



# Propensity Score Diagnostics

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# Background

- Commonly used in observational data to deal with confounding bias.

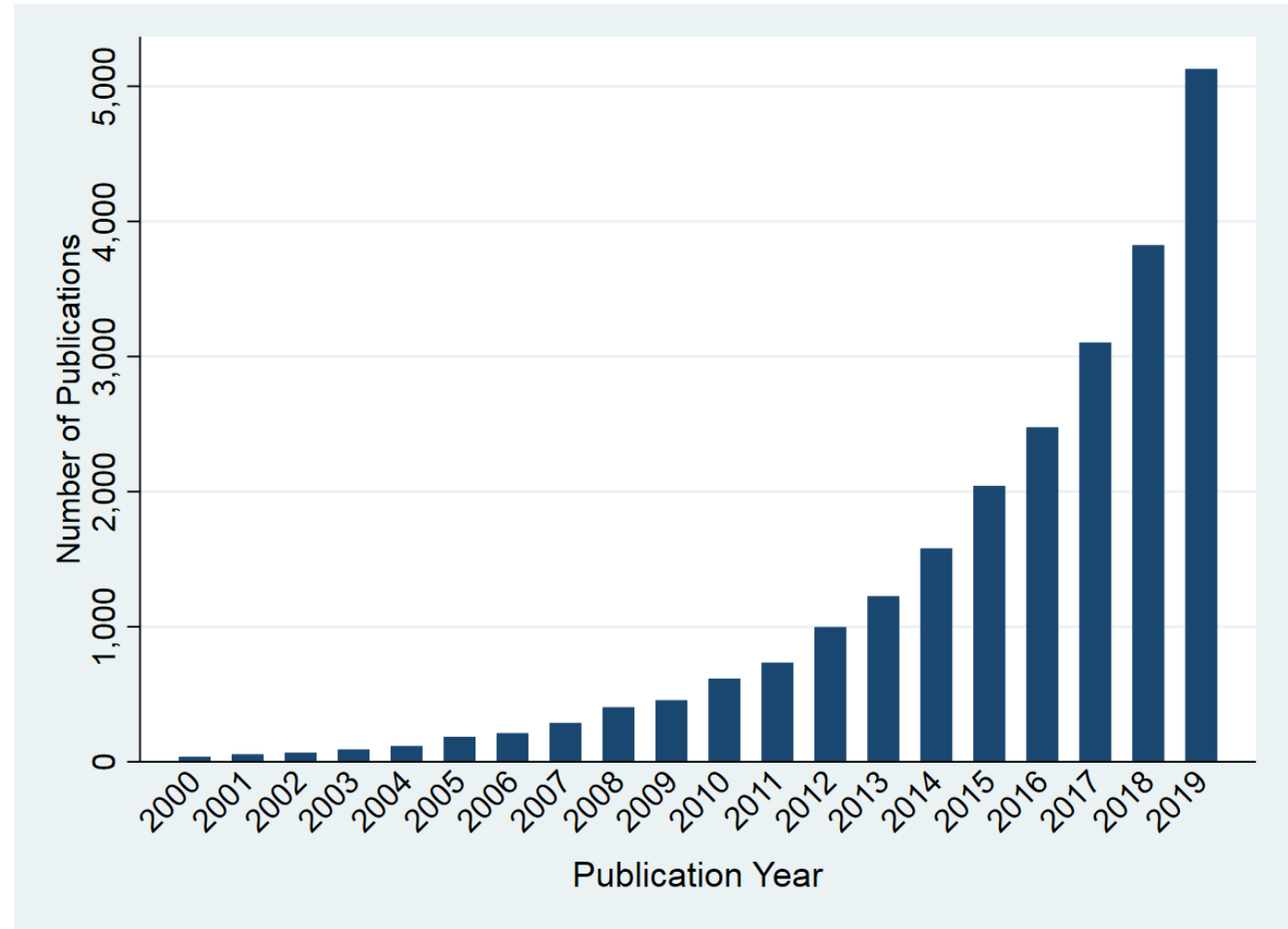


Figure 1: Number of propensity score publications in medical research by year

# Background

- Commonly used in observational data to deal with confounding bias.
- Poorly estimated propensity scores may lead to biased estimates.
- Use of diagnostics to assess propensity scores is important.
- Currently unknown how best to assess propensity scores.

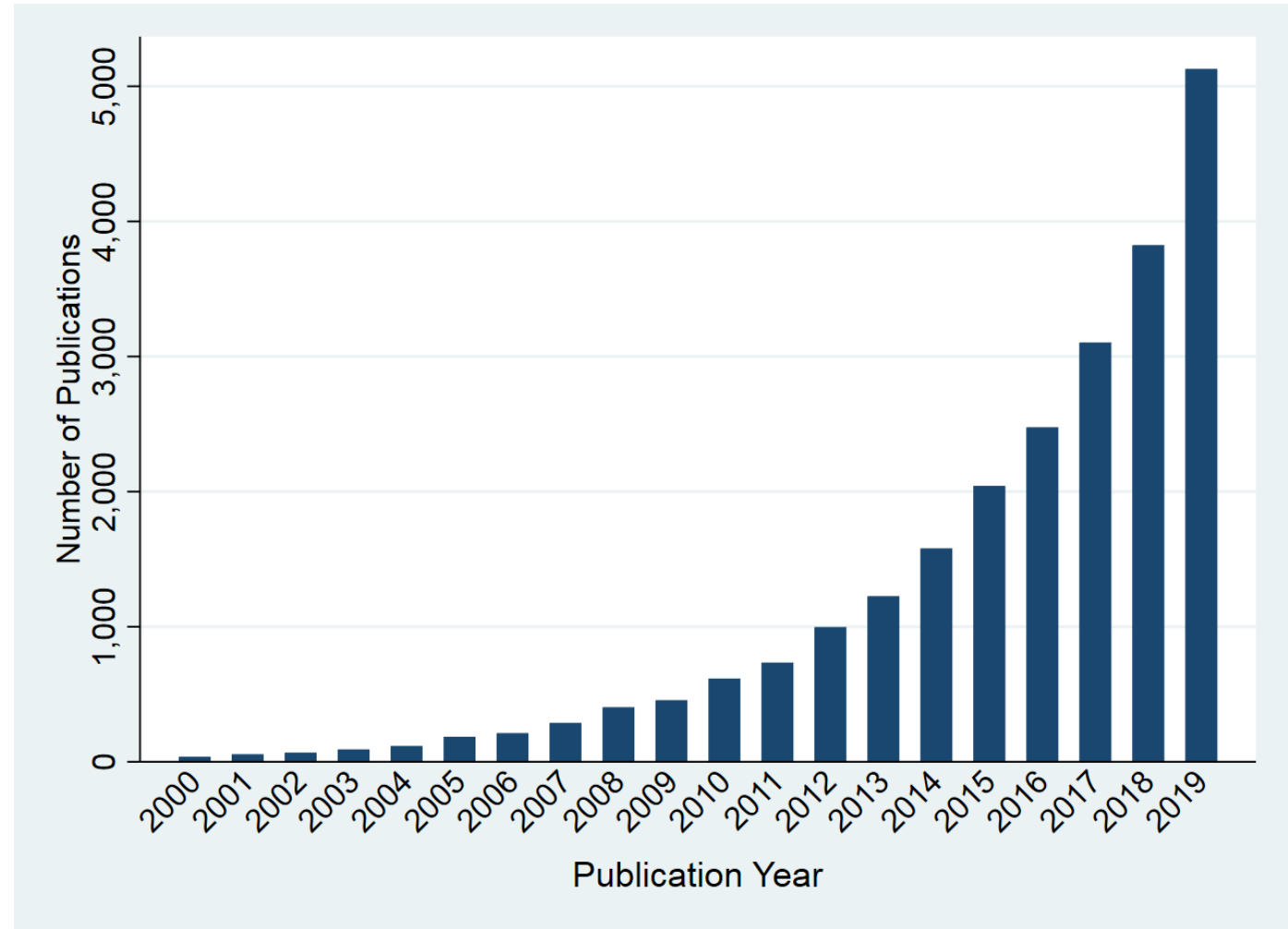
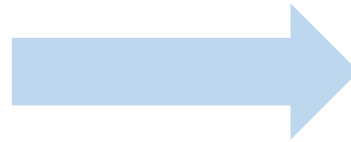


Figure 1: Number of propensity score publications in medical research by year

# Aims of research

## **Aim 1:**

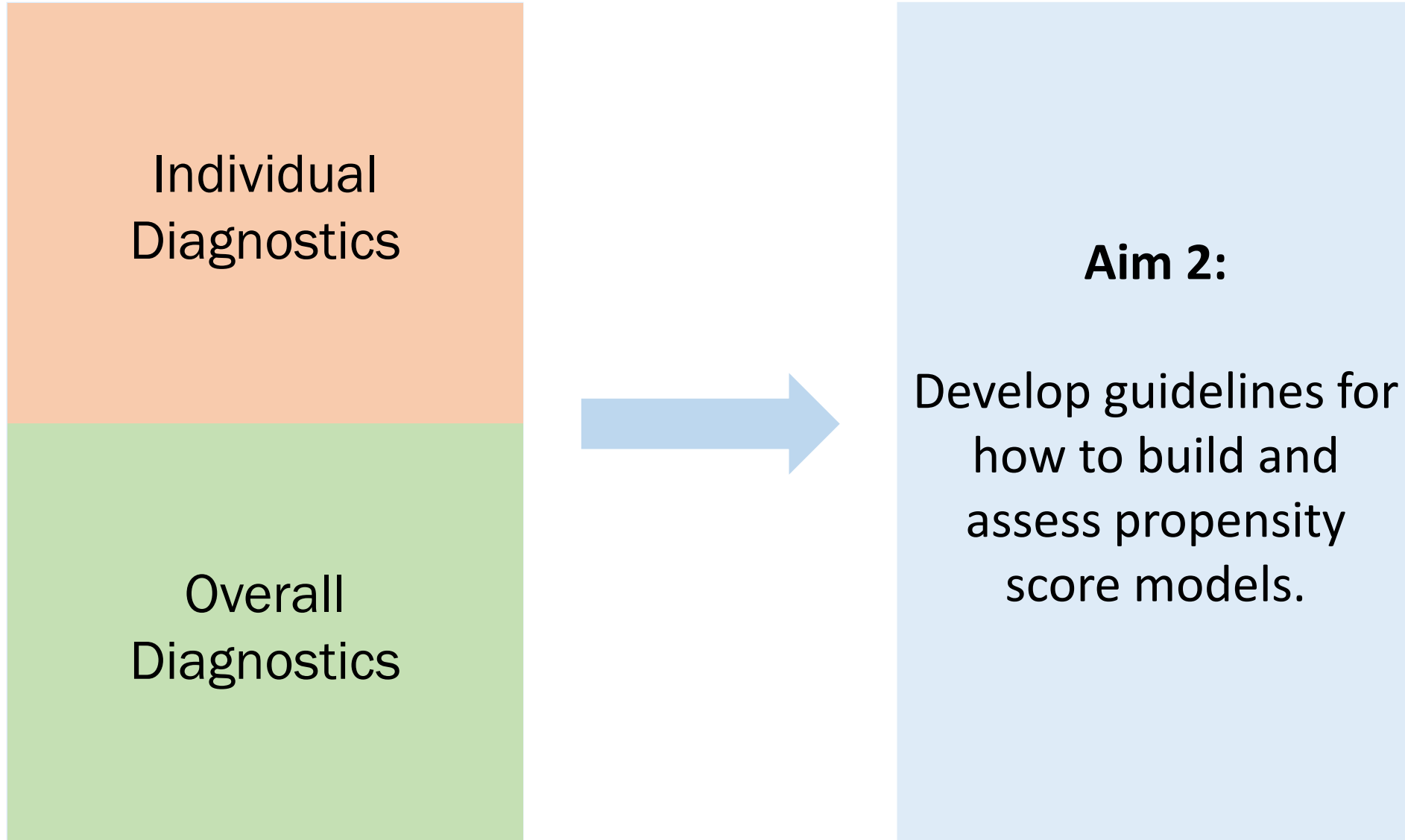
Review and compare  
the existing  
propensity score  
diagnostics.



## **Aim 2:**

Develop guidelines for  
how to build and  
assess propensity  
score models.

# Aims of research



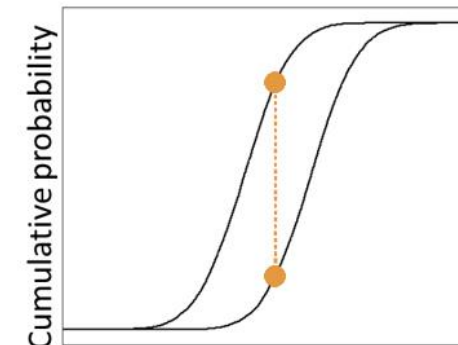
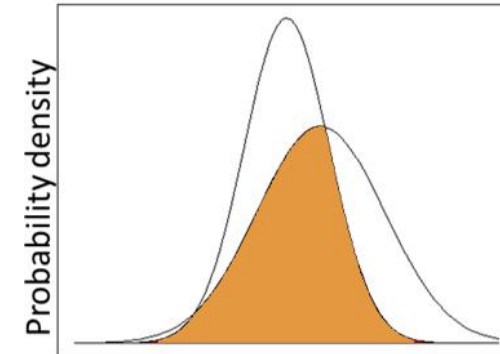
# Individual diagnostics

## Mean-based

- Standardised difference (SD)
- t-test statistic ( $t$ )
- Percent reduction in mean difference (PR)

## Distribution-based

- Overlapping coefficient (OVL)
- Kolmogorov-Smirnov Statistic (KS)

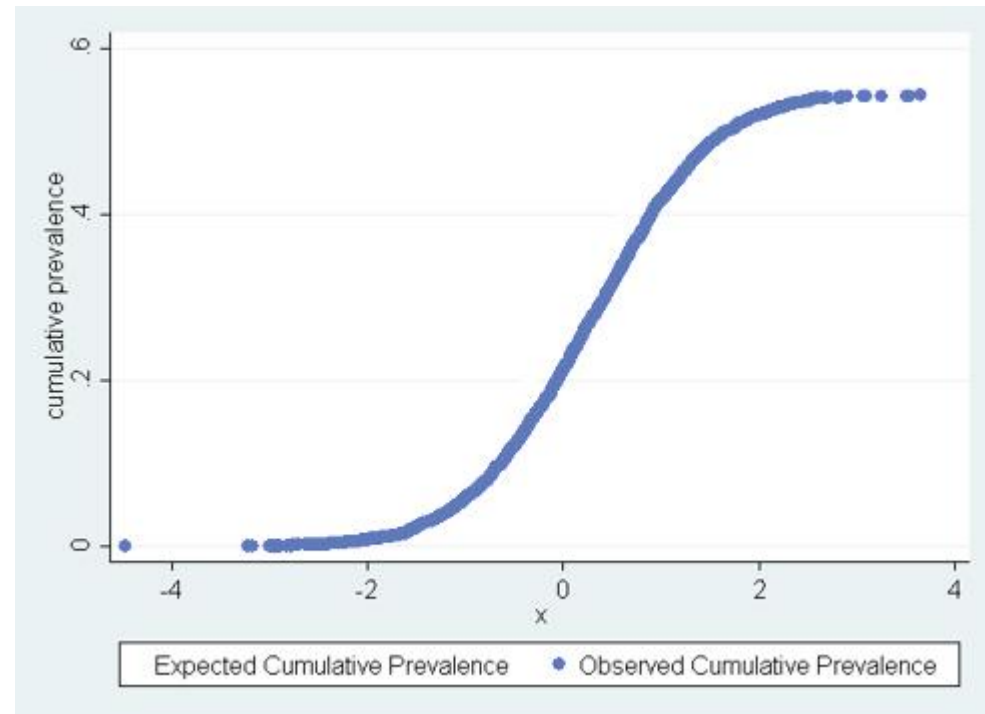


# Cumulative prevalence of exposure

*Notation:* exposure indicator for subject  $i$ :  $E_i$ , propensity score for subject  $i$ :  $PS_i$ , sample size:  $n$ .

For continuous variable  $X$ :

- $OCP_X(X_0) = \frac{1}{n} \sum_{i: X_i \leq X_0} E_i$

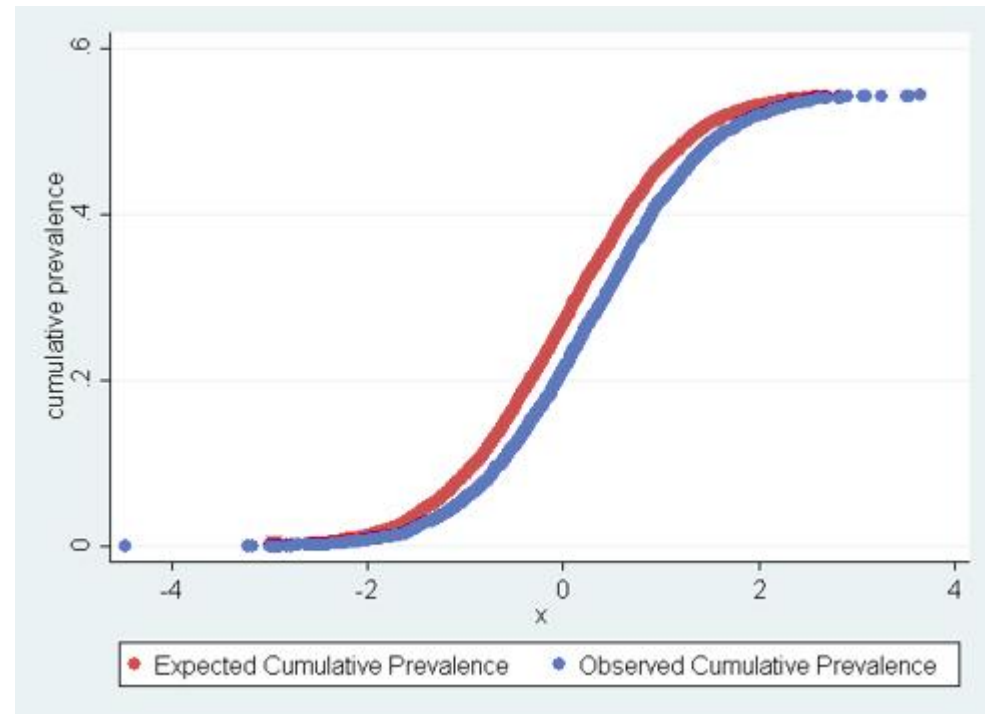


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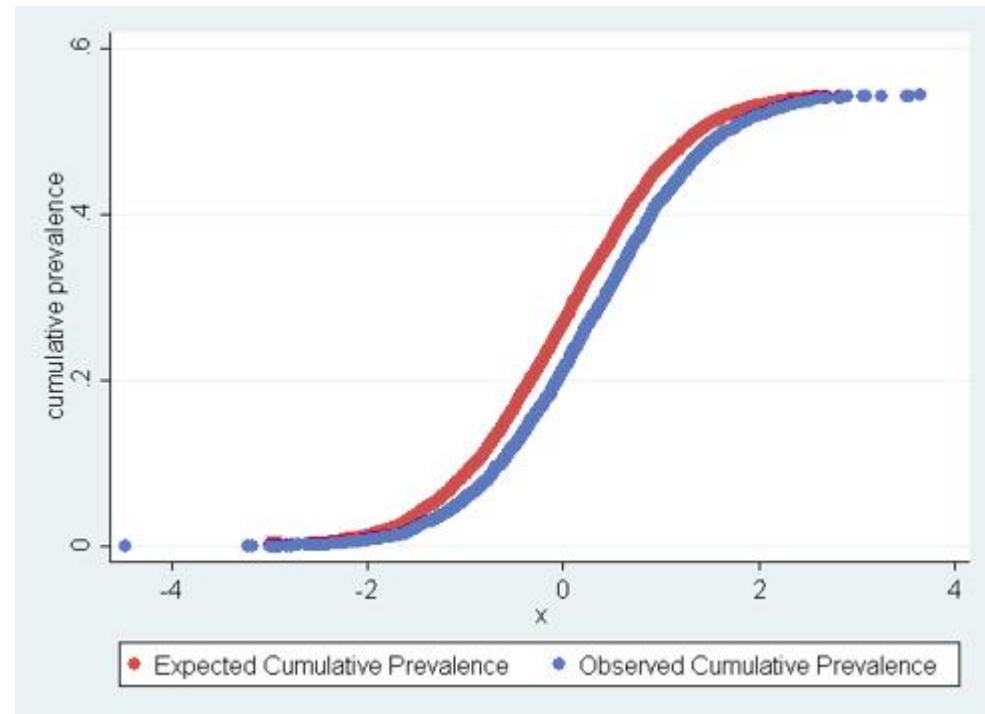


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- $ECP_X(X_0) = \frac{1}{n} \sum_{i: X_i \leq X_0} PS_i$
- $D_X = |OCP_X - ECP_X|$



# Simulated data

## Propensity score model:

- $\text{logit}(\text{PS}) = \alpha_0 + \alpha_1 X_1 + \alpha_2 X_2 + \dots + \alpha_7 X_7 + \alpha_8 X_8$

## Variation between scenarios:

### Correct PS:

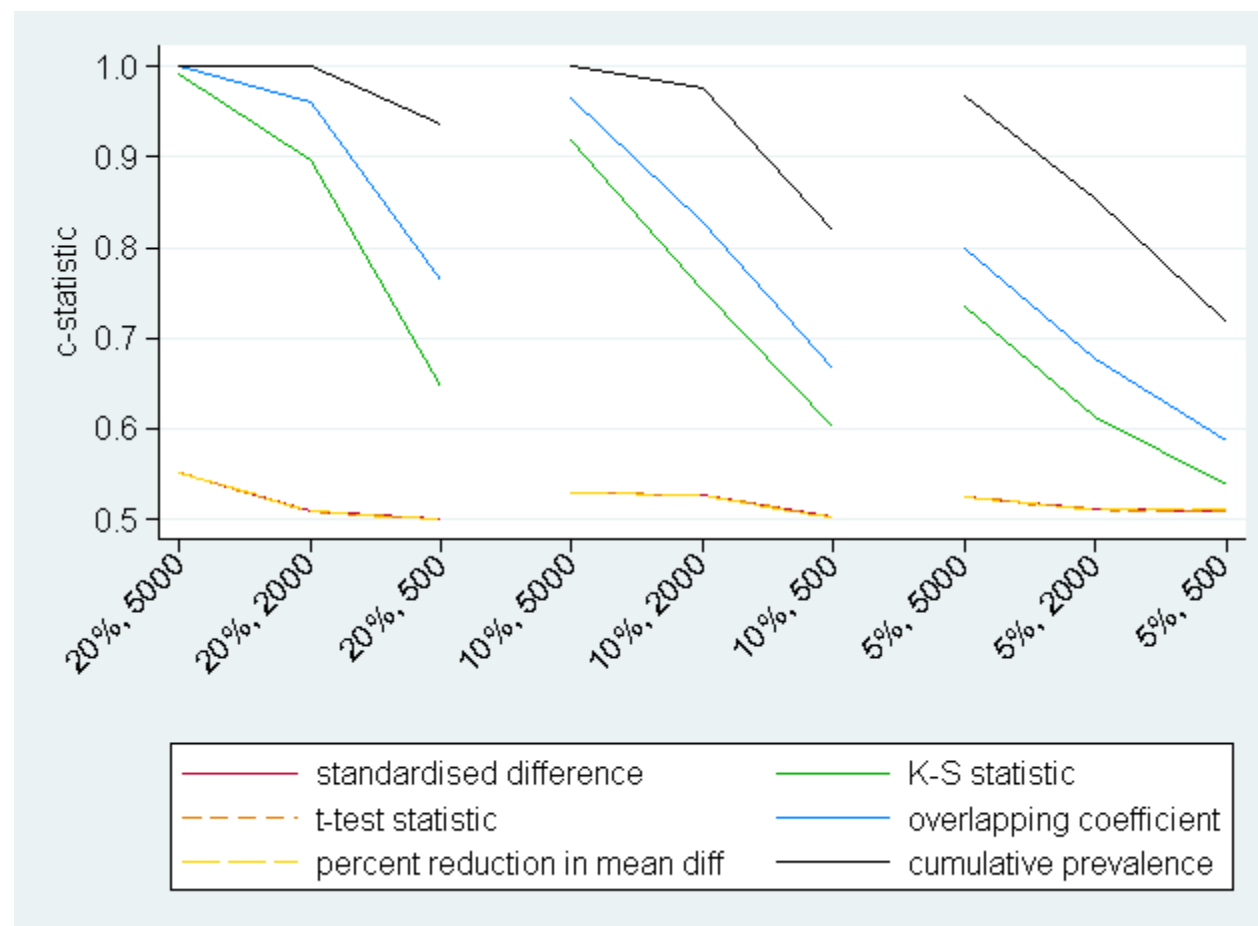
### Incorrect PS:

S1: $X_8 = 0.4(3.5^{X_1} - 1)$	Nonlinearity added (monotonic)	$X_8 = 0$
S2: $X_8 = X_4 X_5$	Binary-binary interaction	$X_8 = 0$
S3: $X_8 = X_4 X_1$	Binary-continuous interaction	$X_8 = 0$
S4: $X_8 = X_1 X_2$	Continuous-continuous interaction	$X_8 = 0$

## Data Analysis:

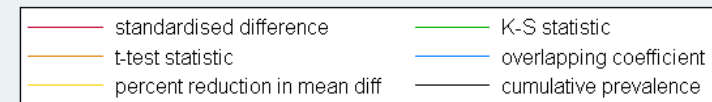
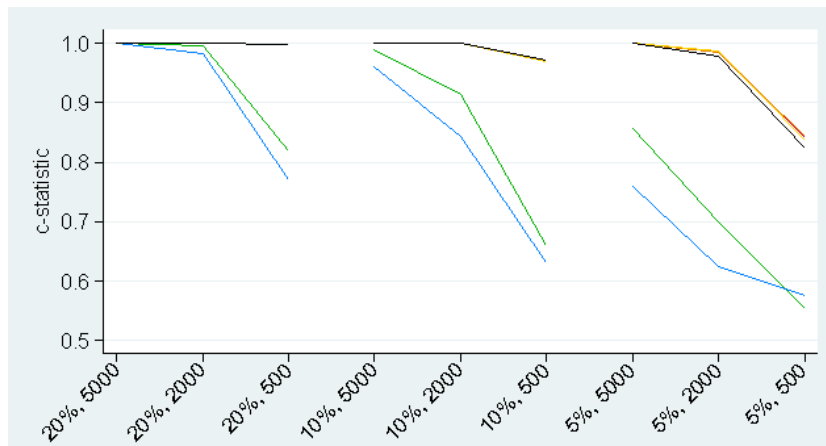
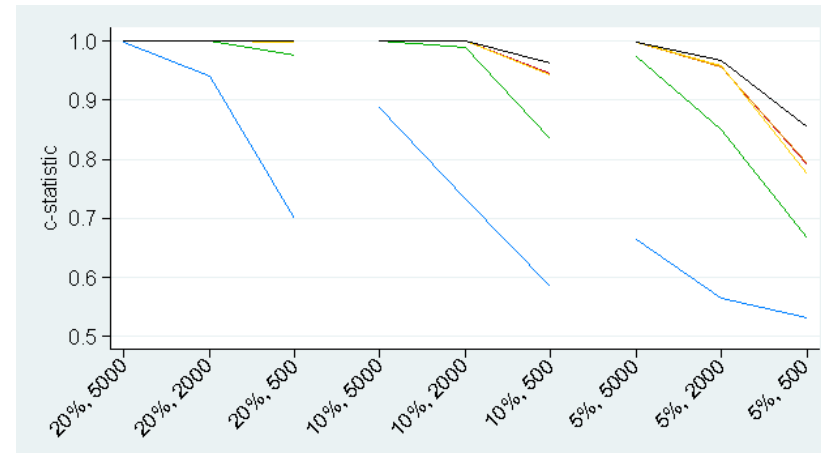
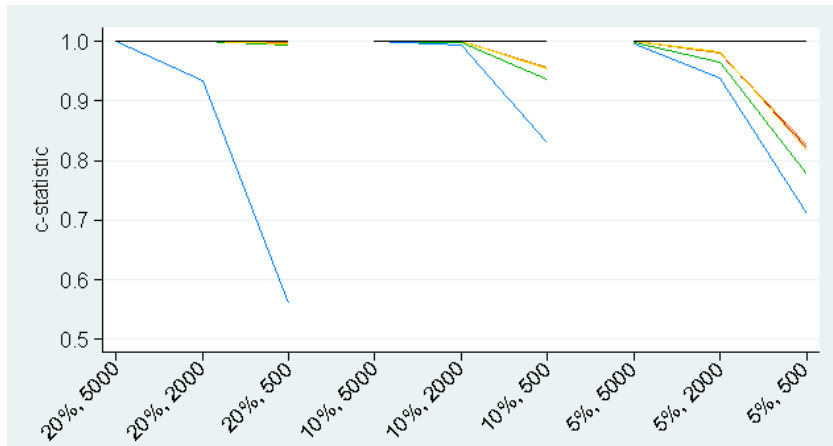
- Logistic Regression: PS indicator ~ PS diagnostic

# Scenario 1: Misspecification of a non-linear term



\*diagnostics used to assess balance/specification the non-linear covariate  $X_1$ ; balance assessed in PS-matched samples.

# Scenarios 2-4: Omission of an interaction term



## Figures:

- Scenario 2 (top left) binary-binary
- Scenario 3 (top right) binary-continuous
- Scenario 4 (bottom left) continuous-continuous

\*diagnostics used to assess balance/specification the interaction term;  
balance assessed in PS-matched samples.

# Overall diagnostics

## Disease risk score

- Disease risk scores (DRS) defined as predicted outcome under the control condition
- Standardised mean difference in DRS as a propensity score diagnostic [Stuart et al. 2013]

## Weighted average of balance

Let  $w_j$  denote the weight for covariate  $X_j$ .  
Then:

- $w_j = \gamma_j \text{Std.Dev}(X_j)$  [Caruana et al. 2015]
  - $\gamma_j$  is the coefficient for  $X_j$  obtained after regressing outcome on  $X_j$ .

Balance measured using either:

- Standardised difference (SD)
- Overlapping coefficient (OVL)
- Kolmogorov-Smirnov Statistic (KS)

# Simulated data

## Propensity score model:

- $\text{logit}(\text{PS}) = \alpha_0 + \alpha_1 X_1 + \alpha_2 X_2 + \dots + \alpha_7 X_7$

## Outcome model:

- $Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_7 X_7 + \beta_8 X_8$

## Variation between scenarios:

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S1:  $X_8 = 0$

Independent baseline covariates

S2:  $X_8 = 0$

Correlated baseline covariates

S3:  $X_8 = 0.2(6.0^{X_1} - 1)$

Monotonic non-linearity

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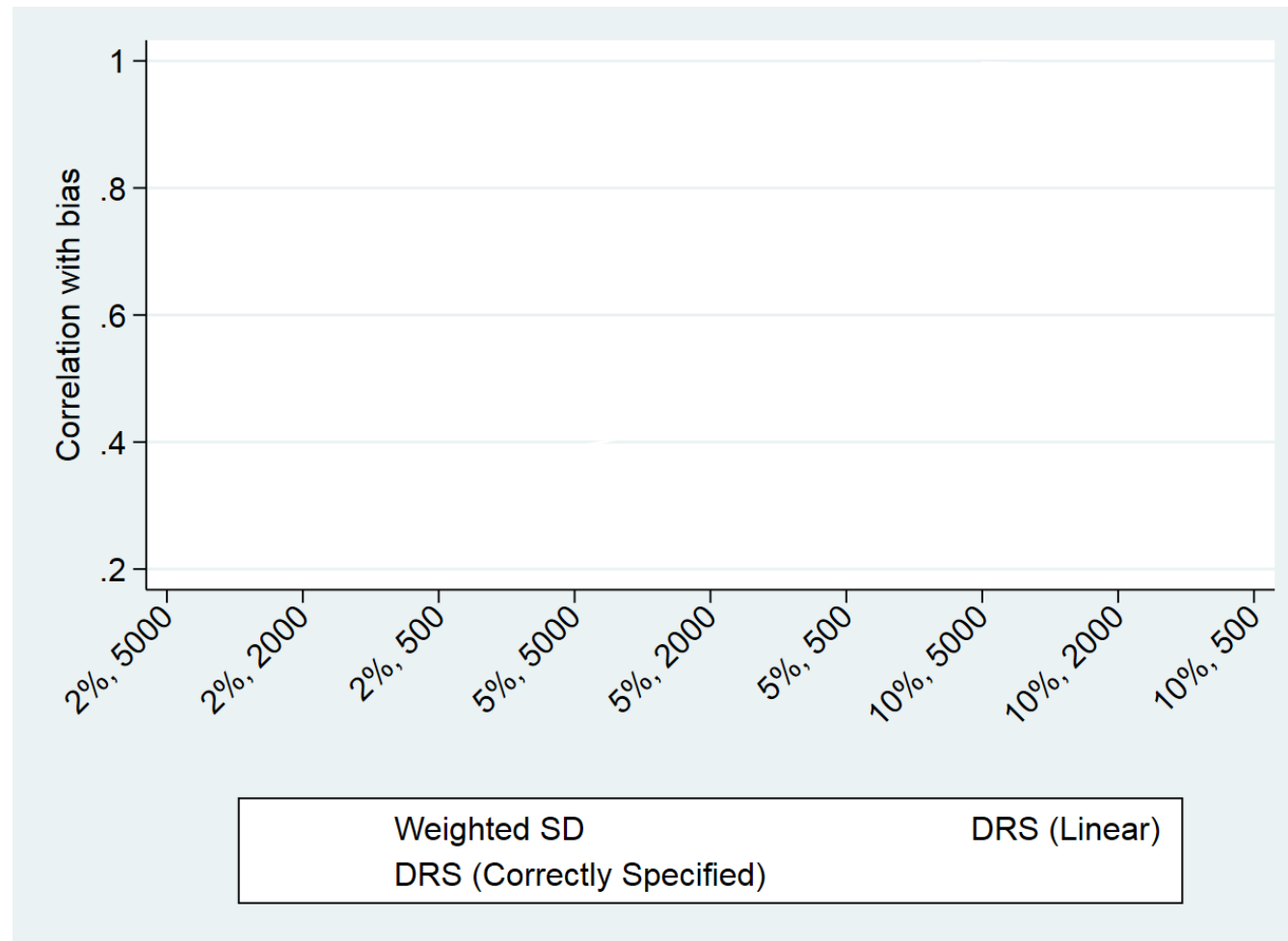
# Scenarios 1 and 2: Linear outcomes

*Table 1: Spearman rank correlation between overall diagnostics and bias (sample sizes 5000)*

Scenario	Weighted SD	Weighted KS	Weighted OVL	SD(DRS)
Scenario 1	0.992	0.137	0.012	1.000
Scenario 2	0.129	0.102	0.031	1.000

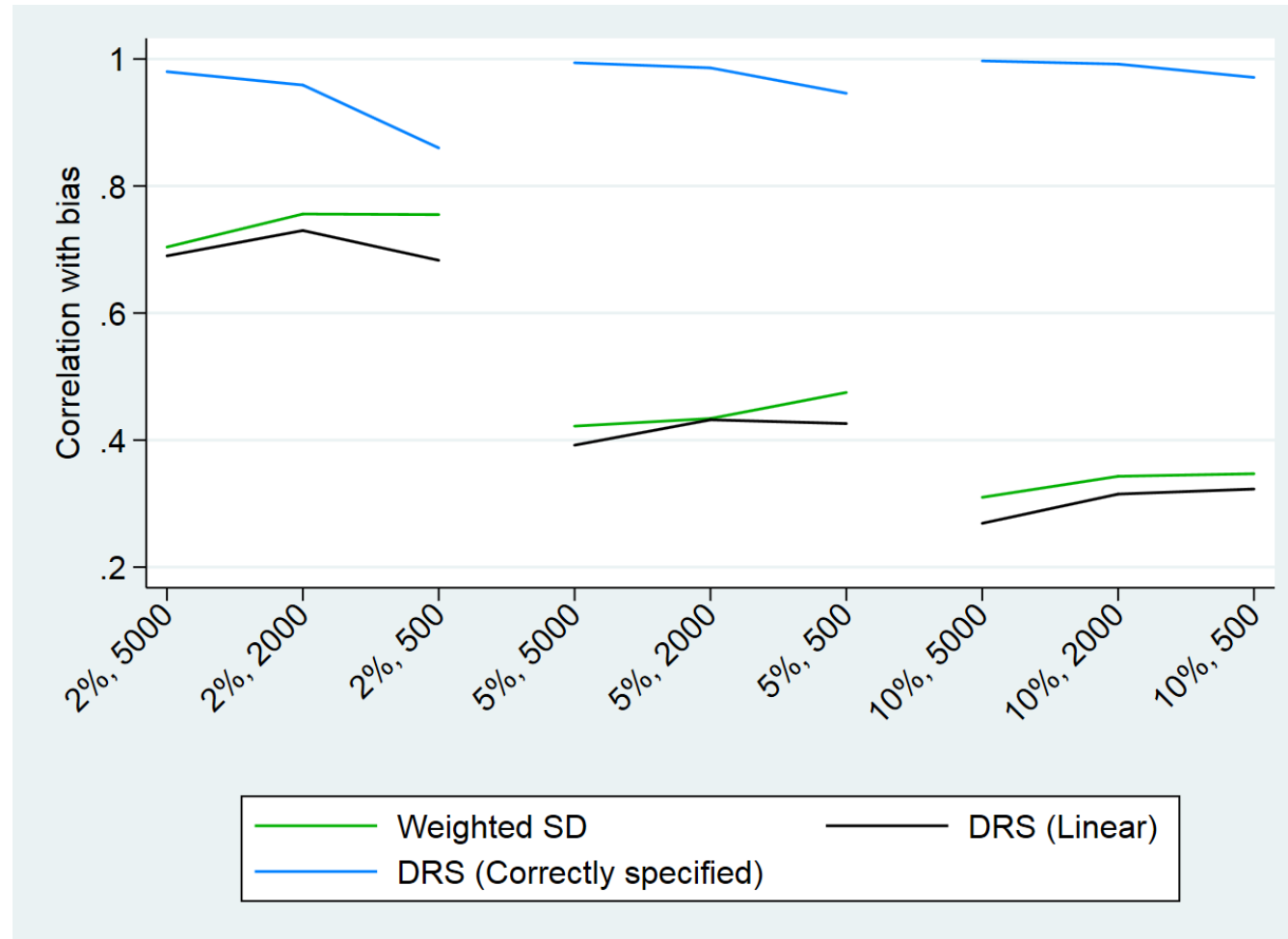
\*SD: Standardised difference; KS: Kolmogorov-Smirnov statistic; OVL: Overlapping coefficient; DRS: Disease risk score

# Scenario 3: Non-linear term in outcome model





# Scenario 3: Non-linear term in outcome model



# Conclusions

- **Cumulative prevalence (CP)** diagnostic most useful for identifying all types of propensity score misspecification.
- Standardised mean difference in the **disease risk score (DRS)** is a promising overall diagnostic.
- Main limitation:
  - DRS not robust to misspecifications in the outcome model
  - Could use CP diagnostics to check specification
  - Future research into different estimation methods for the DRS.

# Aims of research

## **Aim 1:**

Review and compare  
the existing  
propensity score  
diagnostics.



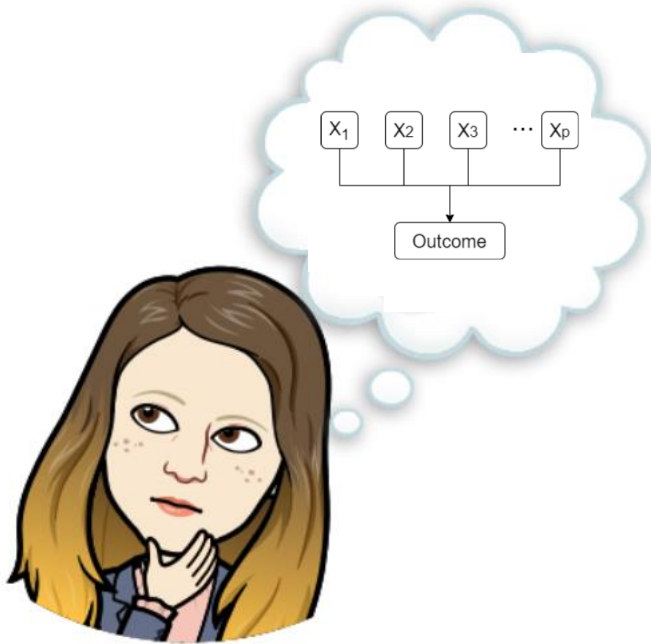
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# Proposed guidelines for propensity score assessment

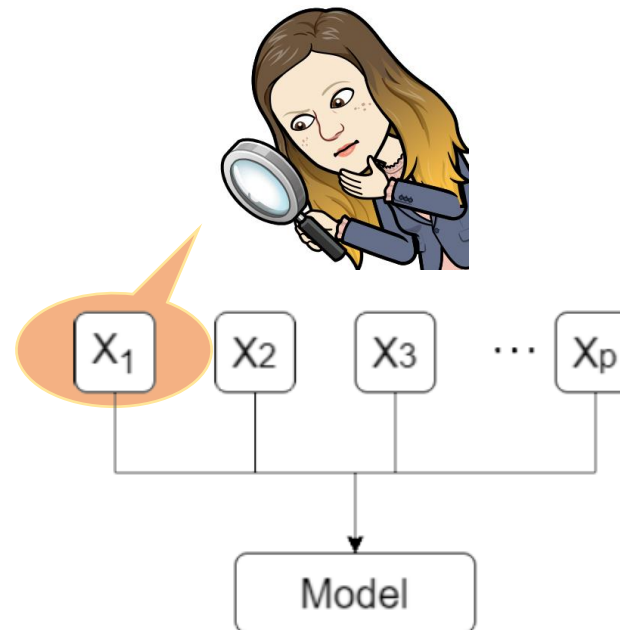
## STEP 1:

Choose variables



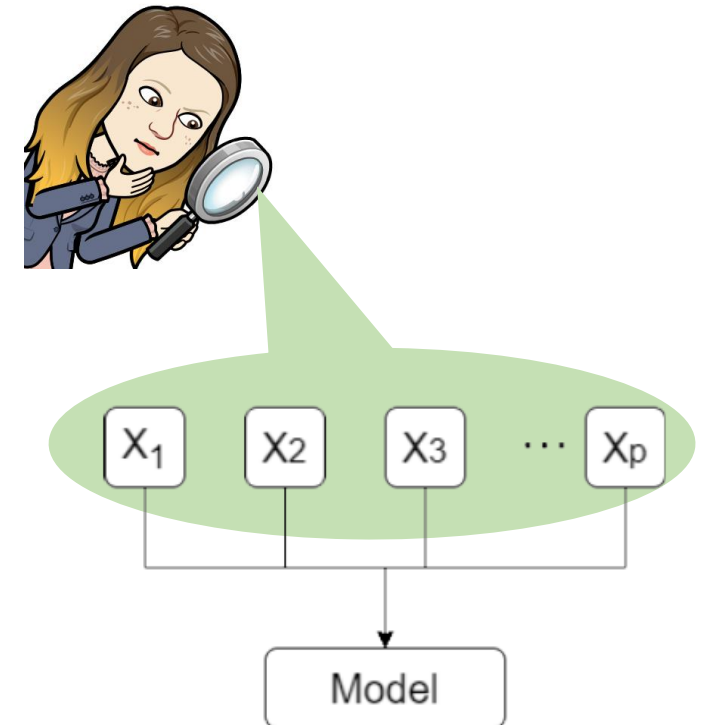
## STEP 2:

Check individual covariates using **CP diagnostics**



## STEP 3:

Check overall balance using **DRS**



# References

- [1] Stuart, EA et al. Prognostic score-based balance measures for propensity score methods in comparative effectiveness research. *Journal of Clinical Epidemiology*. 2013
- [2] Caruana, E et al. A new weighted balance measure helped to select the variables to be included in a propensity score model. *Journal of Clinical Epidemiology*. 2015.
- [3] Belitser, SV et al. Measuring balance and model selection in propensity score methods. *Pharmacoepidemiology and Drug Safety*. 2011.
- [4] Ali, MS et al. Reporting of covariate selection and balance assessment in propensity score analysis is suboptimal: a systematic review. *Journal of Clinical Epidemiology*. 2008.
- [5] Granger, E et al. A review of the use of propensity score diagnostics in papers published in high-ranking medical journals. *BMC Research Methodology*. 2020.

