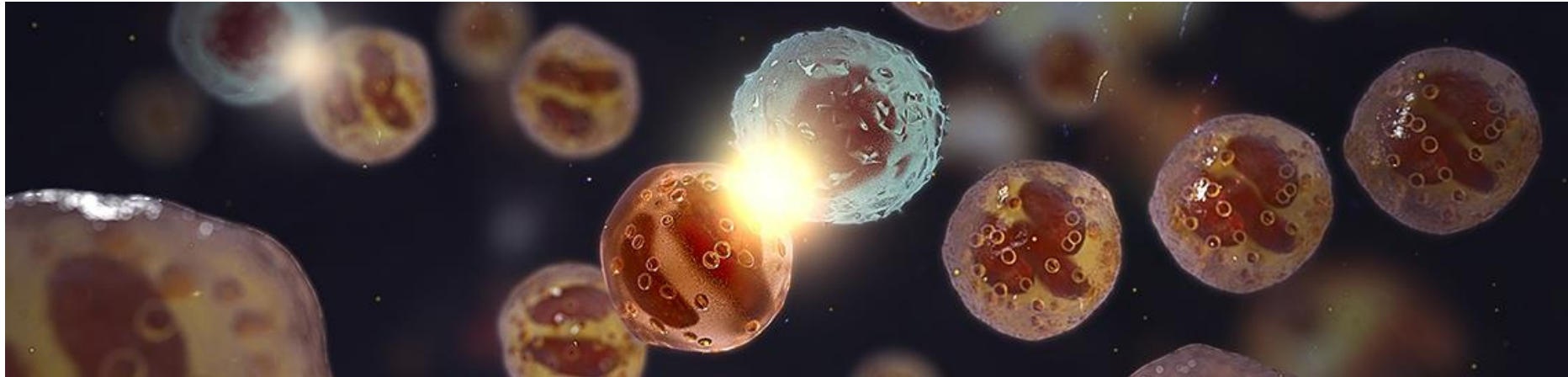


# Consistency of treatment effect across pre-specified subgroups – should we (and, if so, how?) adjust for biases?

**Speaker: David Svensson – Advanced Analytics, AstraZeneca R&D Sweden.**

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# Content & Outline

- A key step in the evaluation of any pivotal Ph III RCT is to make a risk-benefit assessment and **identify the right patient population** to treat.
- This is '**consistency of treatment effect**', a regulatory requirement (e.g., ICHE5, E9 & E17, EMA [13, 21]). A failure might lead to restricted labelling.
- Special case of interest: regions/countries in MRCTs. Overall result versus country-specific results?
- Despite the importance, no detailed regulatory guidance how to statistically demonstrate consistency. Current standard practices: **statistical issues**.
- **Alternative methods** have been proposed: some discussed here.
  - (*industry view on these?*)



## Short remarks:

- We have (at this time) **no** strong preference regarding proposed approaches.
  - Rather: are there **any good alternatives** out there?
- But focusing mostly on **shrinkage** here (as an example of a new approach).
- (Aspects will be illustrated with simulations: **exaggerated** effects for clarity).
- Note: **Overlapping** versus **non-overlapping** subgroups:
  - Mostly non-overlapping here (= *a patient belongs to one subgroup only*).
  - E.g., "country" in MRCT.
- The overlapping (general) case is harder for various reasons (more later).



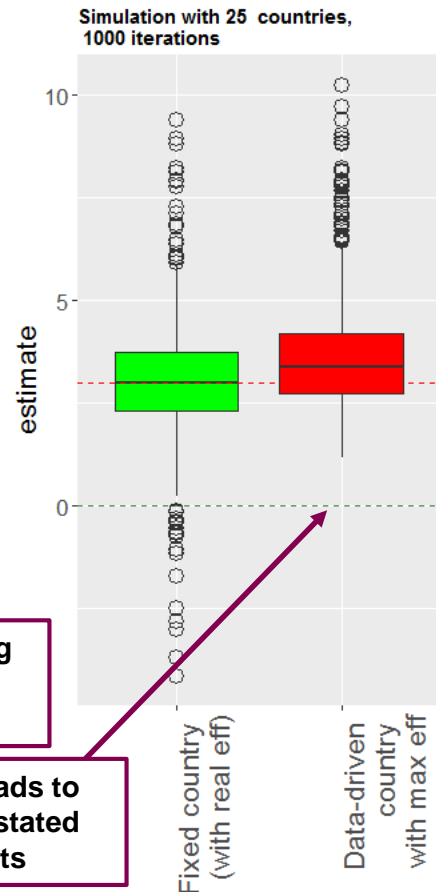
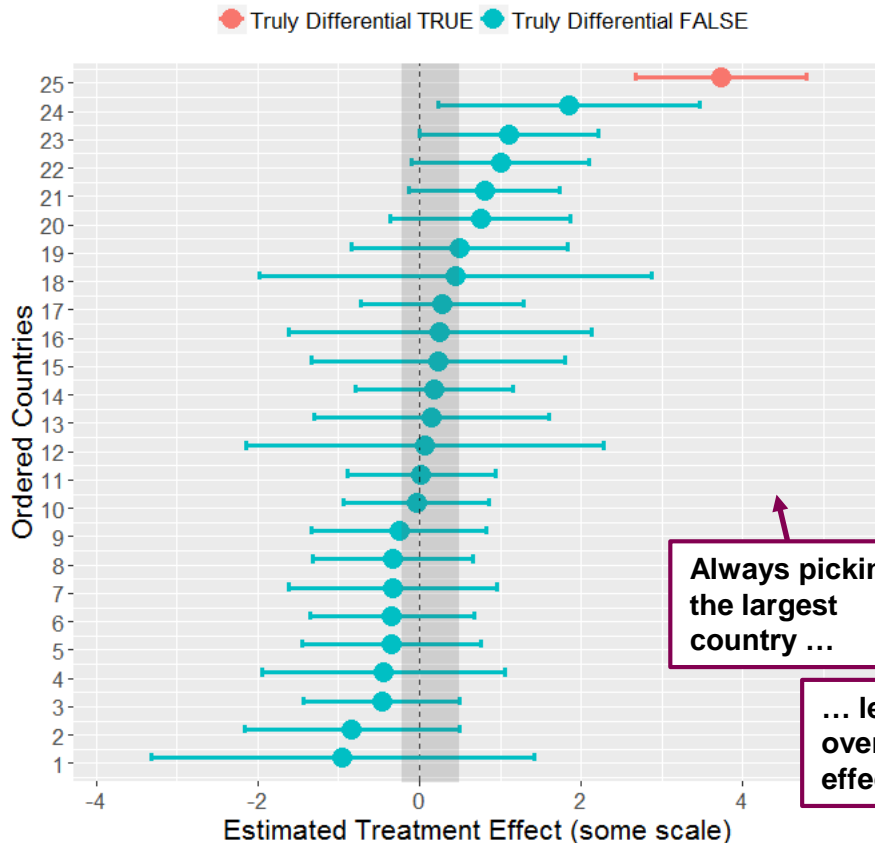
# Are the standard estimates biased?

- Not per se ..
- ... unless the reason you look at it is **data-driven**.

Any data-driven selection: bias. (e.g., [1])

True even if a country truly has larger expected value (simulation illustration).

Illustration: MRCT simulation (1 iteration) with 25 countries, All NULL, except one country with differential effect= 3



Always picking the largest country ...

... leads to overstated effects

# Current Approach to consistency: issues ...

We reckon that consistency assessment is not merely a statistical exercise, (other considerations too, complex ... )

Still, **expect more attention to extremes**; i.e., biases are inherent.

Subgroup estimates = limited data = **large variability**. (Risk of **false positives**).

**Testing** sometimes done: e.g., Global Interaction Test (GIT)?

- **Low power** (trials not sized for it).
- **Multiplicity** (If a test per subgroup factor – general subgroups).
- Not a testing problem. *Absence of evidence vs evidence of absence ...*

Well-known issues [2], [5], [29], and alternatives have been proposed ...



# Any more holistic approach available?

- E.g., **EMA guideline** [13]: useful to assess subgroups together (display full range) although not many details given. Recommendation:
  1. Aim to **pre-group subgroups** into categories (scientific credibility reasons),
    - (A) differential effect plausible, or (B) no differential effect expected.
  2. Use **graphical** methods (e.g., Forest plots).
- **EFSPI Subgroup WG** – White Paper [6]; compared various methods (e.g., simulations); e.g., SEAMOS [8], SIDES [24], Bootstrap Bias Reduction [33], GIT. Recommendation: as EMA, plus:
  - Consider **expected-worst under NULL** [8], to provide a context of the results.
- **Jury still out** on some other methods? (E.g., Bayesian shrinkage?)



# Bayesian Shrinkage has been suggested, e.g.,:

- *Empirical shrinkage estimator for consistency assessment of treatment effects in multi-regional clinical trials.* [30].
- *Multi-regional clinical trial design and consistency assessment of treatment effects.* [31].
- *Treatment effect heterogeneity for univariate subgroup in clinical trials: Shrinkage, standardization, or else.* [35]
- *Exploratory Subgroup Analyses in Clinical Trials.* [32]
- *Bayesian models for subgroup analysis in clinical trials.* [23]
- *Comparing Approaches to Treatment Effects Estimation for Subgroups In Clinical Trials.* [26].
- *Bayesian Assessment of the Influence and Interaction Conditions in Multipopulation Tailoring Clinical Trials.* [27]
- *A Bayesian Approach to Evaluating Regional Treatment Effect in a Multiregional Trial.* [3]





# The idea of Shrinkage

- Well-known **classical** concept, e.g., [11, 12, 26, 36].
  - Stein's classical 'shocking' theorem:
    - *ML estimates can be (dramatically) improved (sometimes).* [11]
- **Widely used** for high-dim data, e.g.,
  - Microarray screening [11], Pharmacovigilance FDA Signal Detection [19].
- **The Core Idea:** seems to be:
  - *True effects less spread out than observed ones...*
  - *Some shared information across subgroups.*
  - *Self-tuning smoothing towards overall estimate...*
- Borrowing information might help with sparse subgroups (e.g., few events)?



# Shrinkage, basic model

$i = \text{countries: standard est. } \hat{\delta}_i \sim N(\delta_i, s_i^2)$

**Prior** for (true) country effects:

$$\delta_i \sim N(\delta, \tau^2)$$

Gives posterior 'estimate' (MAP):

$$\tilde{\delta}_i = w_i \cdot \hat{\delta}_i + (1 - w_i) \cdot \tilde{\delta}$$

with  $\tilde{\delta}$ =overall (RE) estimate, and weights:  $w = \tau^2 / (\tau^2 + s_i^2)$

**Model fit?** MCMC (Full Bayesian [16]), or **Empirical Bayes** (ML fit of prior to data [26, 35]).

Illustration: MRCT simulation with 25 countries, no true differential effects, Forest plot ordered after estimates. Shrunken estimates overlaid (Empirical Bayes)



# Drivers of amount of Shrinkage? (How much modified):

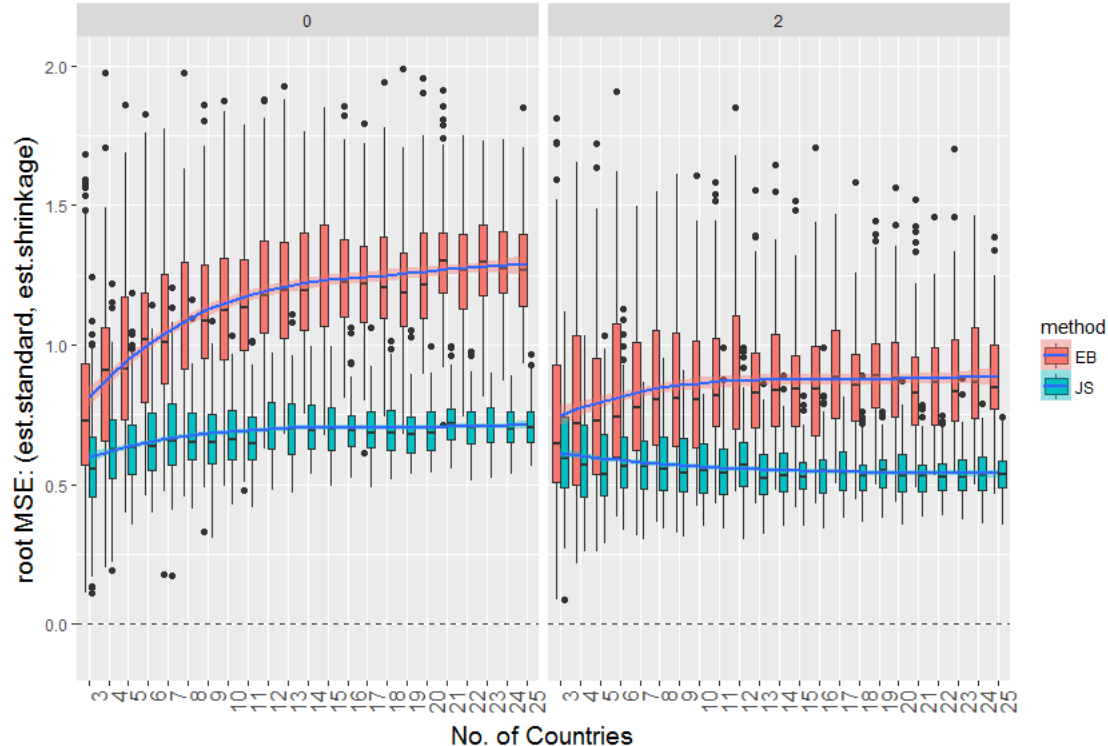
- Between  $\hat{\tau}$  vs within  $s_i^2$

$$s_i^2 \approx 0 \Rightarrow \tilde{\delta}_i \approx \hat{\delta}_i$$

$$s_i^2 \text{ large} \Rightarrow \tilde{\delta}_i \approx \tilde{\delta}$$

- **No. of countries**  
(=x-axis in graph)
- Shrinkage **method** choice  
(several exist –more later).

AVERAGE DISTANCE between Standard Estimates and Shrunken Estimates, as a function of no. of countries. Simulation under two different TAU values: 0 and 2 (heterogeneity)  
100 iterations of MRCT with x countries, NULL effects. Endpoint sd.dev=20. Country sizes: 100 to 1000.



# Some remarks on Emp Bayes estimates

- Remark 1: They are **unbiased** (despite shrinkage towards overall – any confusion?). [under the correct model]
- Remark 2: They also do **bias-adjustments** of 'random-highs':
  - The max.EB (and min) is still biased, but less so.
- Remark 3: Modified ('Improved' or 'manipulated'?) estimates: **controversial?**
  - Assumes 'exchangeability' (  $\delta_i \sim N(\delta, \tau^2)$ : unrealistic?).
  - (Note: an assumption re. the *unobservable, true* country effects).

Real differential effects will be shrunken - but recall, noise shrunken too.

– Does it make it **easier or harder** to detect truly differential subgroups?

Question: *shrinkage useful as a kind of secondary 'sensitivity analysis'?*

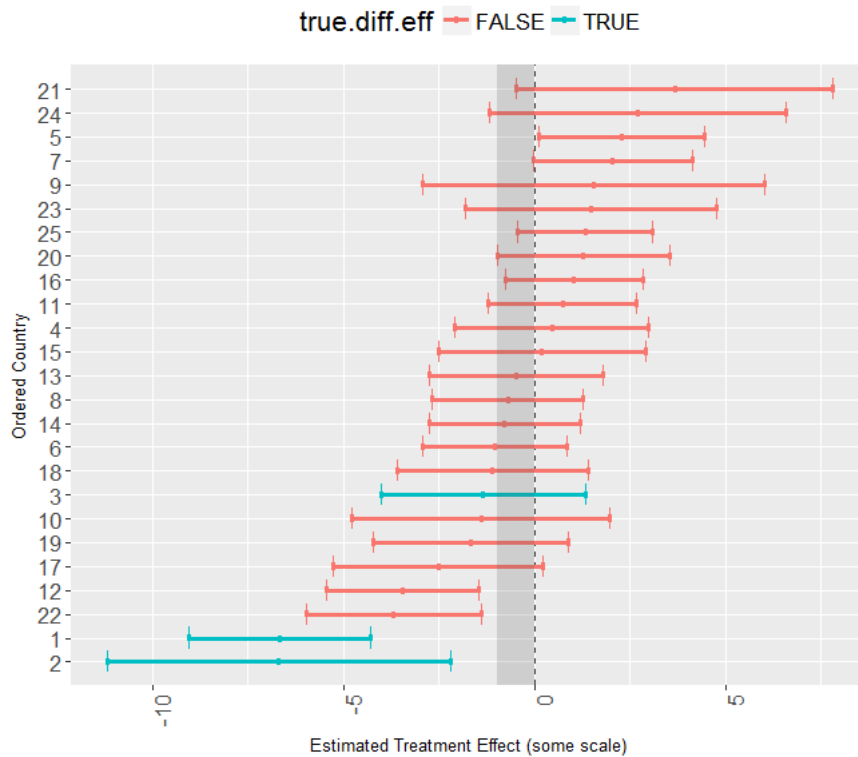


# Simulation illustration, under broken assumption:

Illustration: MRCT simulation with 25 countries, many NULL countries, but three with true differential effect (expected value) equal to -3 Classical Estimates, overall CI (shaded)

## What if exchangeability is violated?!

- Iterations of simulation of
  - MRCT with 25 countries.
  - 22 with **NULL** effects,  $\delta_i=0$ .
  - 3 **DIFF** countries with  $\delta_i=x$ . (-3 in graph)
- **QUESTION:** our ability to detect this, with/without shrinkage?
- Tracking MSE & BIAS & Interval Coverage.
- (Also overlaps: country-specific intervals relative to the overall interval).



## Some results, broken assumption (highlights only):

(Full results in Back-Up section)

Considerably **lower MSE for** Emp.Bayes, overall seen.

– (but not uniformly so)..

EmpBay **estimates biased:**

– differential countries were **under-estimated**.

– The null countries (zero effect) were **slightly over-estimated**.

But what does it mean, in terms of ability to detect differential countries?



# CI overlaps? (under broken assumption).

What if **CI overlap** was the key aspect? (Discovery Rates).

- **QUESTION:** ability to detect differential countries via 'non-overlaps', BEFORE/AFTER shrinkage?

1.  $\hat{\delta}_j^{EB} \rightarrow$  center (estimates moving).
2.  $\|CI_{ac}^{EB}\| \geq \|CI_{ac}\|$  (width, allcomer).
3.  $\|CI_j^{EB}\| \leq \|CI_j\|$  (width, countries j).

How will it play out? (1,2 vs 3)

Illustration: MRCT simulation with 25 countries many NULL countries, some with true differential effect equal to -2  
Classical Estimates, overall CI (shaded)

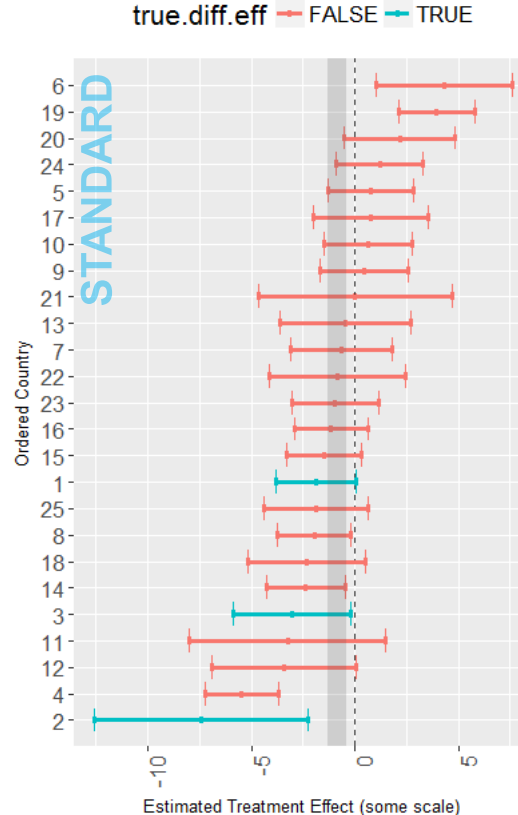
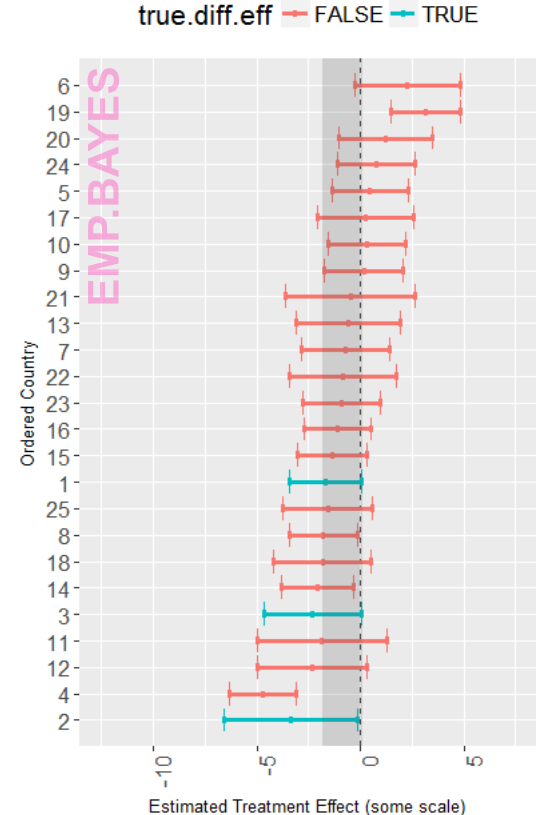


Illustration: MRCT simulation with 25 countries many NULL countries, some with true differential effect equal to -2 EmpBayes Estimates and overall CI from R.E. meta (shaded)

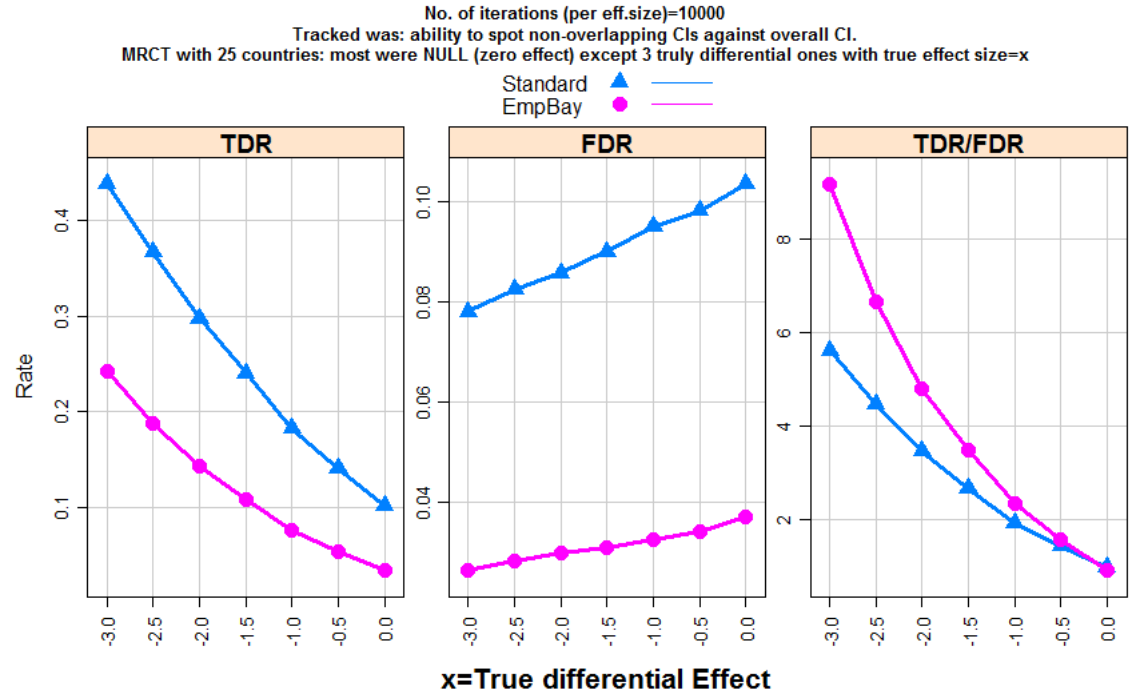


# Easier or harder: preliminary results:

Many iterations,  
each set with true  
differential effect  $\delta_i = x$ .

Tracked (for both methods):

- **True Discovery Rate:**
  - (true countries detected?)
- **False Discovery Rate:**
  - (NULL countries detected?)



Conclusion:

Ratio better "after shrinkage", but driven by lower FDR.





# Shrinkage: not just Emp.Bayes; there is more ...

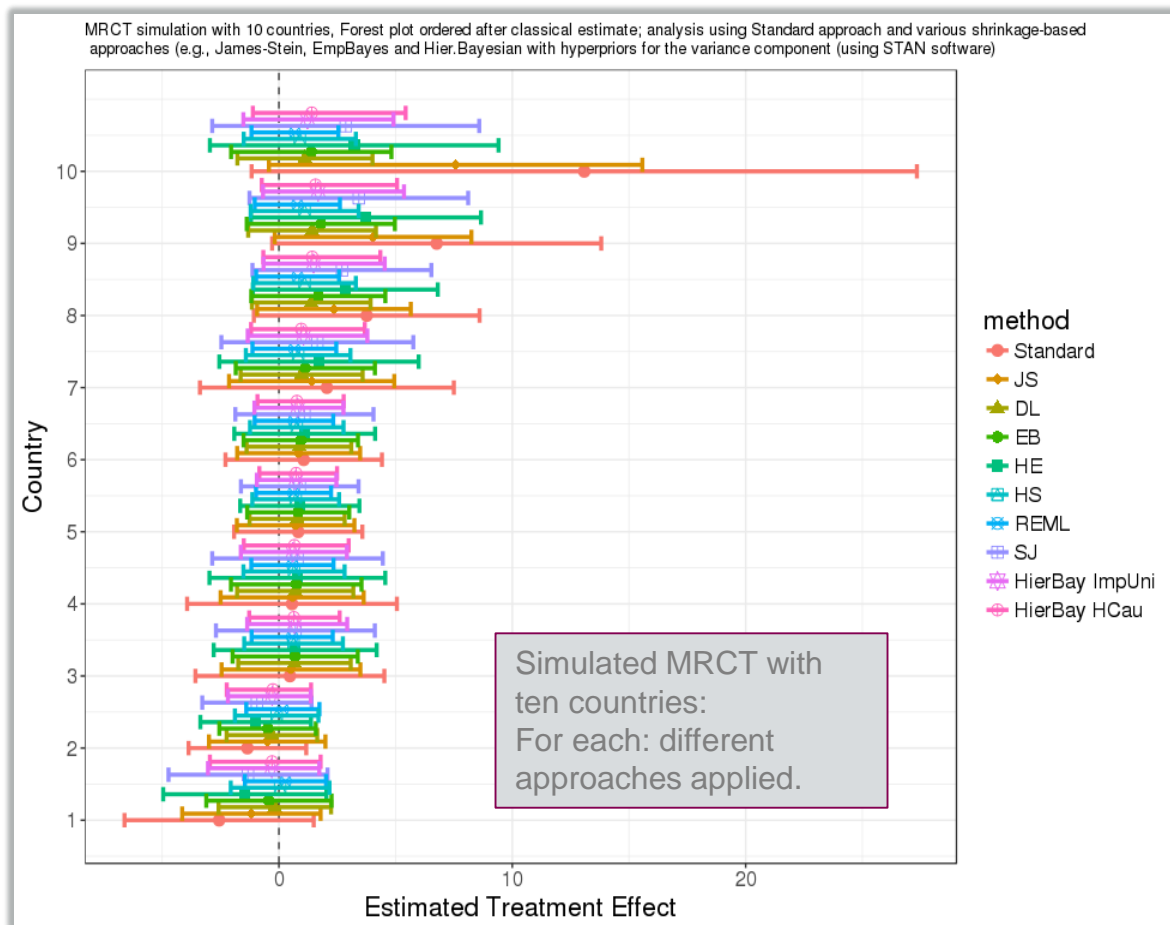
- Empirical Bayes is only one of the possible forms of Shrinkage.
- **Meta Analysis** approaches (same model, but different ' $\tau^2$  approaches' [28,37])
- Other well-known instances: **James-Stein** (frequentist approach) [30].
- **Full Bayesian Hierarchical Model** [16]:
  - Don't fit the prior to the data (as EB did): instead, 'let the data speak'.
  - But requires hyperpriors for variance component
  - Computational intensive fitting (e.g., STAN, hamiltonian chains [18]).
- (One motivation: EB ignores uncertainty in prior estimation, FB doesn't).



# Shrinkage: many flavours, different results...

(RE approach, various options [37, 27],  
Hier. Bayes [16,17], Emp.Bayes [37],  
James-Stein [30]).

- Which is 'Vanilla'?
- Sponsor cherry-picking?
- Note:
- Amount of shrinkage.
- Interval width.

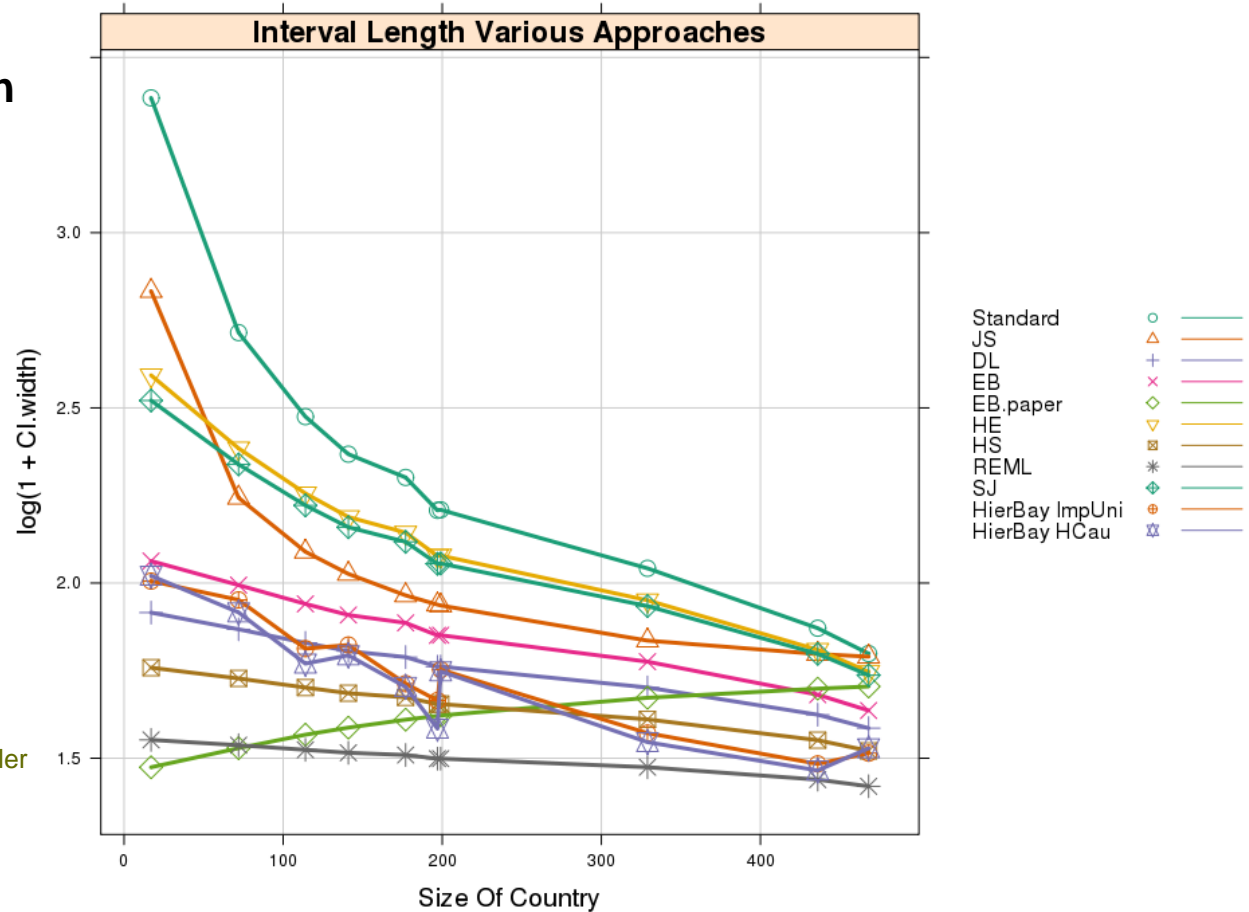


# Shrinkage flavours; operating char. differ ...

- Interval Length as a **function of subgroup size** – across different shrinkage flavours.

(RE approach, various options [37, 27], Hier. Bayes [16,17], Emp.Bayes [37, 30], James-Stein [30]).

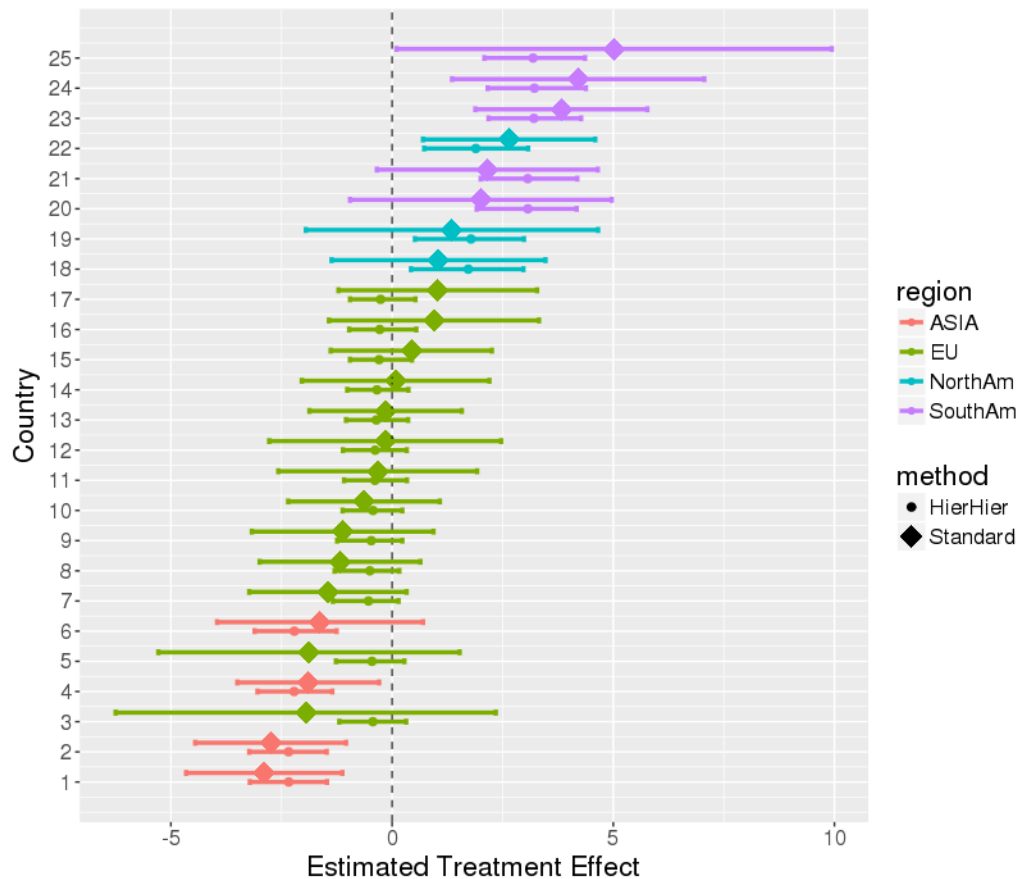
(EmpBayes Variance formula wrong in [30]: smaller regions got more precise CIs ...)



# Multi-level hierarchical (e.g., country within region?).

- In the spirit of the EMA guideline recommendation [13, 7]; **shrinkage within pre-specified stratas** useful?
  - Could be 'region'.
  - Or, other categories:
    - Cat1: differential eff. not expected
    - Cat2: differential eff. plausible
- Possible using a Full Bayesian model.
- Assumptions:
  - Relaxed assumption re. country exchangeability.
  - Prior for Regional eff.,
  - Prior for country-eff. within Regions,
  - Hyper-priors for variance components.
  - Unknown operating characteristics?

MRCT simulation with 25 countries, nested within region. Standard approach, and double-hierarchical Full Bayesian Model (using STAN software)



# Shrinkage; overlapping subgroups:

The general (overlapping) case (e.g., Disease Status, Conc.Med, Gender):

- Hierarchical Bayesian models after splitting patients into **mutually exclusive** (disjoint) subgroups (e.g., [9, 26, 38]).
  - Sometimes model fitting instability due to 'sparse cells'?
- Special case of Model selection: **Model averaging** of subgroup-specific models; [1]. Primarily developed for bias-reduction of *selected* subgroups.
  - But does shrinkage for all subgroups (via BIC model uncertainty).
  - Technically different: ensemble of models instead of a single model (such as e.g., with the EB)



# Shrinkage; some question marks:

- Shrinkage **assumes what we want to demonstrate?**
- - e.g., FDA [34] – “*apriori assuming exchangeability*”.
  - (“*Perhaps this can be helpful in limiting [...] random highs*”).

(Stratified assumption/mixture more realistic?).

- **Normality** assumption of the true effects:? Some **skewness** and/or heavier tails more realistic?
- Full Bayes **requires hyper-prior**:
  - No non-informative exists (variance component). [17]. So, which is it?
  - Diagnostics – **operator dependent?**



# Conclusions:

- The current approach has **flaws**, and novel approaches have question marks too.
  - Which **flavour of Shrinkage**?
  - Hierarchical **model fitting** issues with overlapping subgroups?
  - **Exchangeability** assumption unrealistic?
  - **Unclear** trade-offs by shrinkage when true differential effects are present.
  - **Stratified** (pre-specified?) approach might be worth considering?
  - Technical **model fitting** issues sometimes with overlapping subgroups?
- Several **other approaches** suggested ([1], [32, 33], [4]):
  - Some only for selected subgroup? some only for non-overlapping subgroups?
  - Some further head-to-head comparisons of operating characteristics needed?
  - Some question marks on the handling of prognostic factors for some permutation approaches [15]?

Regulator's & other sponsor's view on this entire topic? **What do you think?**



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In the event of errors or false claims, the main author takes the full responsibility for it.





## References (1).

- [1] **Bornkamp, B., Ohlssen D., Magnusson B., Schmidli, H.;** Model Averaging for Treatment effect estimation in subgroups. *Phar. Statistics*, 2017, vol 16.
- [2] **Brookes ST., Whitely E., Egger M., Smith GD., Mulheran PA., Peters TJ.;** *Subgroup analyses in randomized trials: risks of subgroup-specific analyses; power and sample size for the interaction test.* *J Clin Epidemiol.* 2004, Vol 57(3).
- [3] **Chen, YH., Wu, YC., Wang M.;** *A Bayesian approach to evaluating regional treatment effect in a multiregional trial.* *J. Biopharm Stat.*, 2009, 19(5).
- [4] **Chen J., Zheng H., Quan H., Li G., Gallo P., Ouyang SP., Binkowitz B.;** Ting N, Tanaka Y, Luo X, Ibia E; *Graphical assessment of consistency in treatment effect among countries in multi-regional clinical trials.* *Clin Trials.* **2013**;10(6).
- [5] **Cui L, Hung JHM, Wang SJ, Tsong Y.** *Issues related to subgroup analysis in clinical trials.* *Journal of Biopharmaceutical Statistics* 2002, Vol 12.
- [6] **David, C. E., Leffingwell, D. P.;** *Empirical Bayes estimates of subgroup effects in clinical trials.* *Controlled Clinical Trials* 11, 1990.
- [7] **Dane, A., Spencer, A., Rosenkranz, G., Lipkovich, I., Parke, Tom.** *Subgroup Analysis and Interpretation for Phase 3 Confirmatory Trials: White paper of the EFSP/PSI Working Group on Subgroup Analysis.* [Submitted 2018], *Pharmaceutical Statistics.*
- [8] **Dane, A., Spencer, A., Stone, A., Svensson, D.;** *SEAMOS: The use of a Resampling Based Graphical methods to Present Standardised Effects Adjusted for Multiple Overlapping Subgroups.* [Submitted 2017] *Statistics in Biopharmaceutical Research.*
- [9] **Dixon D., Simon R.;** *Bayesian Subset Analysis.* *Biometrics* 47.



## References (2).

- [10] **Dmitrienko, A., Millen, B., Lipkovich, I.**; *Multiplicity considerations in Subgroup Analysis*. Stat. In Medicine, 2017.
- [11] **Efron, Bradley**. *Large-scale inference: Empirical Bayes methods for estimation, testing, and prediction*. Cambr. Univ. Press 2010.
- [12] **Efron, Bradley**. *Empirical Bayes Methods for Combining Likelihoods*. 1996, J. Am. Stat. Assoc. 91.
- [13] **EMA Guideline on the investigation of subgroups in confirmatory clinical trials**  
[http://www.ema.europa.eu/docs/en\\_GB/document\\_library/Scientific\\_guideline/2014/02/WC500160523.pdf](http://www.ema.europa.eu/docs/en_GB/document_library/Scientific_guideline/2014/02/WC500160523.pdf).
- [14] **Fletcher C, Chuang-Stein C, Paget MA, Reid C, Hawkins N.**; *Subgroup analyses in cost-effectiveness analyses to support health technology assessments*. Pharm.Stat. 2014 Jul-Aug;13(4):265-74
- [15] **Foster J., Nan B., Shen L., Kaciroti N., Taylor J.**; *Permutation Testing for Treatment-Covariate Interactions and Subgroup Identification*. Stat. Biosci. 2016 Jun;8(1):77-98
- [16] **Gelman A, Carlin J., Stern H.** Bayesian Data Analysis, Chapman Hall, 3<sup>rd</sup> Ed., 2013.
- [17] **Gelman A.** *Prior Distributions for Variance Parameters In Hierarchical Models*. Bayesian Analysis Vol 1(3), 2006.
- [18] **Gelman A., & The Stan Development Team**; STAN software: <http://mc-stan.org/>
- [19] **Harpaz R., DuMouchel W., LePendou P., Bauer-Mehren A., Ryan P., Shah N.**; *Performance of Pharmacovigilance Signal Detection Algorithms for the FDA Adverse Event Reporting System*. Clin Pharmacol Ther. 2013 June; 93(6).



## References (3).

- [20] **Hauck, W.W., Anderson, S., Marcus SM.** *Should we adjust for covariates in nonlinear regression analyses of randomized trials?* Control Clin. Trials 1998, 19(3), 249-56.
- [21] **ICH Efficacy Guidelines** <http://www.ich.org/products/guidelines/efficacy/article/efficacy-guidelines.html#16>
- [22] **Jackson D., Veroniki A., Law M., Tricco A., Baker R.** *Paule-Mandel estimators for network meta-analysis with random inconsistency effects.* Research Synthesis Methods, Vol 8 (4), 2017.
- [23] **Jones H., Ohlssen D., Neuenschwander B., Racine A., Branson M.;** *Bayesian models for subgroup analysis in clinical trials.* Clinical Trials 2011; 8: 129–143.
- [24] **Lipkovich I. , Dmitrienko A., Denne J., Enas G.;** *Subgroup Identification based on Differential Effect Search (SIDES) – a recursive partitioning method for establishing response to treatment in patient subpopulations.* Stat. In. Medicine, 2011.
- [25] **Marius T., Borncamp, B.;** *Comparing Approaches to Treatment Effects Estimation for Subgroups In Clinical Trials.* Stat. in Bioph. Research, 9:2, 2016.
- [26] **Parametric Empirical Bayes Inference: Theory and Applications.** Morris C., Journal of the American Statistical Association, Vol. 78, No. 381 (Mar., 1983)
- [27] **Miller B., Dmitrienko A., Song G.;** *Bayesian Assessment of the Influence and Interaction Conditions in Multipopulation Tailoring Clinical Trials.* J. of Biopharm. Stat., Vol 24, 2014.
- [28] **Normand T.;** *Tutorial in Biostatistics. Meta-analysis: formulating, evaluating, combining, and reporting by S-L.* Statistics in Medicine, 18 (1999).



## References (4).

- [29] **Paget M., Chuang-Stein C., Fletcher C., Reid C.;** *Subgroup analyses of clinical effectiveness to support health technology assessments*. Pharm. Stat., Vol 10 (6) 2011.
- [30] **Quan H., Li M., Shih WJ., Ouyang, SP., Chen J., Zhang J., Zhao PJ.;** *Empirical shrinkage estimator for consistency assessment of treatment effects in multi-regional clinical trials*. Stat. In Med., Vol 32, Issue 10, 2013.
- [31] **Quan H., Li M., Shih WJ., Ouyang SP., Chen J., Zhang J., Zhao PJ.;** *Multi-regional clinical trial design and consistency assessment of treatment effects*. Stat. In Med., Vol 33, Issue 13, 2014.
- [32] **Rosenkranz G.;** *Exploratory Subgroup Analyses in Clinical Trials*. ISCB Pre-Conference Training Course, Birmingham, August 21, 2016
- [33] **Rosenkranz G.;** *Bootstrap corrections of treatment effect estimates following selection*. Computational Statistics and Data Analysis, Vol 69, 2014.
- [34] **Russek-Cohen., E.;** [http://www.ema.europa.eu/docs/en\\_GB/document\\_library/Presentation/2015/03/WC500183614.pdf](http://www.ema.europa.eu/docs/en_GB/document_library/Presentation/2015/03/WC500183614.pdf)., *Comments from the FDA working group on SUBGROUPS*, 2015.
- [35] **Varadhan R., Wang SJ.;** Treatment effect heterogeneity for univariate subgroup in clinical trials: Shrinkage, standardization, or else. Biometrical Journal 58, 2016.
- [36] **Stein C.,** *Inadmissibility of the usual estimator for the mean of a multivariate normal distribution*. Proceedings of the Third Berkeley Symposium on Mathematical Statistics and Probability, Berkeley and Los Angeles, University of California Press, 1956, Vol. 1.
- [37] **Viechtbauer W.,** METAFOR R package, <https://cran.r-project.org/web/packages/metafor/index.html>, 2017.

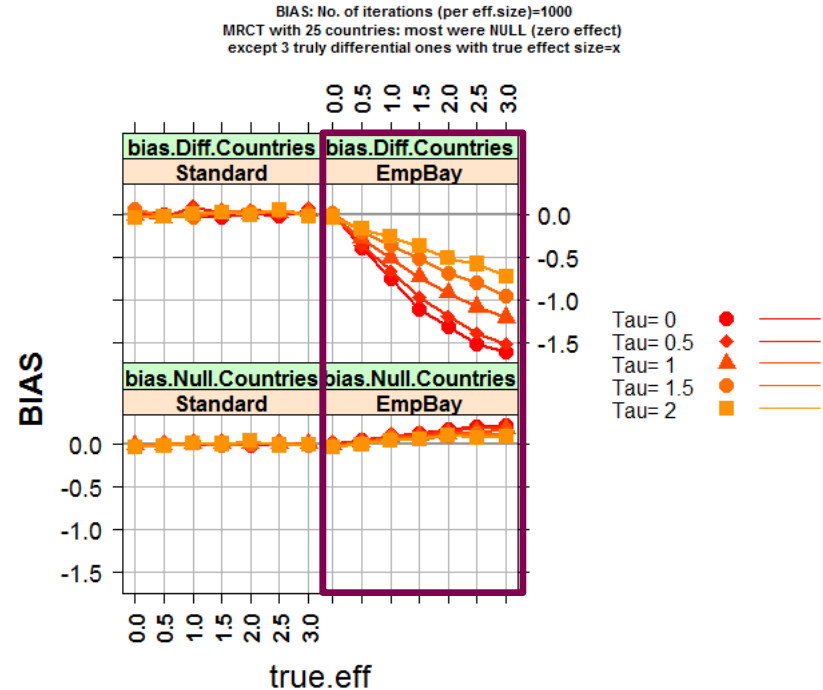
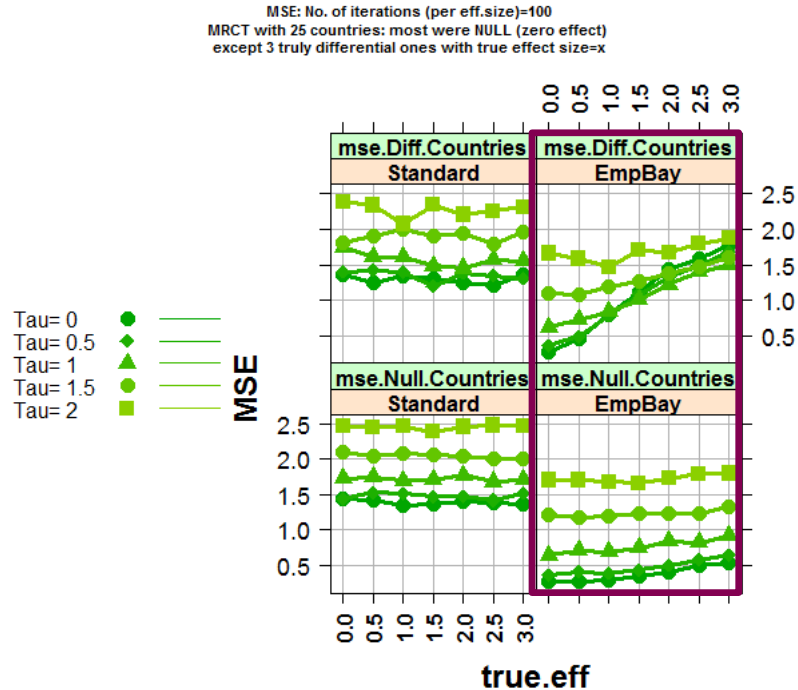


## Back-Up Slides:



# MSE & Bias (exchangability violated).

X-axis= True Effect ( $x$ ) in the differential countries.



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