

Posterior Probabilities in Meta-Analysis:

An Intuitive Approach of Dealing with Publication Bias

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Meta-analysis & publication bias

- Meta-analysis: combining the results of multiple primary studies to:
 - Estimate effect size
 - Estimate heterogeneity
- Publication bias:
 - Overestimated effect sizes
 - Over- and underestimated levels of heterogeneity

Current methods often have problems providing accurate and intuitive results

Bayesian Meta Analytic Snapshot (BMAS)

Key message

- BMAS is an intuitive method that shows which models are most likely
- BMAS shows how much evidence there is for a model
- Provides insight in the level of (un)certainty
- Corrects for publication bias
- Sensitivity analysis to assess publication bias

Overview of today's presentation

- Introduce example
- BMAS without publication bias correction
- BMAS with correction

Example dataset

McCall & Carriger (1993)

Infant habituation & recognition memory performance as predictors of later IQ.

- 31 samples
- N differs between 11 and 143
- Correlation between $r = .01$ and $r = .66$
- 26 studies are statistically significant if tested two-sided
- 28 if tested one-sided

Random effects meta-analysis

Effect size: Medium to large

$r = .407$ [.341 : .469], $p < .0001$

Heterogeneity: Small to medium amount

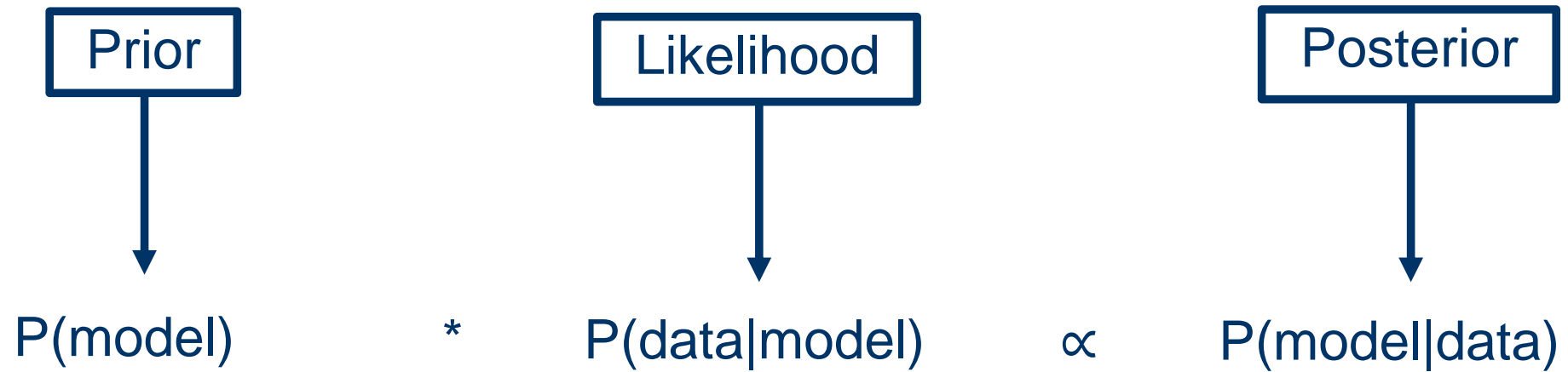
$Q(30) = 49.57$, $p = .0137$

$I^2 = 39.5\%$ [2.6% : 55.3%]

Bayesian Meta Analytic Snapshot (BMAS)

Bayesian Meta Analytic Snapshot

Bayes theorem

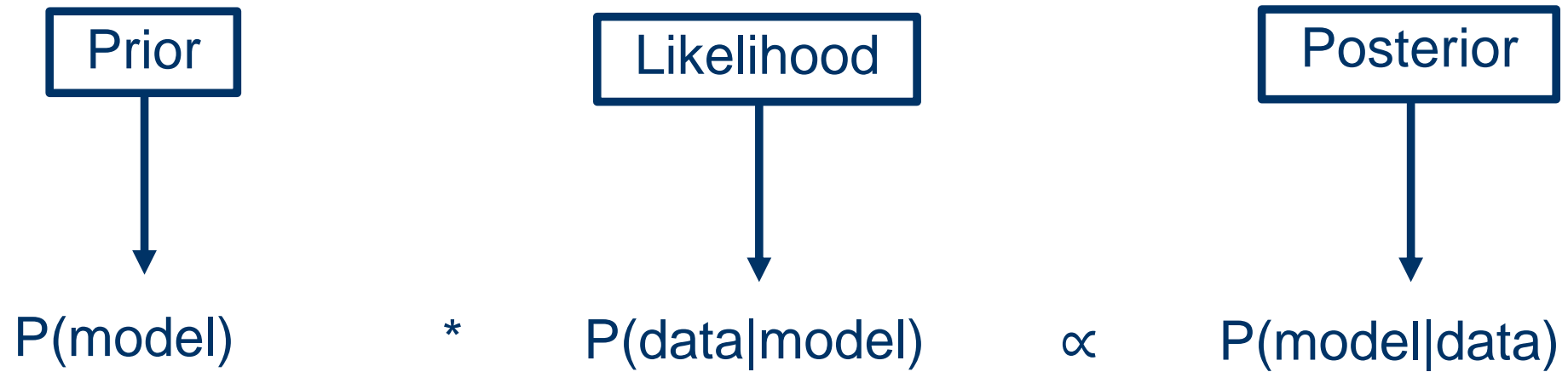


Bayesian Meta Analytic Snapshot

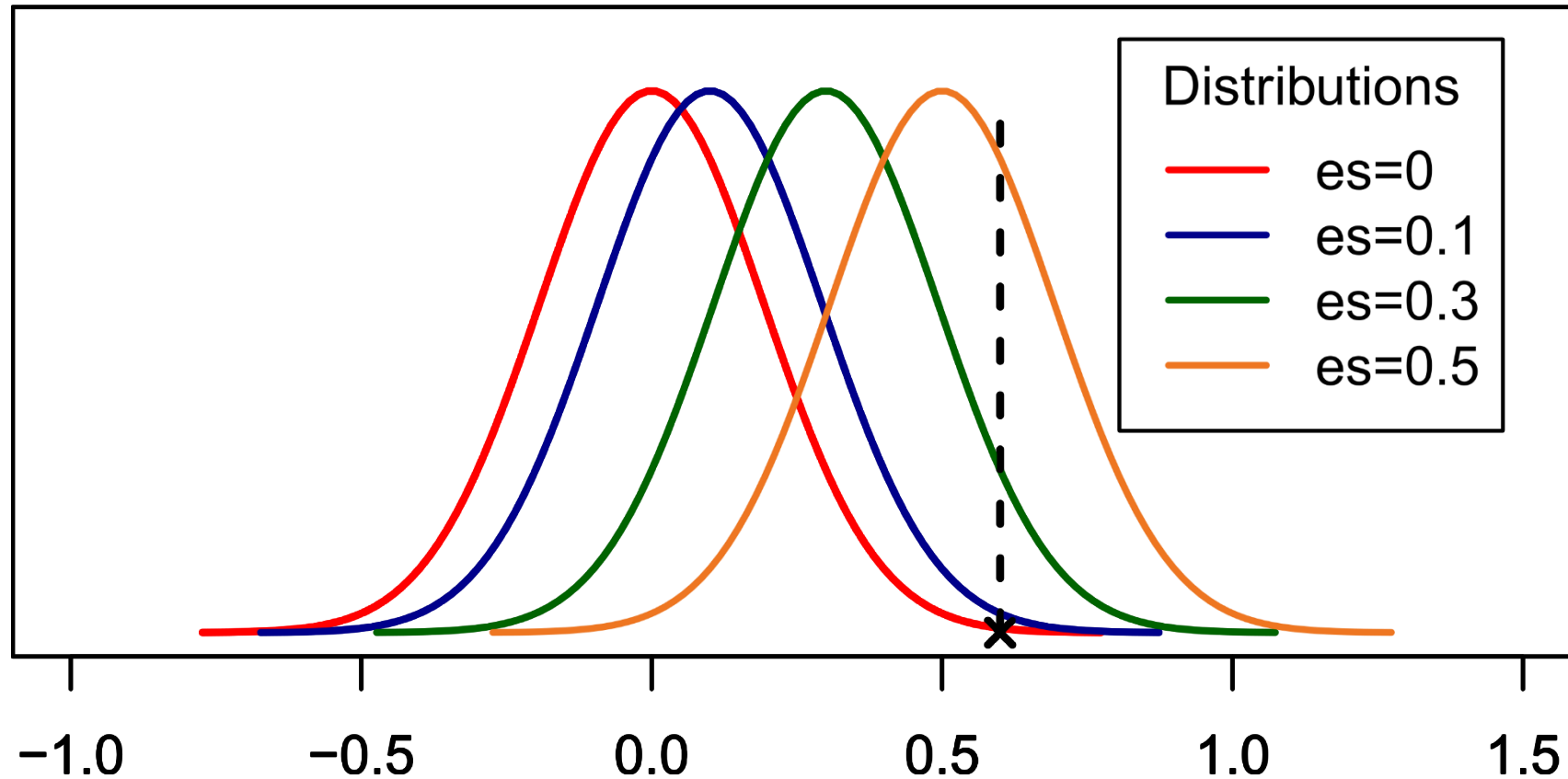
- Provide the posterior probability of 16 (4x4) models
 - Effect size (none, small, medium, large)
 - Heterogeneity (none, small, medium, large)

Bayesian Meta Analytic Snapshot

Bayes theorem

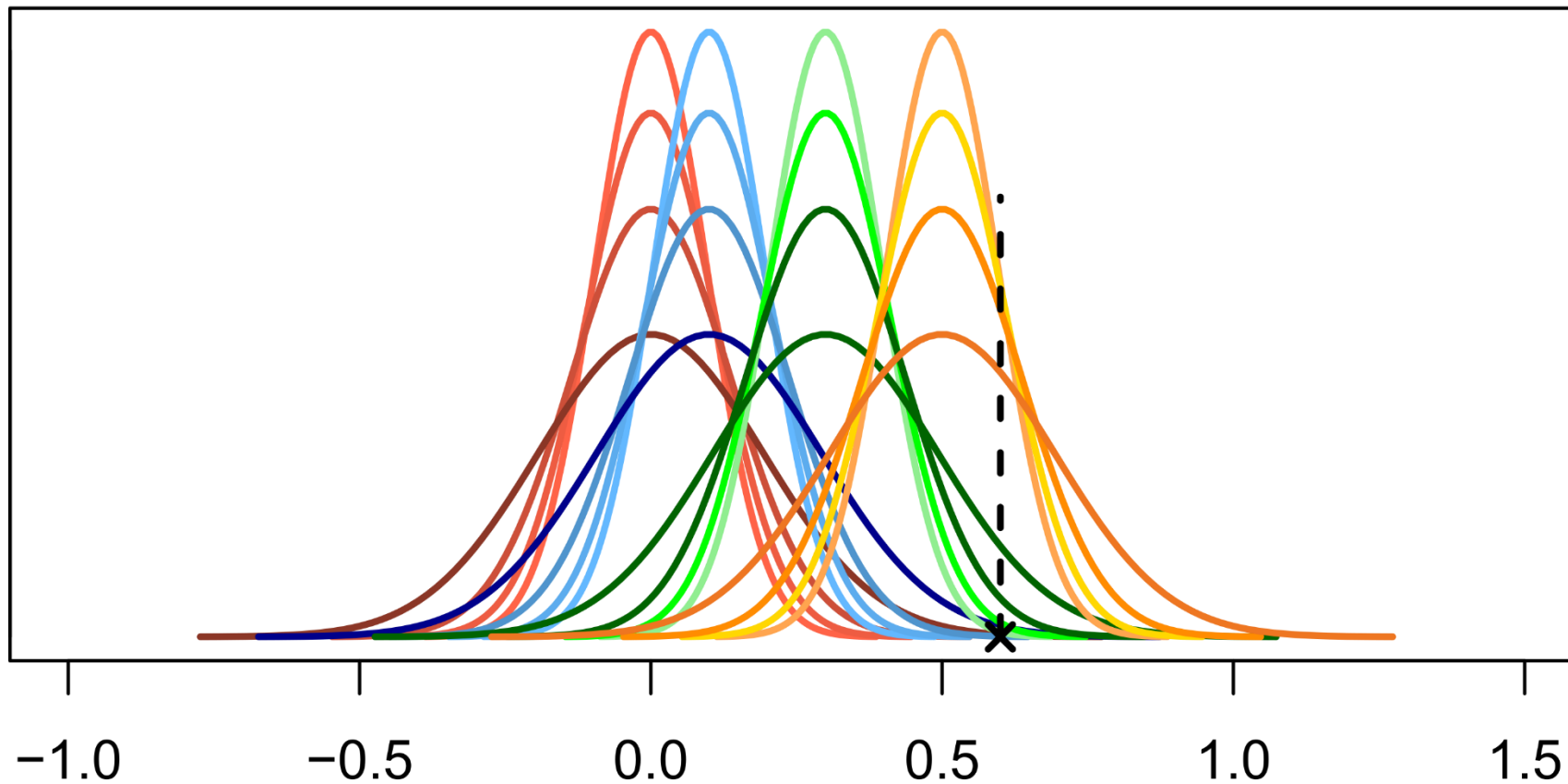


Comparison of Distributions



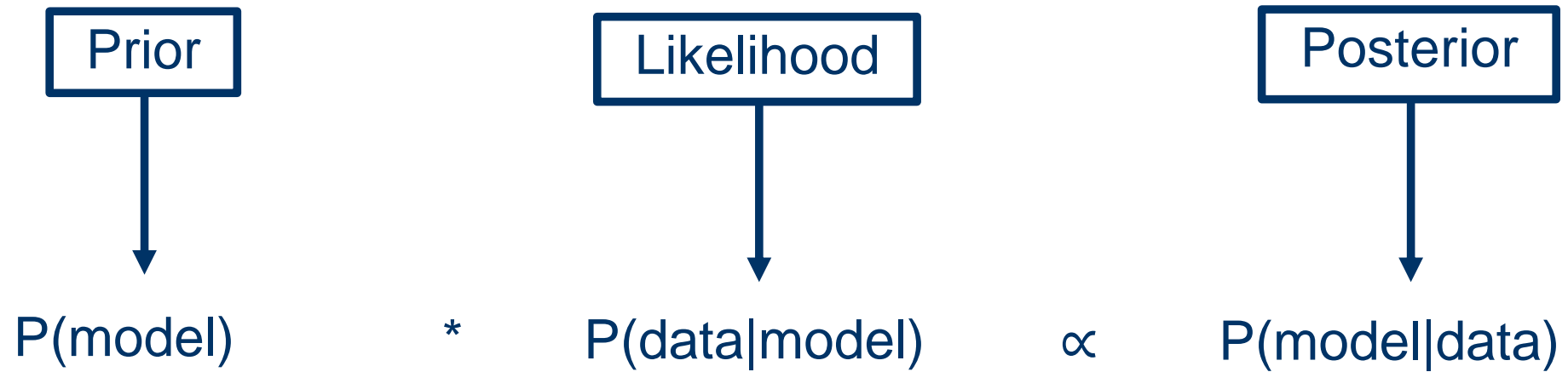
Bayesian Meta Analytic Snapshot

Comparison of Distributions



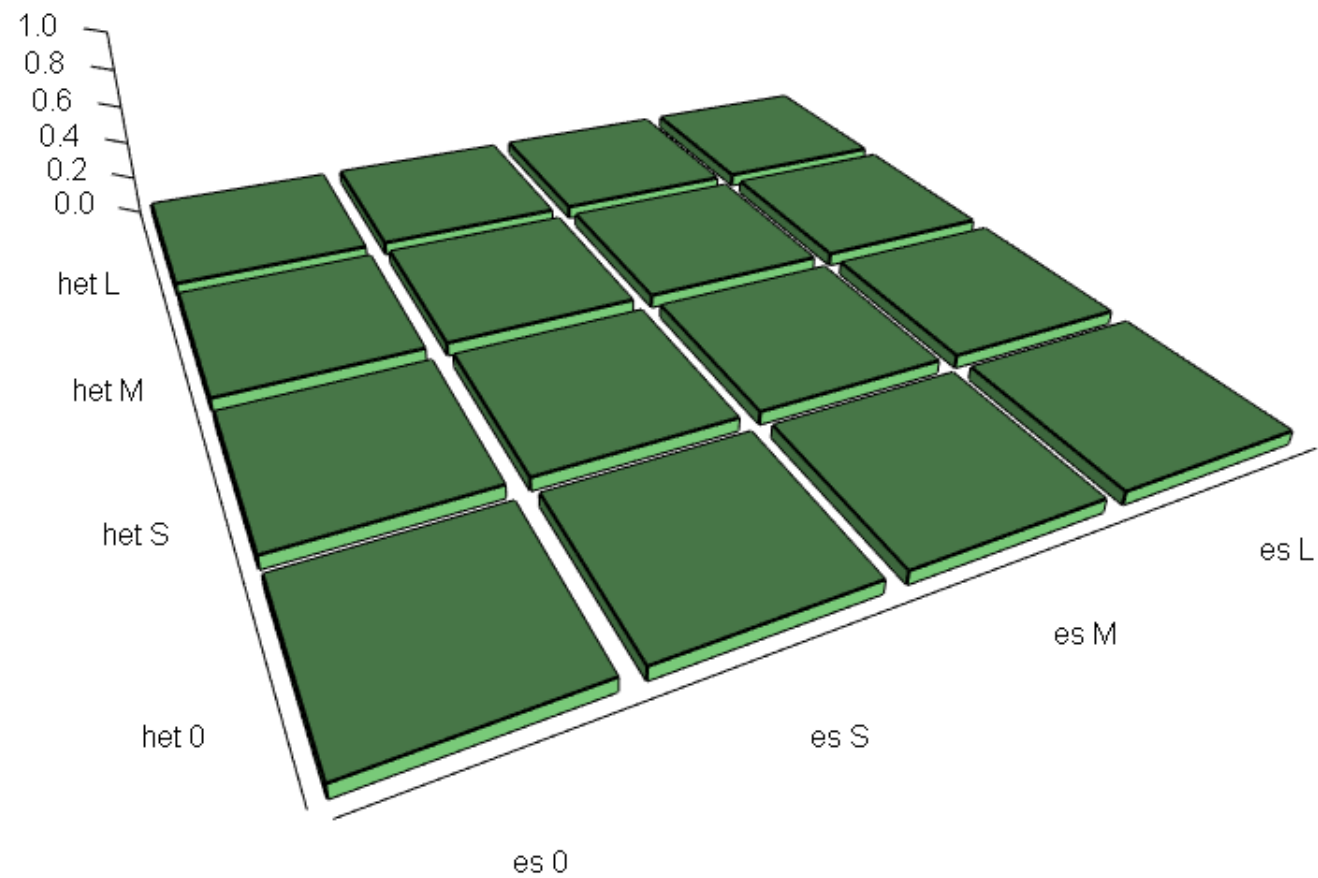
Bayesian Meta Analytic Snapshot

Bayes theorem



Start: no data

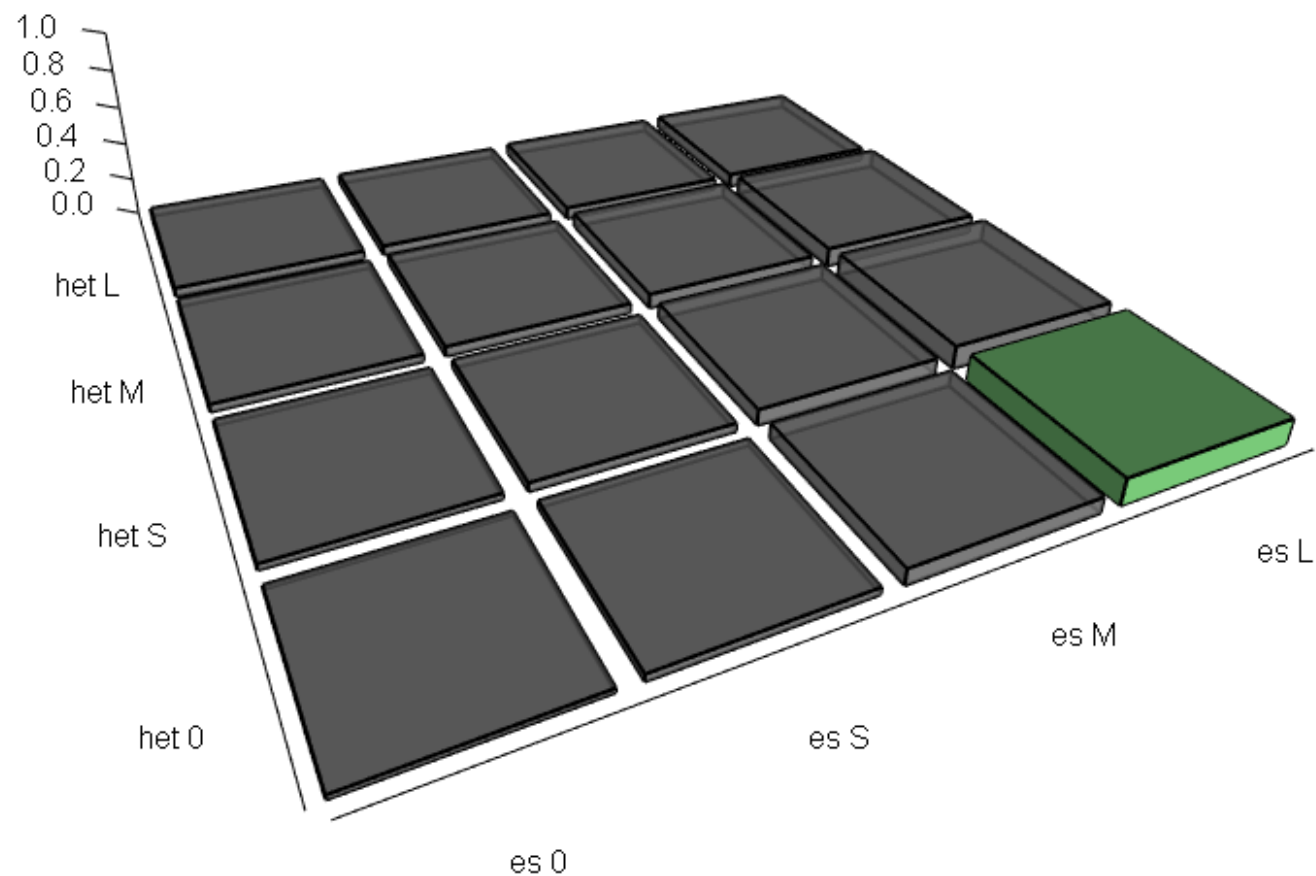
uncorrected BMAS



First datapoint:

$$r = 0.61, N = 11$$

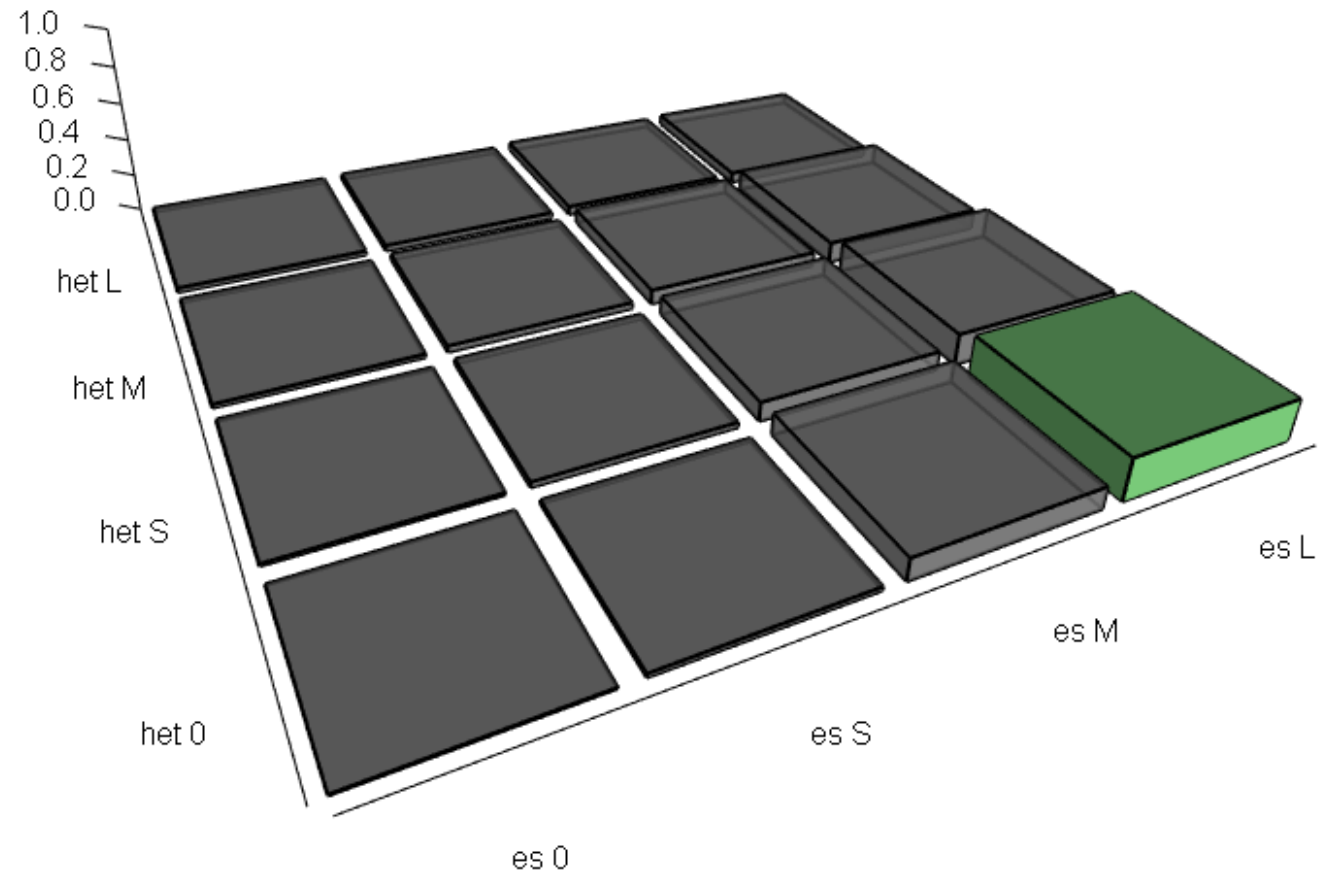
uncorrected BMAS



Second datapoint:

$$r = 0.43, N = 21$$

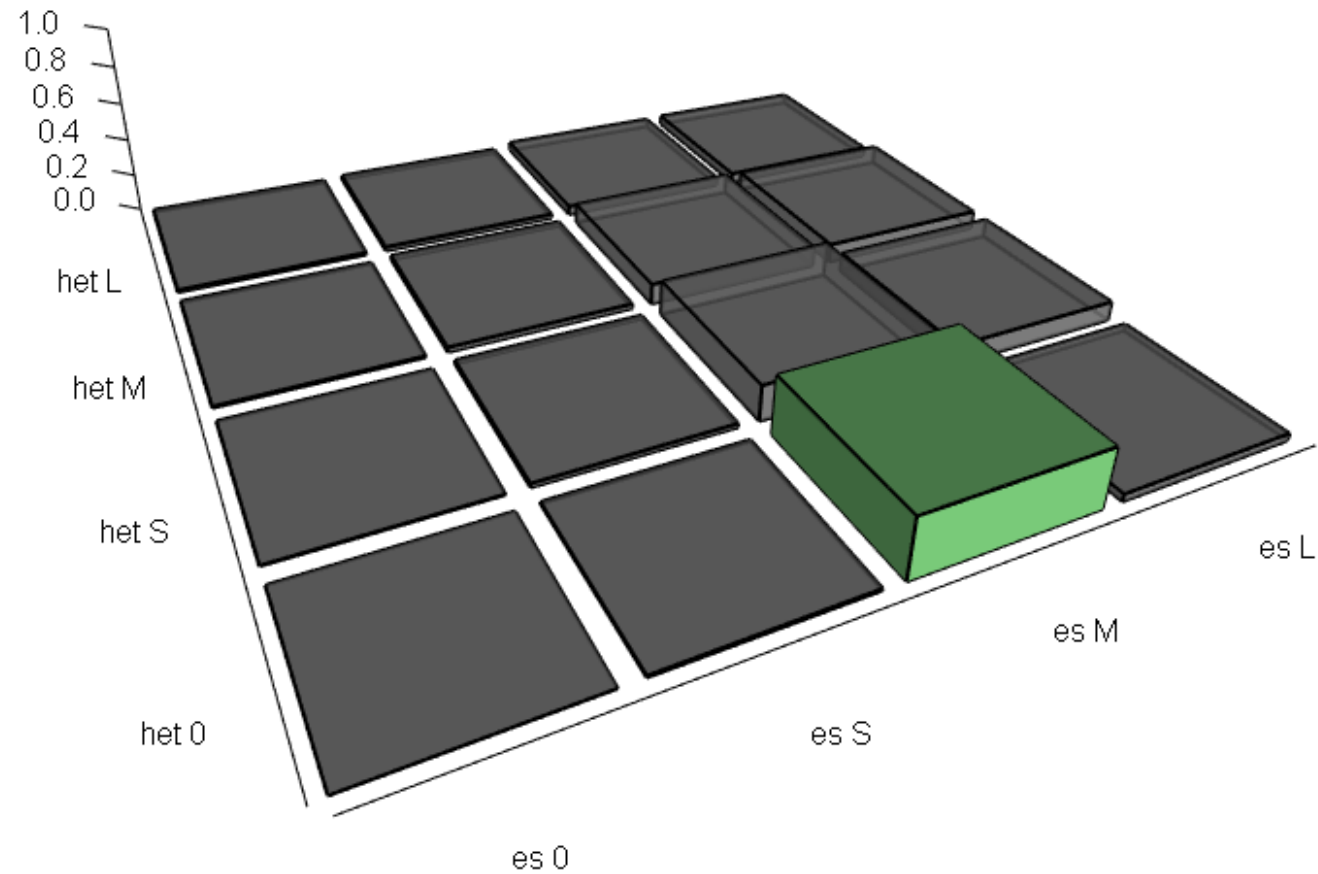
uncorrected BMAS



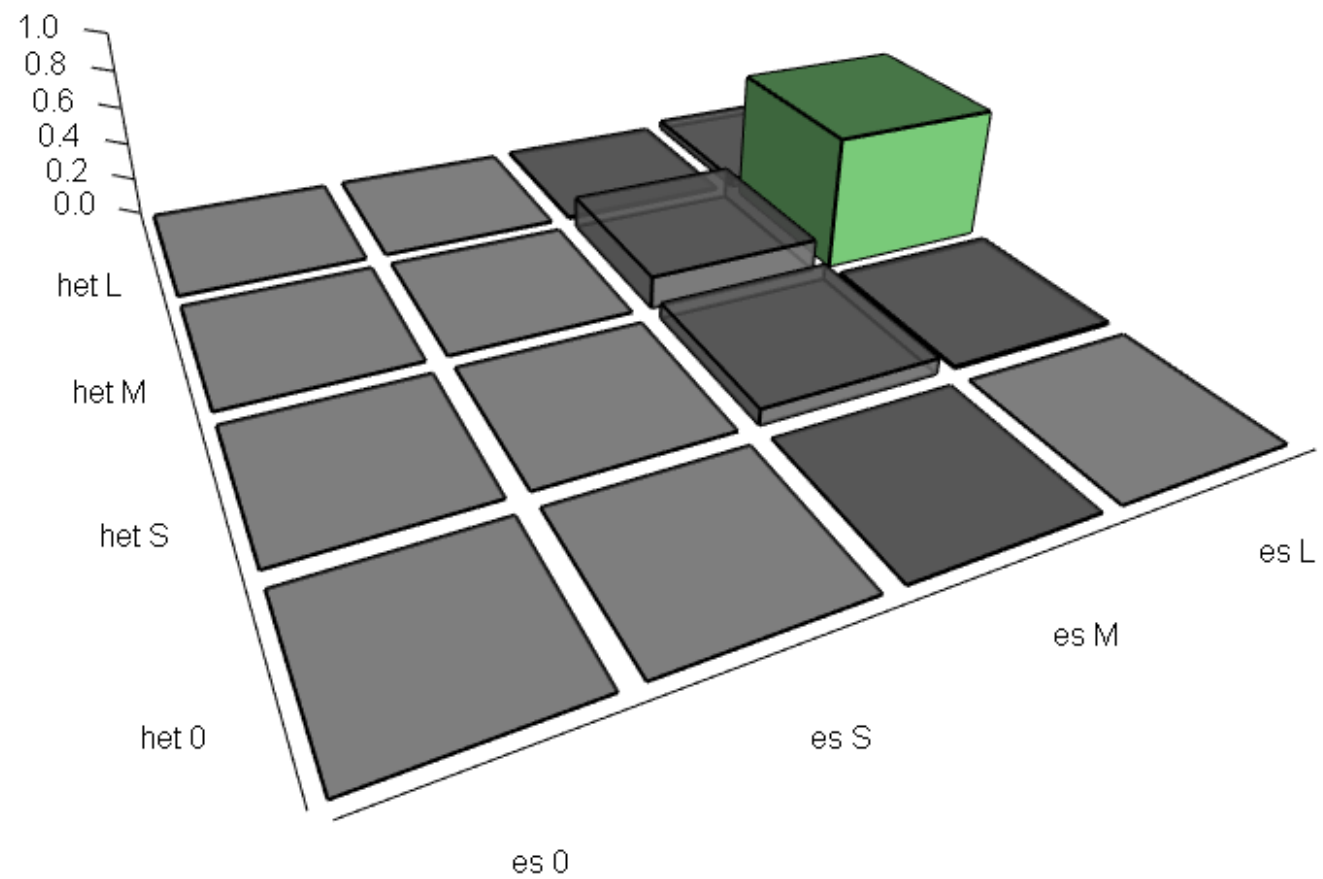
Third datapoint:

$r = 0.29$, $N = 91$

uncorrected BMAS

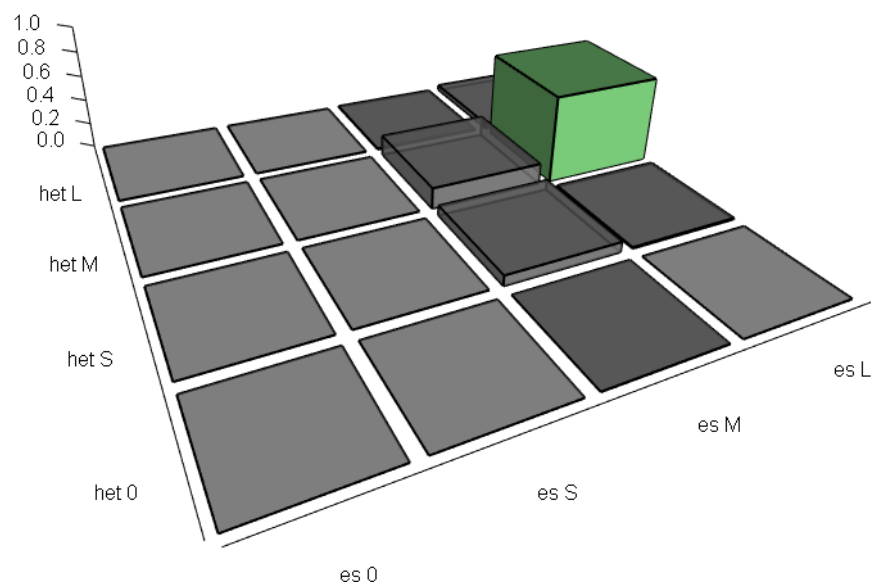


uncorrected BMAS



Bayesian Meta Analytic Snapshot: Uncorrected

uncorrected BMAS



uncorrected BMAS

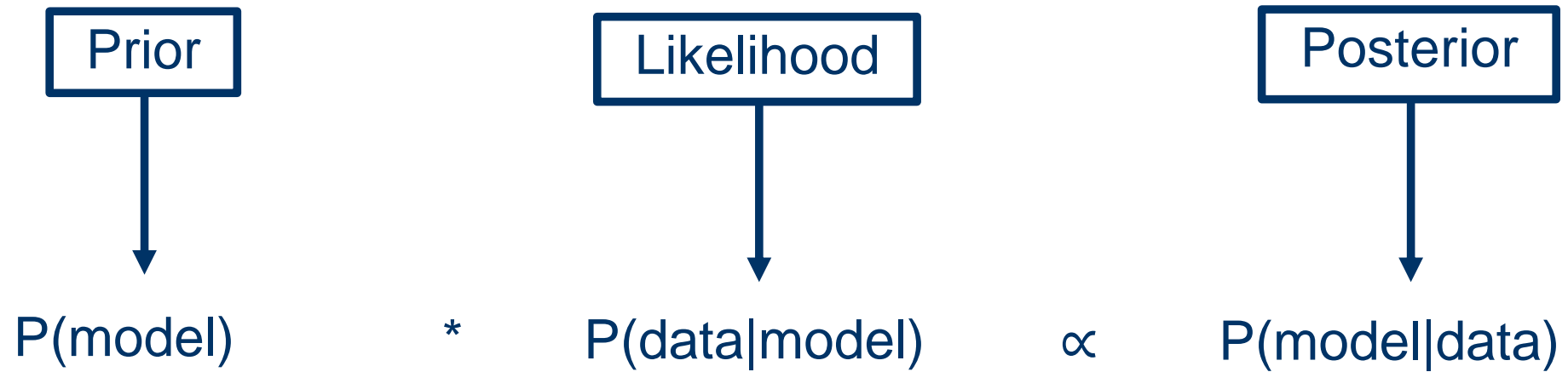
model	es0	esS	esM	esL	cummulative
het 0	0	0	0.0017	0	0.0017
het S	0	0	0.0796	0.0099	0.0894
het M	0	0	0.1662	0.6932	0.8594
het L	0	0	0.0066	0.0429	0.0494
cummulative	0	0	0.2541	0.7459	1

Publication bias

- Is there publication bias?
- Overestimated effect size?
- Incorrect estimate of heterogeneity?
- Correction needed!

Bayesian Meta Analytic Snapshot

Bayes theorem



Bayesian Meta Analytic Snapshot: Corrected

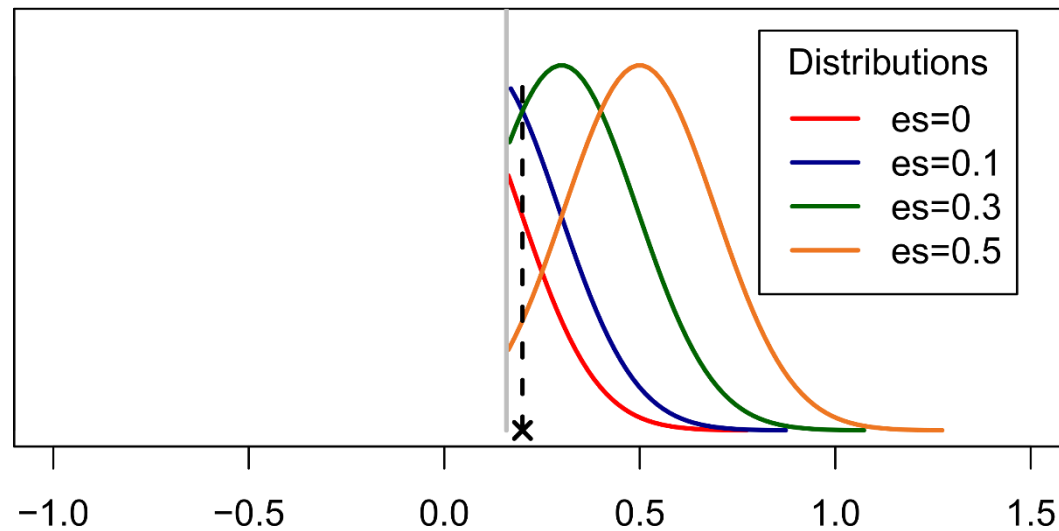
$$r = .20$$

Significant

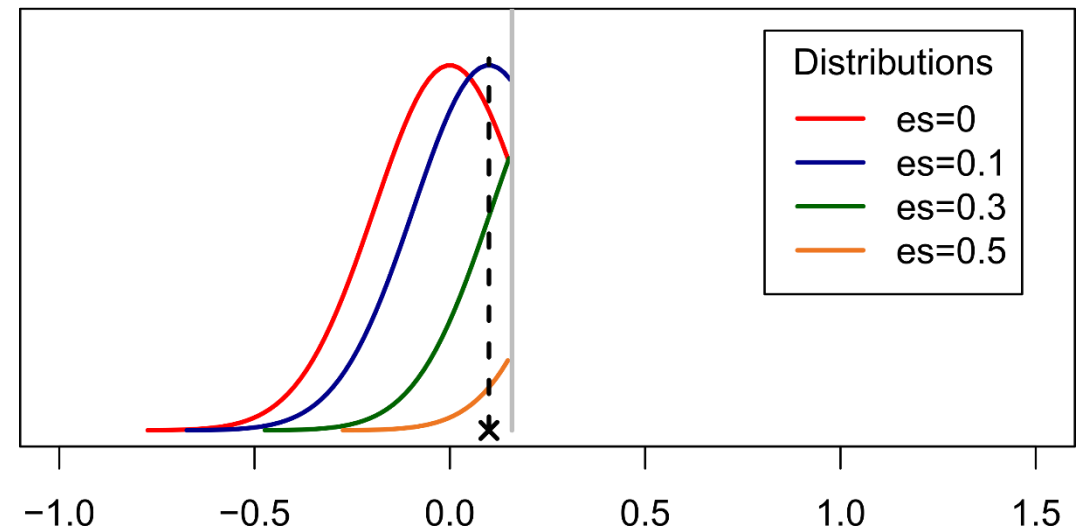
$$r = .10$$

Non-significant

Comparison of Distributions

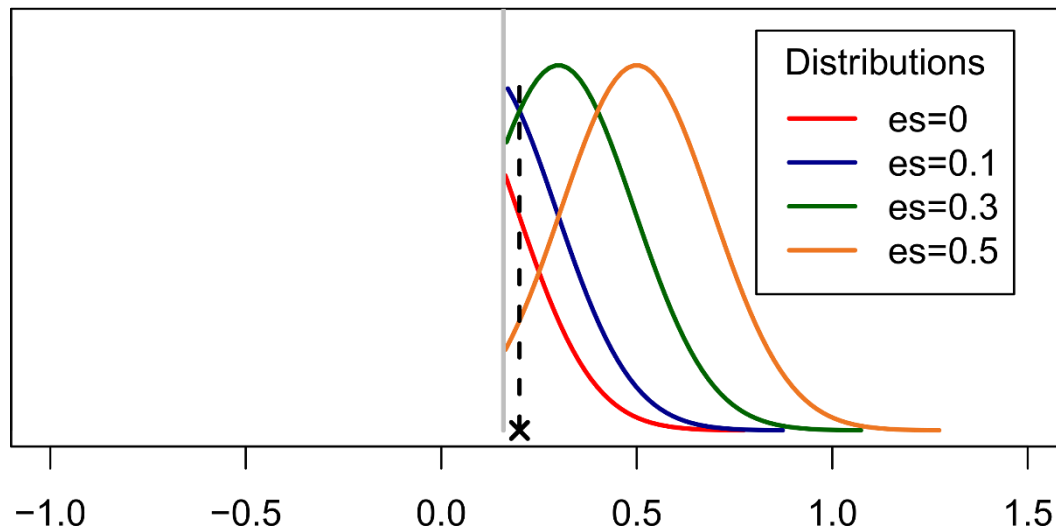


Comparison of Distributions

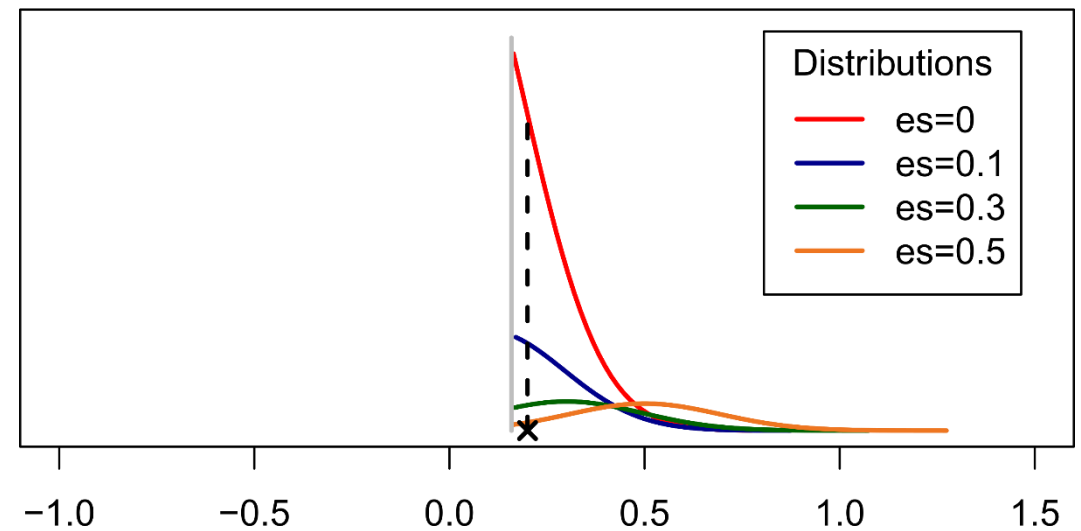


Bayesian Meta Analytic Snapshot: Corrected

Comparison of Distributions

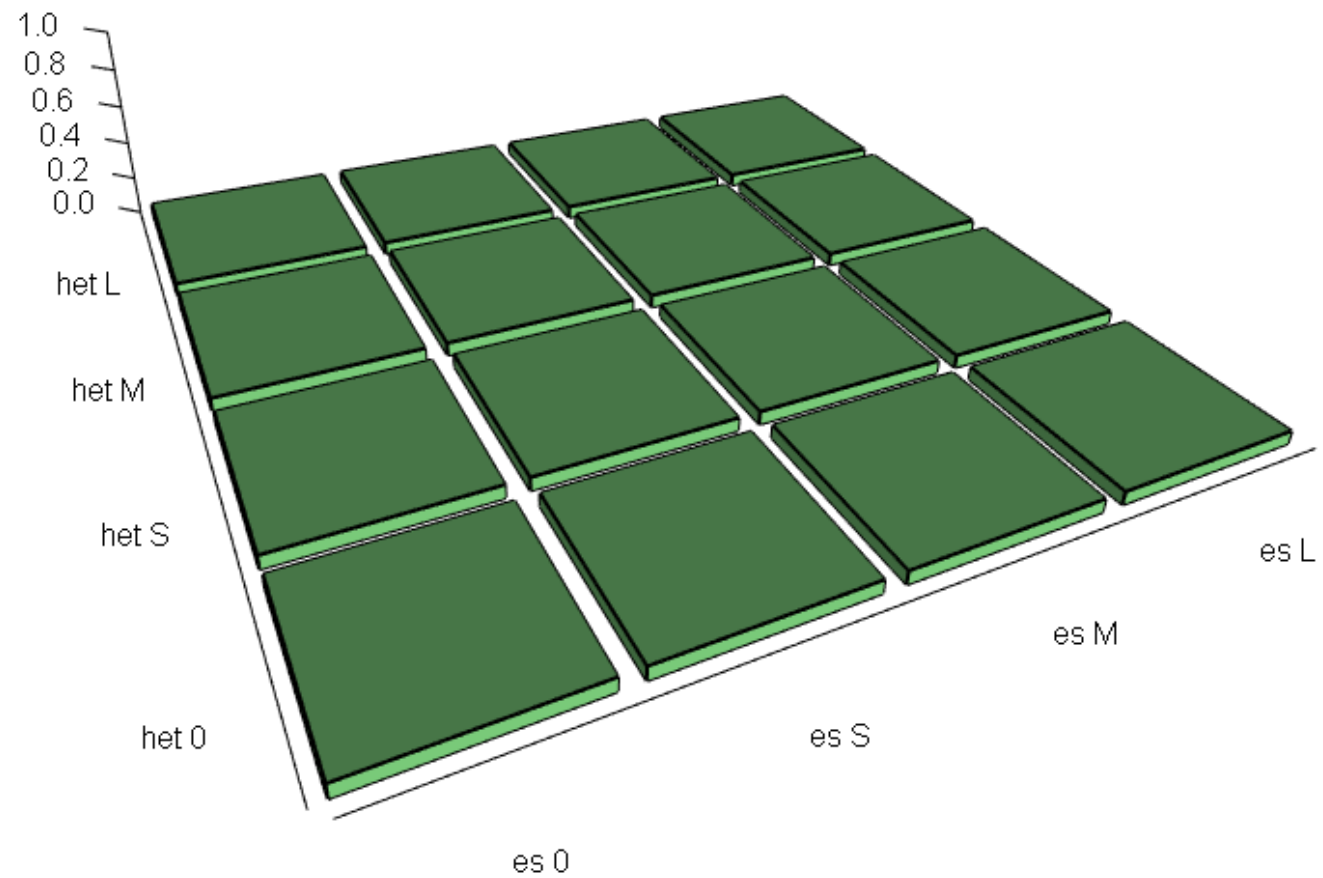


Comparison of Distributions



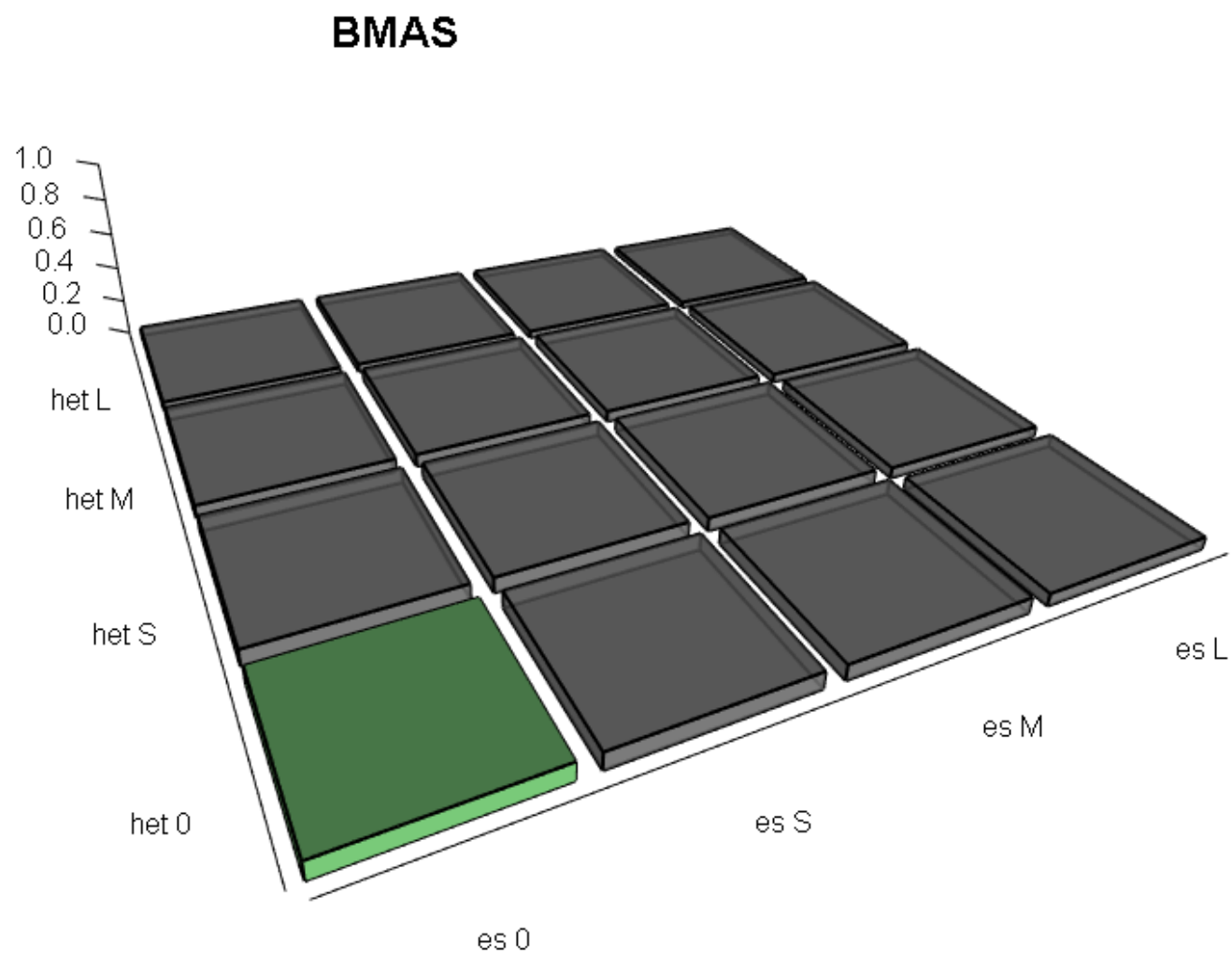
Start: no data

BMAS

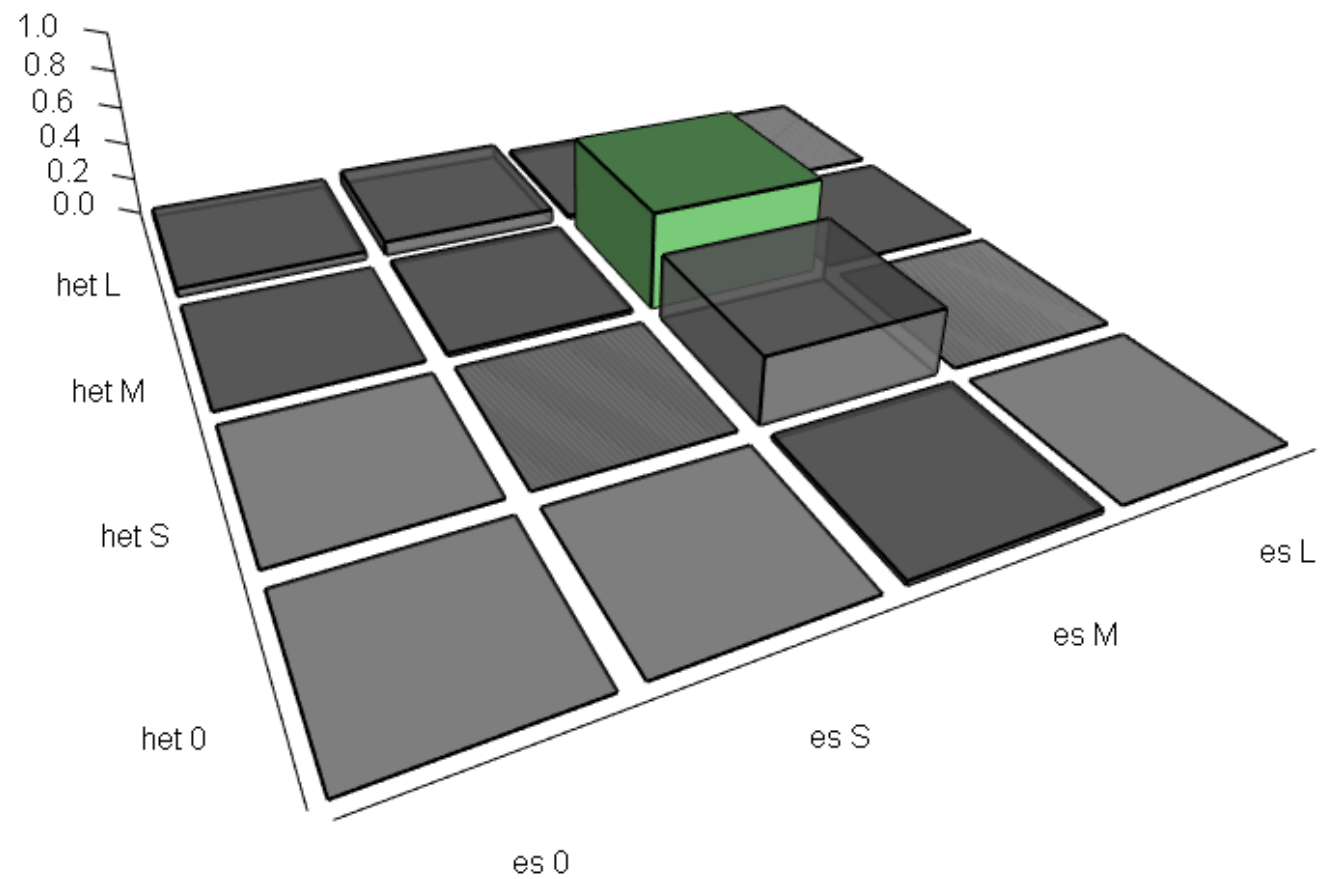


First datapoint:

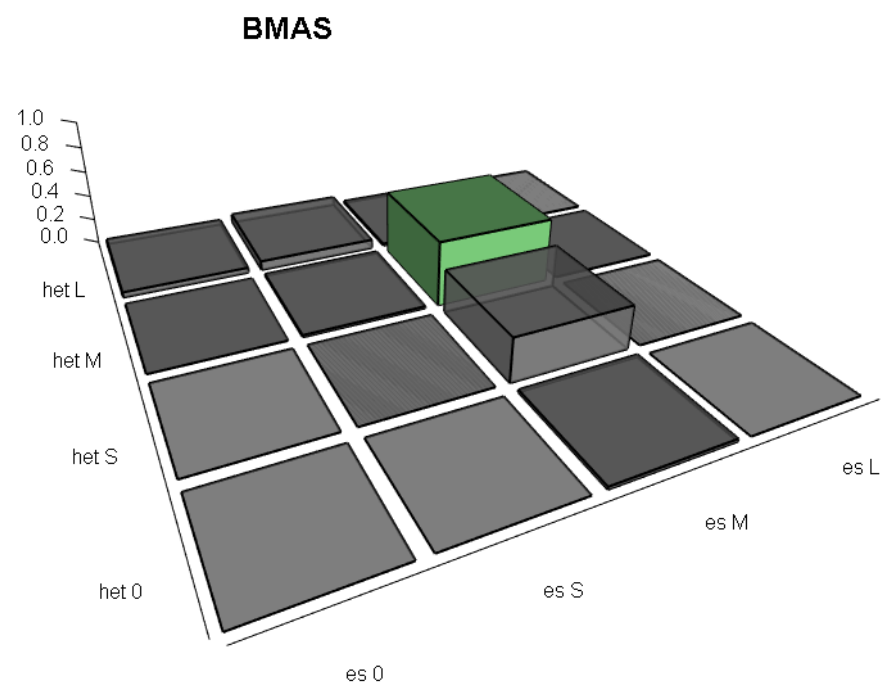
$$r = 0.61, N = 11$$



BMAS



Bayesian Meta Analytic Snapshot: Corrected



model	BMAS				cummulative
	es0	esS	esM	esL	
het 0	0	0	0.0199	0	0.0199
het S	0	0	0.3349	0	0.3349
het M	2e-04	0.0147	0.5015	1e-04	0.5166
het L	0.0401	0.0717	0.0167	0	0.1285
cummulative	0.0403	0.0864	0.8731	2e-04	1

Result overview

Random effects:

- Medium to large effect ($r = .41$)
- Small to medium amount of heterogeneity ($I^2 = 39.5\%$)

Uncorrected BMAS:

- Large effect size
- Medium amount of heterogeneity

Corrected BMAS:

- Medium effect size (large effect very unlikely)
- Small to medium amount of heterogeneity

Key message

- BMAS is an intuitive method that shows which models are most likely
- BMAS shows how much evidence there is for a model
- Provides insight in the level of (un)certainty
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Future work

- Paper on properties of BMAS
- R package / Shiny app
- Other (non-uniform) priors
- Continuous version using 16 intervals?

