

# Portland CAR: next steps

2016-08-05

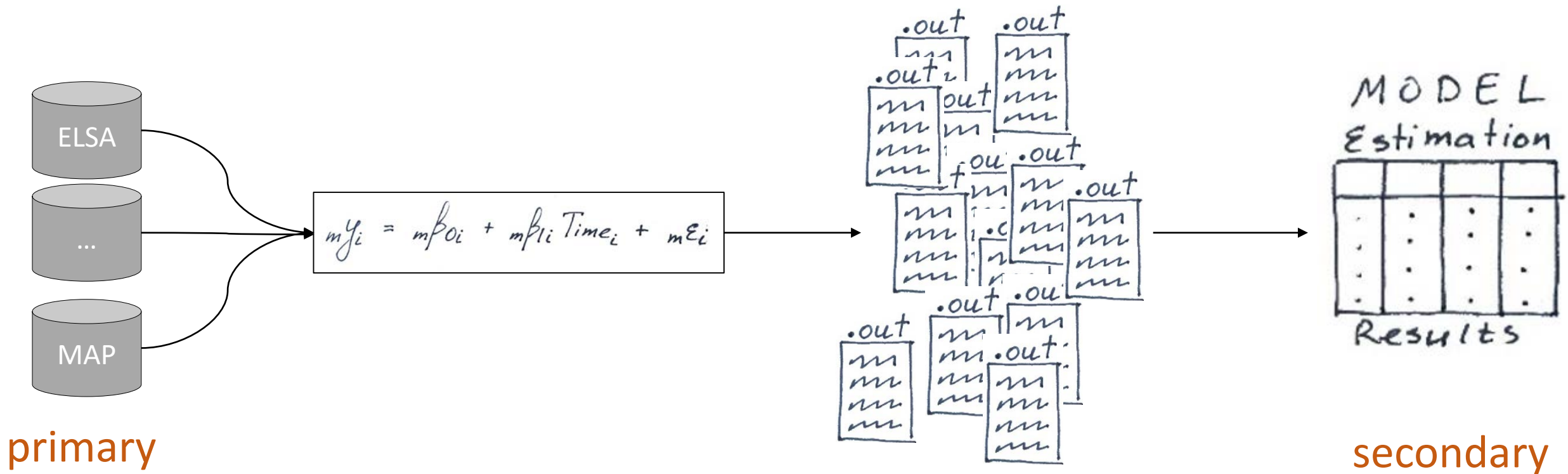
# Contents

- Preamble : context so far
- CPA poster : Gen2 scripts
- Next steps

# Original vision

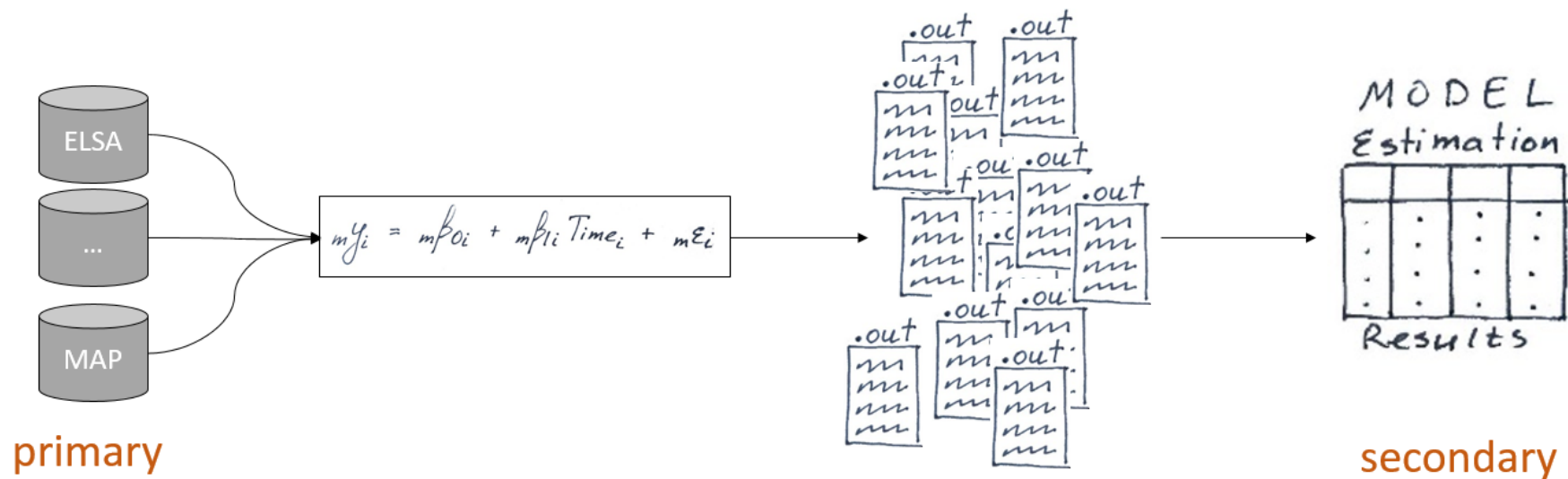
Aggregation Phase: Coordinate statistical analysis of the **primary** data

Analysis Phase: Facilitate the analysis of the **secondary** data



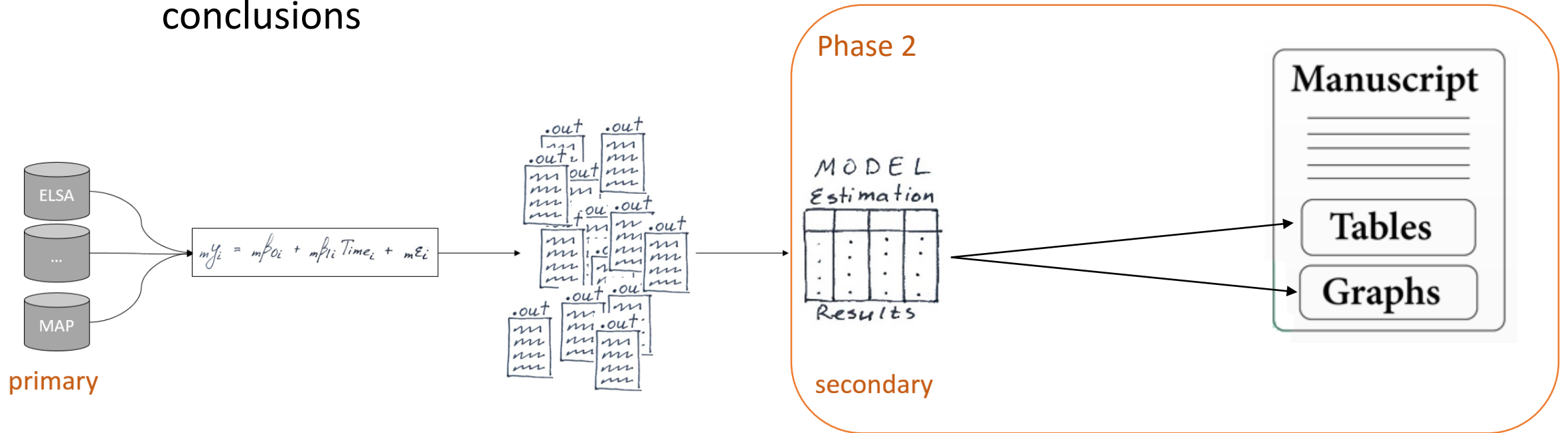
# Original vision

1. Coordinate statistical analysis of the primary data
  - A. Standardize input data
  - B. Standardize syntax for fitting models
  - C. Remove human error during extracting indices from outputs
2. Facilitate the analysis of the secondary data



# Original vision

1. Coordinate statistical analysis of the primary data
2. Facilitate the analysis of the secondary data
  - Originally we thought a simple BISR correlation will be all we need
  - But there were many unexpected specifics that threatened the validity of conclusions



# Current progress in the Analysis Phase

- Table of BISR correlation
- Table of Growth processes
- Study-specific tables for manuscript seeds

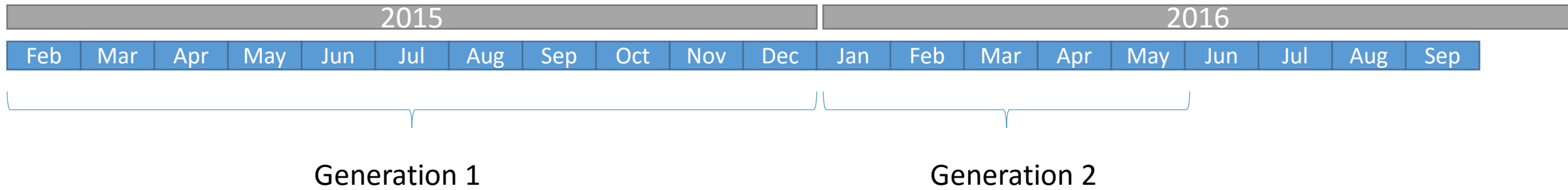
target process a : gait

	Fixed effects	animals	bnt	mmse	gait
a	intercept	18.93(2.41),**	18.93(2.41),**	18.93(2.41),**	18.93
a	slope	0.80(0.47)*	0.80(0.47)*	0.80(0.47)*	0.08
a	age	-0.15(0.16)**	-0.15(0.16)**	-0.15(0.16)**	-0.15
a	education	0.88(0.26)***	0.88(0.26)***	0.88(0.26)***	0.88
a	height	-0.01(0.11)	-0.01(0.11)	-0.01(0.11)	0
a	smoking	1.53(1.44)	1.53(1.44)	1.53(1.44)	1.15
a	cardio	-0.38(2.44)*	-0.38(2.44)*	-0.38(2.44)*	0
a	diabetes	-4.39(2.59)***	-4.39(2.59)***	-4.39(2.59)***	-4.39
a	slope*age	-0.04(0.03)**	-0.04(0.03)**	-0.04(0.03)**	-0.04
a	slope*education	-0.07(0.05)	-0.07(0.05)	-0.07(0.05)	0
a	slope*height	-0.01(0.02)	-0.01(0.02)	-0.01(0.02)	0
a	slope*smoking	-0.10(0.26)*	-0.10(0.26)*	-0.10(0.26)*	0
a	slope*cardio	-0.07(0.73)***	-0.07(0.73)***	-0.07(0.73)***	-0.07
a	slope*diabetes	0.18(0.40)	0.18(0.40)	0.18(0.40)	0.01
b	intercept	38.58(2.78)*	38.58(2.78)*	38.58(2.78)*	
b	slope	-0.78(0.72)***	-0.78(0.72)***	-0.78(0.72)***	
b	age	-0.35(0.19)***	-0.35(0.19)***	-0.35(0.19)***	
b	education	0.72(0.29)	0.72(0.29)	0.72(0.29)	
b	height	-0.10(0.14)	-0.10(0.14)	-0.10(0.14)	
b	smoking	0.78(1.34)	0.78(1.34)	0.78(1.34)	
b	cardio	1.06(3.11)	1.06(3.11)	1.06(3.11)	
b	diabetes	-5.08(2.30)**	-5.08(2.30)**	-5.08(2.30)**	
b	slope*age	-0.03(0.04)**	-0.03(0.04)**	-0.03(0.04)**	
b	slope*education	0.07(0.06)	0.07(0.06)	0.07(0.06)	
b	slope*height	0.00(0.03)	0.00(0.03)	0.00(0.03)	
b	slope*smoking	0.08(0.39)*	0.08(0.39)*	0.08(0.39)*	
b	slope*cardio	0.00(0.76)	0.00(0.76)	0.00(0.76)	
b	slope*diabetes	0.01(0.66)	0.01(0.66)	0.01(0.66)	
<b>Variance components</b>					
a	Intercept	6.13(.52)**	6.13(.52)**	6.13(.52)**	
a	Slope	.85(.12)*	.85(.12)*	.85(.12)*	
a	Residual	1.23(1.02)***	1.23(1.02)***	1.23(1.02)***	
b	Intercept	3.34(.52)***	3.34(.52)***	3.34(.52)***	
b	Slope	.66(.08)*	.66(.08)*	.66(.08)*	
b	Residual	2.08(.82)*	2.08(.82)*	2.08(.82)*	
<b>Covariance Components</b>					
	Cov(IPhys-ICog)	0.076(.051)	0.076(.051)	0.076(.051)	
	Cov(SPhys-SCog)	0.076(.051)	0.076(.051)	0.076(.051)	
	Cov(IPhys-SCog)	0.076(.051)	0.076(.051)	0.076(.051)	
	Cov(ICog-SPhys)	0.076(.051)	0.076(.051)	0.076(.051)	
	AIC	44259.46	44259.46	44259.46	
	BIC	44345.68	44345.68	44345.68	

This presentation is about phase of  
**mechanical aggregation.**

The subjective analysis phase is for another meeting

# Two generation of software scripts

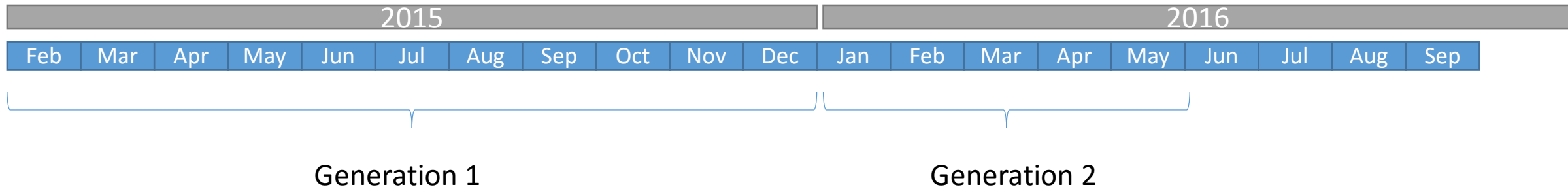


## Limitations

- Difficult to re-run models
- Humans create syntax files
- No certainty that data described are data modeled



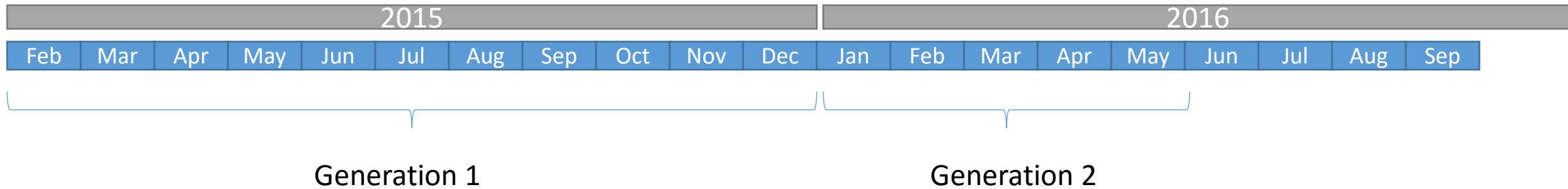
# Two generation of software scripts



Uncertainty about:

- Right subjects? (e.g. wrong subgroup filter)
- Misspecified models? ( e.g. relied on filename for model shape)
- Violated convention that were suggested to the drivers?

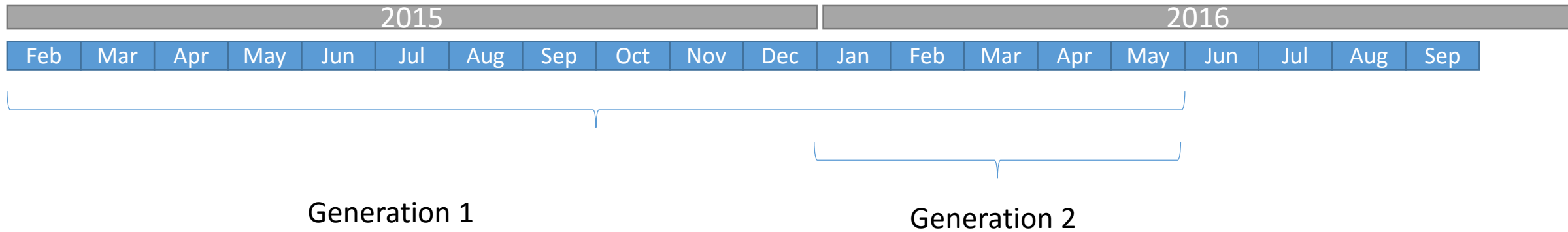
# Two generation of software scripts



Creates Threats to Statistical Conclusion Validity (Analysis Phase):

1. Many models have not estimated correlations
  - (instead, computed post hoc)
2. We cannot rely ONLY on the value of BISR correlations. Here's why:
  - A. Variance of slope
  - B. Sample size due to subgroup split
    - A. What if we drop models with insufficient sample size?  
(e.g.  $N < 100$ )
  - C. Number of included waves
  - D. Untraced human errors during estimation

# Two generation of software scripts



## Generation 1

### Limitations

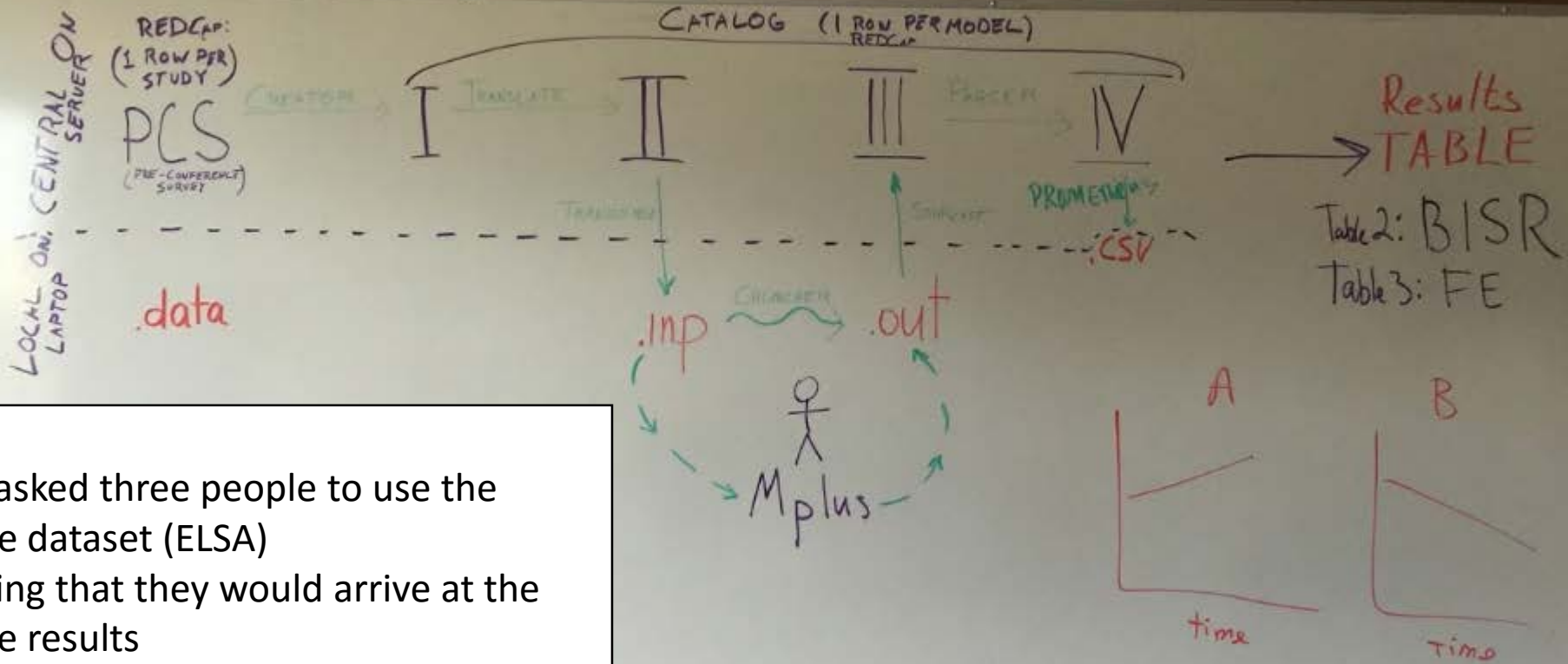
- Difficult to re-run models
- Humans create syntax files
- No certainty that data described are data modeled

## Generation 2

### Must improve

- Easy to re-run models
- Automatic generation of syntax files for each study
- Ensure the data described is the data modeled
- Greater trust in replication

On May 26, 2016 we have successfully tested aggregation phase or Gen 2 scripts using ELSA data



- We asked three people to use the same dataset (ELSA)
- Testing that they would arrive at the same results
- Focus on the software ergonomics more than the analysis

# POSTER PRESENTATION

# (Automated) Chain of Custody

- evidence must be documented, otherwise it can't be used in courtroom
- cannot be vouched for
- can be contaminated during investigation

Gen 2 has the evidence under control the entire time from the crime scene to the courtroom

- Adds transparency and reproducibility to the process
- Videotaping the entirety of the investigation
- No assurance that the knife found on the crime scene is the murder weapon.
- But solid confidence that the knife presented in the courtroom is the knife found at the crime scene

# (Automated) Chain of Custody

Start:

- Pre-Conference Survey + data
- No human intervention after that:
  - no subjective decision
  - only click-and-run
  - oversight of script execution

Without ACC we cannot be certain about:

- Right subjects? (e.g. apply subgroup filter)
- Misspecified models? (relied on filename for model shape)
- Violated convention that were suggested to the drivers?

# Bottlenecks

## 1 generation

- generating and aggregating the output ( ~12 months)
- comprehending the aggregated output ( ~ 5 months into it)

## 2 generation

- ~~generating and aggregating the output (~ 1 day)~~
- comprehending the aggregated output



# Advantages of Gen 2

- Lower cost of collaboration during coordinated analysis
- Alleviates the disheartening difficulty/length of result extraction
- If each workshop takes too long to process then you will be tempted to swing for the fences (become more aggressive to achieve results here and now, b/c you don't want to wait for another 18 months.
- greater focus on achievable goals

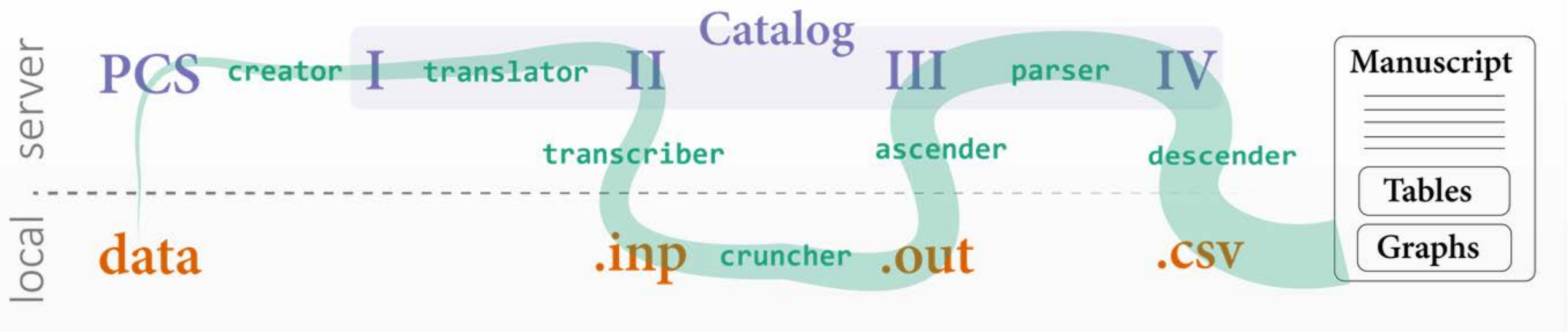
# Advantages of Gen 2

## Focus changes

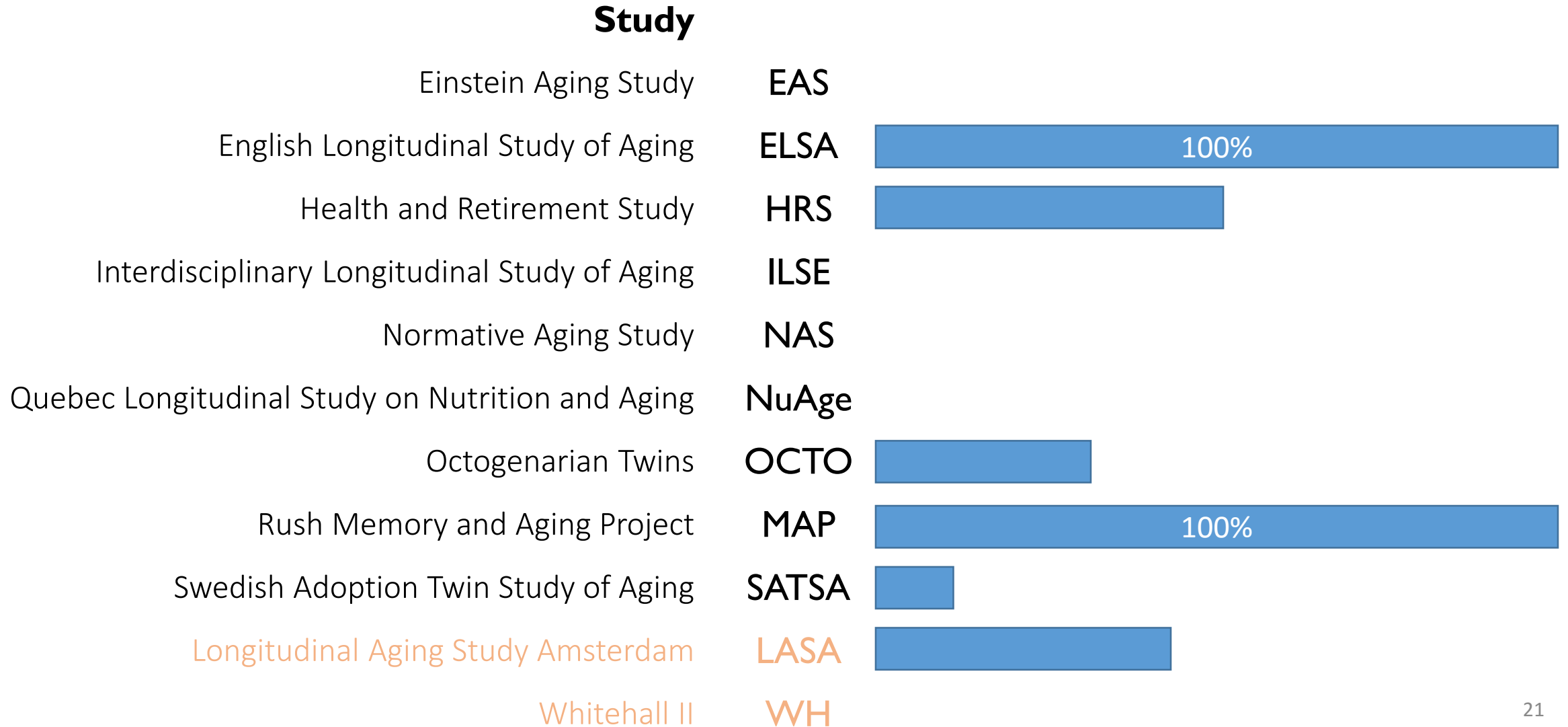
- From :How many models can we bring together?
- To: How can we organize the results?
- To : How many results can we make sense of?
  
- Will take less time
- Can do remotely
- More frequent
- More focused workshop
- Greater emphasis on Phase 2

# Future directions

1. iLifeSpan-based:
  - Groom available studies to fit Portland needs = standard for a “general grooming”
2. Same model (BISR), new workshop with new studies or/and variables
  1. Keep variables, change studies
  2. change variables, change studies
3. New statistical model
  1. Old studies
  2. New studies



# IALSA-study-curator project



# New options in Gen 2

- Unlike Gen 1 that offered only option 1, Gen 2 offers different types of workshops:
  - together ( 4 days + travel costs)
  - remote completely (regular, spaced out meetings online)
  - hybrid (muscle happens quickly, more time dedicated to the interpretation and writing), happens at the conference workshop ( ~6 hours)