

Fast Permutation Tests, Especially for Multiple Comparisons and Even When One Sample is Large, that Efficiently Maximize Power Under Conventional Monte Carlo and Allow for Simultaneous Permutation-Style P-Value Adjustments

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Contents

- 1. Goal and Rationale
- 2. Permutation Sampling, Duplicate Samples, & Power
- 3. Maximizing Power Under Conventional Monte Carlo
- 4. Efficiently "Oversample" Based on Expected Runtime
- 5. Approximate the Optimal Number of Samples
- 6. How Much Power Gain...?
- 7. ...At What Cost?
- 8. Speed Premium for Multiple Comparisons
- 9. Increased Power for Permutation-Style P-Value Adj



1. Goal and Rationale

GOAL:

Quickly implement many non-parametric permutation tests, even when one sample in a pair is large, with maximum power under conventional Monte Carlo

- WHY MANY TESTS &/OR ONE LARGE SAMPLE?
 - "Parity Testing" in Regulatory Telecom OSS Reports
 - Medical studies using MRI data
 - clinical trials with large controls and many and smaller studies
 - Any multiple comparisons context requiring permutation-style p-value adjustments of permutation test p-values (and thus, computationally intensive nested sampling loops)



1. Goal and Rationale

- WHY CONVENTIONAL MONTE CARLO?
 - Faster, more efficient sampling techniques (e.g. various methods of importance sampling) are not always implementable
 - when such methods can be implemented but their results are suspect, conventional Monte Carlo can be a useful verification
- WHY MAXIMUM POWER?
 - best test, all else equal

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2. Permutation Sampling, Duplicate Samples, & Power

- PROC PLAN, PROC MULTTEST, PROC NPAR1WAY, & PROC TWOSAMPL® can sample without replacement within a sample, as required of permutation tests
- None can sample without replacement across samples (i.e. none can avoid drawing duplicate samples)
- Duplicate samples → loss of power due to increased variance of estimated p-value



3. Maximizing Power Under Conventional Monte Carlo

- Use "oversampling" to efficiently obtain a unique set of samples (no duplicates)
 - a. draw more samples than desired (r)
 - b. delete duplicates
 - c. randomly select the desired number (T) of samples from the remainder
 - d. recall PROC PLAN if fewer than T samples remain
- "Oversampling" preserves the uniform distribution sampling assumption of nonparametric permutation tests



4. Efficiently "Oversample" Based on Expected Runtime

- Draw just enough "extra" samples (r-T) to minimize expected runtime
- Expected Runtime = g(n₁, n₂, r, T) = PROC PLAN RunTime * expected # of Calls To PROC PLAN = PPRT(r, [n₁+n₂]) * CTPP(r, T, [n₁+n₂]!/[n₁!n₂!])
- Choose optimal r, r*, such that $\partial g/\partial r = 0$



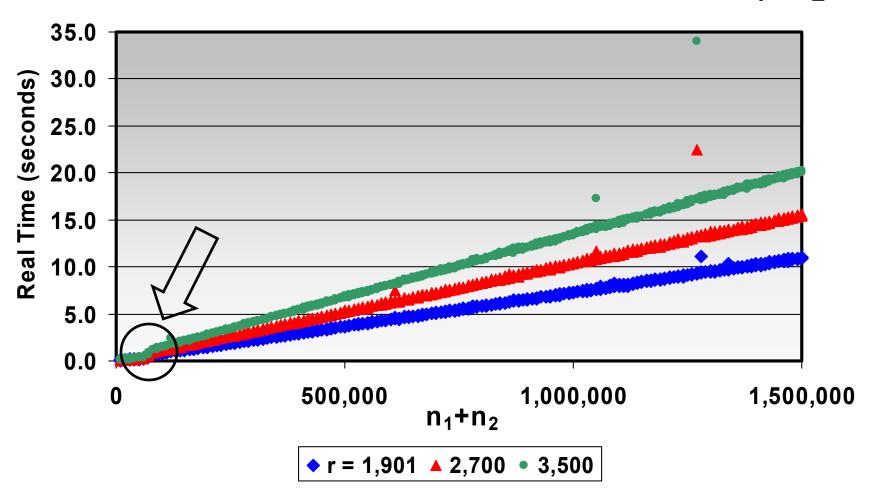
4. Efficiently "Oversample" Based on Expected Runtime

- PPRT(r,[n_1+n_2]) $\approx \beta_0 + \beta_1*(n_1+n_2) + \beta_2*r + \beta_3*r*(n_1+n_2)$
- CTPP(r, T, $[n_1+n_2]!/[n_1!n_2!]$) =

$$\left(\frac{1}{p}\right) = \left[\sum_{j=T}^{r} \left[\frac{\frac{(n_1 + n_2)!}{n_1! n_2!}}{j! \left(\frac{(n_1 + n_2)!}{n_1! n_2!} - j\right)!} \sum_{i=0}^{j} \frac{(-1)^i j! (j-i)^r}{i! (j-i)! \left(\frac{(n_1 + n_2)!}{n_1! n_2!}\right)^r}\right]\right)^{-1}$$

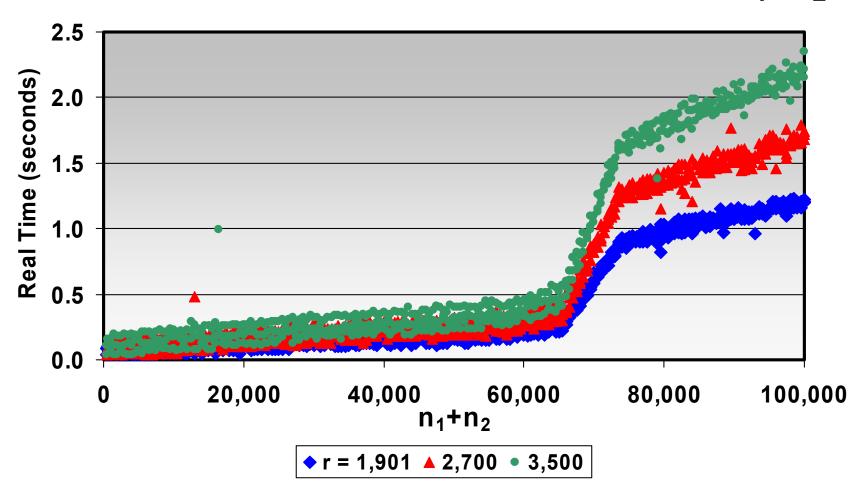


GRAPH 1: PROC PLAN Runtime by r by n₁+n₂

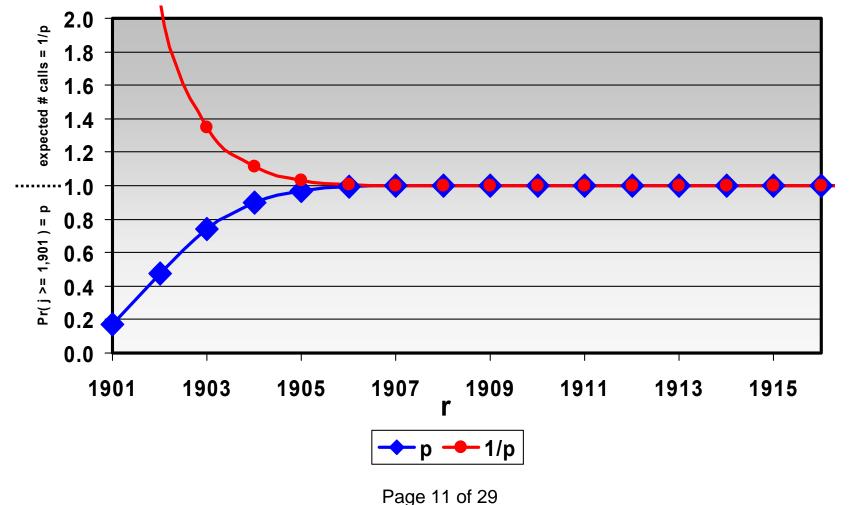




GRAPH 2: PROC PLAN Runtime by r by n₁+n₂

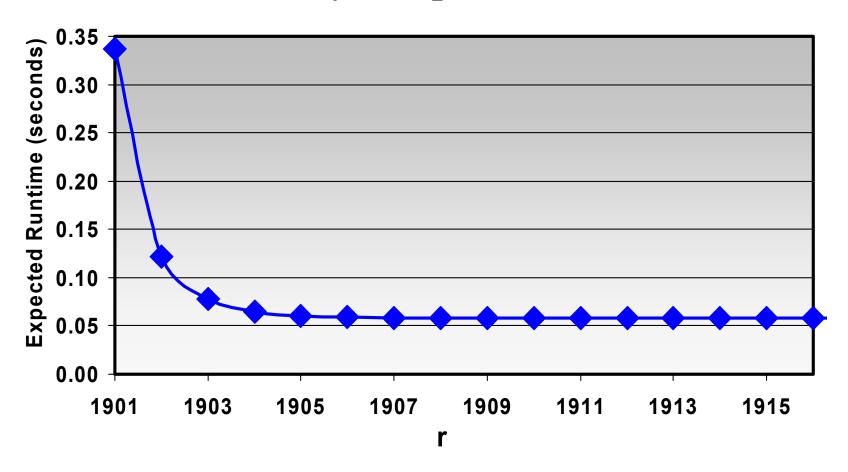


GRAPH 3: Probability of At Least T Unique Samples (p) & Expected Number of Calls to PROC PLAN (1/p) by r (for n₁=68, n₂=4, and T=1,901)



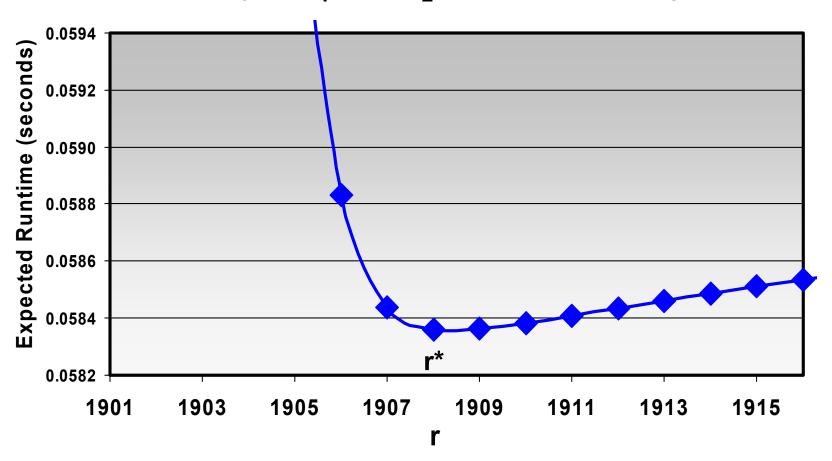


GRAPH 4: Expected Runtime (1/p * one runtime) by r (for $n_1=68$, $n_2=4$, and T=1,901)





GRAPH 5: Expected Runtime (1/p * one runtime) by r (for n_1 =68, n_2 =4, and T=1,901)





5. Approximate Optimal r*

- Precision required to numerically calculate r* is too high to do "on the fly" in SAS[®] for every n₁ & n₂
- However, since $\partial g/\partial r \approx 0$ for r slightly > r*, we can approximate:
 - a. Define ranges based on combinations $(n_1+n_2)!/n_1!n_2!$
 - b. Pick suboptimal r*s corresponding to each lower bound to obtain largest r* for each range
 - c. Runtime of suboptimal $r^* \approx runtime$ of r^* because $\partial g/\partial r \approx 0$ for r slightly > r^* (see Graphs 4 & 5)



5. Approximate Optimal r*

| $C = (n_1 + n_2)!/n_1!n_2!$ | "low-end" r* | p (lower bound) | 1/p (lower bound) |
|---------------------------------|-----------------|-------------------------|----------------------|
| C < 10,626 | С | 1.0 (assuming C ≥ T) | 1.0 |
| 10,626 ≤ C < 52,360 | 2,138 | 0.997929320330667 | 1.002074976280530 |
| 52,360 ≤ C < 101,270 | 1,956 | 0.999058342955471 | 1.000942544598290 |
| 101,270 ≤ C < 521,855 | 1,934 | 0.999429717692296 | 1.000570607715190 |
| 521,855 ≤ C < 1,028,790 | 1,912 | 0.999726555240808 | 1.000273519551680 |
| $1,028,790 \le C < 10,009,125$ | 1,908 | 0.999512839120371 | 1.000487398321020 |
| 10,009,125 ≤ C < 25,637,001 | 1,904 | 0.999961594180711 | 1.000038407294350 |
| 25,637,001 ≤ C < 100,290,905 | 1,903 | 0.999944615376581 | 1.000055387691050 |
| 100,290,905 ≤ C < 5,031,771,045 | 1,902 | 0.999839691379204 | 1.000160334323770 |
| 5,031,771,045 ≤ C | 1,901 | 0.999641154940541 | 1.000358973875460 |



- Permutation test p-values relying on any type of sampling will have actual size level (asl) > α
- ∴ either p-values or critical value (c_α) should be adjusted
- Smaller variance of no replacement sampling (NR) \Rightarrow smaller asl \Rightarrow larger $c_{\alpha}^* \Rightarrow$ larger power

•
$$\sigma^2_{NR} < \sigma^2_{WR}$$
 \Rightarrow $asl_{NR} < asl_{WR}$ \Rightarrow $c_{\alpha NR}^* > c_{\alpha WR}^*$ \Rightarrow power_{NR} > power_{WR}



- σ^2_{WR} is based on the binomial, $\sigma^2_{bin} = n_p pq$ σ^2_{NR} based on hypergeometric, $\sigma^2_{hyp} = n_p pq(N-n_p)/(N-1)$ (N = # possible samples, n_p = # permutation samples)
- $\sigma^2_{bin} > \sigma^2_{hyp} \Rightarrow \sigma^2_{WR} > \sigma^2_{NR}$

•
$$\operatorname{asl}_{\mathsf{WR}} = \mathsf{Pr}(\mathsf{S} \leq \mathsf{n_p}^* \alpha \mid \mathsf{p}) = \frac{1}{n_p} \sum_{i=0}^{n_p} \sum_{k=0}^{\lfloor n_p \alpha \rfloor} \binom{n_p}{i} \left(\frac{i}{n_p}\right)^k \left(1 - \frac{i}{n_p}\right)^{(n_p - k)}$$

• asl_{NR} = Pr(S \le n_p*\alpha | p) =
$$\frac{1}{n_p} \sum_{i=0}^{N} \frac{\sum_{k=0}^{N} \binom{i}{n_p} \binom{n_p}{n_p}}{\sum_{k=0}^{N} \binom{N-S}{n_p}}$$
•
$$\frac{1}{n_p} \sum_{s=0}^{N} \sum_{k=0}^{N} \frac{\binom{N-S}{n_p}}{\binom{N}{n_p}}$$

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- if (asl / α) is essentially constant close to α = 0.05, then $c_{\alpha}^* \times (asl / \alpha) = \alpha$ $c_{\alpha}^* = \alpha^2 / asl$
- $\therefore c_{\alpha WR}^* = \alpha^2 / asl_{WR} and c_{\alpha NR}^* = \alpha^2 / asl_{NR}$
- power can only be obtained via simulation, but by CLT we know that asymptotically:

power =
$$_{1-\Phi}\left(z_{\alpha}-\frac{\delta\sqrt{n}}{\sigma}\right)$$
 where δ = effect, z_{α} = $\Phi^{-1}(1-\alpha)$

• power_{NR}
$$\approx 1 - \Phi\left(z_{c_{\alpha_{NR}}^*} - \frac{\delta\sqrt{n}}{\sigma}\right)$$
, power_{WR} $\approx 1 - \Phi\left(z_{c_{\alpha_{WR}}^*} - \frac{\delta\sqrt{n}}{\sigma}\right)$

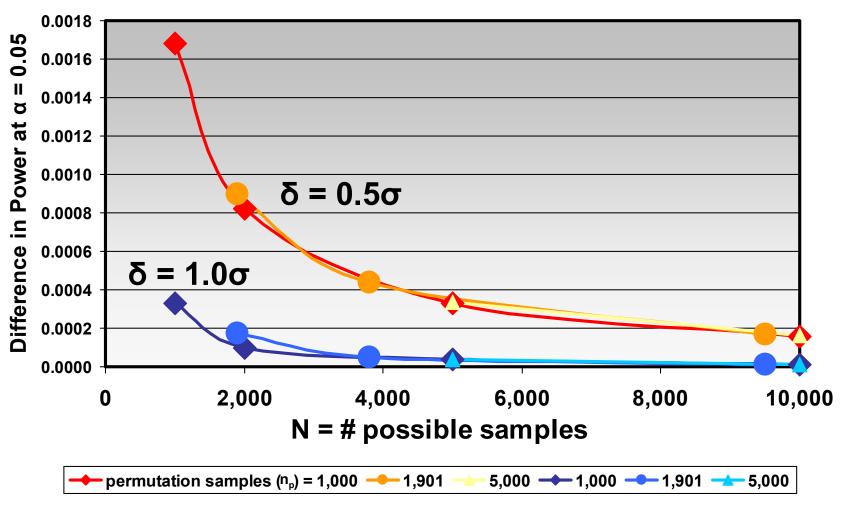


| # Permutation Samples, n _p | # Possible Samples, N | asl _{NR} | asl _{WR} | Ca* | c _a * |
|--|--------------------------|-------------------|-------------------|---------|------------------|
| 1,000 | 1,000 | 0.05100 | 0.05144 | 0.04902 | 0.04859 |
| 1,000 | 2,000 | 0.05122 | 0.05144 | 0.04880 | 0.04859 |
| 1,000 | 5,000 | 0.05136 | 0.05144 | 0.04868 | 0.04859 |
| 1,000 | 10,000 | 0.05140 | 0.05144 | 0.04863 | 0.04859 |
| 1,901 | 1,901 | 0.05050 | 0.05073 | 0.04951 | 0.04927 |
| 1,901 | 3,802 | 0.05062 | 0.05073 | 0.04939 | 0.04927 |
| 1,901 | 9,505 | 0.05069 | 0.05073 | 0.04932 | 0.04927 |
| 5,000 | 5,000 | 0.05020 | 0.05028 | 0.04980 | 0.04971 |
| 5,000 | 10,000 | 0.05024 | 0.05028 | 0.04976 | 0.04971 |



| # Permutation Samples, n _p | # Possible Samples, N | Power _{NR} δ=0.5σ | Power _{WR} δ=0.5σ | ΔPower δ=0.5σ | Power _{NR} $\delta = \sigma$ | Power _{WR} $\delta = \sigma$ | |
|--|--------------------------|-------------------------------|-------------------------------|----------------------|---------------------------------------|---------------------------------------|---------|
| 1,000 | 1,000 | 0.53093 | 0.52925 | 0.00168 | 0.96483 | 0.96450 | 0.00033 |
| 1,000 | 2,000 | 0.58483 | 0.58401 | 0.00082 | 0.98147 | 0.98137 | 0.00010 |
| 1,000 | 5,000 | 0.58434 | 0.58401 | 0.00033 | 0.98141 | 0.98137 | 0.00004 |
| 1,000 | 10,000 | 0.63373 | 0.63357 | 0.00016 | 0.99040 | 0.99039 | 0.00001 |
| 1,901 | 1,901 | 0.53283 | 0.53193 | 0.00090 | 0.96519 | 0.96502 | 0.00017 |
| 1,901 | 3,802 | 0.58708 | 0.58664 | 0.00044 | 0.98173 | 0.98168 | 0.00005 |
| 1,901 | 9,505 | 0.63628 | 0.63611 | 0.00017 | 0.99058 | 0.99056 | 0.00001 |
| 5,000 | 5,000 | 0.58864 | 0.58830 | 0.00034 | 0.98191 | 0.98187 | 0.00004 |
| 5,000 | 10,000 | 0.63787 | 0.63771 | 0.00016 | 0.99068 | 0.99067 | 0.00001 |

GRAPH 6: Permutation Sampling With vs. Without Replacement: Approximate Difference in Power at $\alpha = 0.05$ by N by n_p by δ





7. ... At What Cost?

- Even for small N, n_p , and δ , approximate power gains from NR sampling are relatively small
- However, runtime cost also is small typically less than 1%
- ∴ use NR permutation sampling unless cost of 1% of runtime is high and cost of Type II error is low

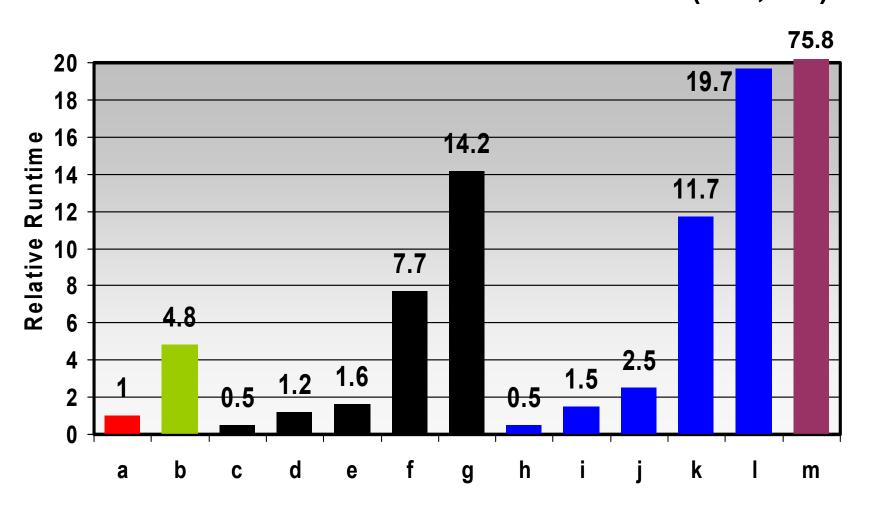


8. Speed Premium for Multiple Comparisons

- For multiple study groups per control group (and/or multiple comparisons), code merges each group of original data to the PROC PLAN sampling output separately
- Separate merges avoids multiple outputting of control records for each corresponding study group (or each multiply-compared group)
- ⇒ Huge runtime savings on sorts/merges of large datasets (see Graph 7)
- PROC MULTTEST, PROC NPAR1WAY, and PROC TWOSAMPL® do not have this option (Graph 7)



GRAPH 7: Relative Start-to-Finish Runtime (T=1,901)





Graph 7 Key

| Column | Method |
|--------|--|
| а | PROC PLAN with "oversampling" |
| b | Cytel's PROC TWOSAMPL® |
| С | PROC NPAR1WAY, study/control=1, (n ₁ +n ₂)<10 ⁴ |
| d | PROC NPAR1WAY, study/control>1, (n ₁ +n ₂)<10 ⁴ |
| е | PROC NPAR1WAY, study/control=1, 10 ⁴ <(n ₁ +n ₂)<10 ⁵ |
| f | PROC NPAR1WAY, study/control=1, 10 ⁵ <(n ₁ +n ₂)<1.5*10 ⁶ |
| g | PROC NPAR1WAY, study/control=1, 10 ⁶ <(n ₁ +n ₂)<1.5*10 ⁷ |
| h | PROC MULTTEST, study/control=1, (n ₁ +n ₂)<10 ⁴ |
| i | PROC MULTTEST, study/control>1, (n ₁ +n ₂)<10 ⁴ |
| j | PROC MULTTEST, study/control=1, 10 ⁴ <(n ₁ +n ₂)<10 ⁵ |
| k | PROC MULTTEST, study/control=1, 10 ⁵ <(n ₁ +n ₂)<1.5*10 ⁶ |
| I | PROC MULTTEST, study/control=1, 10 ⁶ <(n ₁ +n ₂)<1.5*10 ⁷ |
| m | Looping in SAS® (see Jackson affidavit) |



9. Increased Power for Permutation-Style P-Value Adj

- Take a single step resampling method adjustment
- No-replacement sampling ⇒ increased power from:
 - ❖ a) smaller variance of each p_i*

$$\Rightarrow \min_{1 \le j \le k} p^*_{j_{NR}}$$
 is stochastically larger than $\min_{1 \le j \le k} p^*_{j_{WR}}$

$$\Rightarrow \Pr\left(\min_{1 \leq j \leq k} p_{j_{NR}} \leq p_i \mid H_0^C\right) < \Pr\left(\min_{1 \leq j \leq k} p_{j_{WR}} \leq p_i \mid H_0^C\right)$$

$$\Rightarrow \tilde{p}_{i_{NR_a}} < \tilde{p}_{i_{WR_a}} \Rightarrow power_{NR_a} > power_{WR_a}$$



9. Increased Power for Permutation-Style P-Value Adj

❖ b) previous Monte Carlo error p-value adjustment

$$\Rightarrow p_{i_{NR}} < p_{i_{WR}}$$

$$\Rightarrow \Pr\left(\min_{1 \leq j \leq k} p_j \leq p_{i_{NR}} \mid H_0^C\right) < \Pr\left(\min_{1 \leq j \leq k} p_j \leq p_{i_{WR}} \mid H_0^C\right)$$

$$\Rightarrow \tilde{p}_{i_{NR_b}} < \tilde{p}_{i_{WR_b}}$$

$$\Rightarrow power_{NR_{b}} > power_{WR_{b}}$$



9. Increased Power for Permutation-Style P-Value Adj

 use NR sampling for both permutation tests and permutation-style p-value adjustments to maximize power gain

$$\Pr\left(\min_{1 \leq j \leq k} p_{j_{NR}} \leq p_{i_{NR}} \mid H_0^C\right) < \Pr\left(\min_{1 \leq j \leq k} p_{j_{WR}} \leq p_{i_{WR}} \mid H_0^C\right)$$

$$\Rightarrow \tilde{p}_{i_{NR}} < \tilde{p}_{i_{WR}}$$

$$\Rightarrow power_{NR} > power_{WR}$$

Same rationale applies to stepwise adjustments



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