

Thinking About and Approaching Statistics

Waylon Howard, PhD



Slides available at <http://tinyurl.com/pharmerit-slides>

PDF slides at <http://tinyurl.com/pharmerit-pdf>



Global Director of Biostatistics
and Data Analytics



 @waylon-howard

 @wwaylon

 www.waylon.github.io

Education

Ph.D., Quantitative Psychology
University of Kansas (2012);

Advisor: [Todd D. Little](#)

Currently

Lead a [10-person](#) team

Oncology focused RWE analytics

- [Clinical: EMR curation](#)
- Humanistic: PRO data
- Economic: claims

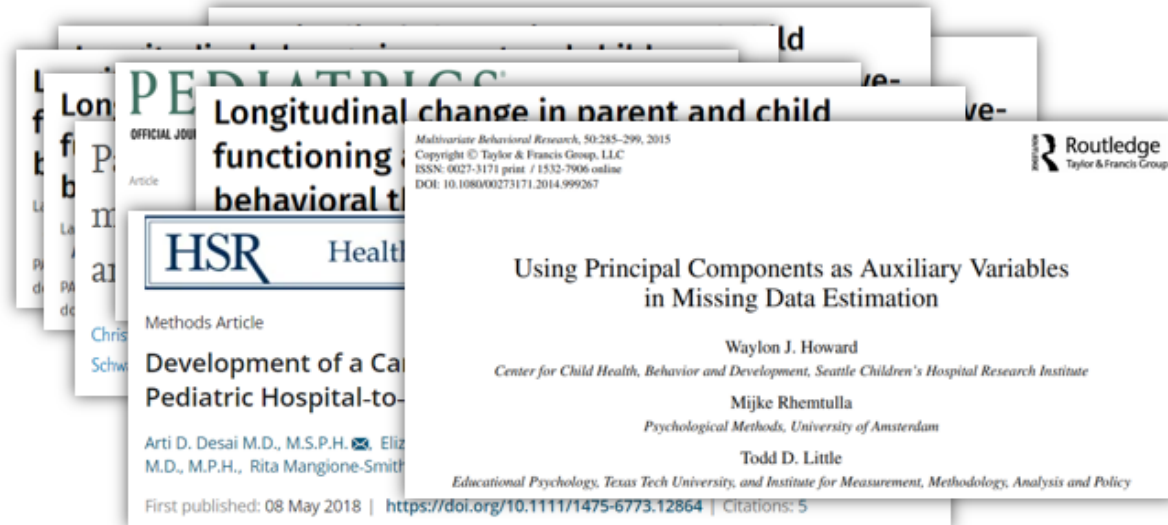
[SymphonyAI](#) acquisition

- [More than 60%](#) growth
- Developed 15+ SOP, WI, GD

Table of Contents

- Research and Leadership
- Motivating example
- Modern modeling highlights
- Summary

Interdisciplinary-oriented collaborator



My work has appeared in **22** different peer-reviewed journals garnering **518** citations, with an h-index of **13** and an i-10 index of **14**.

Grants and contracts

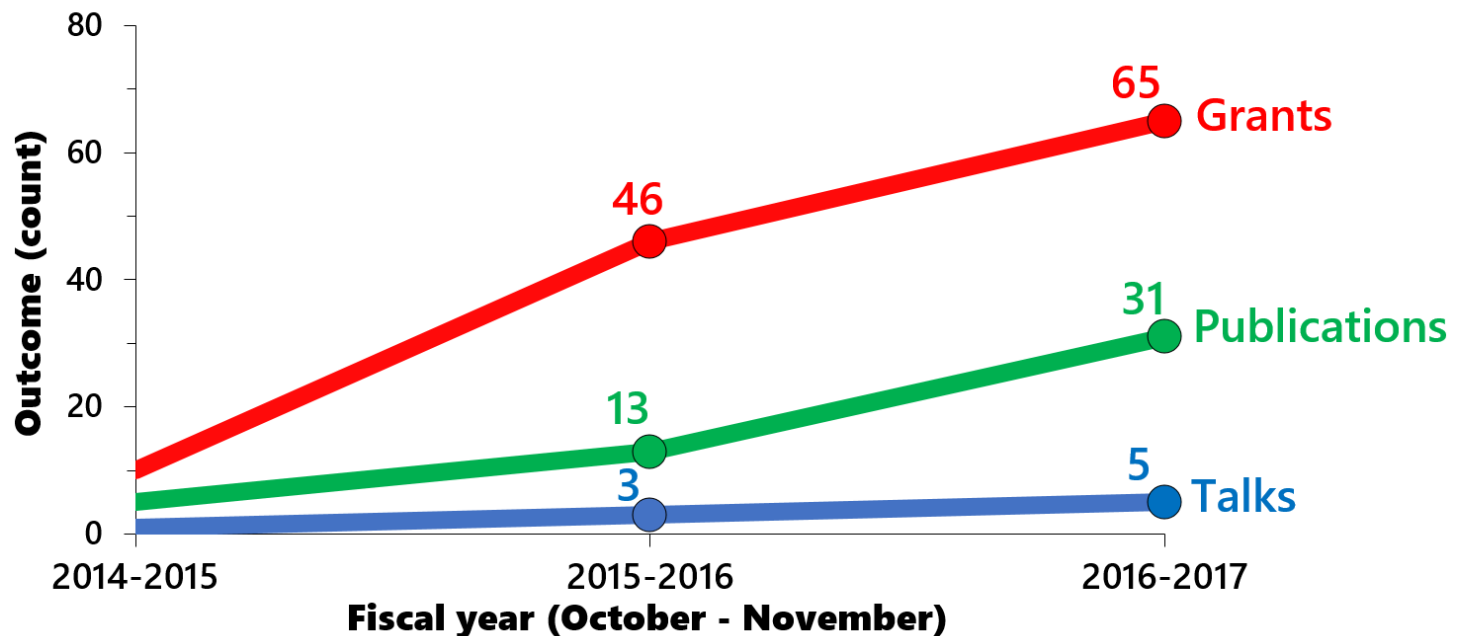
I have directly collaborated with researchers to attain **more than \$5 million** in new research funding and developed considerable experience applying diverse best-practice methods to complex problems in new areas.

Training

Organized, managed, and taught an 8-course advanced methodology workshop for faculty and graduate students (N = 60) and a 6-course basic methodology training series for residents (N = 150).

Mentoring

The coalescence of a new vision for team leadership within the organization that resulted in new funding, partnerships, and additional value offered by the team.

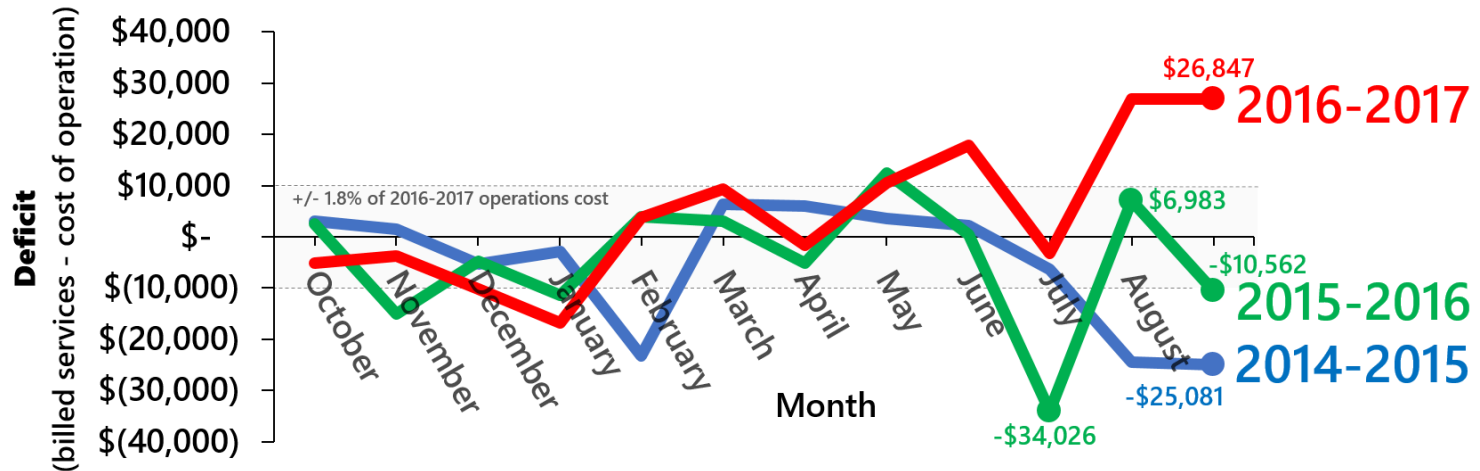


Is this change *significant*?

Different question:

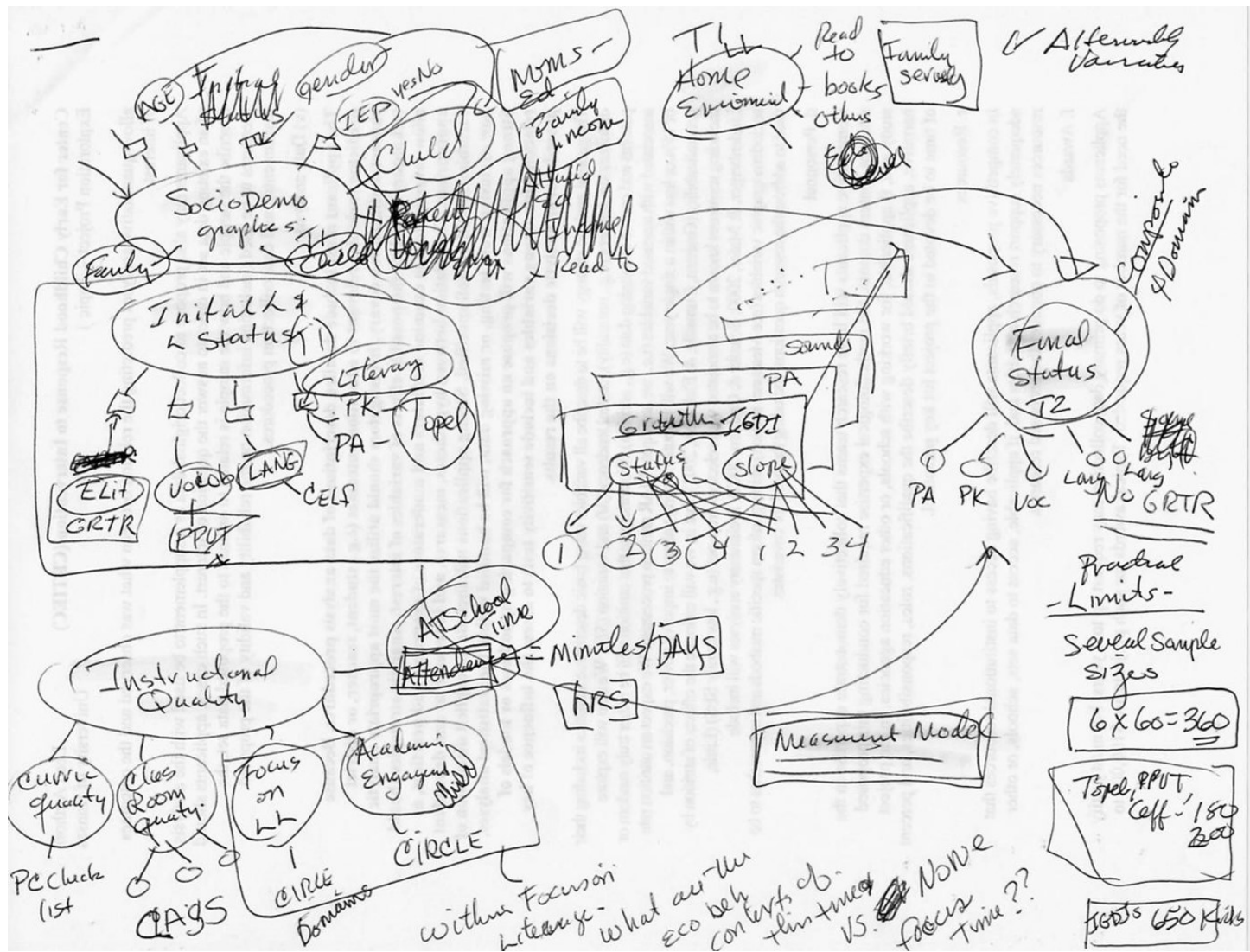
Is this change *sustainable*?

Remaining sustainable

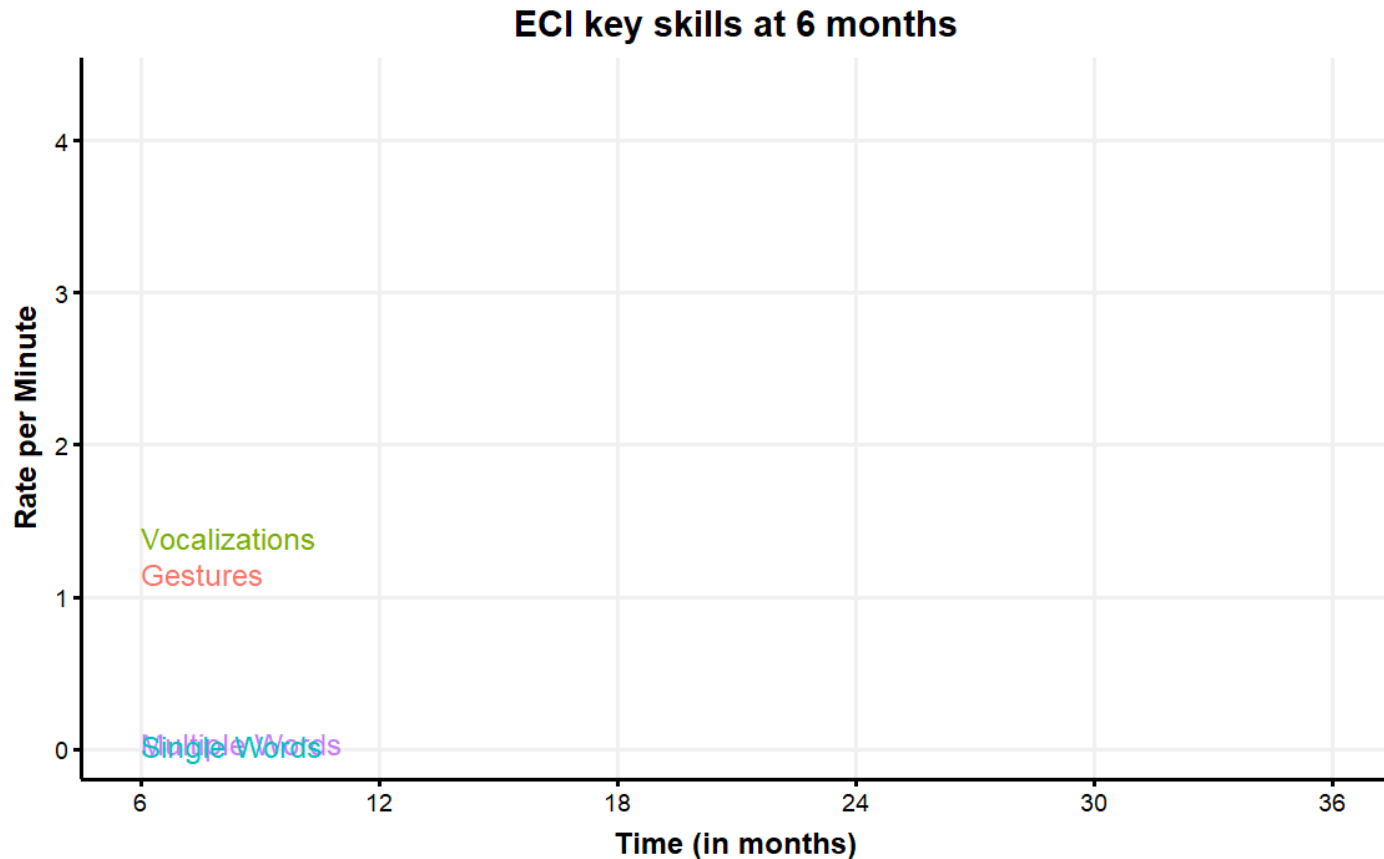


- Services billed increased by 38.6%
- Team averaged 78.3% billable time
- Compound annual growth rate +9.2%

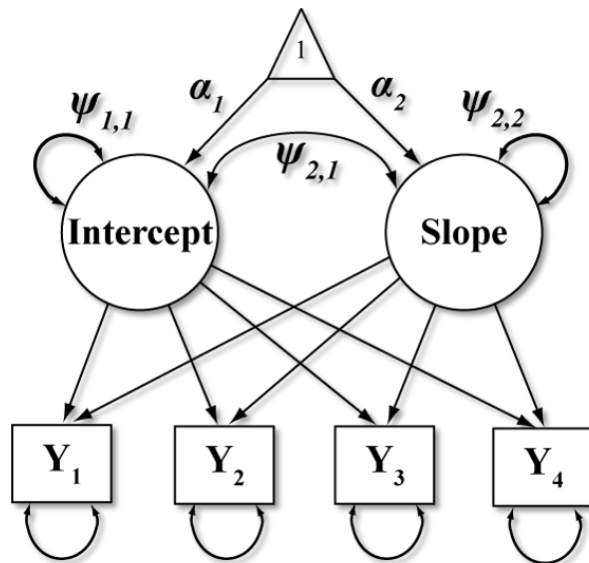
Motivating example



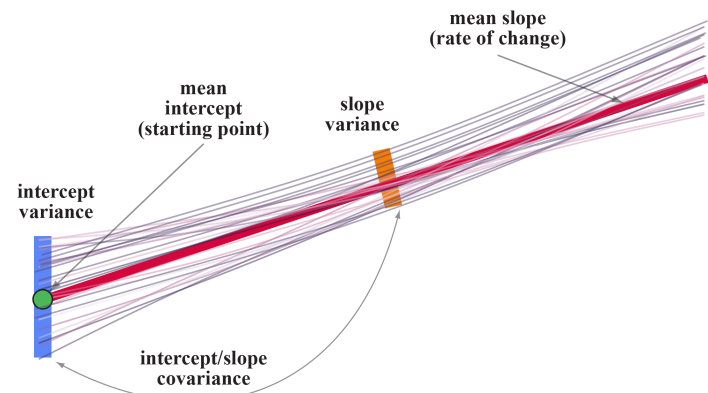
First motivating example for today



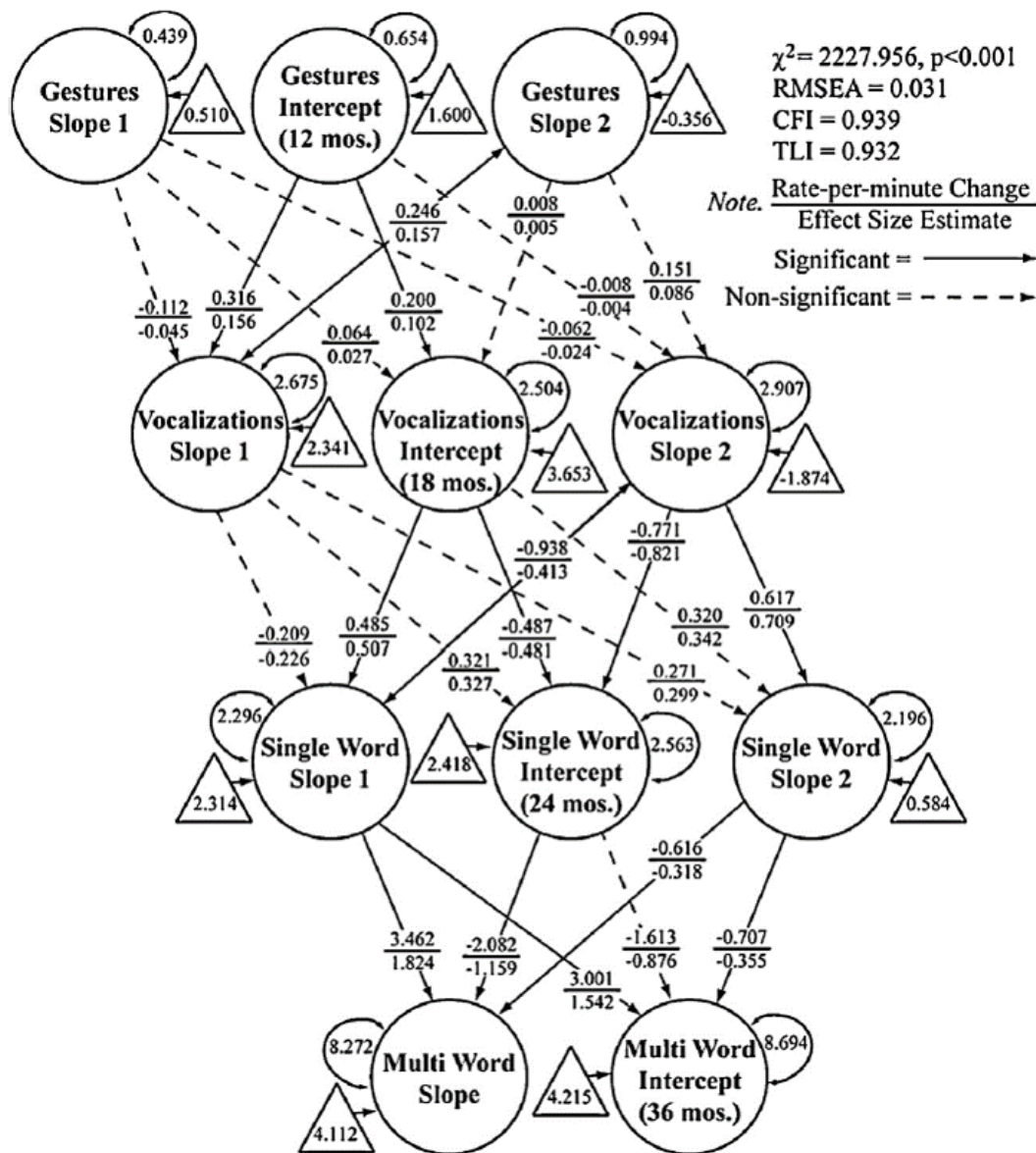
Exemplary LGM model



Path diagram



Parameter illustration

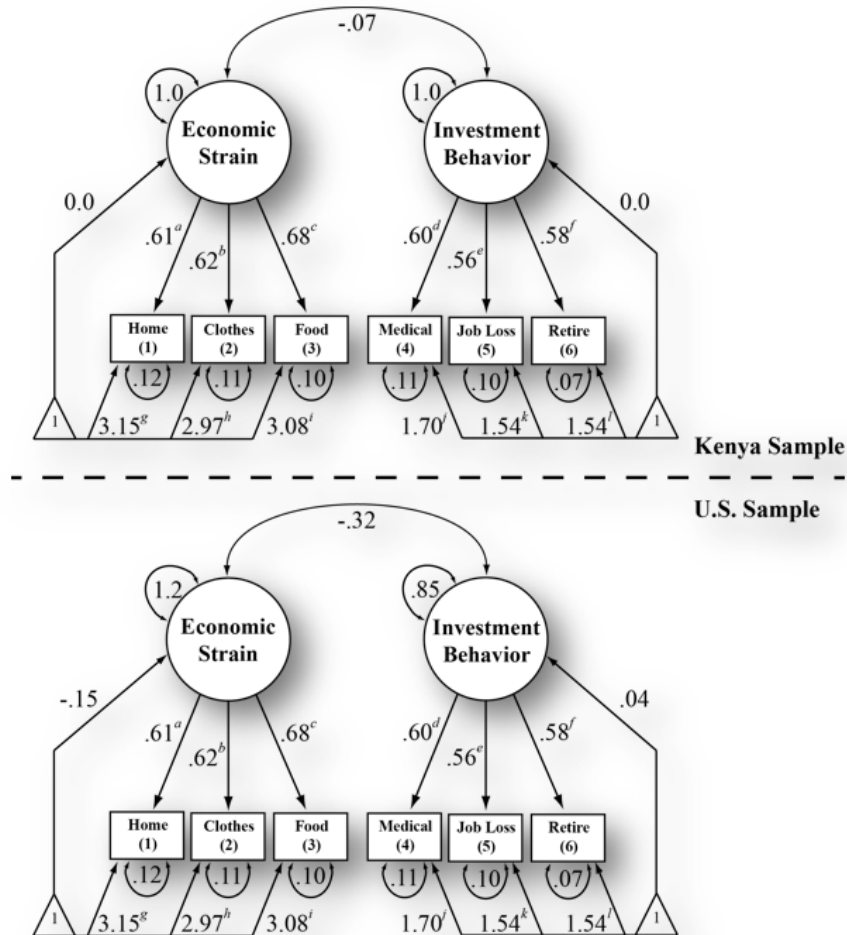


Highlights: Measurement invariance

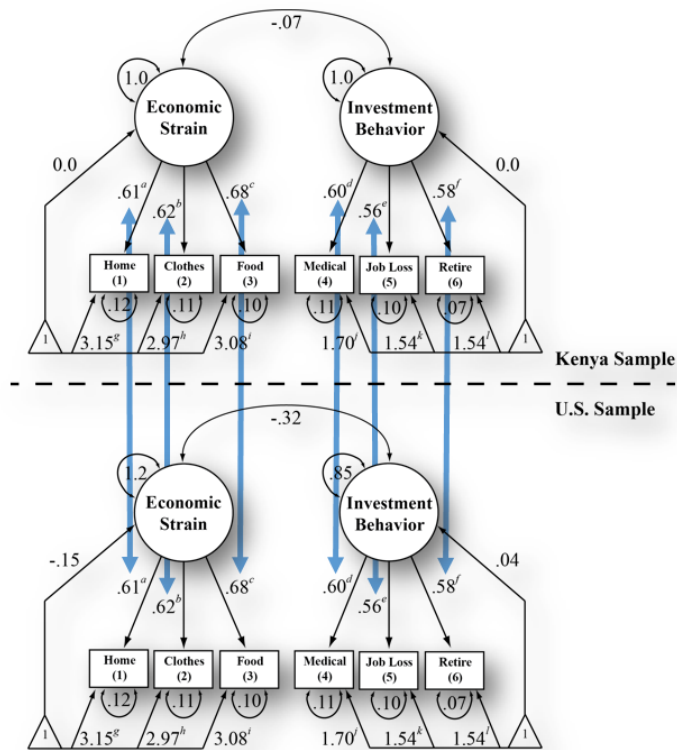
Why?

To ensure we are measuring the same constructs across groups and time - allows for structural invariance testing

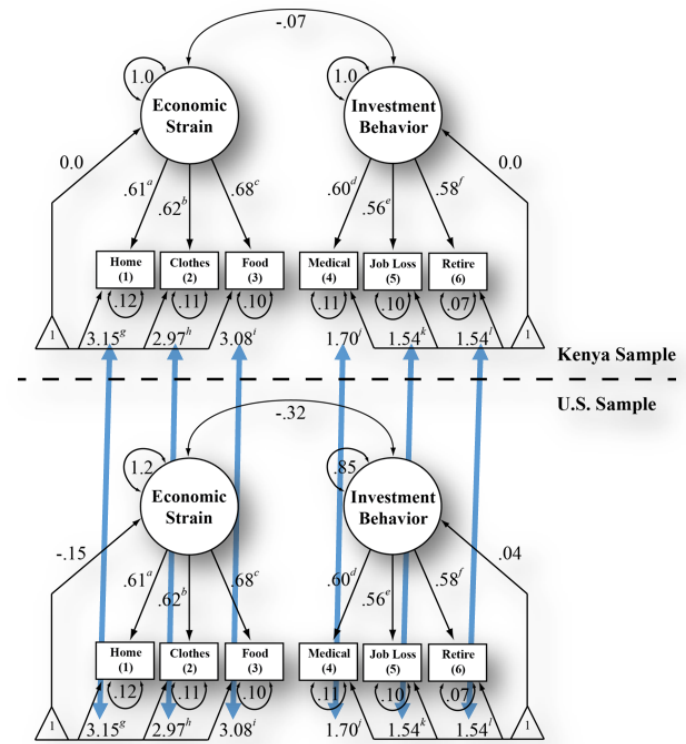
1. Configural invariance



2. Weak invariance

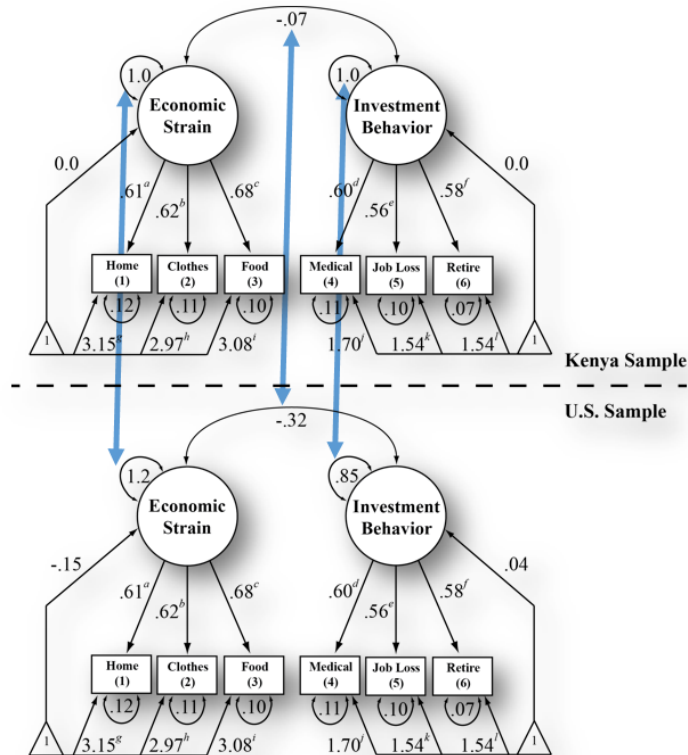


3. Strong invariance

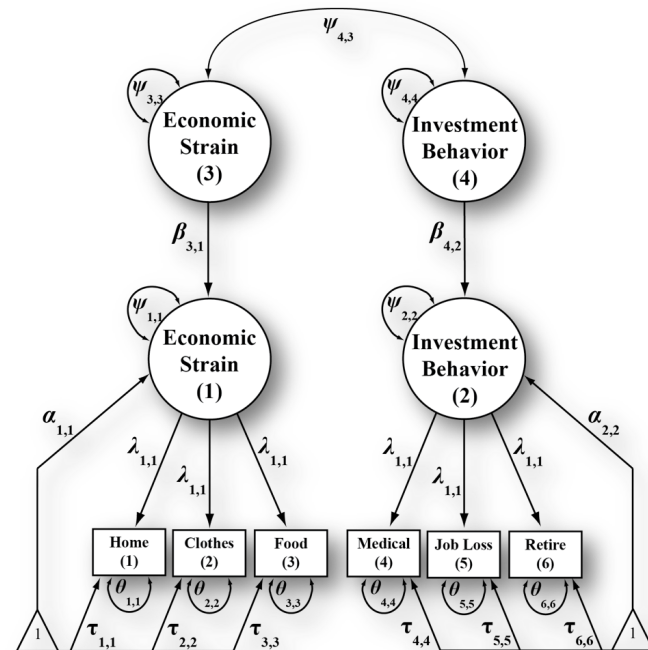


Structural invariance

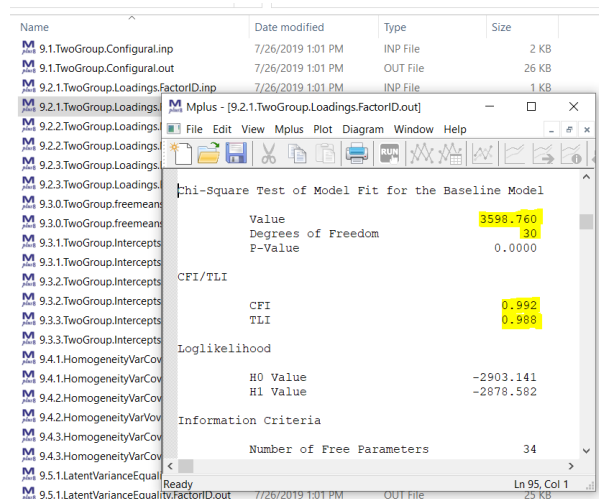
Latent correlations



Phantom variables



A note about efficiency tools

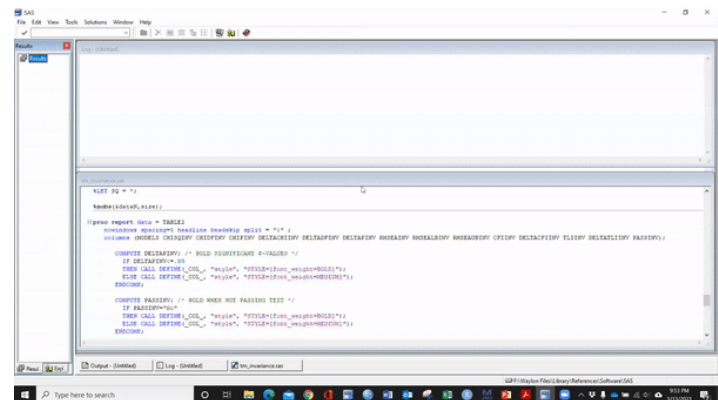


The image shows a file explorer window with a list of Mplus output files. The files are organized by model name and type (INP File or OUT File). The file sizes range from 1 KB to 26 KB. The file names include model numbers (e.g., 9.1, 9.2, 9.3, 9.4, 9.5) and model descriptions (e.g., TwoGroup, Loadings, HomogeneityVarCov, LatentVarianceEqual).

Name	Date modified	Type	Size
9.1.TwoGroup.Configural.inp	7/26/2019 1:01 PM	INP File	2 KB
9.1.TwoGroup.Configural.out	7/26/2019 1:01 PM	OUT File	26 KB
9.2.1.TwoGroup.Loadings.FactorID.inp	7/26/2019 1:01 PM	INP File	1 KB
9.2.1.TwoGroup.Loadings.FactorID.out	7/26/2019 1:01 PM	OUT File	25 KB
9.2.2.TwoGroup.Loadings.FactorID.inp	7/26/2019 1:01 PM	INP File	1 KB
9.2.2.TwoGroup.Loadings.FactorID.out	7/26/2019 1:01 PM	OUT File	25 KB
9.2.3.TwoGroup.Loadings.FactorID.inp	7/26/2019 1:01 PM	INP File	1 KB
9.2.3.TwoGroup.Loadings.FactorID.out	7/26/2019 1:01 PM	OUT File	25 KB
9.3.0.TwoGroup.freemean	7/26/2019 1:01 PM	OUT File	25 KB
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9.3.1.TwoGroup.Intercepts	7/26/2019 1:01 PM	OUT File	25 KB
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9.4.1.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
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9.4.2.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
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9.4.3.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
9.4.3.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
9.5.1.LatentVarianceEqual	7/26/2019 1:01 PM	OUT File	25 KB
9.5.1.LatentVarianceEqual	7/26/2019 1:01 PM	OUT File	25 KB

Chi-square Test of Model Fit for the Baseline Model	
Value	3598.760
Degrees of Freedom	30
P-Value	0.0000
CFI/TLI	
CFI	0.992
TLI	0.988
Loglikelihood	
H0 Value	-2903.141
H1 Value	-2878.582
Information Criteria	
Number of Free Parameters	34

Numerous **Mplus** output files



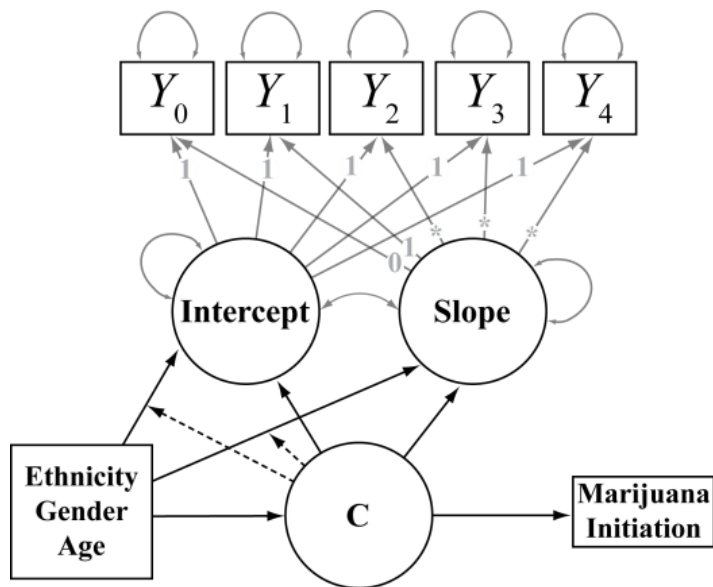
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9.4.1.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
9.4.2.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
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9.4.3.HomogeneityVarCov	7/26/2019 1:01 PM	OUT File	25 KB
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Loglikelihood	
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Information Criteria	
Number of Free Parameters	34

Example model fit table
automation

A note about organizational tools



Σ_k	# of classes (K)	LL	npar	AIC	BIC	CAIC	AWE	LRTS	Adj LMR p-value	BF _{K,K+1}	cmP _K
Most Restricted Model Class-invariant, diagonal $\Sigma_k = \Sigma$ Variances freely estimated BUT constrained across groups - covariances fixed to zero (psidiaginv.lnp)	1	-1079.65	4	2167.29	2179.22	2183.22	2211.16	11.96	---	4.46	---
	2	-1073.66	7	2161.33	2182.21	2189.21	2238.10	13.39	0.014	2.18	0.1690
	3	-1066.97	10	2153.94	2183.77	2193.77	2263.61	2.80	0.027	435.03	0.0775
	4	-1065.57	13	2157.14	2195.92	2208.92	2299.71	5.09	0.199	138.16	0.0002
	5	-1063.02	16	2158.04	2205.78	2221.78	2333.52	---	0.709	---	0.0000
Class-varying, diagonal Σ_k Variances freely estimated BUT covariances fixed to zero (psidiagvar.lnp)	1	-1079.65	4	2167.29	2179.22	2183.22	2211.16	23.65	---	1.88	---
	2	-1067.82	9	2153.64	2180.49	2189.49	2252.34	10.29	0.002	1499.69	0.4000
	3	-1062.67	14	2153.35	2195.12	2209.12	2306.89	5.35	0.160	1471.76	0.0003
	4	-1060.00	18	2156.00	2209.70	2227.70	2353.41	---	0.000	---	0.0000
Class-invariant, unrestricted $\Sigma_k = \Sigma$ Variances freely estimated BUT covariances equated across groups (psionddiagvar.lnp)	1	-1077.56	5	2165.11	2180.03	2185.03	2219.95	9.07	---	18.94	---
	2	-1073.02	8	2162.05	2185.91	2193.91	2249.78	15.08	0.312	0.94	0.0266
	3	-1065.49	11	2152.97	2185.79	2196.79	2273.61	3.62	0.026	289.00	0.0283
	4	-1063.68	14	2155.35	2197.12	2211.12	2308.89	4.61	0.603	176.34	0.0001
	5	-1061.37	17	2156.75	2207.47	2224.47	2343.19	---	0.221	---	0.0000

a transparent, traceable tool

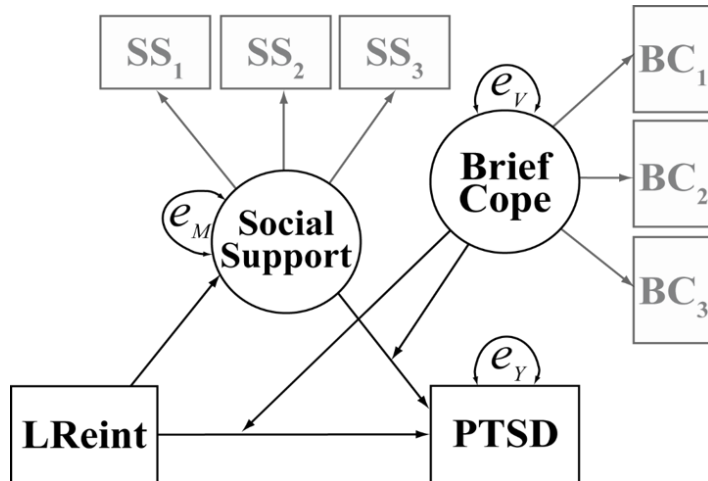
a complex, iterative process...

Highlights: moderated mediation

Why?

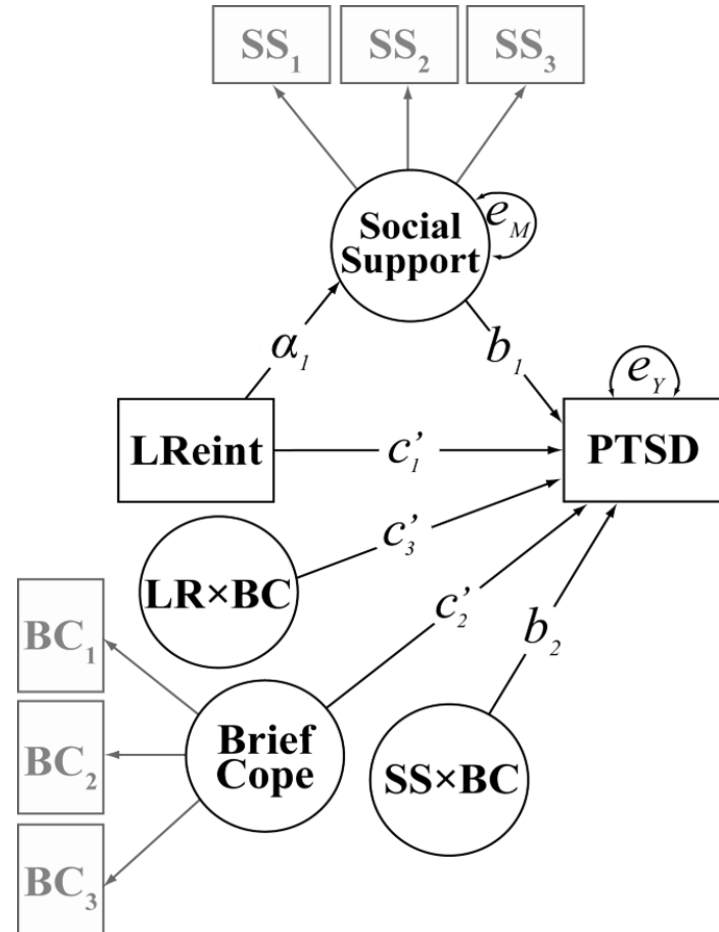
We often want to know how (mediation) and when (moderation) predictive relations hold or are strong versus weak

Conceptual diagram



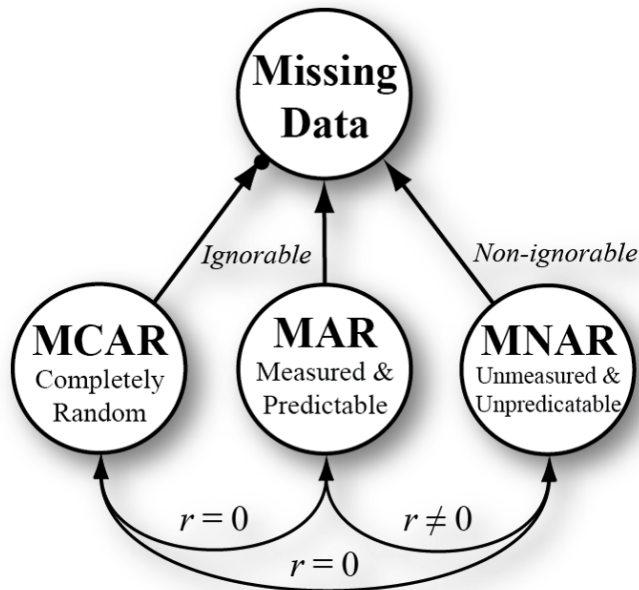
See [Full-text available](#)

Statistical diagram



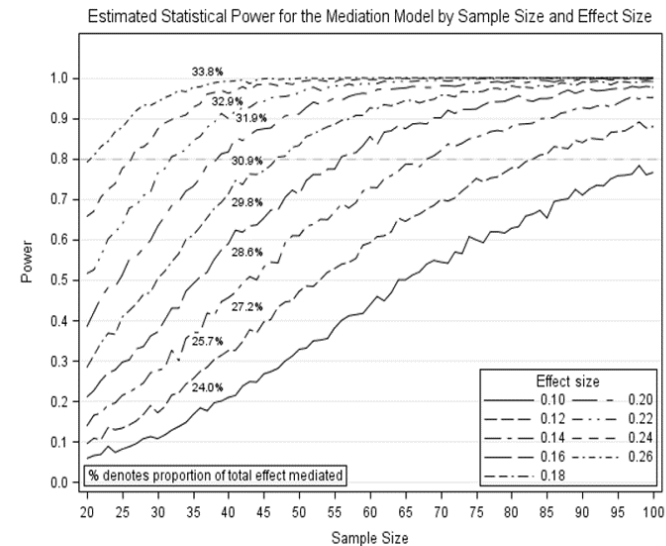
Highlights: Missing data and power

Missing Data Analysis



Simulation in MBR **Full-text**
Reporting practices **Full-text**

Monte Carlo power simulations



Software wrangling
Statistical library and templates

Summary

1. Leading technical experts for the past 10 years
2. Extensive real-world experience applying diverse best-practice methods to complex problems in new areas
3. Readily pivot from the strategic to hands on
4. Thrive in fast-paced, changing work environment

Thanks!

Any further questions?

- Slides created via the R package `xaringan` by Yihui Xie
- Slides' source code at <https://www.waylon.github.io/assets/ph-2021/>
- R code from throughout the slides as an R script as `slide_code.R`

Appendix

Previous research appointments



Biostatistics Manager



Dir. of Biostatistics



Dir. of Biostatistics Core



Faculty Res. Methodologist



Dir. of Res. & Evaluation



Senior Res. Data Analyst

KU Quantitative Analyst

- Position details [here](#)

Full-text research content

- Peer-reviewed research articles
- Talks, workshops, posters

Exemplar research tools

- Consultant-based **effort estimator**
- Timeline visualizations for **project management and communication**
- Rmarkdown automation of CV, NIH biosketch, NSF biosketch, etc.

Example standard process development

Training	Document ID	Document Title
Course-015	GD-OSS015-R01	Statistical Programming Best Practices
Course-016	GD-OSS016-R01	Analysis Datasets Results Verification
Course-017	GD-OSS017-R01	Reusable Code Validation
Course-020	GD-OSS020-R01	Procedures for Annotation and QC of TLFs
Course-001	CHKLST-OSS001	Source Code Version Control
Course-002	CHKLST-OSS002	Programming Quality Control Checklist
Course-003	CHKLST-OSS003	Disk Space Management Checklist
Course-004	CHKLST-OSS004	AWS Import/export Checklist
Course-005	CHKLST-OSS005	Data Delivery Checklist
Course-006	CHKLST-OSS006	Chemo Master List
Course-006	WI-OSS001	De-identification and Data Transfer

Software summary (not all-inclusive)

- SAS 9.4 (STAT, IML)
- Mplus 8.4 (base, mixture, multilevel)
- IBM SPSS statistics 25 (base, missing values, AMOS)
- R (data manipulation: dplyr, tidyr, stringr, RMySQL, RSQLite; data visualization: ggplot2, htmlwidgets; reporting: shiny, rmarkdown; analysis: lavaan, psych, lme4/nlme, survival)

Methodology interest summary

- Design and measurement issues in longitudinal research, panel designs, latent growth curve analysis, latent class and finite mixture modeling, multi-level SEM with longitudinal data, moderation and mediation, missing data analysis and power, and measurement invariance testing.