

# Use of Data-Driven Simulations to inform Real-World Survival Analyses

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# Simulation Panel's Mission & Objective of the Current Study

## ■ STRATOS Simulation Panel's Mission:

**Promote more widespread and more accurate use of simulations in both methodological and applied statistical research, through enhancing their:**

- (i) Validity (lack of bias, neutrality) [Boulesteix et al., 2018]
- (ii) Reproducibility (accurate reporting, software availability) [Morris et al., 2019]
- (iii) Practical Relevance (plausibility)

## ■ Current Study's Goal (focus on Applied research):

To stimulate use of data-driven simulations  
to assess the impact of specific imperfections in the available data  
on the results of real-world time-to-event analyses



# Outline

- Background / Rationale
- Overview of the proposed Approach to Data-driven Simulations
- 2 Real-World Illustrations:
  - 1) *Omitting an important risk factor (potential Confounder)*
  - 2) *Imprecise timing of (interval-censored) events associated with a time-varying exposure*
- Conclusions



## Background: Need to be *more pro-active* when *dealing with imperfections of real-world data*

- Most real-world clinical/epi **papers recognize** (in Discussion) some **imperfections in the available data** and/or limitations of the study design **that may affect the accuracy (or even validity?) of the results**
- Traditionally, this was **limited to a lip service**, possibly with **vague qualitative comments aimed at minimizing the problem\*\***, e.g.:  
*“Lack of data on disease severity might have affected some of our estimates, but similar problems are common to this area of research.”*

\*\* Applies also to many papers co-authored by members of our team 😊

# Background: Main existing approaches for Quantitative Bias Analysis (QBA)

- Recent studies incrementally rely on QBA to get a *Quantitative assessment of the potential impact of selected common data imperfections* (e.g., unmeasured confounder or exposure measurement error) [Lash et al., 2009]
- **2 main existing Alternative QBA approaches** [Banack et al., 2021]:
  - 1) Analytical correction formulas for selected, relatively simple analyses, including e.g. E-values for unmeasured confounding [Vanderweele & Ding, 2017], **OR**
  - 2) Simulating Synthetic data, with data structure generally similar to the real-world data used in a given empirical study

Lash, Fink, Fox, Springer 2009.

Banack, Hayes-Larson, Mayeda, *Epidemiol Rev* 2021.

Vanderweele, Ding, *Ann Int Med* 2017.



# Limitations of existing QBA approaches

- (A) Neither approach accounts fully for the **complex Multivariable data structure** actually encountered in a given real-world dataset (relationships of different covariates with (i) each other, (ii) exposure, and (iii) outcome)
- (B) **Not well developed for Time-to-Event analysis** (for which the outcome is often dichotomized, i.e. Event Times are ignored) [e.g., Barberio et al., 2021]
- (C) **Not clear if/how to handle Time-Varying Exposures** (or time-varying covariates) ?



# Main features of Our approach: QBA via Data-Driven Simulations

- To address limitation (A), **we Combine:** (A1) **Observed Multivariable Real-world data\*\*** with (A2) **Simulating Additional Data items** (outcomes and/or covariates) based on carefully defined assumptions
- To address limitations (B) and/or (C), **we rely on the dedicated, validated, “Permutational Algorithm”** to simulate bivariate survival outcomes (follow-up duration & status) that (B) **copy the observed real-world distribution of the event times**, and (C) reflect the **assumed association(s) with time-varying exposure(s)/covariate(s)** [Sylvestre & Abrahamowicz, 2008]

\*\* Note this **Contrasts with traditional Methods-driven Simulations** (in *statistical papers*) that often assess or compare performance of selected methods across a range of (usually hypothetical) plausible data structures



# Implementation of Data-Driven Simulations: Preliminary Steps 1-3

## 3 Preliminary Steps:

*(see later slides for implementation in 2 illustrative examples)*

- 1) **Identify relevant Data Imperfection(s)** in your Available Real-World data & (if relevant) carry related Initial Data Analyses
- 2) Perform relevant, usually **Multivariable, Analyses of the Available Data to get 'naïve' estimates** (Not corrected for the Imperfection(s) identified in step 1) of the relationships between exposure, outcome, and covariates
- 3) Based on substantive knowledge and/or literature, **Formulate Assumption(s)** regarding how the *available data* can be modified or expanded to create the *oracle dataset* that is corrected for the expected impact of the imperfection identified in step 1 \*\*

(\*\* Several plausible alternative scenarios may be considered here, each implying repeating further steps 4-7)





# Implementation of Data-Driven Simulations: Main Steps 4-7

Data Simulations & Analyses (Steps 4-6) to be independently repeated across  $m$  (e.g., 1000) replications\*:

- 4) **Generate the 'Oracle data'** (Free of the imperfections of interest) that combine relevant empirical estimates from step 2 with additional data simulated according to the assumptions from step 3
- 5) **Modify the 'Oracle data'** from step 4 **to account for imperfection(s)** identified in step 1
- 6) **Analyze (6a) the 'Oracle' and (6b) the Modified (Imperfect) data** (from steps 4 *and* 5, respectively), *using the same methods*, and **contrast the corresponding results**
- 7) FINAL Step: summarize the results of step 6 across  $m$  replications and formulate the Conclusions regarding the Impact of the Data Imperfection

\* Steps 4-7 must be repeated for each alternative simulated scenario identified in step 3



# Example # 1: Impact of omitting cancer stage in a prognostic study of colon cancer mortality

- Goal of the analyses: estimate the independent (adjusted) association of **obstruction of the colon by a tumour (“exposure”)** with **all-cause mortality (“outcome”)** among patients diagnosed with colon cancer.
- Data source: publicly available dataset from the survival R package [Therneau, 2021], with N = 906 colon cancer patients, 175 (19.3%) with the colon obstructed, and 441 deaths during follow-up [Moertel et al., 1995]. Several time-invariant prognostic factors, measured at cancer diagnosis, are available, some associated with both (i) obstruction exposure and (ii) survival, calling for multivariable analyses.



# Example # 1: steps 1 - 3

- **Step 1 (Imperfection)**: available data do not include cancer stage at diagnosis, a powerful predictor of mortality in colon cancer [Quantin et al, 1999], with higher stage likely associated with both obstruction exposure (i.e. **potential unmeasured confounder**) and some measured covariates
- **Step 2 (Naïve analyses)**: multivariable Cox proportional hazards (PH) model, with adjustments for measured covariates (but NOT stage), yields **HR = 1.33 for colon obstruction (95% CI: 1.06; 1.68)**
- **Step 3 (Substantive Assumptions)**: **higher cancer stage** at diagnosis (dichotomized: stage III-IV versus I-II) assumed to have **HR = 4.0 for mortality**, and **OR = 1.2 for colon obstruction**, as well as associations with selected measured covariates.  
**4 alternative scenarios**: with the true **HR = 1.0, 1.3, 1.5 or 2.0 for colon obstruction**.

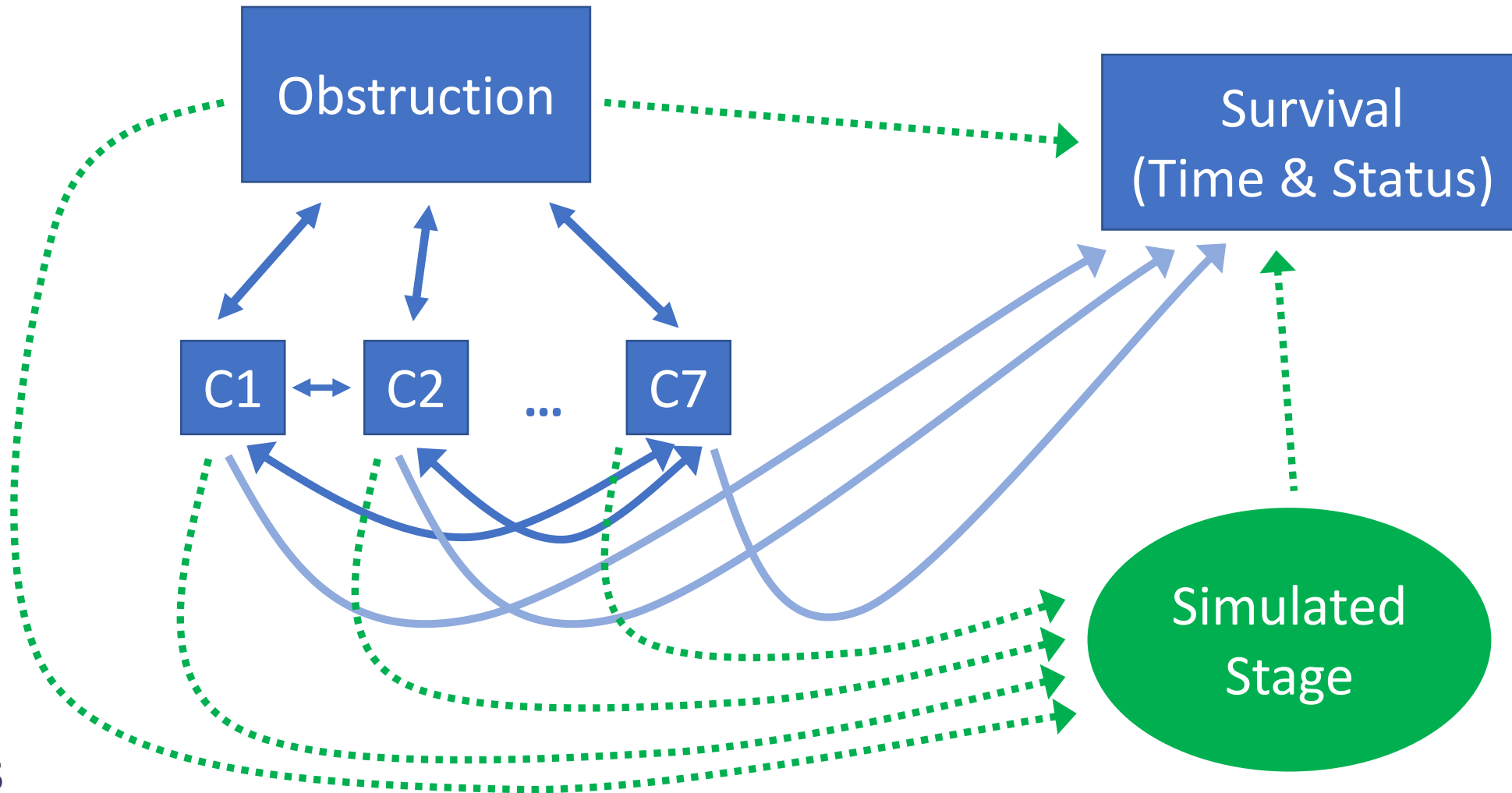


# Example # 1: Simulation Methods (*Step 4: “Oracle data” generation*)

- **Step 4.1:** Across scenarios & 1000 samples use (fixed) real-world data on:
  - (4.1.1) 906 multivariable X vectors (exposure + measured covariates)
  - (4.1.2) Outcomes: Times of 441 events (deaths) + 465 censorings
- **Step 4.2:** **{Stage | exposure, covariates}** generated **independently** in each sample, based on ORs assumed in step 3
- **Step 4.3:** Use **Permutational Algorithm** to assign each of the events or censoring obs. (with times from 4.1.2) to one of the 906 ‘expanded’ X vectors (from 4.1.1 + Stage from 4.2) based on the ‘true’ PH model, with:
  - (i) *Assumed HRs for Stage and Obstruction*, specified in step 3; and
  - (ii) For *measured covariates: ‘empirical’ adjusted HRs estimates* from step 2



# Example 1: Observed & Simulated Data Structure

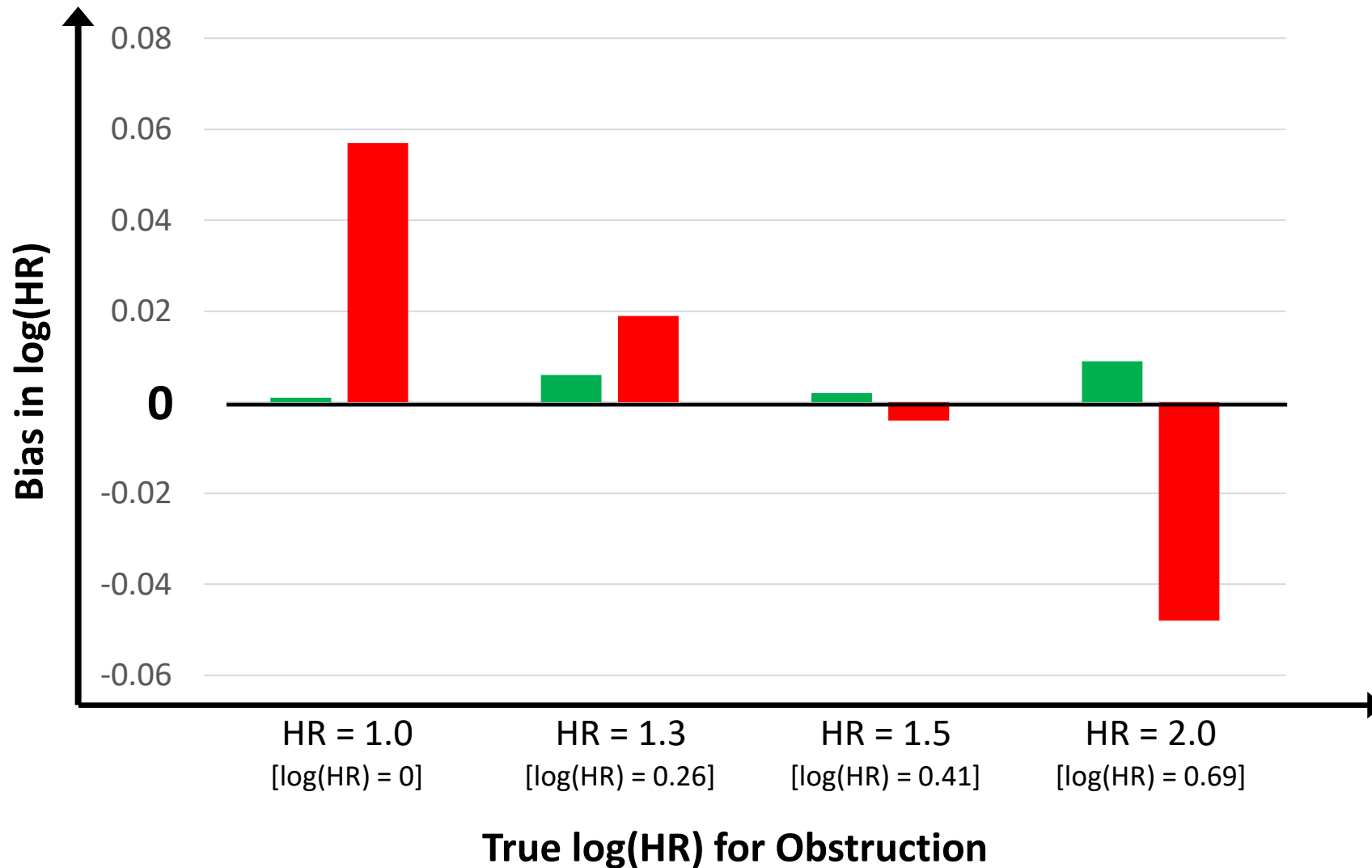


# Example # 1: *Steps 5 - 7*

- *Step 5 (Modifying Oracle data to **Insert the Imperfection**):*  
each of  $m$  samples simulated in step 4 modified by **Deleting “Stage”**
- *Step 6 (Analyses of **(6a) Oracle** vs **(6b) Imperfect data**):*  
**multivariable Cox PH models**, with Colon Obstruction & all Measured Covariates, **Only Difference: Stage (6a) Included vs. (6b) Stage Excluded**
- *Step 7 (Summarizing the results):*  
focus on **BIAS (Mean of 1000 Estimates – True) in Adjusted log (HR) for Colon Obstruction: (6a) vs. (6b)**



# Example # 1: BIAS in $\log(\text{HR})$ for Obstruction as a function of 'true' HR: Oracle vs. Imperfect data



# Example # 1: Conclusions

- **Lack of data on Cancer Stage has likely only a minor impact** on the accuracy of the adjusted log(HR) for Colon Obstruction (absolute Bias < 0.1, coverage rate of 95% CI:  $\geq 90\%$ )
  - Expected Bias varies depending on the strength of the (assumed) true association\*\*:
    - (i) Slight over-estimation of null or weak effects ( $1 \leq \text{HR} \leq 1.3$ ) *versus*
    - (ii) Slight under-estimation of stronger effects ( $\text{HR} \geq 1.5$ )
- \*\* Due to a **Combination** of (i) **Unmeasured Confounding** (OR = 1.2 for Stage-Obstruction) **vs.** (ii) **Non-Collapsibility** (HR = 4.0 for omitted Stage)





## Example # 2: Association of a Time-Varying exposure with an (imprecisely timed) Interval-Censored event

- Goal of the analyses: estimate the association of recent benzodiazepine use with cognitive impairment
- Data source: *synthetic data* based on real-world time-varying patterns of benzodiazepine use [Bartlett et al., 2004], with N= 1250 new benzodiazepine users generating 285 (23%) events of cognitive impairment during up to 3 years of follow-up. 2 measured time-invariant covariates: sex and age. **Binary Time-Varying Exposure (TVE) = Any Benzodiazepine use in the last 2 weeks.**



# Example # 2: Steps 1 - 3

- **Step 1 (Imperfection):** an event of cognitive impairment is detected only at the time of the first clinic visit after its actual occurrence, so the **actual event times remain unknown**, resulting in **Interval-Censored events** (see **Next Slide for Implications for TVE analyses**)
- **Step 2 (Naïve analyses):** 2 Cox PH models (adjusted for age & sex), with **alternative Event Times Imputation:**
  - (i) @ END of the Interval (visit when event 'detected'): HR(TVE) = **1.2**; (95%CI: **0.87**-1.68) vs.
  - (ii) @ MID-Point of the Interval (between 2 adjacent visits): HR(TVE) = **1.47**; (95%CI: 1.09-2.00)
- **Step 3 (Substantive Assumptions):** True (UN-known) event equally likely to occur at any time **within the between-visit interval** at the end of which it was detected.  
4 alternative scenarios: with HR = 1.0, 1.5, 2.0 or 2.5 for TVE (recent benzodiazepine use)

# Impact of inaccurate timing of interval-censored events on the associated 'current' values of time-varying exposure

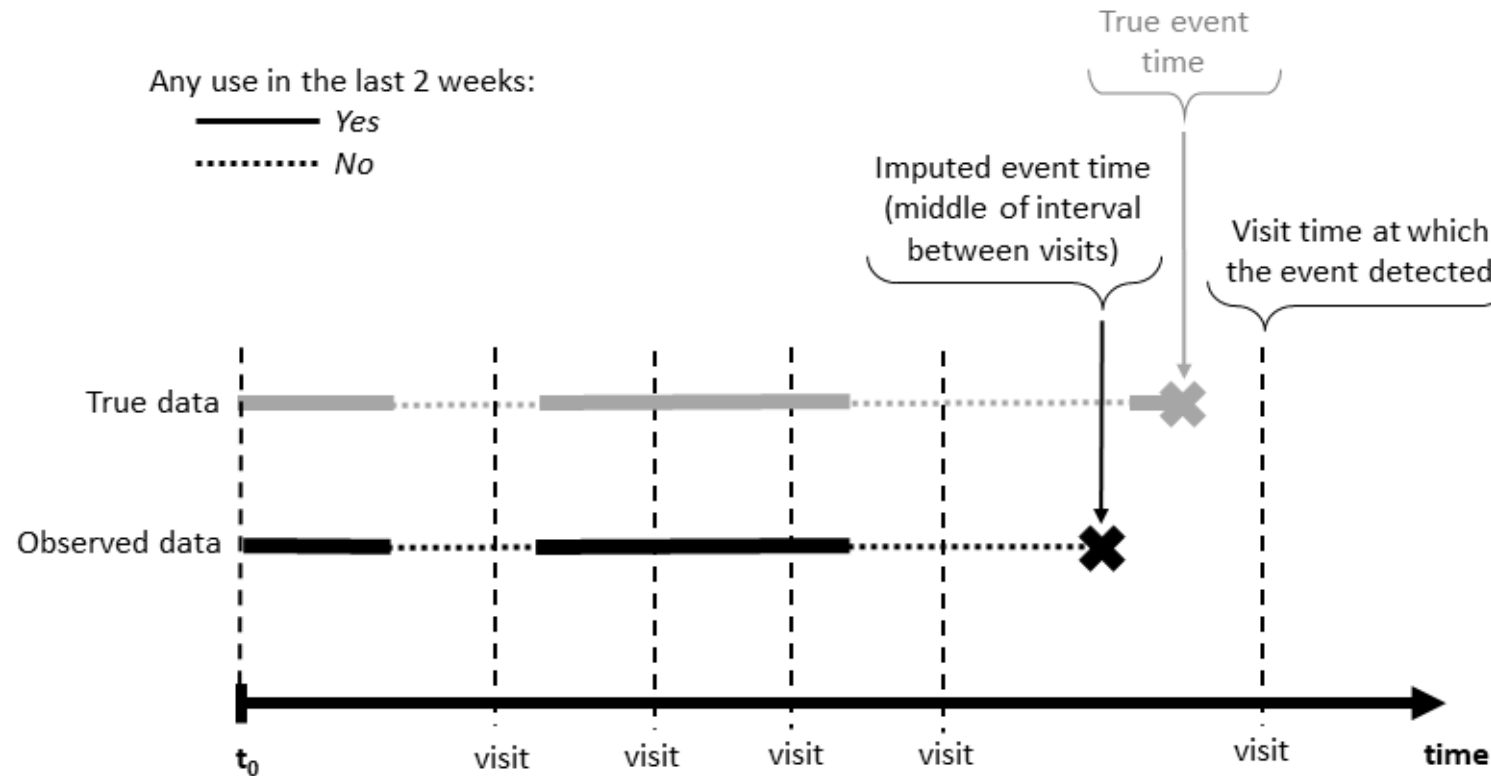
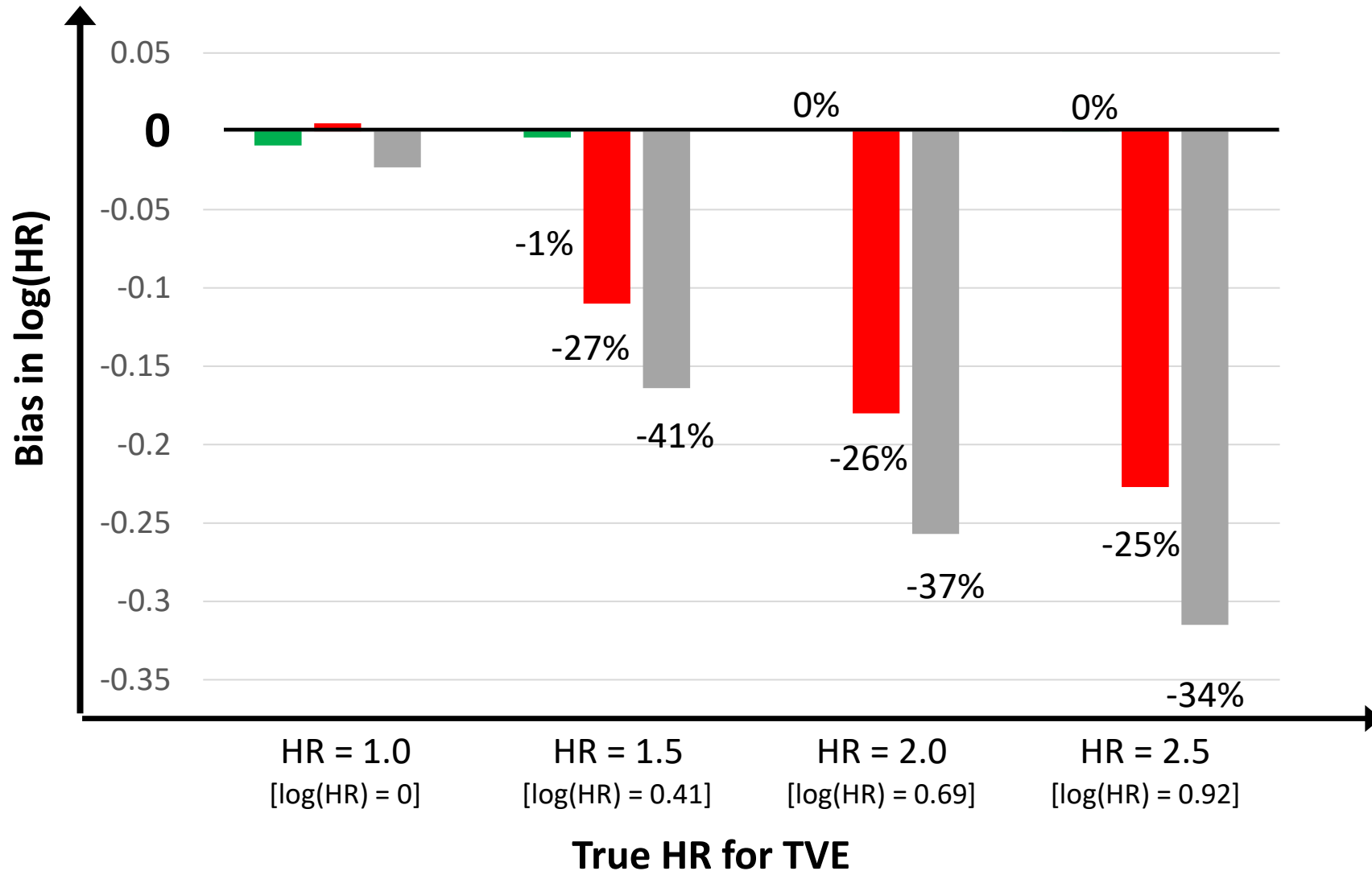


Illustration of the impact of inaccurate timing of interval-censored events for a hypothetical subject: the time-varying exposure metric "any use in the last 2 weeks" value differ between the true event time (exposure = yes) and the imputed event time (exposure = no) at the middle of the intervals between the visits when the event was detected and the preceding visit.

# Example # 2: Steps 4 - 6

- Step 4: (“Oracle data” generation):
  - (4.1) Across scenarios & 1000 samples use (fixed) observed data on: 1250 TVE time-vectors (daily benzo use) with corresponding age & sex
  - (4.2) “True” Time of event  $i = 1, \dots, 285$  generated (independently for each of  $m$  samples) from Uniform  $U \sim [t_{i,(j-1)}; t_{i,j}]$  over the interval between the earlier visit  $t_{i,(j-1)}$  and visit  $t_{i,j}$  when it was detected
  - (4.3) Use **Permutational Algorithm to assign each of the events** obs. (with times from 4.2) **to one of the TVE vectors** (from 4.1) based on the ‘true’ PH model, with: (i) assumed HR for TVE (from step 3) and (ii) empirical HR estimates for age and sex (from step 2)
- **Step 5: (Inserting the Imperfection): Exact (‘true’) event times were Deleted** and only the times of the visits when events were detected were reported
- **Step 6: (Analyses): 3 multivariable Cox models: (6a) Oracle data (True event times) vs. (6b) Event Times Imputed (Imperfect data) at: (6b1) End (detection visit) or (6b2) Mid-point of the interval  $[t_{i,(j-1)}; t_{i,j}]$**

Example # 2: BIAS in  $\log(\text{HR})$  for TVE as a function of 'true' HR: **Oracle** vs. **Imputation @: MID** vs **END**



# Example # 2:

## Summary of Results & Conclusions

- **(i) Imprecise Timing of the events (transient Cognitive Impairment) induces considerable Bias to the Null in the estimated HR for the Time-Varying Exposure (recent use of Benzodiazepines)**
- **(ii) Bias is systematically stronger for Imputing the events at the End (~ 35-40% relative bias) than at the Mid-point (~ 25% relative bias) \*\* of the interval between the adjacent visits**  

(\*\* Also, Root Mean Squared Error (RMSE) of End-imputed estimates is 20%-30% higher than for Mid-point-imputed estimates
- (iii) Given (i) & (ii), the **'naïve' estimate based on Mid-point Imputation** of event times [**HR = 1.47** (95%CI: 1.09-2.00)] provides a solid **evidence of Risk Increase** associated with a recent Benzodiazepines use **but likely Underestimates its strength !**



# Conclusions

- **Carefully designed Data-Driven Simulations** can provide valuable insights regarding the **expected impact of a specific Data Imperfection** or Design Limitation on the results and conclusions of a **particular Real-World study**
- **Our methods extend the QBA toolbox to address complexities of:**
  - *Multivariable* data structures
  - *Time-to-Event* (Survival) analysis
  - *Time-Varying Exposures/Covariates*

**but further real-world applications are necessary to fully assess their practical usefulness/potential...**



Thank you!

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