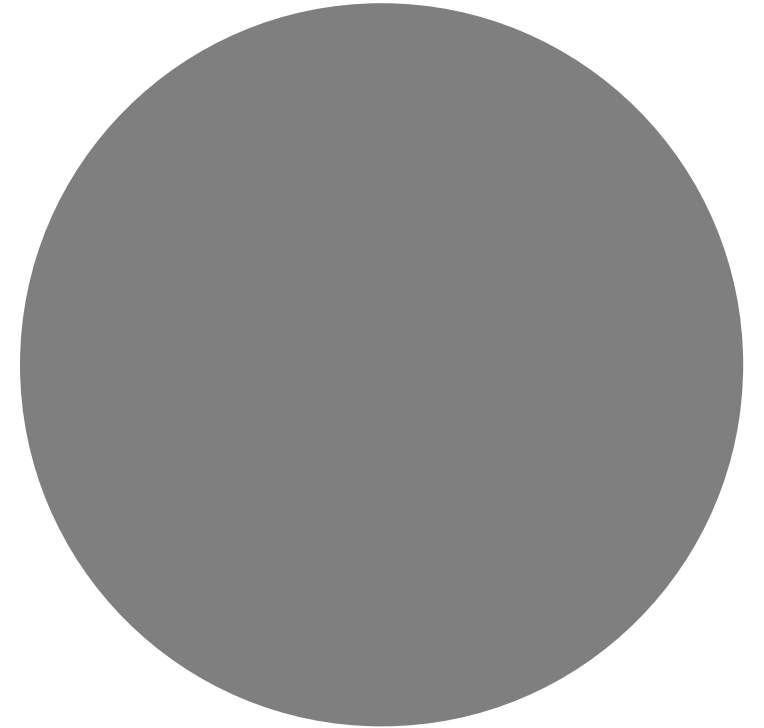


# A Practical Review of Real World Data for Effectiveness Decisions

Case Studies of Using Historical Data in  
Developing Novel Therapies by DIA  
NEED

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Ziliang Li, Chenkun Wang  
Vertex Pharmaceuticals Inc.



# Historical control: not a new concept

Compared to a different group of people with similar situation treated in a different manner – **Historical Control**

Or compared to the same group of people previously untreated – **Self-control**

Randomized clinical trial was widely recognized.

Gold standard of clinical trial

- Remove the potential bias
- Produce compared groups

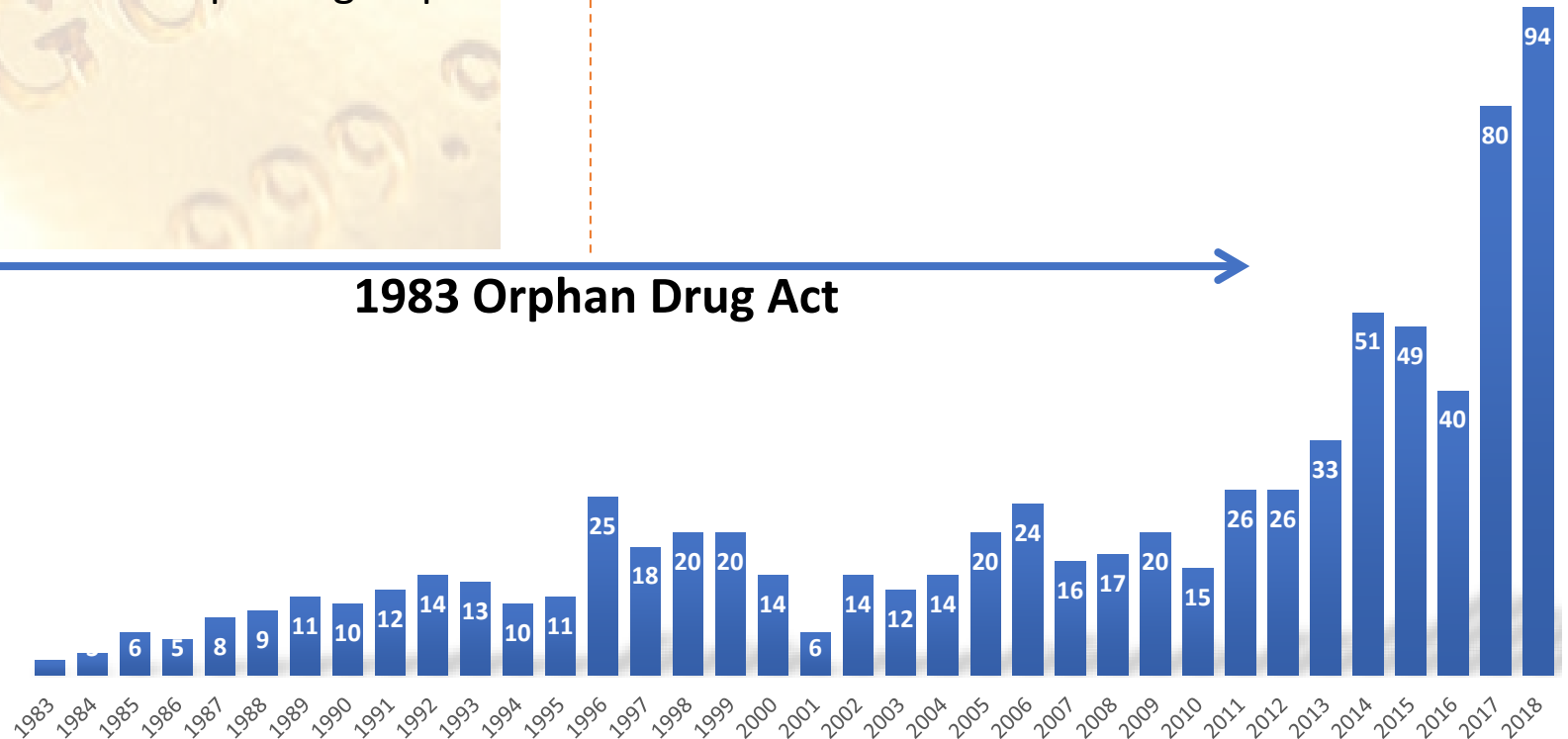
Back to “historical control” ?

- Accessibility to massive historical data – **resources**
- Advances in genetic research for rare disease – **environment**

1950

1983 Orphan Drug Act

## Number of Orphan Indications Approved in the United States 1983–2018:



# DIA-ADSWG Nature and Extent of Evidence Needed for Rare Diseases (NEED) Subteam

## Co-leads:

Rui (Sammi) Tang , Servier Pharmaceuticals

Mercedeh Ghadessi, Bayer

## Team members (alphabetical order):

Chaoqun Mei, University of Wisconsin-Madison

Chenkun Wang, Vertex Pharmaceuticals

ChunQin(C.Q.) Deng, United Therapeutics Corp

Kiichiro Toyozumi, Shionogi Inc.

Jeffrey Schwartz, Pfizer

Joey Zhou , Q2BI

Lihua Yue, Celgene

Lixia Zhang, BERG health

Robert Beckman, Georgetown University

Rong Liu, Celgene

Ziliang Li, Vertex Pharmaceuticals



GEORGETOWN UNIVERSITY



SHIONOGI INC.



WISCONSIN  
UNIVERSITY OF WISCONSIN-MADISON



# A Practical Roadmap

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1 Resources of HD



2 Should we use HC?

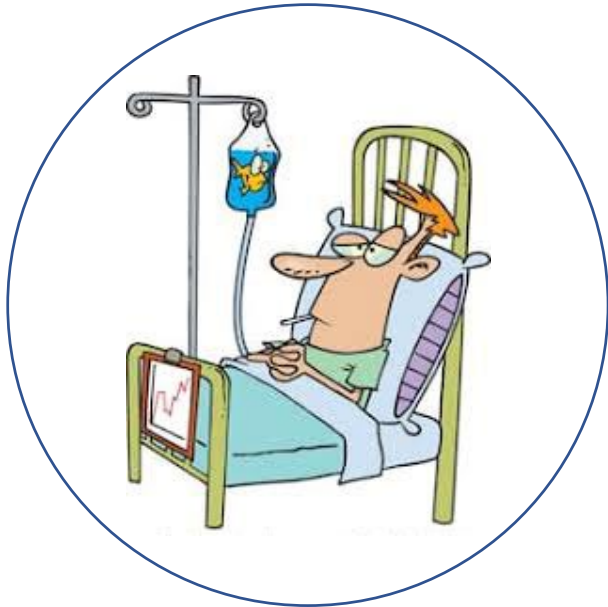


3 Disadvantages of using HC



4 Minimizing disadvantages

Medical chart



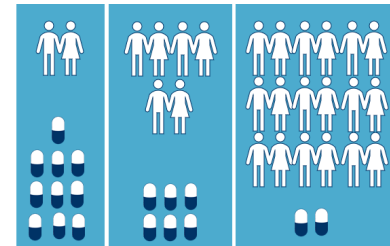
Patient registry



Claim data

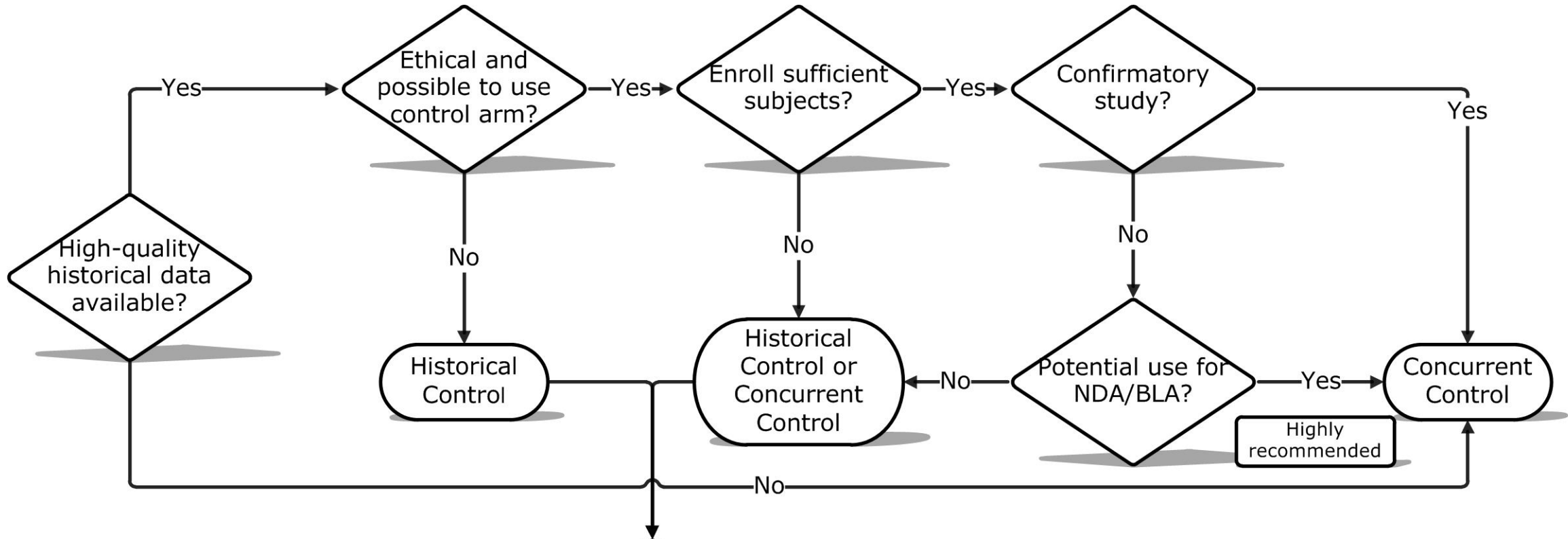


Previous clinical trials



Natural history studies

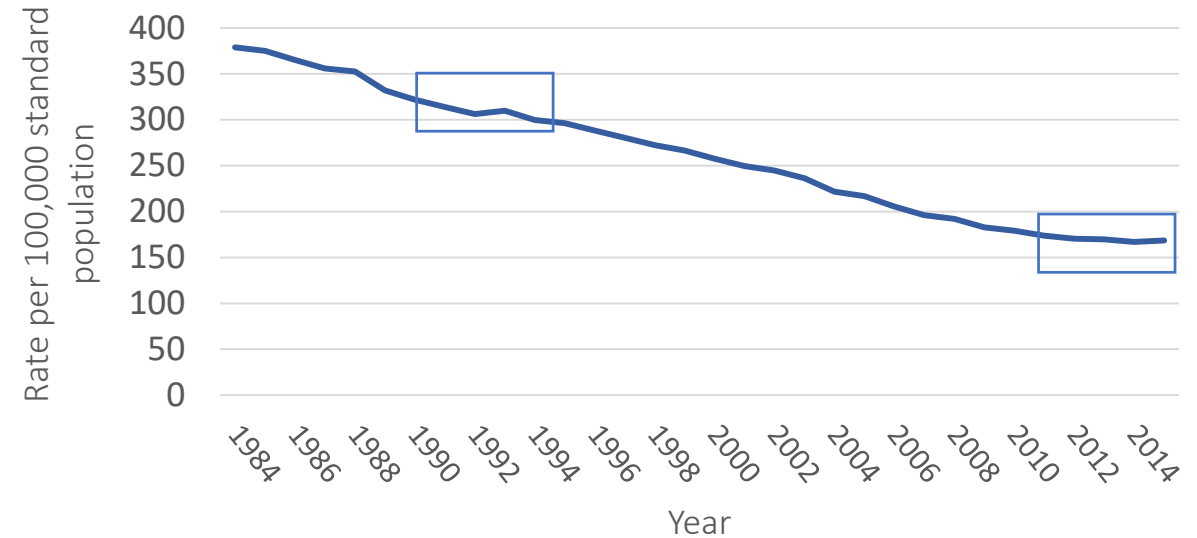




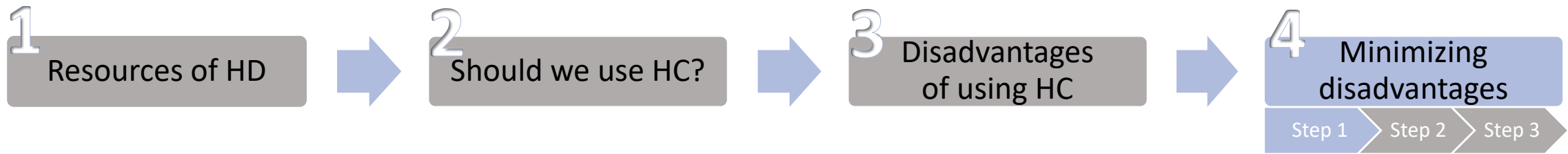


- Not randomized
  - Population between groups are not comparable
- Not concurrent
  - Improved standard care
  - More sensitive technology
- Data collection
  - Lack of blinding
  - Accuracy
  - Missingness

Age-adjusted Death Rates for *Heart Disease*:  
United States, 1980-2015



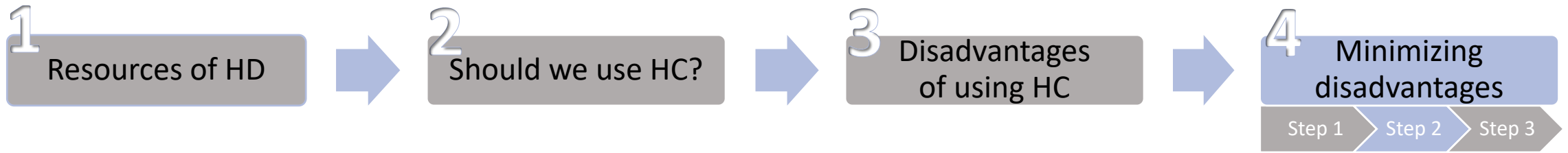
Source: <https://data.cdc.gov/NCHS/NCHS-Age-adjusted-Death-Rates-for-Selected-Major-C/6rkc-nb2q>



## Step 1: At the design stage

- Choose high-quality historical control(s)
  - Checklists to evaluate real-world data
  - Flowcharts by Clinical Trials Transformations Initiative (CTTI) to evaluate registry data
- Design the current study closely to the selected historical control(s)
  - Pocock's (1976) criteria
    - Inclusion/exclusion criteria for patient population
    - Type of study design
    - Exact definition of the outcome
    - Quality of study execution and management;
    - Potential biases due to time trends





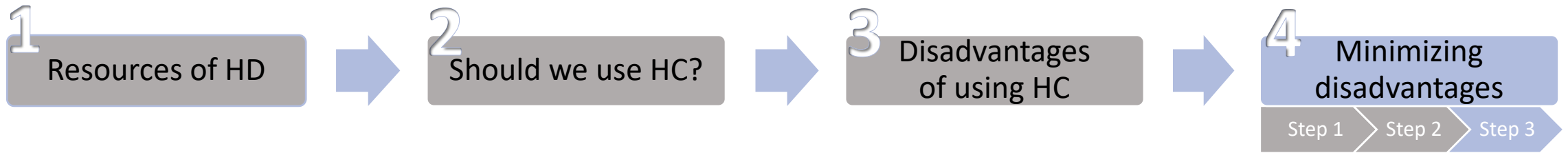
## Step 2: At the analysis stage

Meta-analysis

Casual inference/  
observational study

Historical borrowing

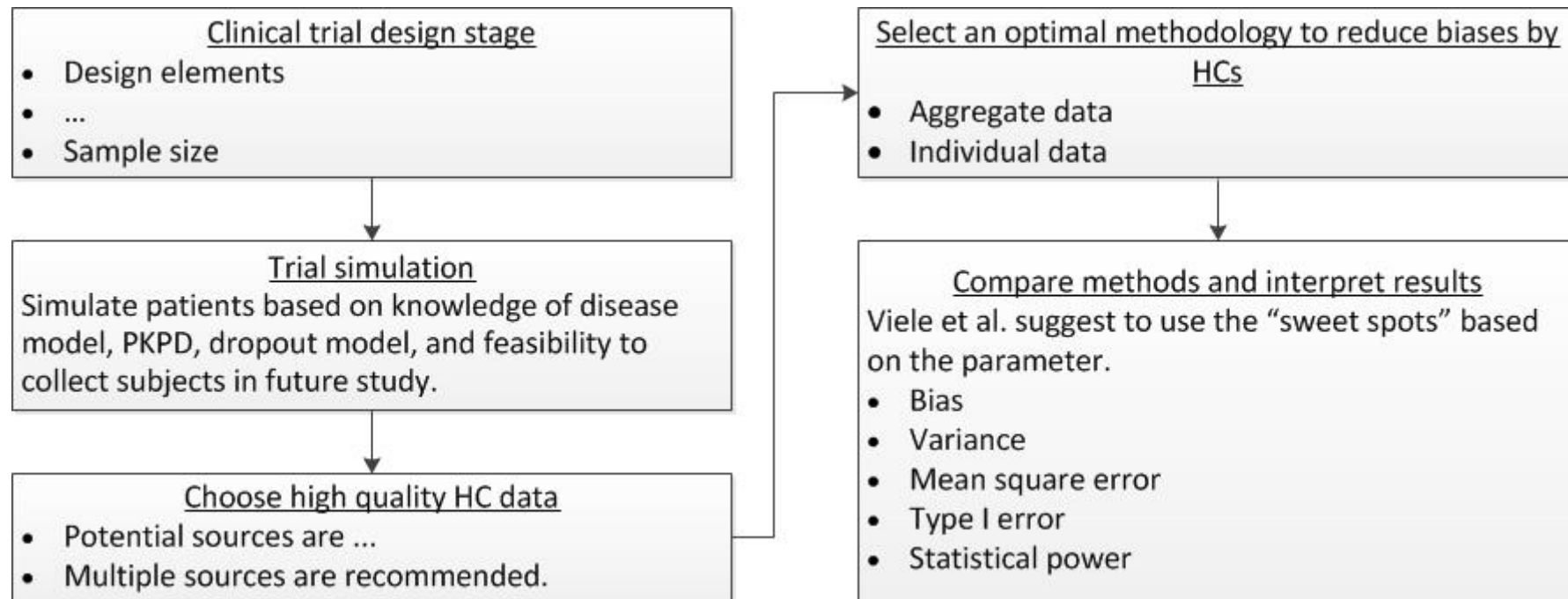
- Network meta-analysis
- Simulated treatment outcome
- Matching-adjusted indirect comparison
- Matching based on covariates or propensity score
- Weighting by the inverse probability of treatment
- Etc.
- Bias-variance (Pocock SJ, 1976)
- Test-then-pool (Viele K, et al, 2014)
- Power prior (Ibrahim JG, et al, 2000)
- Meta-analytic Combined (Neuenschwander B, et al, 2010)
- Meta-analytic Predictive (Neuenschwander B, et al, 2010)
- Etc.



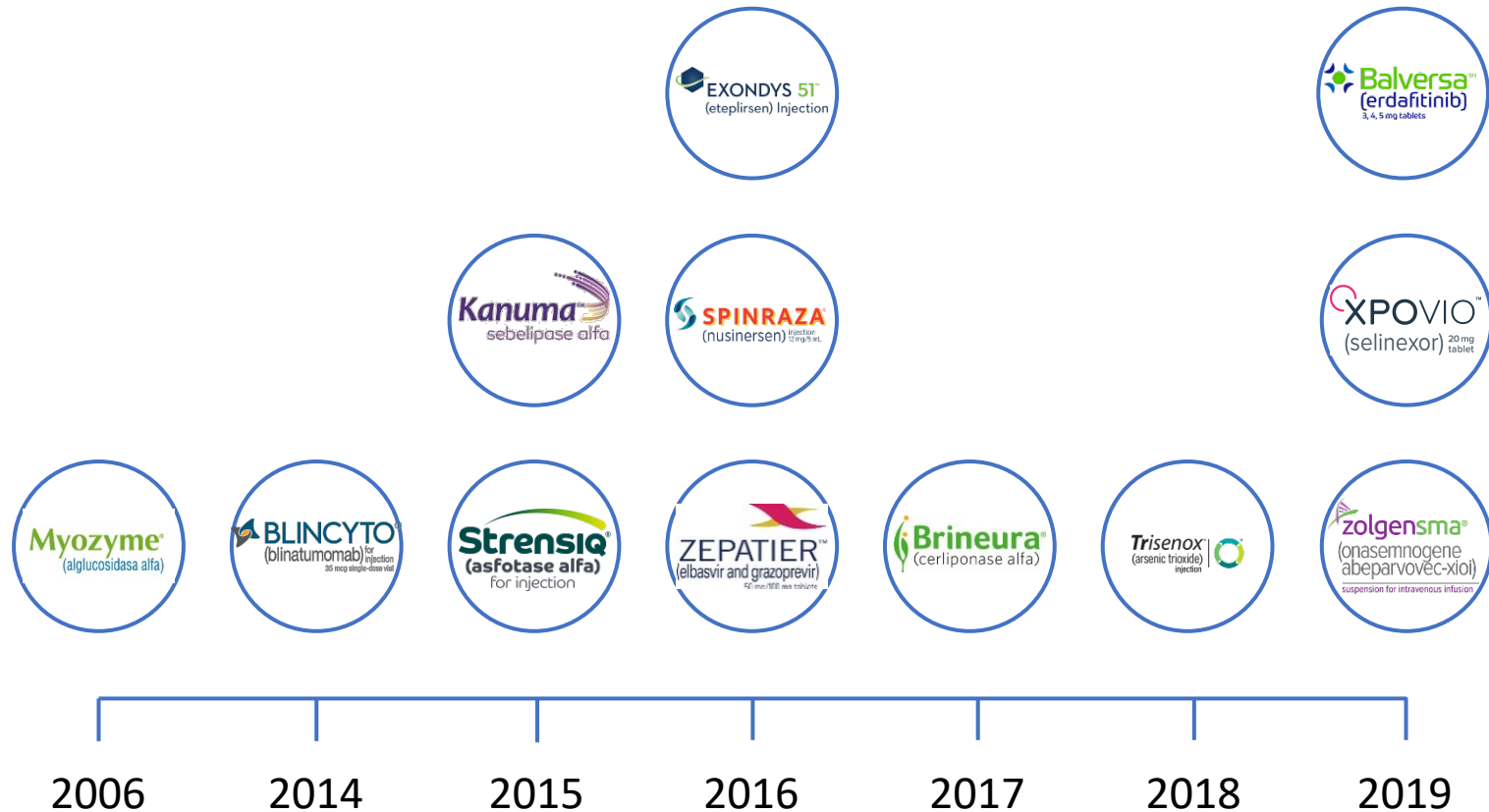
## Step 3: Simulation and sensitivity analysis

No one solution that works in all situations.

The recommendation is to perform simulations as the following process:



# Case Studies



# Case Studies - Disclaimers



This is a **biased** sample of drug development programs



Numerous factors impact how the historical data are utilized in the development



Historical data/RWD have a much broader spectrum of applications than filing

# 1+2

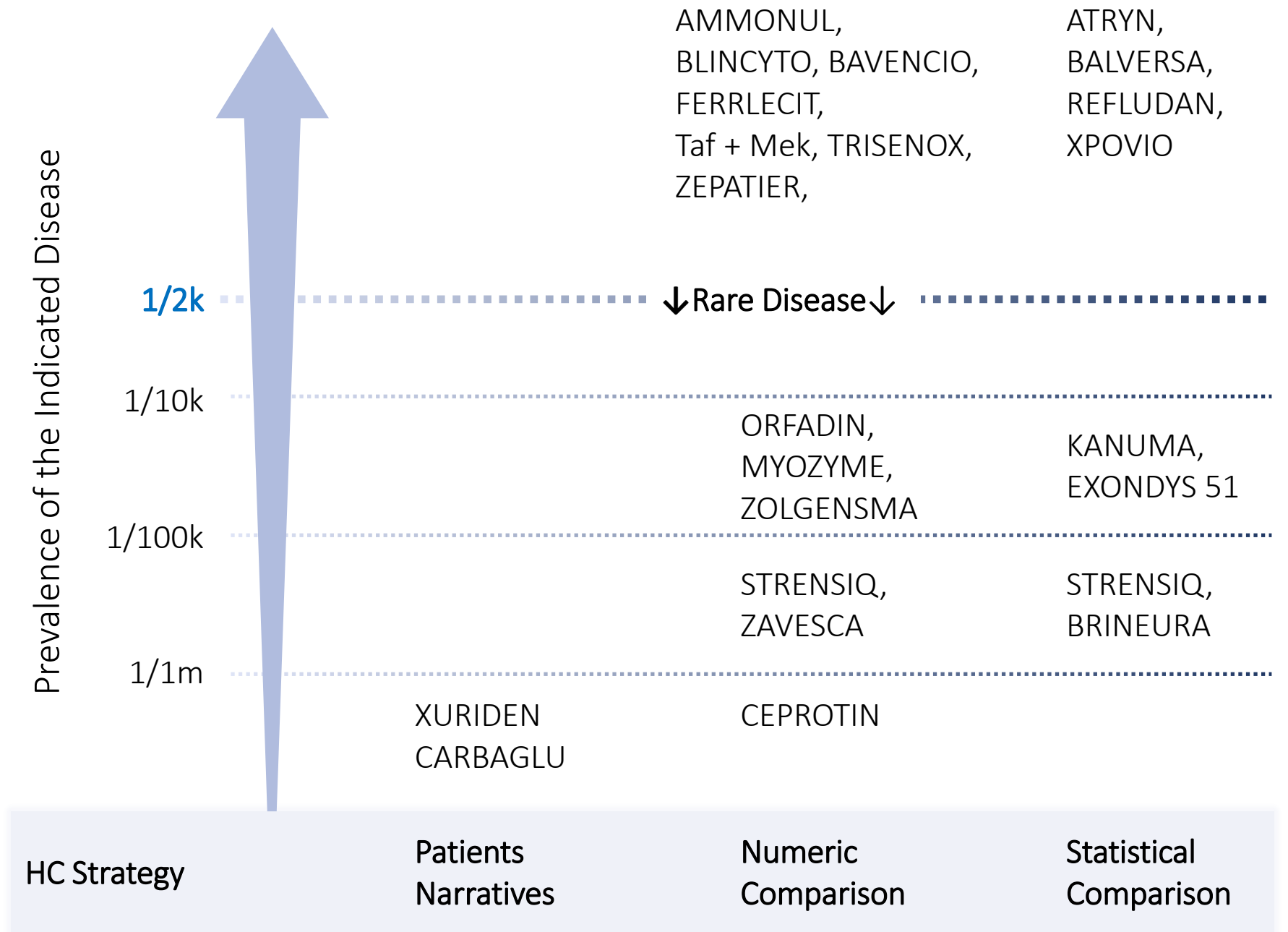
## HC Strategy

NEED team reviewed filings from 22 drugs.

Source of HD:

- Chart review, database, publication (12)
- Registry (6)
- Previous clinical trials (1)
- Natural history (3)

Strategy to utilize historical data (to compare against IMP) varies.



## 1+2

## HC Strategy

Many other factors impact the drug development program, take type 1 spinal muscular atrophy as an example.

| Drug      | US Approval Year | Modality              | Source of HC                                   | Method to utilize HC   |
|-----------|------------------|-----------------------|--|--|
| SPINRAZA  | 2016             | Intrathecal injection | Natural history                                | Served as the control arm in a Ph2 dose-ranging study intended as basis for NDA <sup>1</sup> |
| ZOLGENSMA | 2019             | Gene Therapy          | Prospective Natural history study <sup>2</sup> | Characterized disease progression to support the pivotal study                               |
| Risdiplam | In development   | Oral                  | Natural history                                | Establish an efficacy threshold <sup>3</sup>   |

<sup>1</sup> [https://www.accessdata.fda.gov/drugsatfda\\_docs/nda/2016/209531Orig1s000MedR.pdf](https://www.accessdata.fda.gov/drugsatfda_docs/nda/2016/209531Orig1s000MedR.pdf)

<sup>2</sup> Finkel RS, et al. *Neurology*. 2014;83:810-7.

<sup>3</sup> Reid and Burger. 3rd EFSPi Workshop on Regulatory Statistics, 2018.

## Selection Bias, and How to Minimize It

| Compound                             | BLINCYTO   | MYOZYME  | XPOVIO  |
|--------------------------------------|--|--|---|
| <b>Pivotal Study</b>                 | Ph2 fixed-dose single-arm study using a Simon 2-stage design with extension  | Ph2/3 open-label, dose-ranging study in patients <6 mos with infantile-onset Pompe disease   | Ph2b single-arm, open-label study in patients with relapsed/refractory multiple myeloma   |
| <b>Primary Endpoint</b>              | Complete remission or CR with partial hematological recovery (CR/CRh*) rate within first 2 cycles of treatment   | Ventilator-free survival at 12 mos of age  | Overall response rate (proportion of patients with a partial response or better)  |
| <b>Source of Historical Control</b>  | Retrospective historical database by combining existing databases from EU and US source  | Retrospectively natural history cohort (identified based on entry criteria used in the pivotal study)  | Retrospective historical cohort identified from Flatiron health analytic database   |
| <b>Limitations and Review Issues</b> | <ul style="list-style-type: none"> <li>The estimated CR rate has accounted for important differences between patients in the HCs and MT103-211</li> <li>Each of the historical studies provided large # and variety of patients</li> </ul> | <ul style="list-style-type: none"> <li>Patients were 'healthier' in the clinical study than HC given they need to survive long enough to enter the study and start treatment</li> <li>Natural history study suggested improved survival overtime, the benefit of Myozyme is likely to be over estimated</li> </ul> | <ul style="list-style-type: none"> <li>HC selection criteria systematically ensure that the clinical cohort will have longer expected OS compared to HC cohort (HC has 31% subjects with baseline ECOG missing)</li> <li>Imbalances in baseline characteristics between the HC and clinical cohorts were noted</li> </ul> |

## Assessment Bias, and How to Minimize It

| Compound                             | KANUMA   | BRINEURA   | EXONDYS 51   |
|--------------------------------------|--|--|--|
| <b>Pivotal Study</b>                 | Ph1/2 open-label, single-arm study in infants with rapidly progressive disease due to LAL deficiency | Ph1/2, first-in-human, single-arm, open-label, dose-escalation   | Ph2 randomized, placebo-controlled study followed by OLE   |
| <b>Primary Endpoint</b>              | Time to death from birth up to Month 12  | % of patients with an absence of an unreversed (sustained) 2-point rate (slope) of decline or a score of 0 in the Motor total score over 96 weeks  | % dystrophin positive fibers<br><ul style="list-style-type: none"> <li>• 2<sup>nd</sup> endpoint: 6-minute walk test (6MWT)</li> </ul>   |
| <b>Source of Historical Control</b>  | Retrospective natural history cohort (baseline matched)  | Baseline matched natural history cohort based on registry data   | <i>Post hoc</i> historical cohort identified using 5 relevant baseline factors from 2 DMD patients registries  |
| <b>Limitations and Review Issues</b> | Sponsor took the recommendation from DGIEP and used survival as the primary endpoint                 | The applicant submitted evidence is not sufficiently strong regarding the CLN2 rating scale comparability between HC and pivotal study<br><ul style="list-style-type: none"> <li>• FDA's efficacy assessment focused only on the motor domain and ignored the language domain score</li> </ul> | <ul style="list-style-type: none"> <li>• Performance on 6MWT can be improved by motivation in a clinical trial setting, but limited in real life due to concerns of failing or injury</li> <li>• 6MWT collected in the HC group was not performed in the same investigative site or same investigator</li> </ul> |



3+4

Working with HC

## Missing Values, and How to Minimize It

| Compound                             | ZEPATIER  | BALVERSA   | TRISENOX   |
|--------------------------------------|---|--|--|
| <b>Pivotal Study</b>                 | Two trials with Placebo control and two trials without Placebo  | Ph2 two-arm, open-label efficacy study   | Ph2, single-arm, multicenter study<br>Ph1, single-arm, single institution study  |
| <b>Primary Endpoint</b>              | % of subjects achieving SVR12 after 12-week treatment   | % of subjects with objective response rate (complete response + partial response)  | % of subjects with complete remission  |
| <b>Source of Historical Control</b>  | Previous clinical trials and publications   | Real-world Flatiron-FMI clinic-genomic database  | A case series from medical chart stored by a single institution  |
| <b>Limitations and Review Issues</b> | Using data from previous clinical trials to form HC substantially reduce the impact of missing values | It may be not practical to use a multiple imputation approach due to small sample size (16-25 patients) in the control cohorts, as about one third of subjects did not have an ECOG value. | The data is limited in information and does not include significant information such as bone marrow biopsy results, drugs involved in prior treatments, dosages and laboratory data. |

## Other Statistical Considerations

For time-to-event endpoint, it is important to properly define the index date to enable a fair comparison between HC and clinical cohort

- The start time in clinical trial is often fixed, e.g., date of first dose
- Defining start time in HC could be tricky, e.g., date of diagnosis, last day of prior therapy, the first time a clinically relevant biomarker exceeding a threshold, etc.

➤ Avoid creating the ‘immortal time’ bias!

Slope of decline is often used as a measure of disease progression

- With repeated and scheduled assessments, slope can be estimated with precision in clinical trial setting
- Challenging to get the same precision in HC (irregular timepoint, insufficient duration)

➤ Responder analysis may be more robust

## Some Commonalities in the Statistical Review

General *guidelines* when evaluating the use of historical control as a comparator

- The course of the untreated disease within a patient population is well understood to be uniform with outcomes that can be predicted reliably
- A valid historical control from a natural history study must have the same eligibility requirements, medical workup, and clinical evaluations as the clinical trial
- Using a historical control is most likely to be persuasive when the study endpoint is objective and when the outcome on treatment is markedly different from that of the historical control

## Summary

- A practical roadmap to design and analyze clinical trial for small populations using historical (external) control
- A brief review on case studies which utilized historical control (to support primary efficacy claim)
  - Different strategy to utilize historical data and/or historical control
  - Disadvantages when using HC and how to minimize them
- Future work/topics
  - Putting new methodologies to test
  - Forward looking: how to conduct good natural history/observation study to generate quality evidence