

APBG

AUSTRALIAN PHARMACEUTICAL  
BIostatISTICS GROUP

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# Real World Evidence

*Australian Pharmaceutical Biostatistics Group (APBG)*

*RWE Seminar July 29<sup>th</sup> 2015*

Alan J.M. Brnabic  
Senior Research Scientist , Eli Lilly  
Chair of APBG

The Lilly logo is written in a white, cursive script font, positioned in the bottom right corner of the slide.

# What is the APBG ?

- ◆ A not-for-profit association of pharmaceutical industry statisticians in Australia.
- ◆ Our mission
  - is to ensure high statistical standards within Australia to assist in the decision processes which provide safe, efficacious and cost-effective health care products produced in a regulated environment for the health and quality of life of people
- ◆ Made up of 4 steering committee members:
  - Alan Brnabic ( Chair) Annie Solterbeck (vice-chair), Philip McCloud (treasurer), Joanna Leadbetter (secretary)
- ◆ Currently ~ 55 members

# What is the APBG ?

- ◆ To provide professional educational opportunities of specific relevance to statisticians in the regulated health care industry.
  - Past Workshops/Courses:
    - DMC
    - Do pharmacological interventions reduce drugs-related deaths ?
    - Mixed Treatment Comparisons
    - Adaptive Designs for Clinical Trials
- ◆ If you want to be on our mailing list speak to Myself or Joanna Leadbetter

# Thanks

- ◆ Thank you to all that helped put this event together
  - Joanna Leadbetter, Annie Solterbeck, Matt Slabbert, Philip McCloud, Laurent Billot, Richard Hutchinson
- ◆ Thank you to our speakers
  - David Grainger
  - Laurent Billot
  - John McNeil
- ◆ Eli Lilly for hosting the event

# OUTLINE

## ◆ Introducing TALKS:

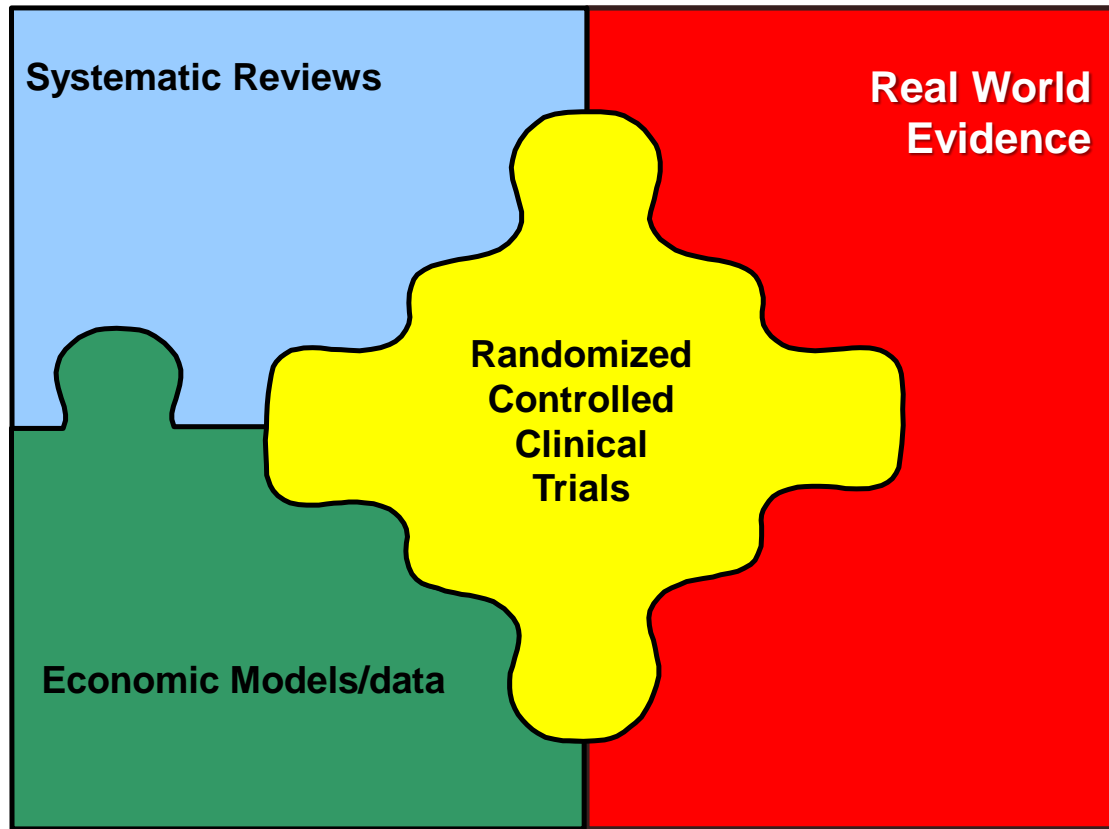
- David Granger: RWE: Policy, Governance and Implications for HTA
- Laurent Billot: Practicalities of using linked data & statistical applications using Propensity Scoring
- John McNeil: Clinical Registries, Opportunity and Limitations

## ◆ Introduction to RWE

- What is RWE
- Why do we need RWE?
- Analytical Challenges of RWE

# What is RWE? - RESEARCH MOSAIC OF EVIDENCE

RWE is complementary to a robust clinical trial program.



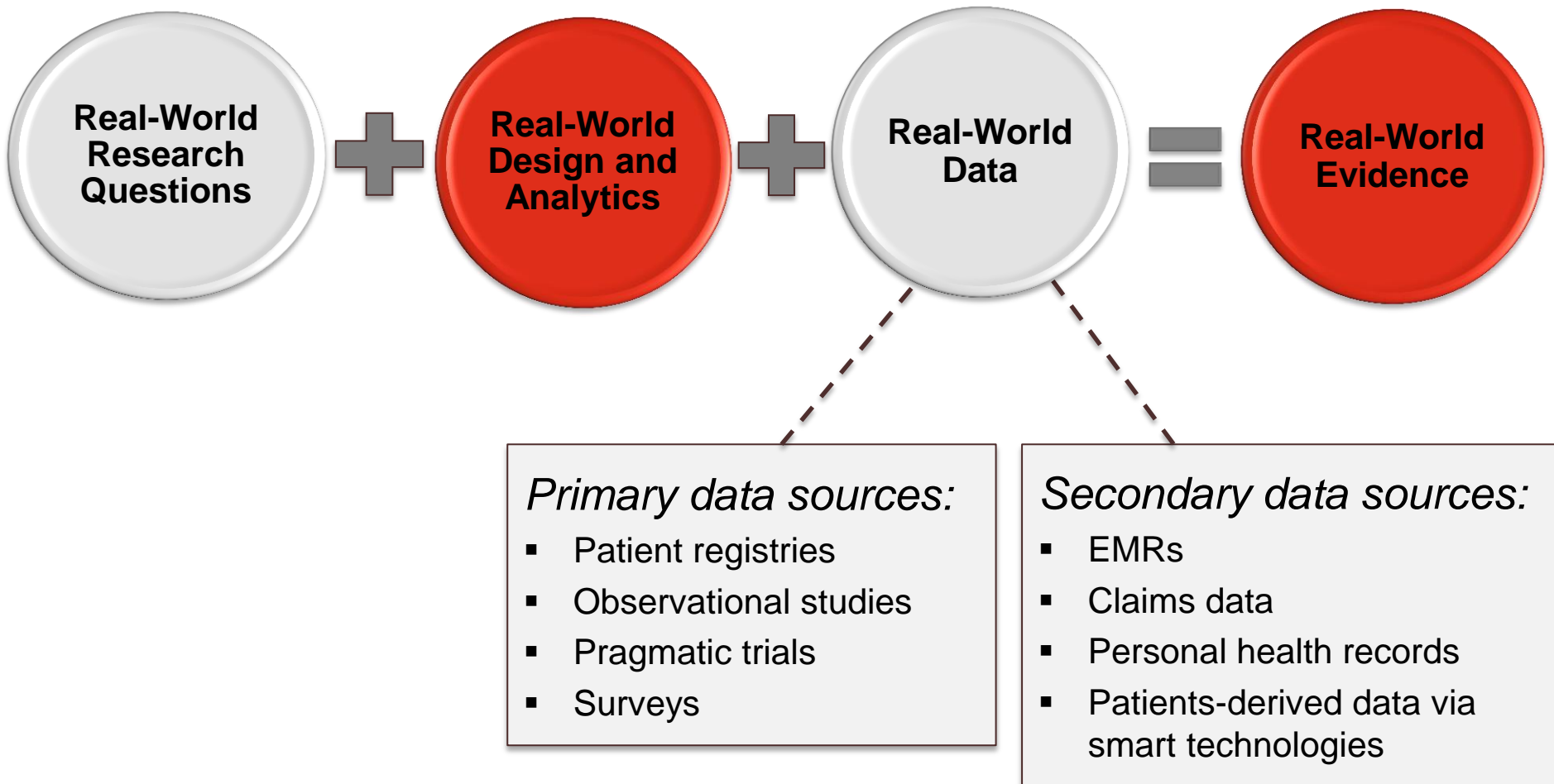
# What is RWE? - Definition

**RWE** is one form of evidence (along with RCT, health economics studies, etc.)

- 1 derived from primary or secondary real world data sources**, with appropriate design/analyses, to
- 2 provide insights** on diseases, medicines, patient populations and healthcare practices, that will
- 3 inform customer and internal decision making.**



# What is RWE? - Definition





# Why RWE? - INCREASING RWE REQUIREMENTS

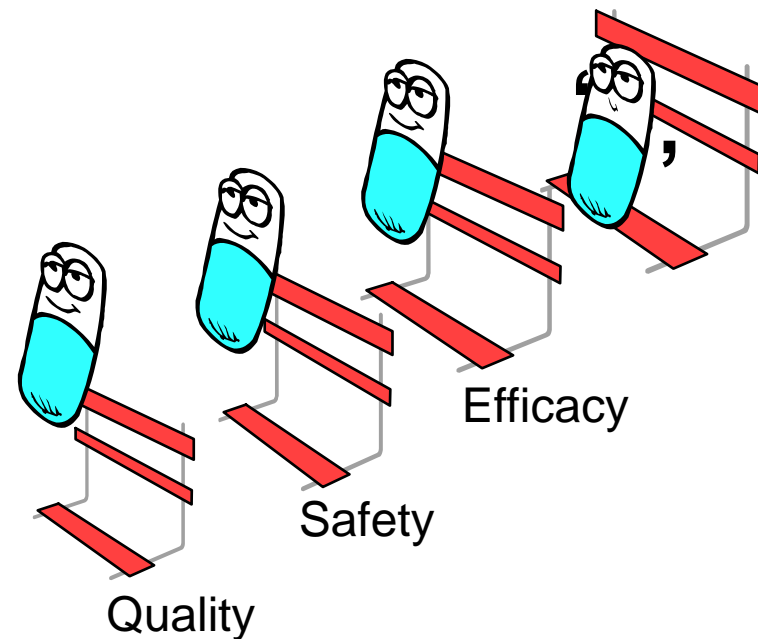
**Globally, RWE Research is increasingly required to gain and maintain reimbursement approval and is linked to which populations get access.**

## Traditional regulatory requirements

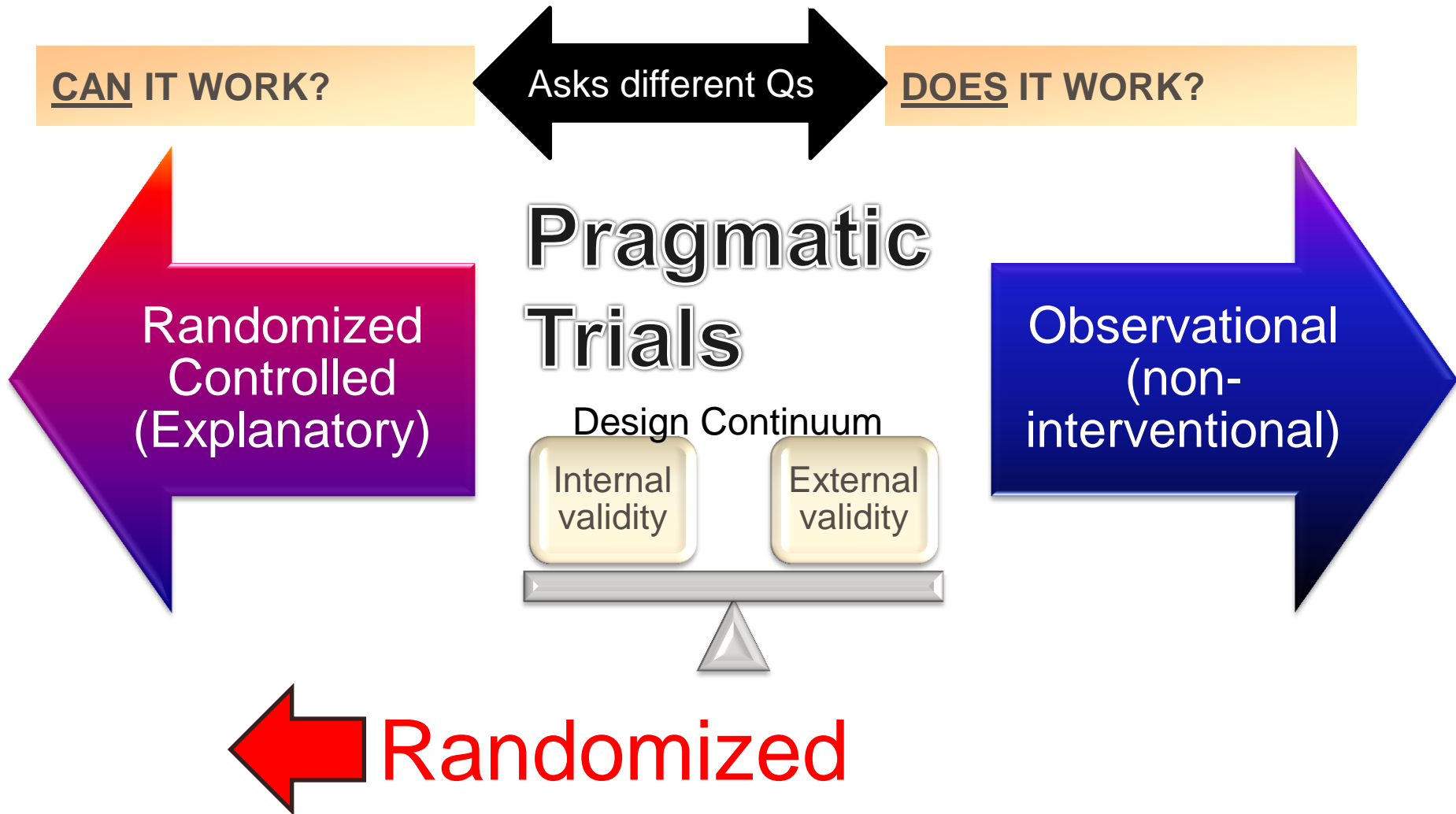
- Quality
- Safety
- Efficacy

## Emerging requirements

- Clinical effectiveness
- Patient outcomes, e.g. quality of life
- Cost Effectiveness
- Budget Impact
- Benefit-Risk
- Patient subtypes/heterogeneity
- Understand unmet clinical need
- Use of drugs in real life



# Why RWE? – A Balance

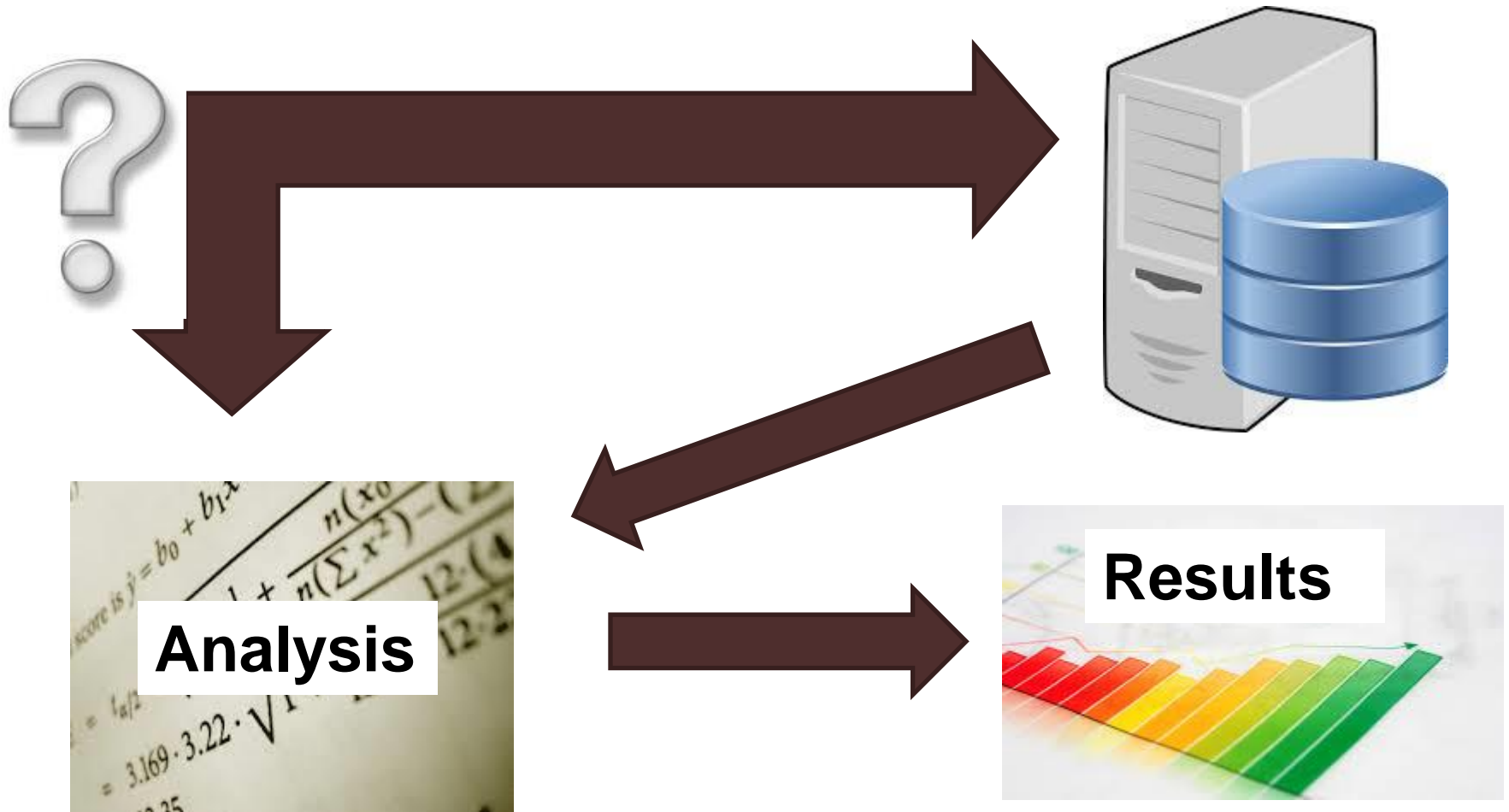


# Application of RWD

## AU Lilly Diabetes/George Institute: EXTEND45 NSW Data linkage Project



# RWE – The question!



# Statistical Challenges in RWD

- ◆ Standard analytic approaches don't work
  - you get biased answers
- ◆ No randomisation
  - requires different methodology
- ◆ Challenges
  - Selection Bias
  - Unmeasured confounding
  - Reliable Data
  - BIG DATA – all of the above!
  - Mixing RCT & RWD

# Statistical Challenges in RWD

## #1 Issue: Selection Bias / Confounding

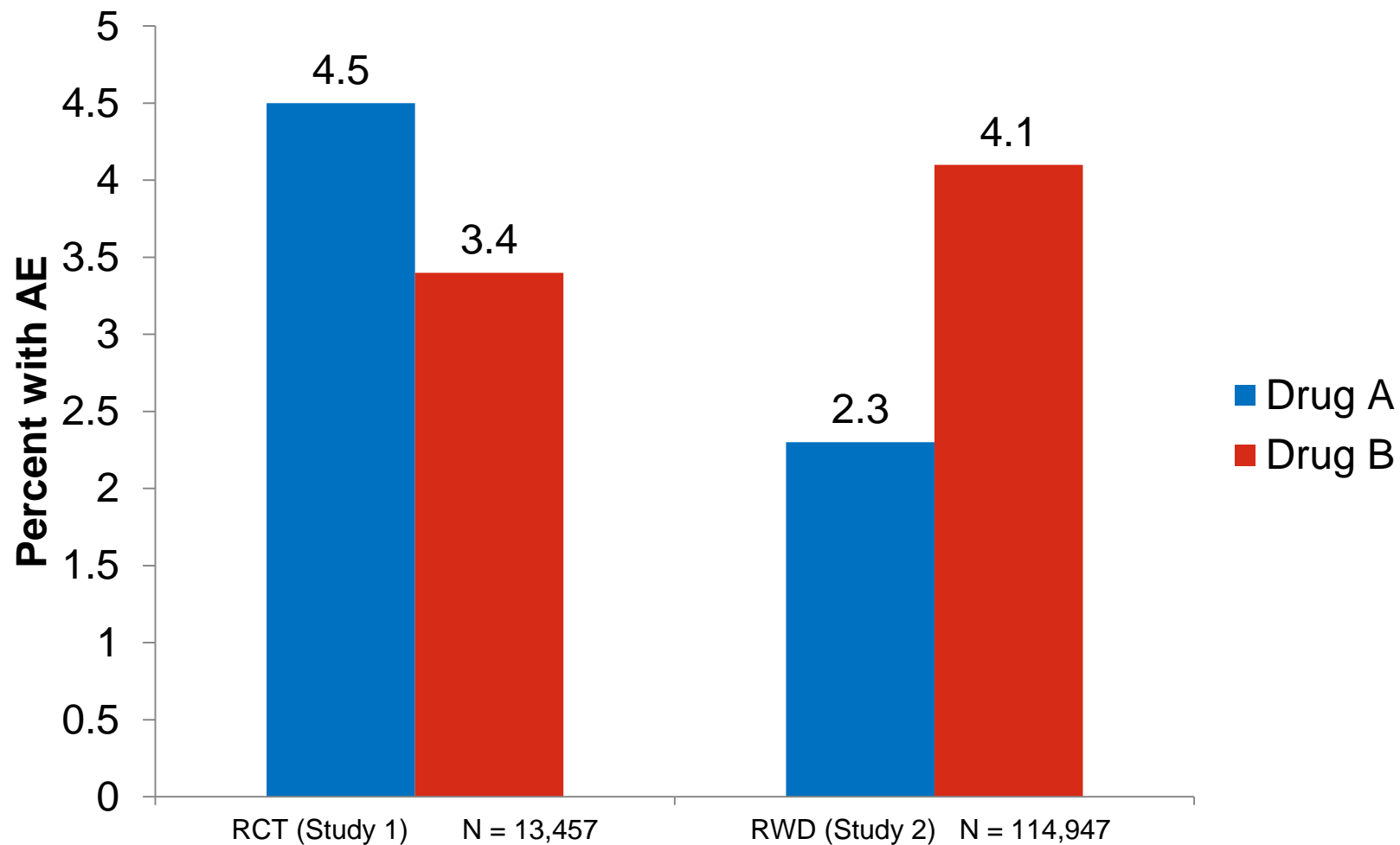


- With randomization – standard methods produce estimates of causal treatment effects



- Without randomization – standard methods produce only ‘associations’ .... Treatment groups are NOT comparable at baseline thus comparisons are BIASED

# Selection bias: AE rates: RCT vs RWD





# Why the difference in Results? =Selection Bias

A confounder is a variable that is associated with both Cohort and Outcome

Study 2 Data Set	Drug A (N=10,448)	Drug B (N=104,499)
Mean Age (years)	57	64
Age > 75 yrs (%)	3.5%	24.9%
Male (%)	75%	66%
Renal Insufficiency	9.3%	19.2%

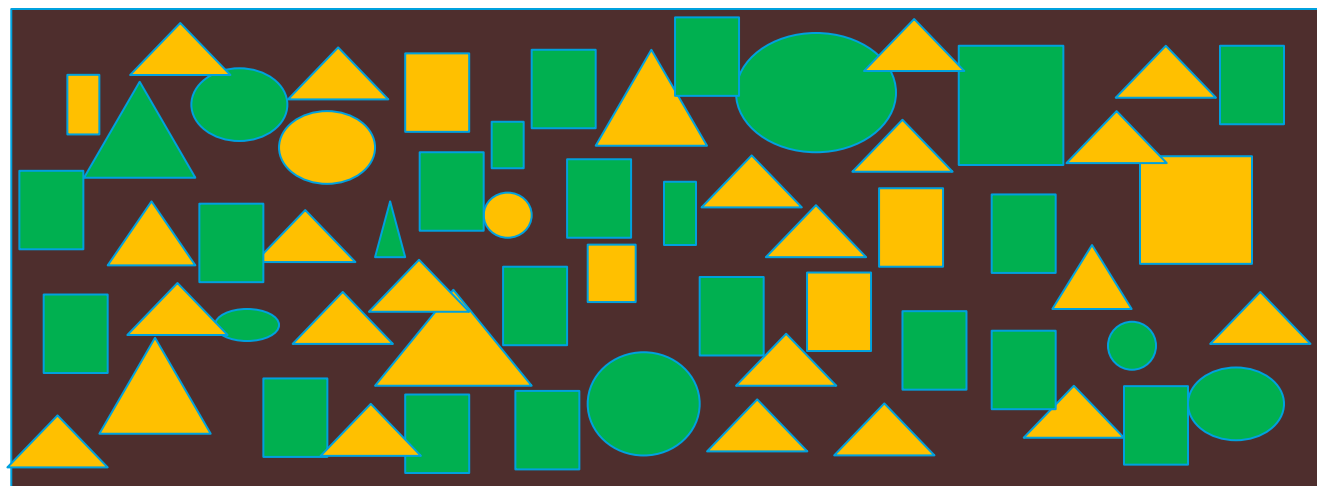
## AE Risk Model

Baseline Factors:

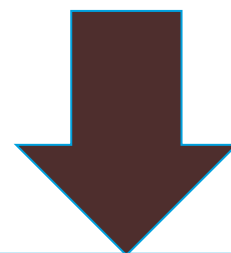
- 1) Older Age
- 2) Female
- 3) Renal Insufficiency
- 4) Blood Pressure
- 5) History of Bleeding

# Selection Bias: A solution

Real World  
Population



Propensity Score:  
Probability of being assigned Treatment A



Propensity  
Matching



# BASIC ASSUMPTIONS FOR CAUSAL INFERENCE

Propensity Score adjustments can provide for estimates of the causal group differences under the following assumptions:

#1

**No Unmeasured Confounders**

All confounders are in the dataset and analysis

#2

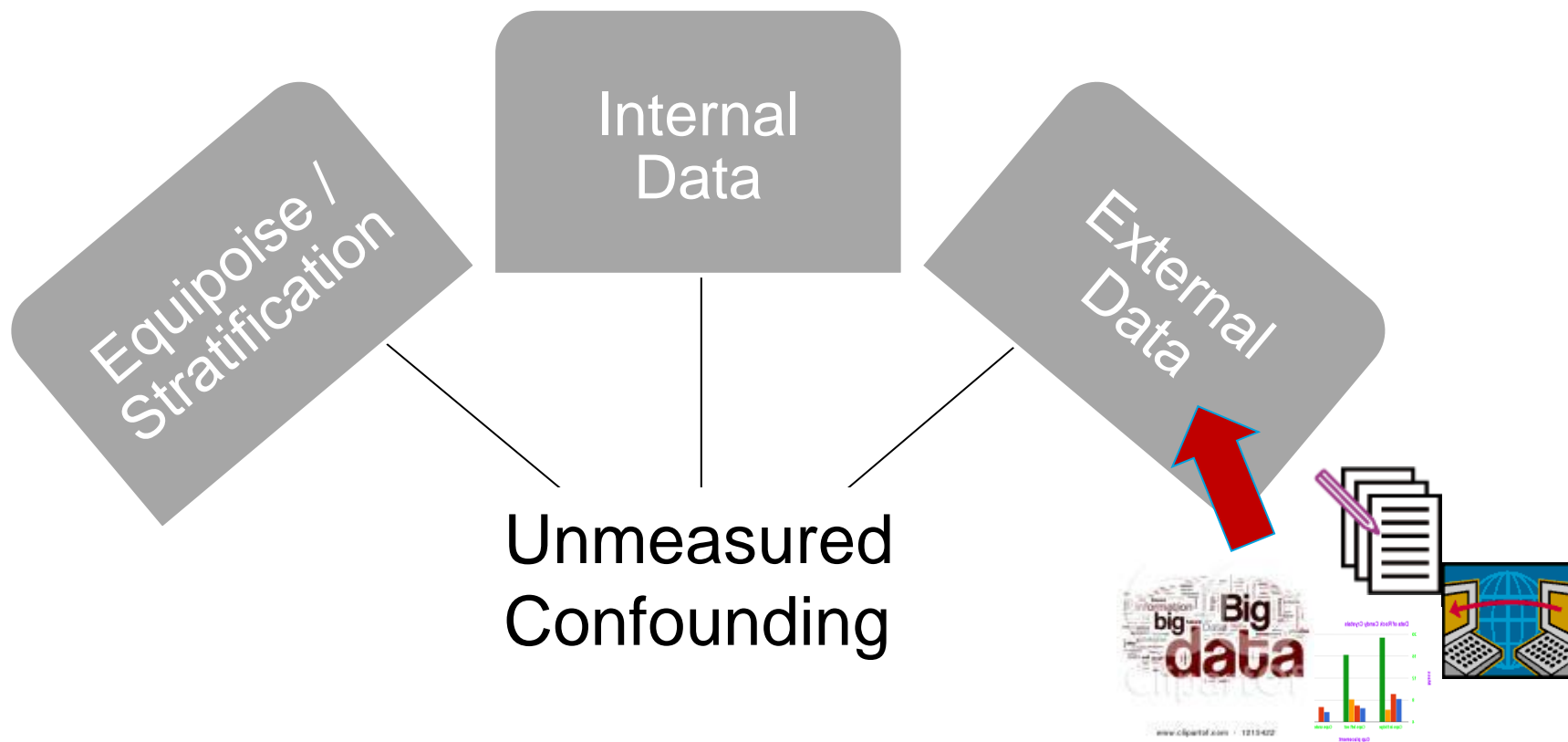
**Sufficient Overlap in Populations**

positivity, no perfect confounding

#3

**Correct Statistical Models**

# UNMEASURED CONFOUNDING



# Challenges with Data

- ◆ Can the data:
  - Answer the question you are asking
    - Collect the information you need e.g. clinical outcomes, capture all costs, QoL
- ◆ Is the data:
  - Representative e.g cover all labs in Australia
  - Complete; % missing of variables reasonable, coding correct/consistent
  - Accurate; can be validated with other sources
  - Up to date
  - Can it be linked to other sources

# Methods for Analysing BIG RWD

- ◆ Focus on Interaction Identification – this identifies relationships that pertain to only certain subgroups
- Traditional DM Machine learning methods include
  - tools used for supervised learning
    - traditional statistical prediction methods like regression models
    - tree-based methods (e.g. classification and regression trees (CART) and random forests)
    - neural networks, discriminant functions and linear classifiers, support vector classifiers and machines (SVM).

# Methods for Analysing BIG RWD

- ◆ Focus on Interaction Identification – this identifies relationships that pertain to only certain subgroups

**CHALLENGE**– How to Apply these methods on BIG RWD given the issues of selection bias & unmeasured confounding?

- neural networks, discriminant functions and linear classifiers, support vector classifiers and machines (SVM).



# Mixing RCT & RWD

- ◆ Using RWD (e.g. prospective observational studies) to generalise RCT results to the “real world” population (via propensity score or entropy balancing weighting)
- ◆ Using IPD and aggregate data (in literature) to match populations better to perform indirect comparisons & network meta analysis (matched adjusted indirect comparisons = MAIC)

# Acknowledgements

- ◆ Doug Faries
- ◆ Tony Zager