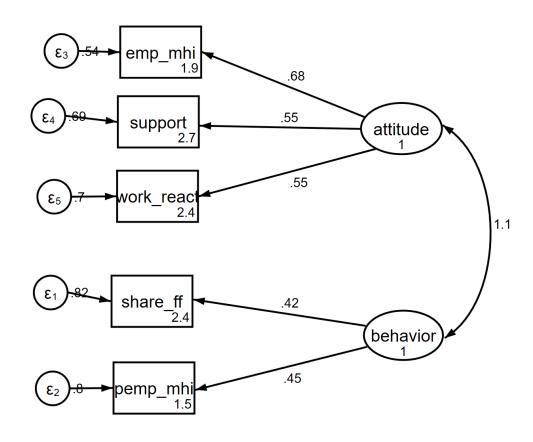
| Factor | Eigenvalue | Difference | Proportion | Cumulative |
|---------|------------|------------|------------|------------|
| Factor1 | 1.39226 | 1.26922 | 1.3405 | 1.3405 |
| Factor2 | 0.12304 | 0.25139 | 0.1185 | 1.4589 |
| Factor3 | -0.12836 | 0.01820 | -0.1236 | 1.3354 |
| Factor4 | -0.14656 | 0.05519 | -0.1411 | 1.1942 |
| Factor5 | -0.20175 | • | -0.1942 | 1.0000 |

LR test: independent vs. saturated: chi2(10) = 172.60 Prob>chi2 = 0.0000

Factor loadings (pattern matrix) and unique variances

| Variable | Factor1 | Factor2 | Uniqueness |
|--|--|---|--|
| work_react support emp_mhi pemp_mhi share_ff | 0.5276 0.5383 0.6289 0.4520 0.4736 | 0.1862 -0.1298 -0.0576 -0.1804 0.1888 | 0.6870 0.6934 0.6011 0.7632 0.7400 |

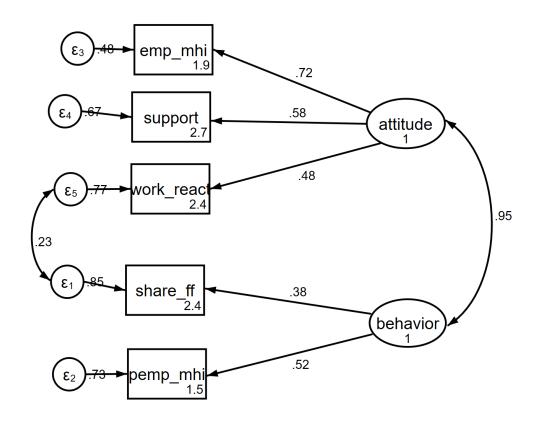
Model 3



Goodness of Fit

| Fit statistic | Value | Description |
|----------------------|----------|--|
| Likelihood ratio | | |
| chi2_ms(4) | 16.669 | model vs. saturated |
| p > chi2 | 0.002 | |
| chi2_bs(10) | 219.029 | baseline vs. saturated |
| p > chi2 | 0.000 | |
| Population error | | |
| RMSEA | 0.095 | Root mean squared error of approximation |
| 90% CI, lower bound | 0.051 | |
| upper bound | 0.144 | |
| pclose | 0.047 | Probability RMSEA <= 0.05 |
| Information criteria | | |
| AIC | 6850.761 | Akaike's information criterion |
| BIC | 6912.579 | Bayesian information criterion |
| Baseline comparison | | |
| CFI | 0.939 | Comparative fit index |
| TLI | 0.848 | Tucker-Lewis index |
| Size of residuals | | |
| CD | 0.667 | Coefficient of determination |

Model 4



Goodness of Fit

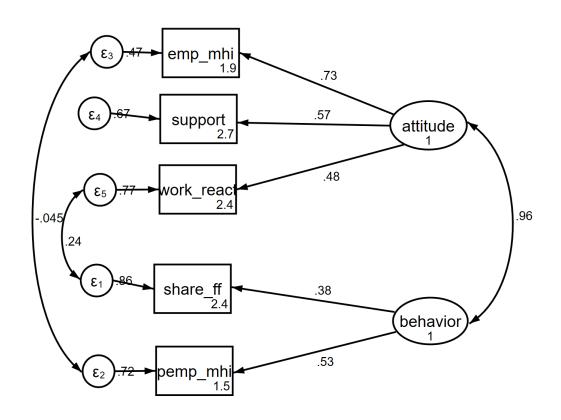
estat gof, stats(all)

Size of residuals

| Fit statistic | Value | Description |
|----------------------|----------|--|
| Likelihood ratio | | |
| chi2_ms(3) | 3.487 | model vs. saturated |
| p > chi2 | 0.322 | |
| chi2_bs(10) | 219.029 | baseline vs. saturated |
| p > chi2 | 0.000 | |
| Population error | | |
| RMSEA | 0.022 | Root mean squared error of approximation |
| 90% CI, lower bound | 0.000 | |
| upper bound | 0.095 | |
| pclose | 0.642 | Probability RMSEA <= 0.05 |
| Information criteria | | |
| AIC | 6839.579 | Akaike's information criterion |
| BIC | 6905.261 | Bayesian information criterion |
| Baseline comparison | | |
| CFI | 0.998 | Comparative fit index |
| TLI | 0.992 | Tucker-Lewis index |
| | | |

Coefficient of determination

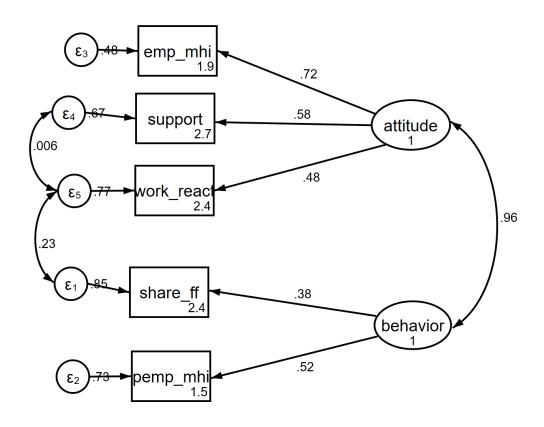
Model 5



Goodness of Fit

| Fit statistic | Value | Description |
|----------------------|----------|--|
| Likelihood ratio | | |
| chi2_ms(2) | 3.370 | model vs. saturated |
| p > chi2 | 0.185 | |
| chi2_bs(10) | 219.029 | baseline vs. saturated |
| p > chi2 | 0.000 | |
| Population error | | |
| RMSEA | 0.044 | Root mean squared error of approximation |
| 90% CI, lower bound | 0.000 | |
| upper bound | 0.124 | |
| pclose | 0.435 | Probability RMSEA <= 0.05 |
| Information criteria | | |
| AIC | 6841.462 | Akaike's information criterion |
| BIC | 6911.007 | Bayesian information criterion |
| Baseline comparison | | |
| CFI | 0.993 | Comparative fit index |
| TLI | 0.967 | Tucker-Lewis index |
| Size of residuals | | |
| CD | 0.718 | Coefficient of determination |

Model 6



Goodness of Fit

| Fit statistic | Value | Description |
|----------------------|----------|--|
| Likelihood ratio | | |
| chi2_ms(2) | 3.480 | model vs. saturated |
| p > chi2 | 0.175 | |
| chi2_bs(10) | 219.029 | baseline vs. saturated |
| p > chi2 | 0.000 | |
| Population error | | |
| RMSEA | 0.046 | Root mean squared error of approximation |
| 90% CI, lower bound | 0.000 | |
| upper bound | 0.125 | |
| pclose | 0.422 | Probability RMSEA <= 0.05 |
| Information criteria | | |
| AIC | 6841.573 | Akaike's information criterion |
| BIC | 6911.118 | Bayesian information criterion |
| Baseline comparison | | |
| CFI | 0.993 | Comparative fit index |
| TLI | 0.965 | Tucker-Lewis index |
| Size of residuals | | |
| CD | 0.707 | Coefficient of determination |

Conclusion

- Confirmatory Factor Analysis demonstrates that the fourth model is better based on a lower AIC and BIC, a higher CFI and TLI
- The RMSEA indicates that there is a very good fit for the fourth model

• Model selection: models do not significantly differ and therefore the fourth model is selected since it is less complicated in comparison (parsimonious) while still demonstrating a very good fit

References

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