Tight, Rigorous, and Automated Type I Error Proofs with Simulation

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Introduction



Figure: Michael Sklar

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Our Goals

- Make the innovation process of trial designs predictable and fast!
- ► New technique: "proof by simulation".
- Proof of Type I Error control not just on simulated points in the null space, but also on the whole null space.
- Enables:
 - Automation of Type I Error mathematical proofs.
 - Rigorous grounding for wide classes of complex designs
 - Fast design and re-design iterations.

Simulation Challenges

- ► Some challenges raised by FDA at the start of the Complex Innovative Design (CID) Pilot Program:
 - How many points in the null hypothesis space to simulate?
 - ► What is the computational complexity? Does it scale with multiple hypothesis testing?
- ▶ Other challenges:
 - Simulation on a finite number of points in the null hypothesis space does not give guarantees for the whole space. How do we deal with composite nulls?
 - Simulation has Monte Carlo error. How do we control Type I Error (on the whole space) accounting for the stochastic error?

Motivation

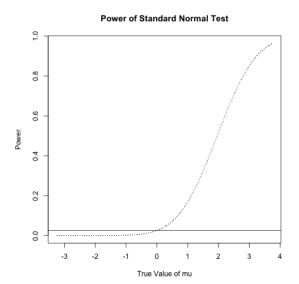
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Simple Example: One-Sided Z-Test Theoretical Results Example Designs

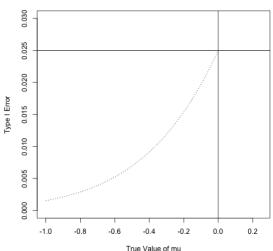
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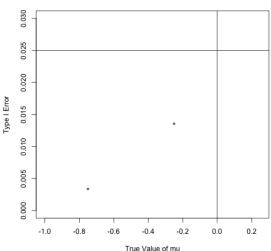
- \triangleright $X \sim \mathcal{N}(\mu, 1)$.
- ► $H_0: \mu \leq 0, H_1: \mu > 0.$
- ▶ Reject if $X > z_{1-\alpha}$.

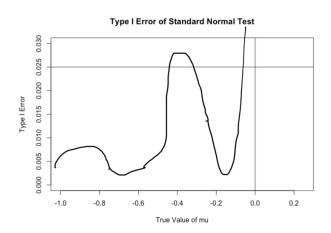




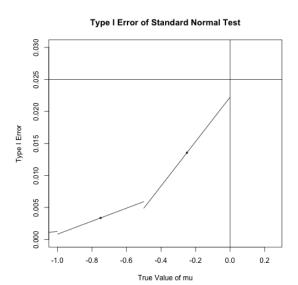




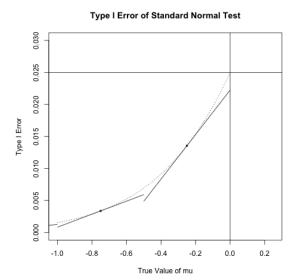




- What if we knew the derivative of the true Type I Error at these points?
- Linear approximation?

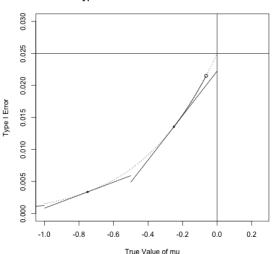


Always under the true Type I Error in this case due to convexity.



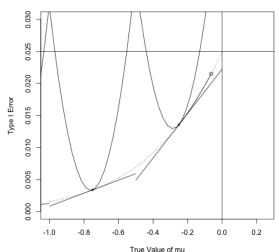
Type I Error of Standard Normal Test

Quadratic approximation?

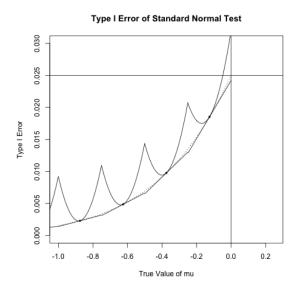


- Conservative estimate using a bound on the second derivative.
- Consequence of Taylor's Theorem.
- This particular bound is pretty bad!

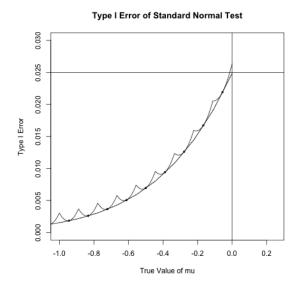
Type I Error of Standard Normal Test



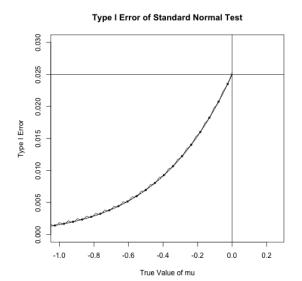
Increase number of simulation points!



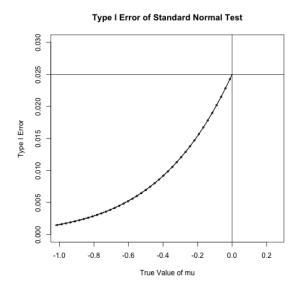
Increase number of simulation points!!



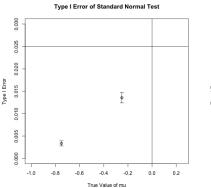
► Increase number of simulation points!!!

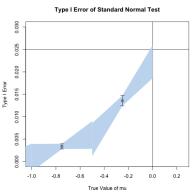


► Increase number of simulation points!!!!



- Why did this quadratic approximation work so well?
- ▶ Problem is 1-dimensional.
 - More dimensions implies orders of magnitude more computation for the same level of approximation.
- Cheated!
 - Assumed knowledge of true Type I Error and its derivatives.





- ► Monte Carlo estimates lie in confidence intervals around true value.
- Derivative estimates produce confidence bands at other values.
- Add a bound on second-order remainder term.



High-Level Sketch

Key steps:

- 1. Simulate design on a finite number of points in the null hypothesis space.
- 2. Construct upper bound estimates of the true Type I Error *on a compact subset of the null space*.
- 3. Prove that such estimates are above the true Type I Error with high confidence, pointwise *on the compact subset*.
- ldea: if the upper bound estimates are under level α , the Type I Error is highly likely to be under α as well.

High-Level Sketch

- Why do we assume compact subset?
- ▶ In practice, it is sufficient to study compact subsets.
- ► Theoretical arguments can often show that Type I Error is small in far regions.

Less High-Level Sketch

- Let Θ_0 denote a compact subset of the null hypothesis space.
- Let $f(\theta)$ denote the true Type I Error of the design if θ were the true parameter.
- ► Key steps:
 - 1. Simulate the design on a (*finite*) set of grid-points in Θ_0 .
 - 2. Construct a process, $\theta \mapsto \hat{U}(\theta)$, that depends on the simulated data.
 - 3. Prove that $\mathbb{P}\left|\left.\hat{U}(\theta)\geq f(\theta)\right|\geq 1-\delta$ for all $\theta\in\Theta_0$.

Setup

- Assume finite max number of patients in each arm.
- Let X, the full patient data, come from an exponential family P_{θ} :

$$dP_{\theta}(x) = \exp\left[T(x)^{\top}\theta - A(\theta)\right]d\mu(x)$$

- ▶ This assumption can be relaxed to log-concave densities.
- Treat a design as a black-box.
 - Adaptive data collection is OK!
 - Censored data is OK!

Polytope Null

- ▶ Assume Θ_0 is a polytope and let θ_0 be a simulation point.
- Perform a Taylor expansion of the true Type I Error: for $v \in \Theta_0 \theta_0$,

$$f(\theta_0 + v) = f(\theta_0) + \nabla f(\theta_0)^{\top} v$$
$$+ \int_0^1 (1 - \alpha) v^{\top} \nabla^2 f(\theta_0 + \alpha v) v \, d\alpha$$

Goal: upper bound each term.

Polytope Null

- Let F(x) denote the indicator that the test rejects with data x.
- ▶ **0th Order**: use $\frac{1}{n} \sum_{i=1}^{n} F(X_i)$ and upper bound with Clopper-Pearson to control $f(\theta_0)$.
- ▶ **1st Order**: use $\widehat{\nabla f}(\theta_0) := \frac{1}{n} \sum_{i=1}^n F(X_i) (T(X_i) \nabla A(\theta_0))$ and upper bound with Cantelli's Inequality to control $\nabla f(\theta_0)^\top v$.
 - Polytope null \implies worst case at one of the corners of $\Theta_0 \theta_0$.
- ▶ 2nd Order: use $U(v) := \frac{1}{2} \sup_{\theta \in \Theta_0} v^{\top} \operatorname{Var}_{\theta} [T(X)] v$ to dominate the remainder term.
- ▶ Combine \implies total upper bound on Θ_0 .

Final case: Compact Null

- Assume Θ_0 is a compact subset of the null space.
- Create a disjoint polytope covering of Θ_0 .
- Apply previous result to each element in the covering.

Possible Workflow Issue?

- ▶ Declaring victory if $\hat{U}(\theta) \leq \alpha$ may still be problematic.
- Due to randomness in \hat{U} , this decision rule does not lead to any meaningful statement about the true Type I Error being under α .

Solution: Tune Threshold

- Tune the rejection threshold!
- ► Key idea: find the critical value, $\hat{\lambda}$, where $\hat{U}(\theta)$ first hits level α exactly for some $\theta \in \Theta_0$.
- ▶ Then, letting $f_{\lambda}(\theta)$ denote the true Type I Error at θ with rejection threshold λ , we have that

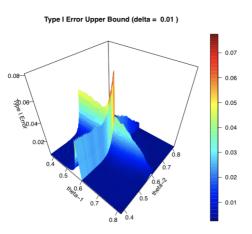
$$\mathbb{P}\left[f_{\hat{\lambda}}(\theta) \leq \alpha\right] \geq 1 - \delta$$
$$\mathbb{E}\left[f_{\hat{\lambda}}(\theta)\right] \leq \alpha + \delta$$

- Interpretation: both a high-probability guarantee for the Type I Error at the selected threshold and an overall guarantee for the Type I Error of this selection procedure.
- ► Guarantees, a priori, that our procedure controls Type I Error at the 2.5% level.

Example: Thompson Sampling

- ► Two-arm trial.
- Outcomes $Y_{ij} \sim Bernoulli(\theta_j)$ (i = 1, ..., 100).
- \vdash $H_0: \theta_1 < 0.6, H_1: \theta_1 \geq 0.6.$
- ightharpoonup Beta(1,1) prior
- ▶ Reject arm 1 at the end if posterior $\mathbb{P}(\theta_1 > 0.6) > 0.7$.

Thompson Sampling



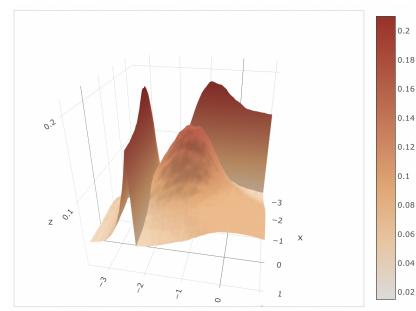
Example: Bayesian Basket Trial from Berry et al. (2013)

Design:

$$Y_j \sim Binom(n_j, p_j)$$
 $j = 1, ..., d$
 $p_j = expit(\theta_j + logit(q_j))$
 $\theta_j \sim \mathcal{N}(\mu, \sigma^2)$
 $\mu \sim \mathcal{N}(\mu_0, \sigma_0^2)$
 $\sigma^2 \sim \Gamma^{-1}(\alpha_0, \beta_0)$

- Let $c \in [0,1]^{d-1}$ be a vector of fixed thresholds.
- ▶ Reject if $\mathbb{P}[p_i > p_0 | Y] > c_i$ for some null (treatment) arm i.

Berry et al. (2013)



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Computation Scale

- Number of parameters to grid should be small (\leq 6) to be tractable.
- Example designs with number of simulations and null points:

Design Type	Number of Sims	Number of Null Points	Total Sims (billion)
Thompson	100,000	16,384	1.6
Exponential Hazard	100,000	16,384	1.6
Binomial Selection	100,000	262,144	26
Berry et al.	10,000	5,308,416	530

Computation Bottleneck

- Most of the bottleneck is in speeding up the trial simulation itself.
- Lots of practical designs use Bayes.
- ► Traditional methods to obtain Bayes quantity is through MCMC or analytical formulas (conjugacy).
- ► Found Integrated Nested Laplace Approximation (INLA) to be efficient and accurate.
- ► Homegrown INLA library in development.

Main Contributor of INLA Library + Application



Figure: Ben Thompson

Simulations are Fast!

- ▶ JAX Python library and homegrown C++ codebase.
- Benchmark on modern Apple M1 Macbook (CPU) and cloud machine with NVIDIA V100 (GPU):

Design Type	Number of Sims	Number of Null Points	Total Sims (billion)	Hardware	Time (m)
Exponential Hazard	100,000	16,384	1.6	CPU	1.57
Binomial Selection	100,000	262,144	1.6	CPU	2.37
Thompson	100,000	16,384	26	CPU	4
Berry et al.	10,000	5,308,416	530	CPU	150
Berry et al.	10,000	5,308,416	530	GPU	1.16

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Acknowledgements



(a) Alex Constantino



(b) Gary Mulder



(c) Daniel Kang

Further Developments

- Extend to bound FDR, Bias, MSE.
- Give similar bounds for importance sampling.
- Bring simulations into cloud computing.

Recap

- Our methodology gives provable control of Type I Error via simulation.
- ▶ Allows complex designs to be studied seamlessly.
- Our software can make simulations extremely fast.