



What can Biostatistics do for



Time-Domain Astronomy ?

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Classical (Bio)Statistics

Jerzy Neyman

(1894 – 1981)



- Polish / American Statistician
- Founder of **Hypothesis Testing**
- Co-Inventor of **Confidence Intervals**
- *Neyman-Pearson Lemma*
(with his advisor Egon Pearson)

" The trouble is that what we [statisticians] call modern [orthodox] statistics was developed under strong pressure on the part of **biologists**. As a result, there is practically nothing done by us which is directly applicable to problems of **astronomy**." -- Jerzy Neyman (late in his life)

Modern Biostatistics

“In academia, the **Bayesian revolution** is on the verge of becoming the majority viewpoint, which would have been unthinkable 10 years ago.”

from The New York Times, January 20th 2004

- **Bradley P. Carlin**

Mayo Professor of Public Health
Head of Division of Biostatistics
University of Minnesota



Modern Biostatistics



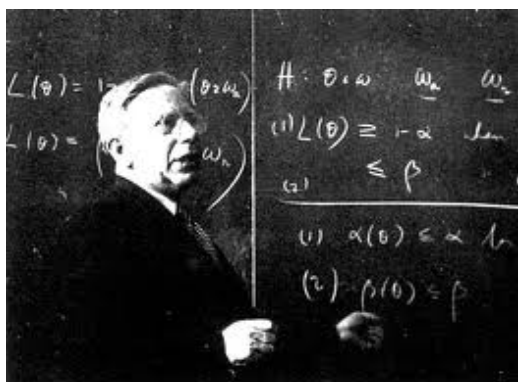
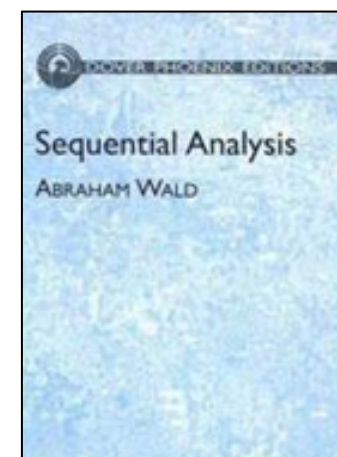
- Pathologies of Orthodox Statistics are more acknowledged nowadays
- Tests of *statistical significance* and design of *clinical trials* are now increasingly Bayesian
- Journals **banning** use of *p-values*
 - The Lancet
 - Medical J. of Australia
 - The British Heart Journal
 - American J. of Public Health
 - Int'l Committee of Medical Journal Eds

Sequential Analysis

Abraham Wald
(1902 - 1950)



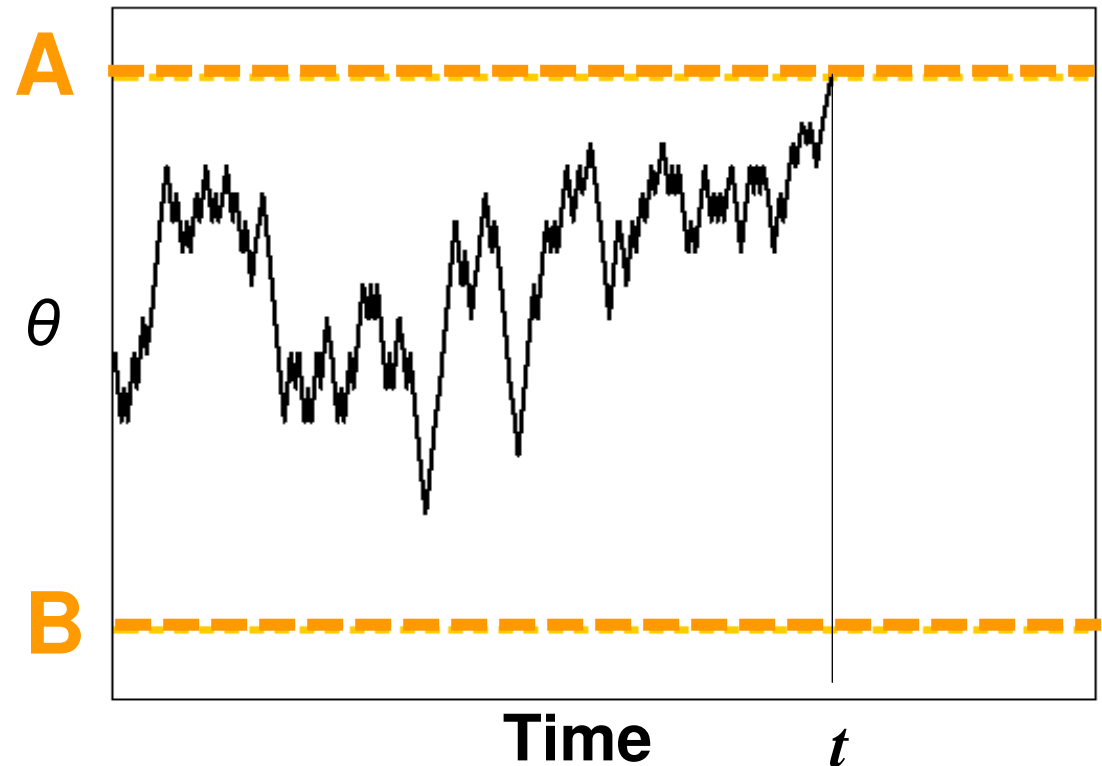
- Hungarian mathematician
- Fled Nazi Europe in late 1930s
- Invented Sequential Analysis
doing QC for Allied bombers in WW2



- Wrote classic text in 1947
- Sequential Probability Ratio Test (SPRT)

Optimal Decision Policy

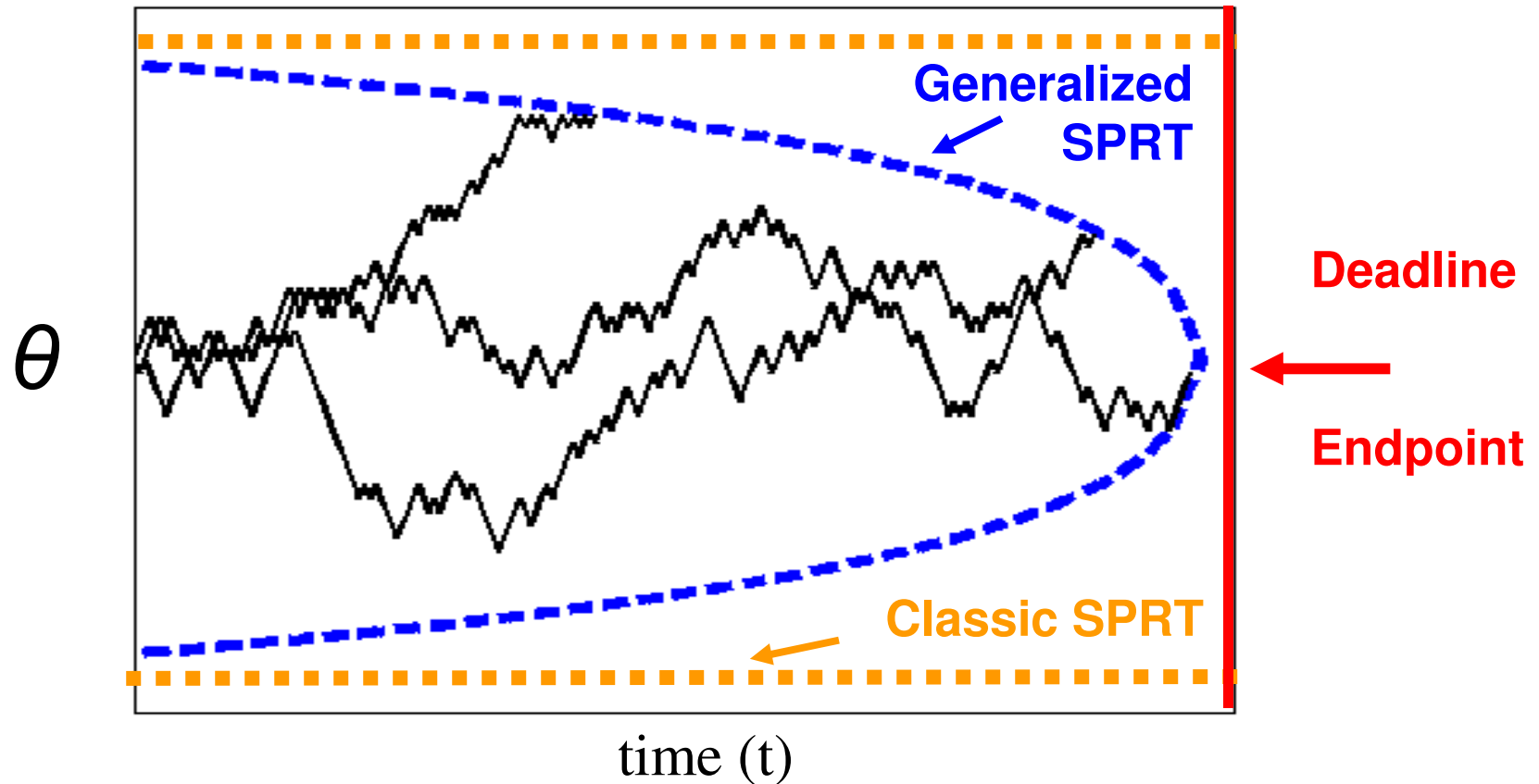
Wald & Wolfowitz (1948) showed that the optimal policy is to stop when θ exits an interval $[A, B]$ and to choose the more likely hypothesis



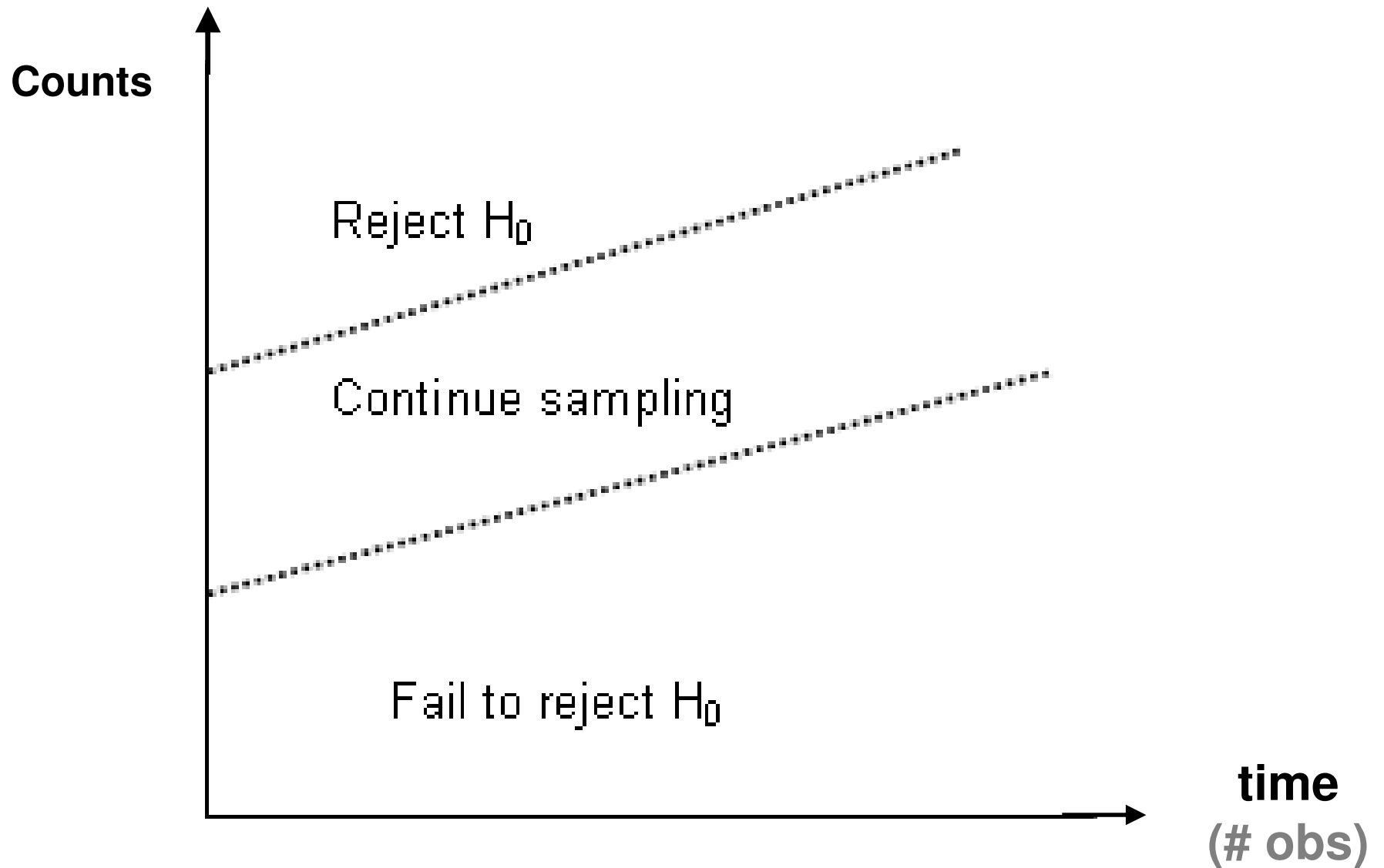
Sequential Probability Ratio Test (SPRT)

Optimal Decision Policy

The optimal policy with a **time-penalty** is to stop as soon as θ exits a region that narrows with time.

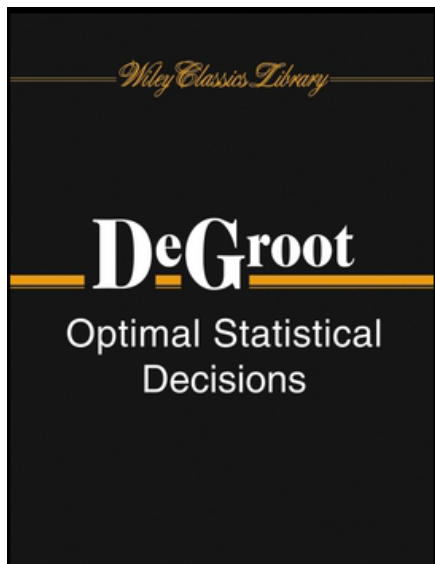


Sequential Probability Ratio Test



Optimal Decisions

Morris DeGroot
(1931 – 1989)

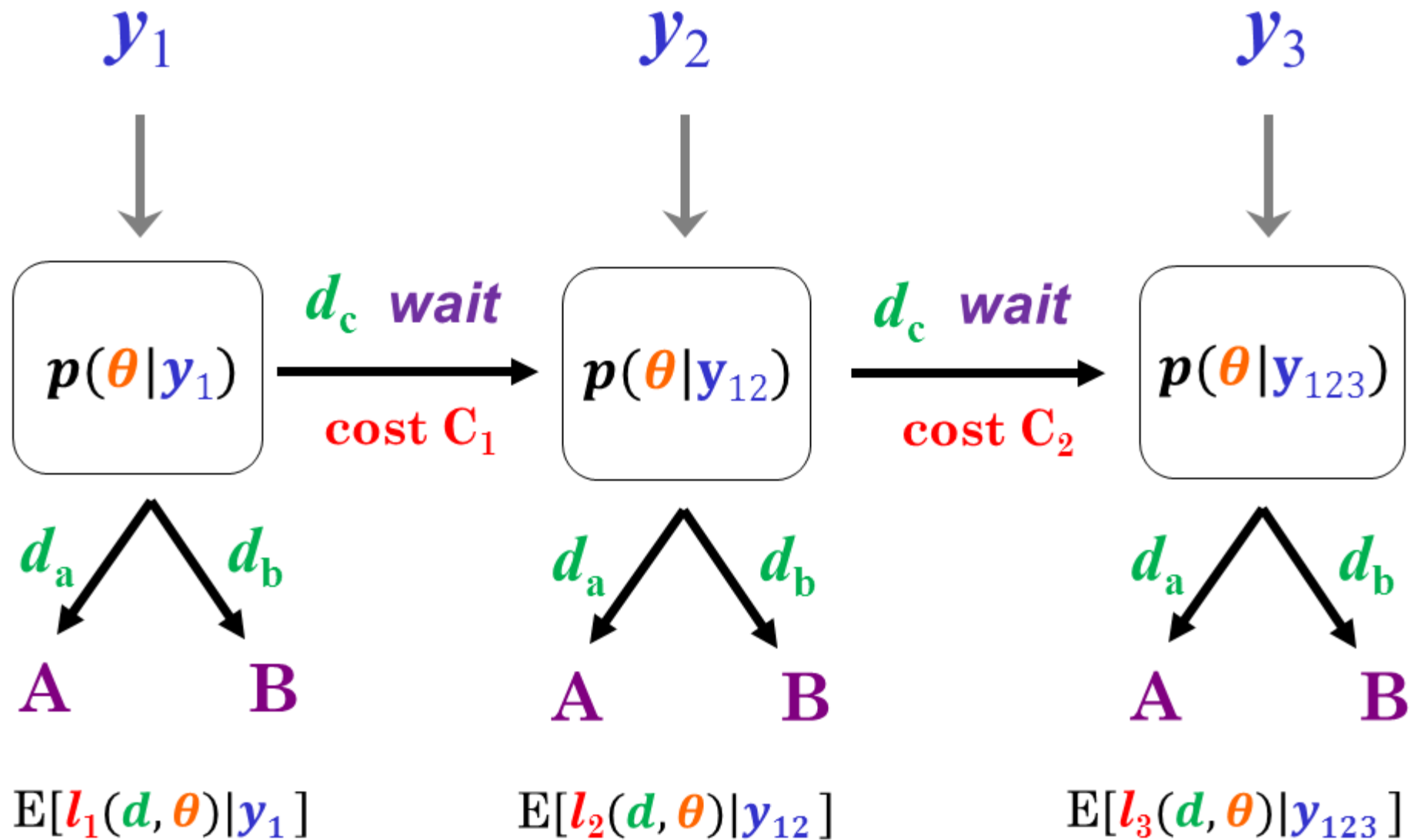


- American Statistician / CMU Professor
- Extended and formalized Wald's SPRT
- Bayesian Sequential Decision Making
- Wrote seminal text book in 1970
- Optimal decision algorithms
- For example, **Backward Induction** (BI)
 - solves the problem "backward in time"
 - also used in Dynamic Programming
 - unfortunately has **exponential complexity**

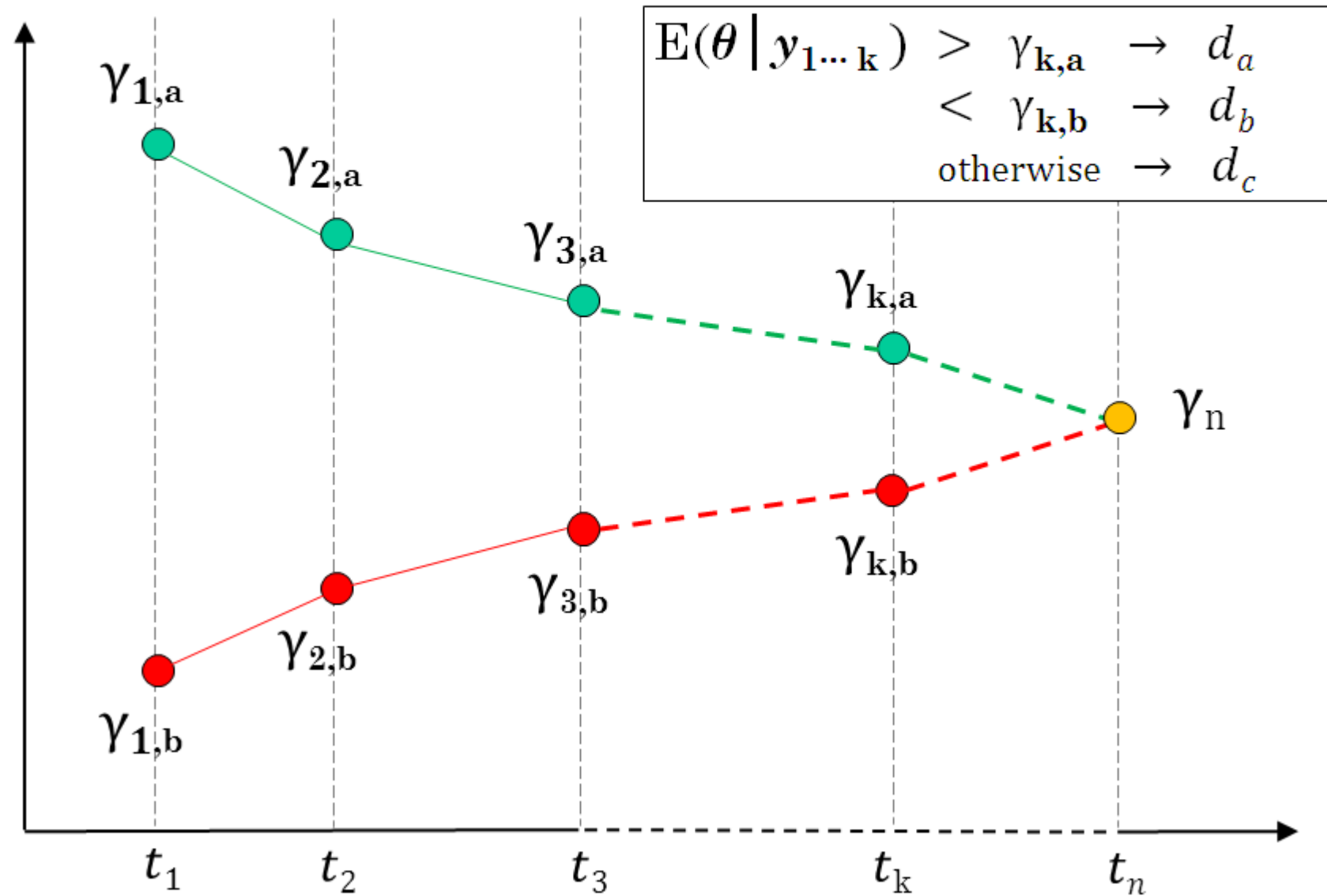
Sequential Decision Theory

- Observations “trickle-in” $\{ \mathbf{y}_i \} \quad i = 1 \dots k$
- Update current beliefs $p(\boldsymbol{\theta} \mid \mathbf{y}_1, \dots, \mathbf{y}_k)$
- Must “act” (make decisions) $\mathbf{d}_a, \mathbf{d}_b, \mathbf{d}_c$
- Which leads to losses L_a, L_b, L_c
 - timeliness (efficiency)
 - decision-induced costs
 - various domain-specific trade-offs
- Need an optimal policy (usually beforehand)
 - minimize **posterior expected loss**

Sequential Hypothesis Tests



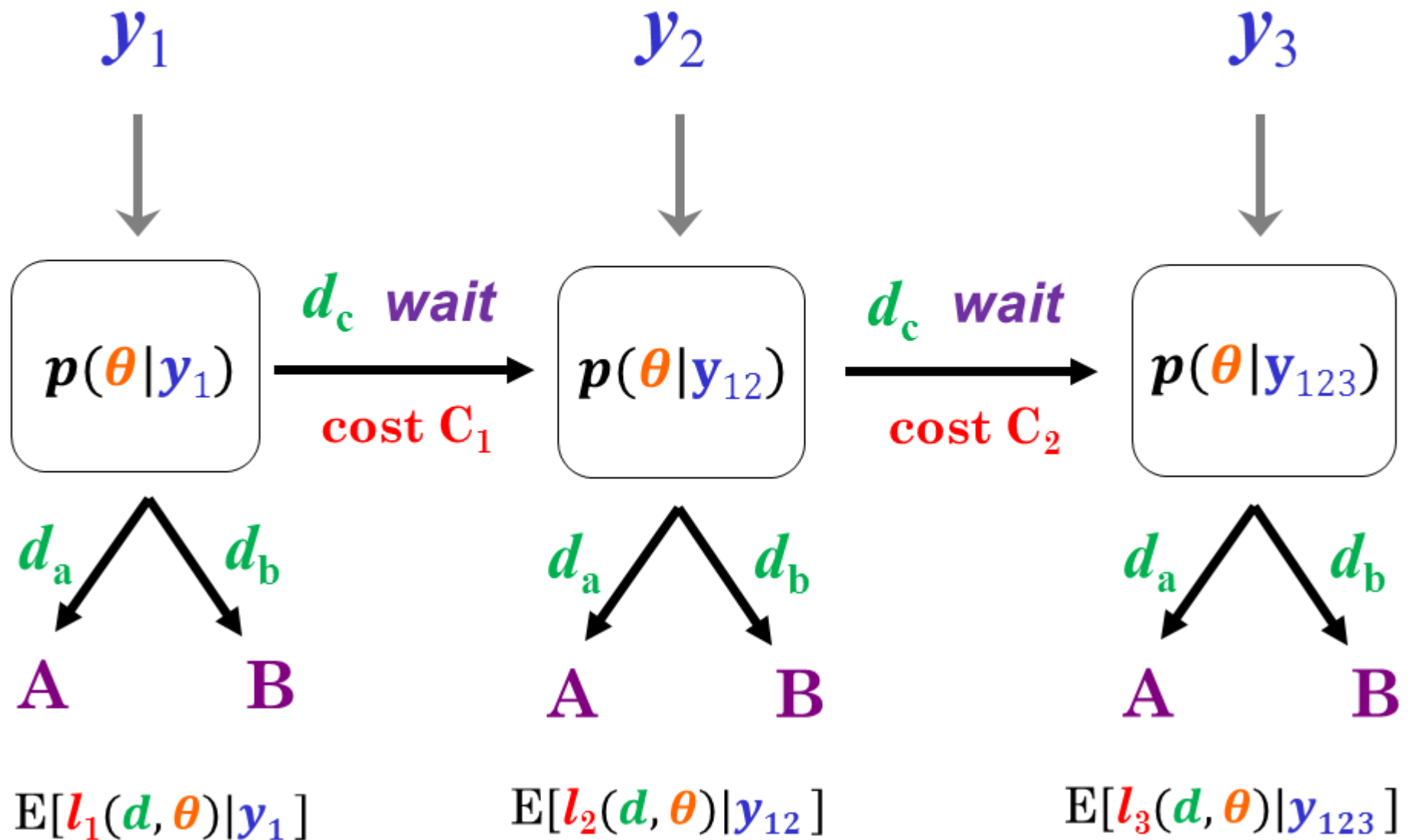
Sequential Hypothesis Tests



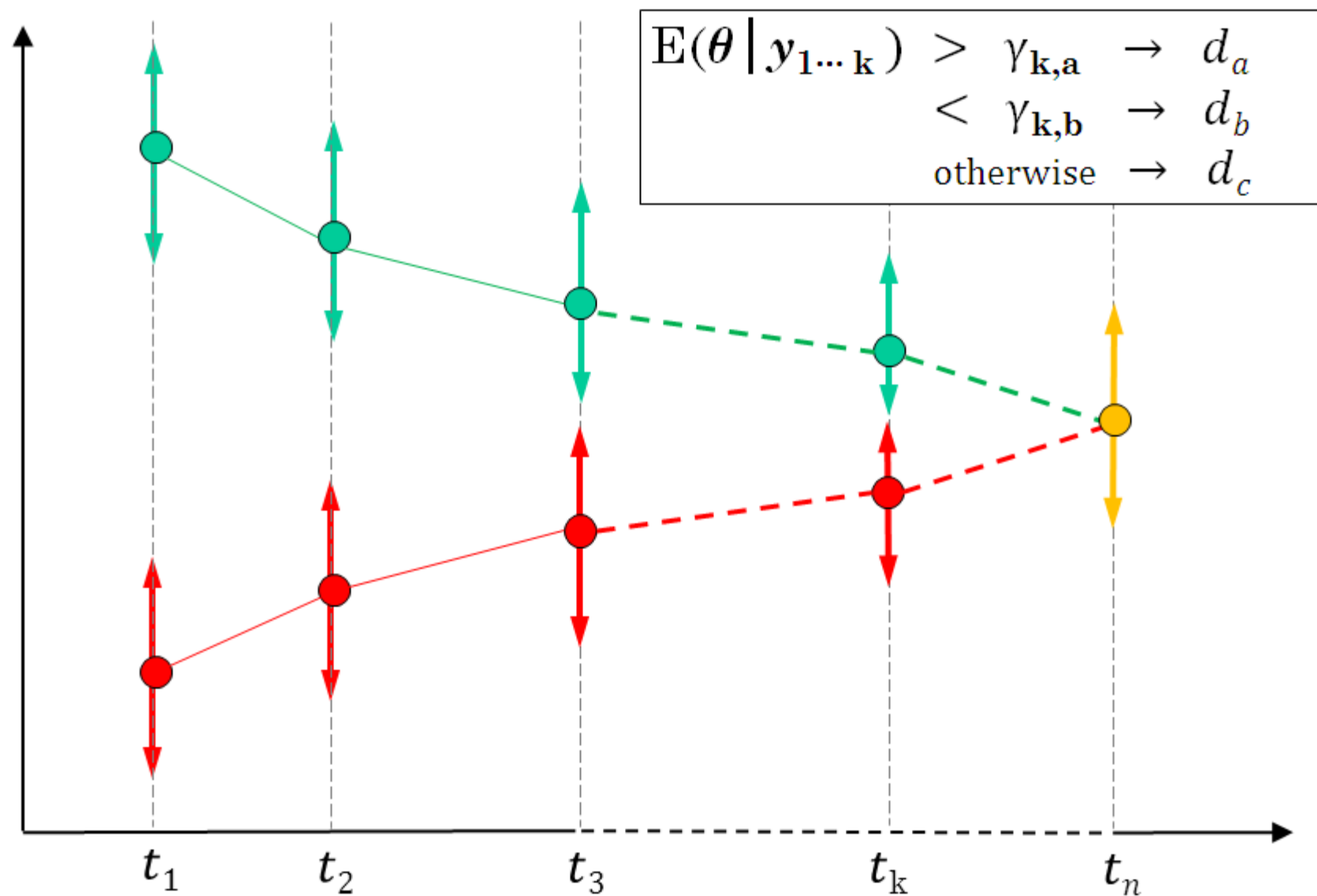
Alternative to BI

- For ~ 30 years **BI** was the best solution
- Then a new method: **Forward Sampling**
 - Carlin, Kadane and Gelfand, “Approaches to Optimal Sequential Decision Analysis in Clinical Trials,”
Biometrics, 54, 964-975, 1998
- Forward (generative) Monte Carlo sampling
- No complex integration required
- Handles arbitrary loss functions
- Complexity is **linear** in # of decision points

MC sample : $\{\theta, y_1, y_2, y_3\} \sim p(\theta, y_1, y_2, y_3)$



min $E[\text{total loss}]$ wrt $\{\gamma_{k,d}\}$



Forward Sampling vs. BI

- Complexity is **linear** vs. **exponential** for BI
- FS finds the **optimal** policy (same as BI)
 - for single-parameter exp-family models, *etc*
- Unlike BI, FS is a (parallelizable) *continuous* optimization problem (grid search, ICM, *etc*)
- FS offers far greater flexibility in terms of loss functions & probability distributions

for further inspiration ...

