

# COMPARISON BETWEEN TREE GATEKEEPING PROCEDURE AND GRAPHICAL APPROACH IN A PIVOTAL CLINICAL TRIAL WITH MULTIPLE OBJECTIVES AND MULTIPLE ENDPOINTS

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**1. Introduction**

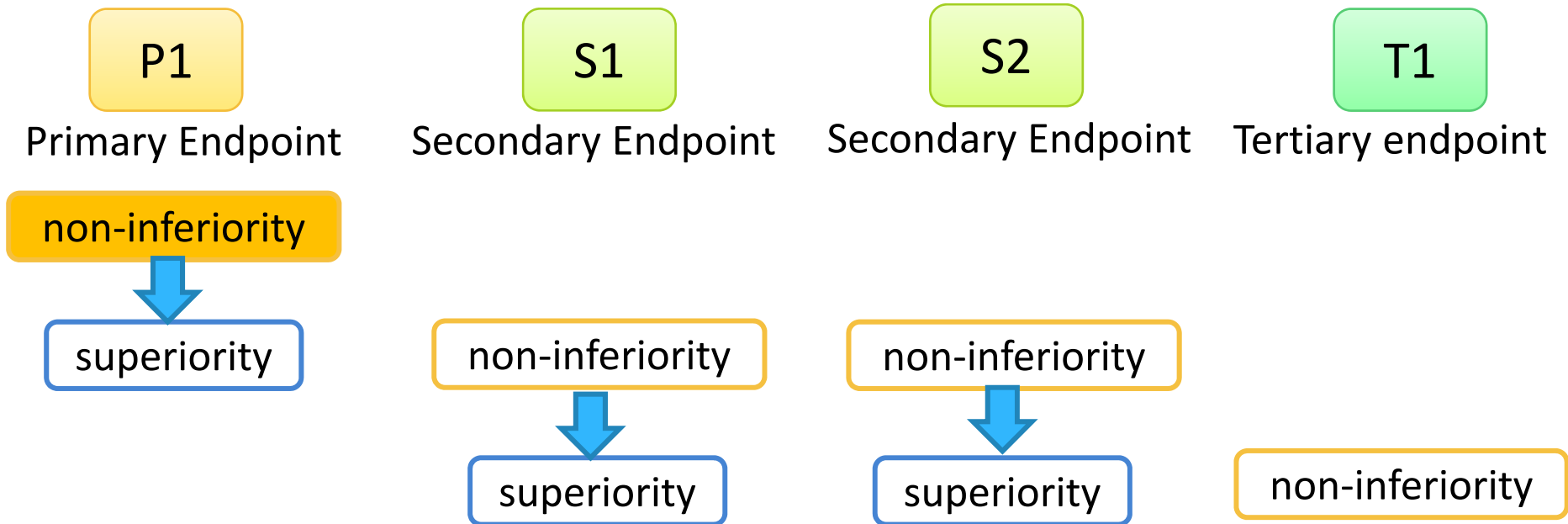
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# 1. INTRODUCTION

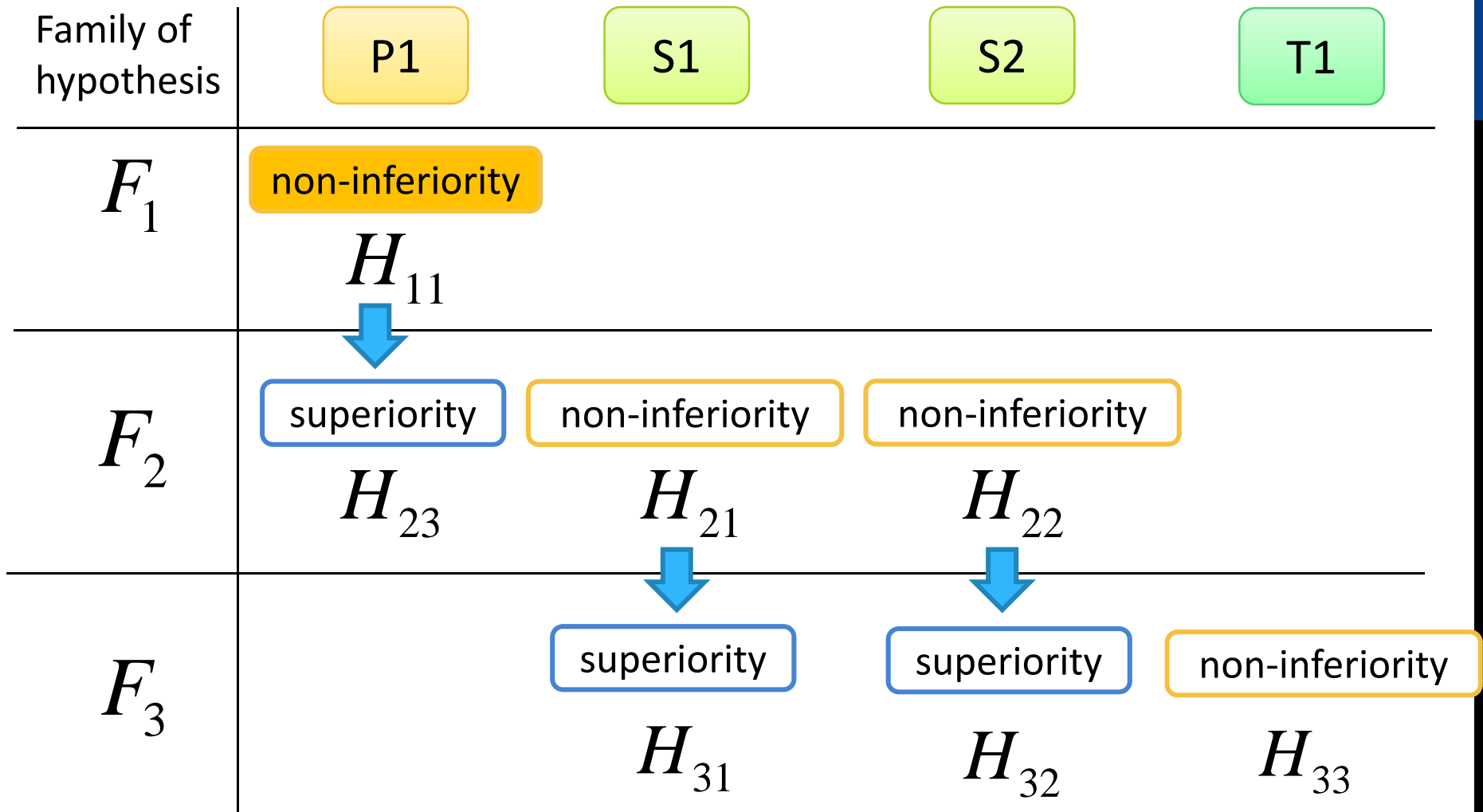
- We assume a pivotal clinical trial for HIV with 2 arms (Drug/Control)



- As to each of primary and secondary endpoints, only when non-inferiority can be shown, superiority is tested

When we would like to include these results *in labeling*, ***FWER (Family-Wise Error Rate) should be controlled.***

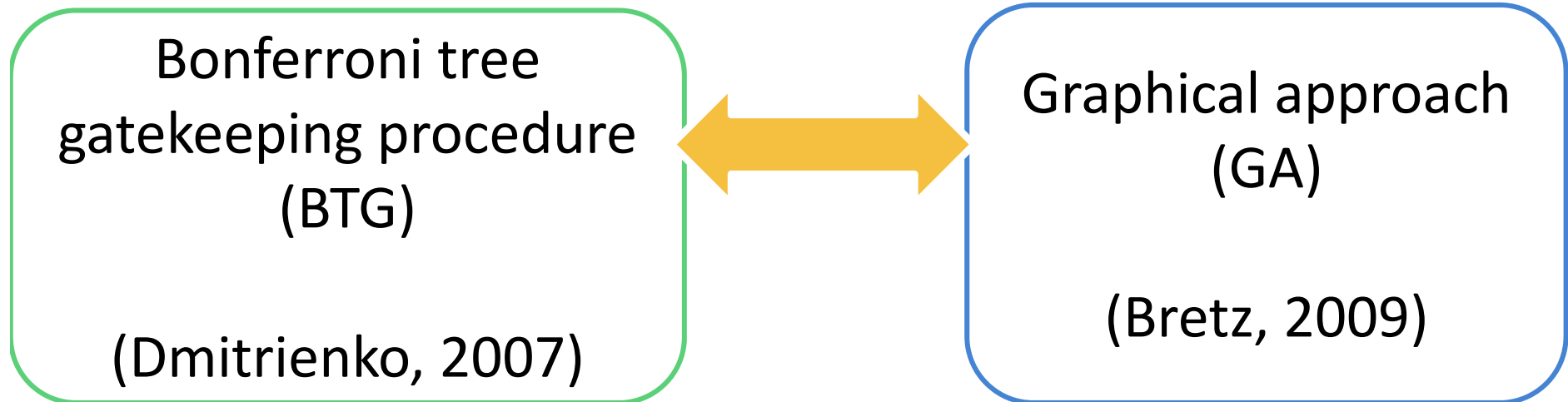
# SETTING (INDIVIDUAL HYPOTHESIS)



We assume the importance of hypotheses in each family are equal

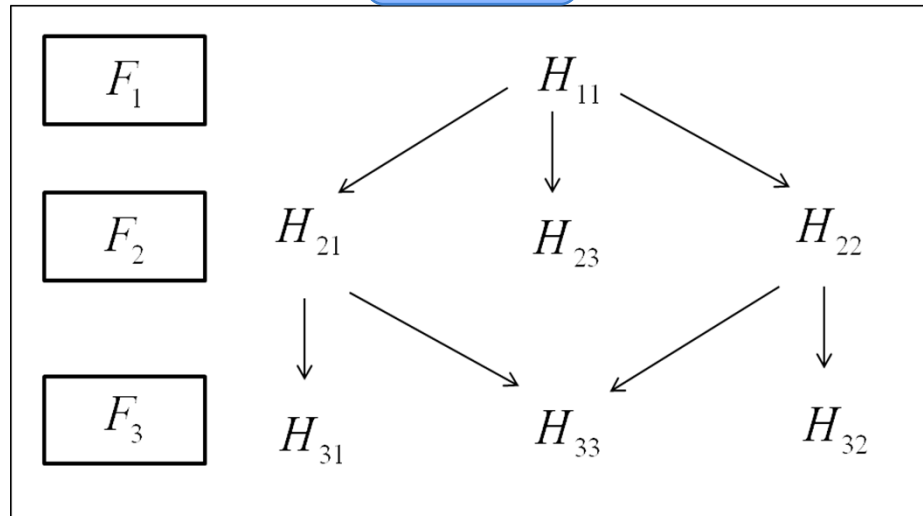
# 2. OBJECTIVE

To compare the performance between Bonferroni tree gatekeeping procedure and Graphical approach for assumed pivotal study

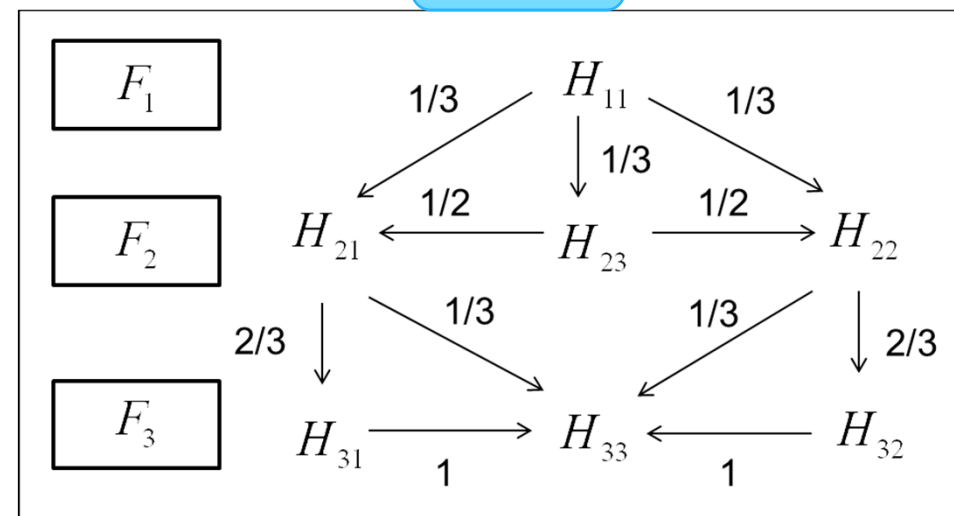


# OVERVIEW OF BTG AND GA

## BTG



## GA



Elements influencing on each procedure's performance

1. Rejection set

2. Weights for each individual hypothesis

1. Transition matrix

2. Default Alpha  $\alpha_{ij}$  for each individual hypothesis

There may be a difference of performance between BTG and GA

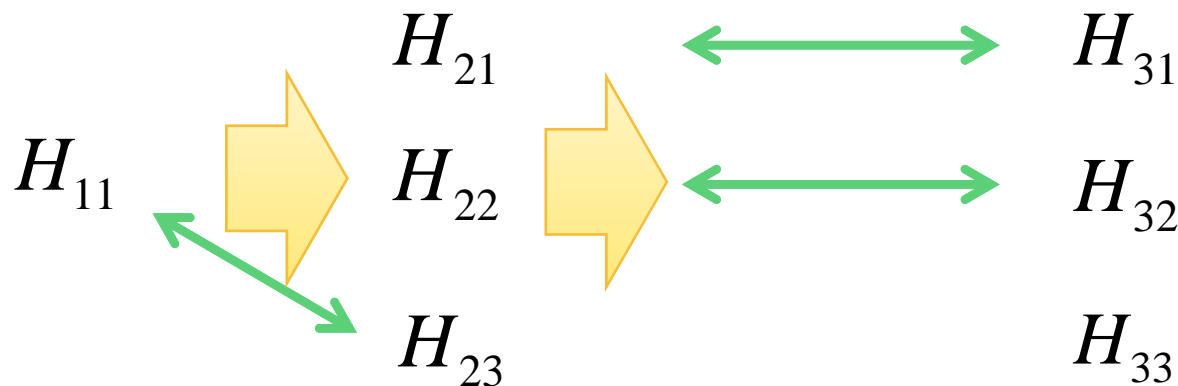
# WHY DID WE CHOOSE BTG AND GA IN THIS PIVOTAL STUDY ?

- Two reasons

1. Hierarchical structure between hypotheses

2. Test scheme

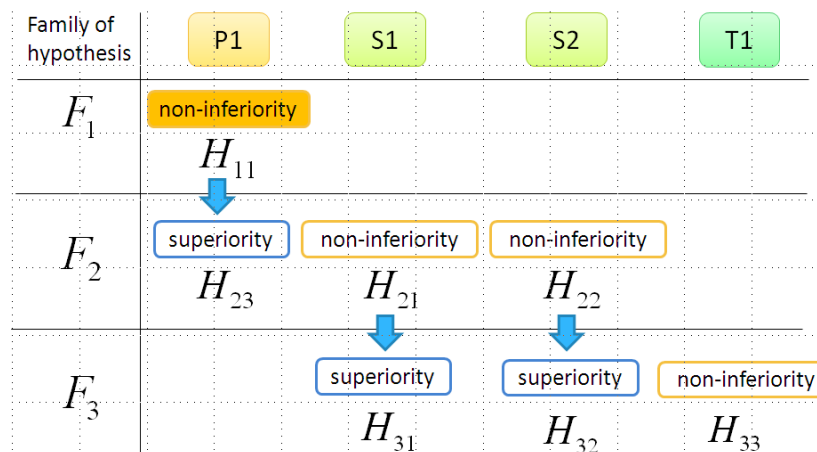
- Only if the non-inferiority for the control drug is verified for P1, S1 and S2, then we can test superiority for these endpoints
- So, we need the framework that *only if specific hypothesis in a hypothesis family is rejected, then we can test the particular hypothesis in the next hypothesis family*



# 3. ASSUMPTION OF SIMULATIONS

Case	Effect size for each hypothesis , Power	Correlations among endpoints
<b>1-1</b>	$H_{11}$ $H_{21}$ $H_{22}$ 0.265      90% $H_{23}$ $H_{31}$ $H_{32}$ $H_{33}$ 0.203      70%	Not considered
<b>1-2</b>	Same as 1-1	Considered
<b>2-1</b>	$H_{11}$ 0.265      90% $H_{21}$ $H_{22}$ $H_{23}$ $H_{33}$ 0.203      70% $H_{31}$ $H_{32}$ 0.160      50%	Not considered
<b>2-2</b>	Same as 2-1	Considered

2arms, Total Sample Size : 600

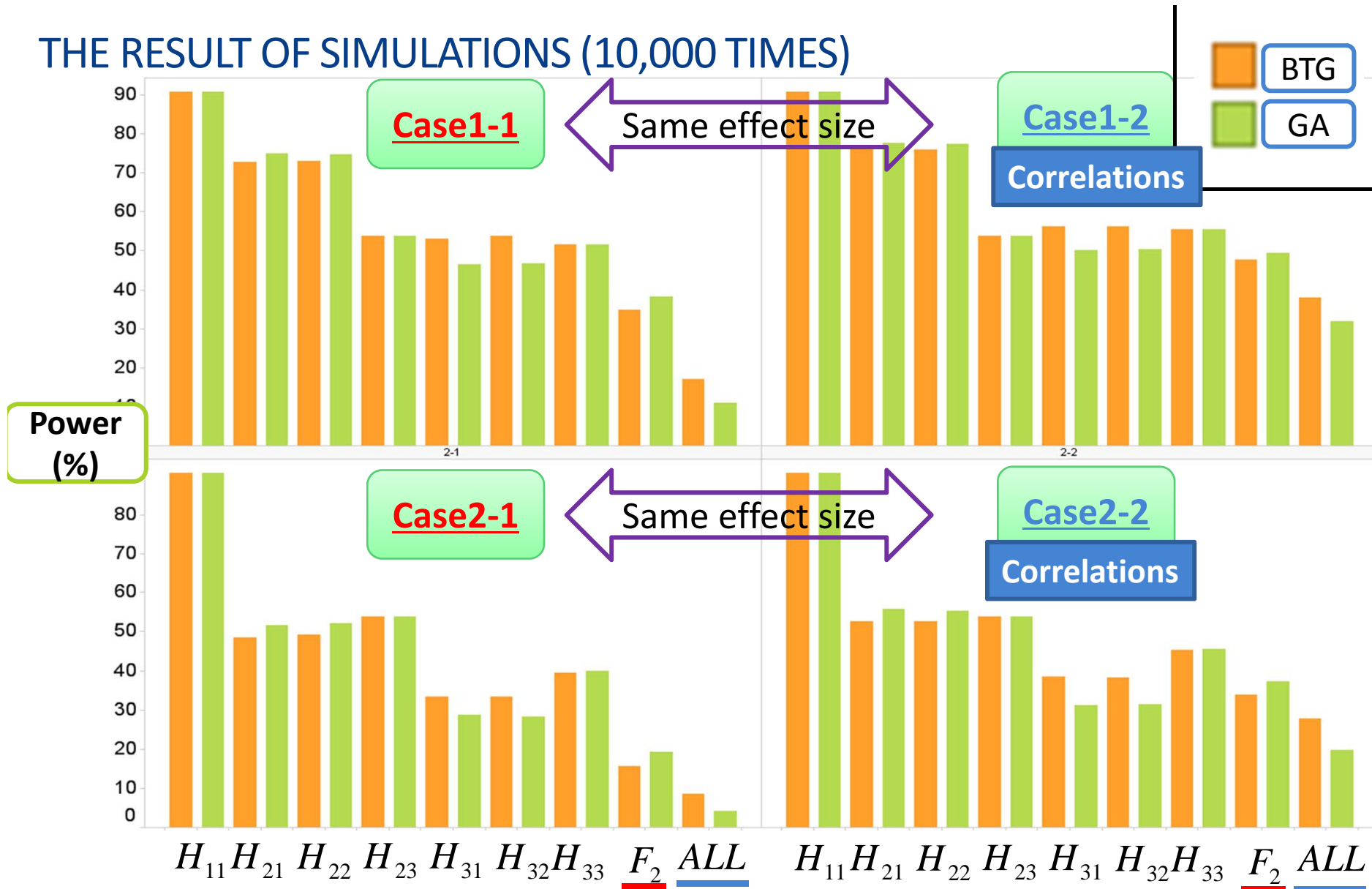


Correlation coefficient matrix

$$(\rho) = \begin{matrix} & \begin{matrix} P1 & S1 & S2 & T1 \end{matrix} \\ \begin{matrix} P1 \\ S1 \\ S2 \\ T1 \end{matrix} & \begin{pmatrix} * & 0.6 & 0.6 & 0.6 \\ * & * & 0.8 & 0.6 \\ * & * & * & 0.6 \\ * & * & * & * \end{pmatrix} \end{matrix}$$

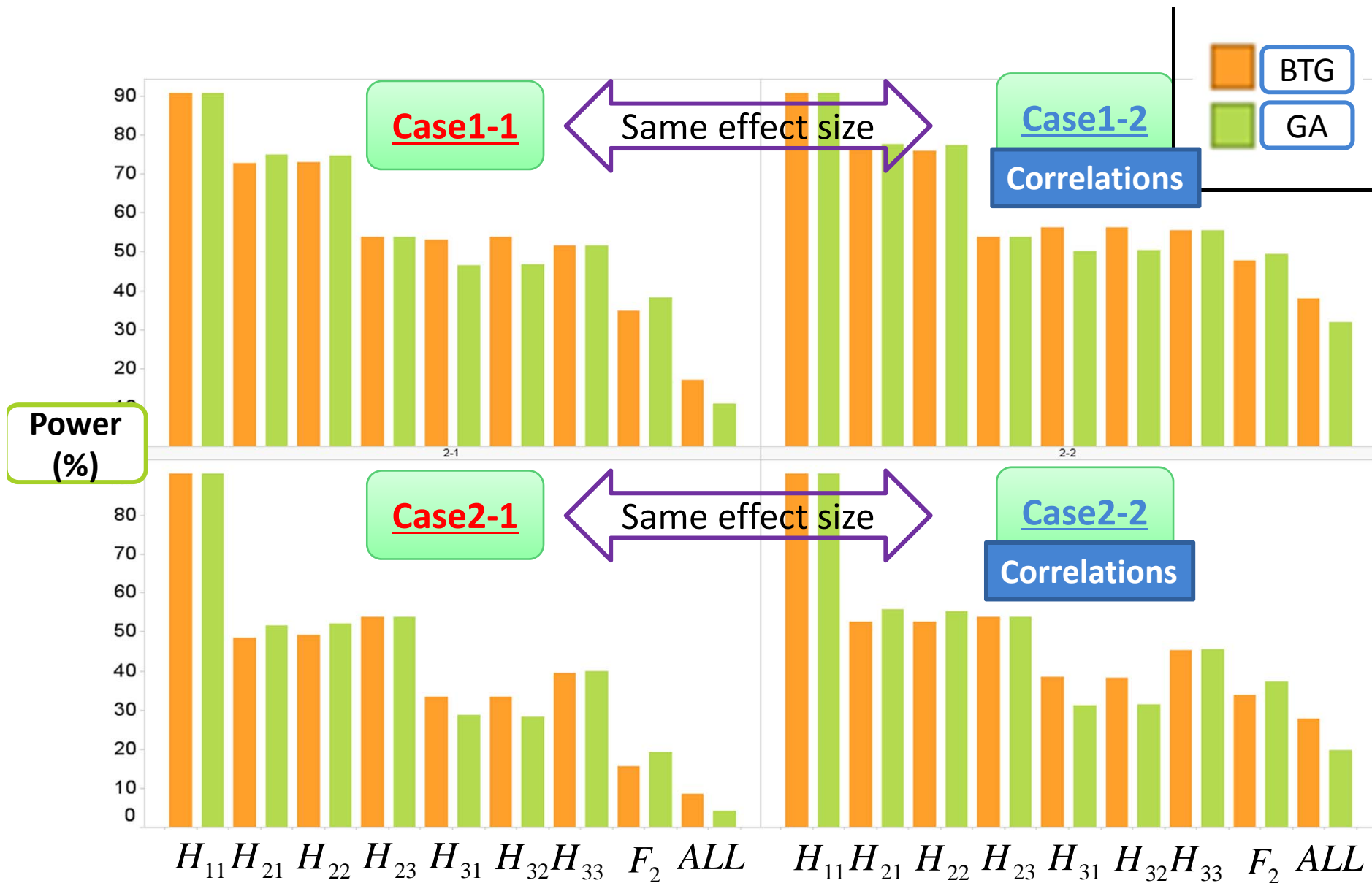


# THE RESULT OF SIMULATIONS (10,000 TIMES)

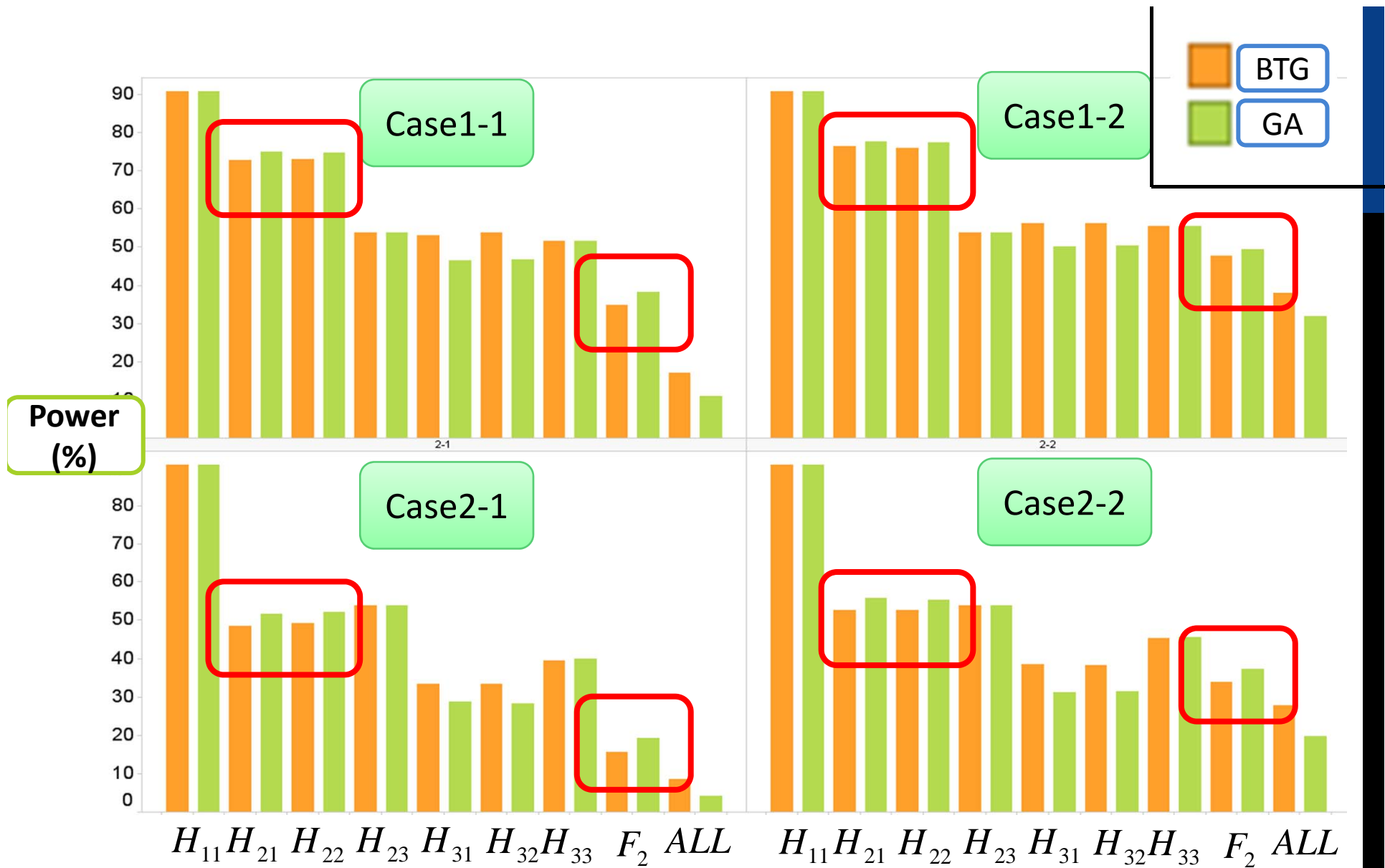


$F_2$  : the power that all hypotheses in  $F_2$  were rejected.

$ALL$  : the power that all hypotheses were rejected.



For both methods, the powers in considering correlations were higher than those in not considering, because the power for primary endpoint was 90%



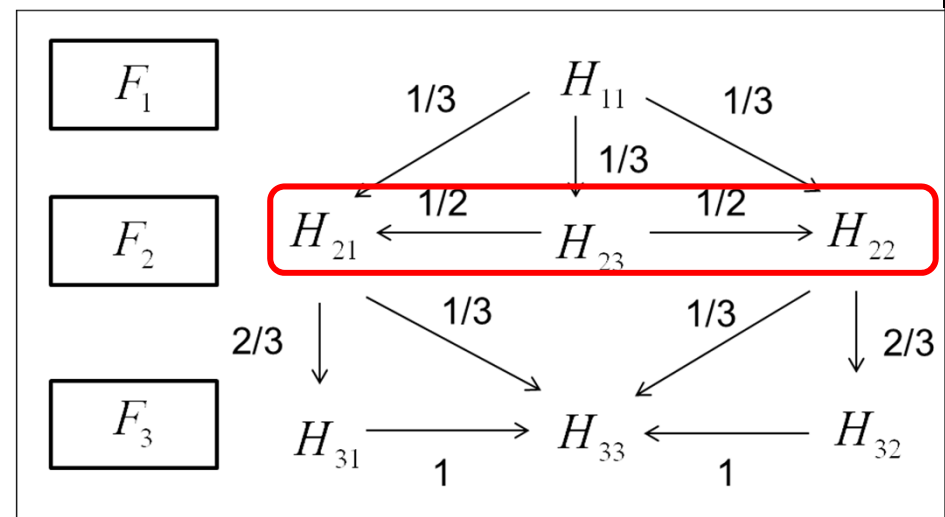
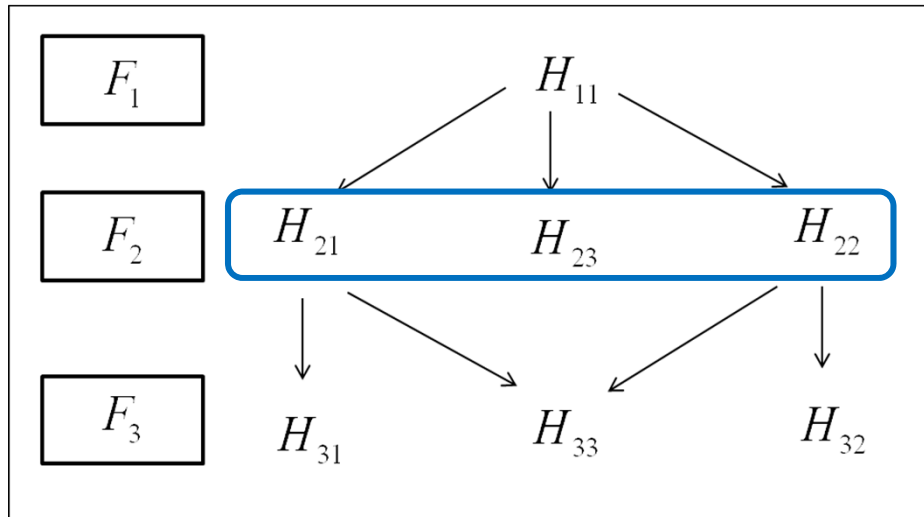
In all cases, the powers of GA for rejecting  $H_{21}$ ,  $H_{22}$  and all hypotheses in  $F_2$  were slightly higher than those of BTG

# FOCUSED ON $F_2$

$H_{23}$  was more easily rejected than  $H_{21}, H_{22}$  under a given significance level  $\alpha \dots$

BTG

GA



Adjusted p-values for  $H_{21}, H_{22}$

$3p_{21}, 3p_{22}$

$2p_{21}, 2p_{22}$



If  $H_{23}$  was rejected, then GA allocated the significant level efficiently for  $H_{21}, H_{22}$  within  $F_2$



In all cases, the powers of BTG for  $H_{31}$ ,  $H_{32}$  and "ALL" were higher than those of GA

# FOCUSED ON $H_{31}$

**BTG**

$p(H)$  : Bonferroni adjusted p-value for each intersection hypotheses  $H$

$\tilde{p}_{ij}$  : adjusted p-value for each individual hypotheses  $H_{ij}$

Intersection hypotheses	$v_{31}(H)$
$H_{31,32,33,22,23}$ $H_{31,32,22,23}$	1/6
$H_{31,32,33,23}$	2/9
$H_{31,32,33,22}$ $H_{31,33,22,23}$ $H_{31,33,22}$ $H_{31,33,23}$ $H_{31,32,33}$ $H_{31,32,23}$ $H_{31,22,23}$	1/3
$H_{31,32}$ $H_{31,33}$	1/2
$H_{31,32,22}$ $H_{31,22}$ $H_{31,23}$	2/3
$H_{31}$	1

If all of  $H_{32}, H_{33}, H_{22}, H_{23}$  were not rejected,

$\Rightarrow p(H) = \min_{i,j} \{ p_{31} / v_{31}(H) \} = 6p_{31}$ 
 $\Rightarrow \tilde{p}_{31} = \max_H p(H) = 6p_{31}$

$\Rightarrow$  If some of  $H_{32}, H_{33}, H_{22}, H_{23}$  were rejected,  $\tilde{p}_{31}$  was getting lower

$\Rightarrow$  Minimum  $\tilde{p}_{31} = p_{31}$

$\Rightarrow$   $\alpha$ -exhaustive in  $F_3$

# FOCUSED ON $H_{31}$

GA

Intersection hypotheses	$v_{31}(H)$
$H_{31,32,33,22,23}$ $H_{31,32,22,23}$ $H_{31,32,33,23}$ $H_{31,33,22,23}$ $H_{31,33,23}$ $H_{31,22,23}$ $H_{31,23}$	2/9
$H_{31,32,33}$ $H_{31,32,23}$ $H_{31,33,22}$ $H_{31,32,33,22}$ $H_{31,32}$ $H_{31,33}$ $H_{31,22}$ $H_{31,32,22}$ $H_{31}$	1/3

If all of  $H_{32}, H_{33}, H_{22}, H_{23}$  were not rejected,  $\tilde{p}_{31} = 4.5 p_{31}$

Even if  $H_{32}, H_{33}, H_{22}, H_{23}$  could be rejected,  $\tilde{p}_{31}$  was minimum  $3 p_{31}$

BTG showed a tendency to get lower adjusted p-values in  $F_3$  than GA, because BTG had  $\alpha$ -exhaustive property in  $F_3$

# SUMMARY OF SIMULATIONS

In this simulation study,

- Powers of GA to reject the all hypotheses in  $F_2$  were slightly higher than those of BTG
  - GA used the significant level efficiently in the family  $F_2$
  
- Powers of BTG to reject all hypotheses were higher than those of GA
  - BTG had  $\alpha$ -exhaustive property in the last family  $F_3$



## 4. CONCLUSION

### Consideration from the simulation results

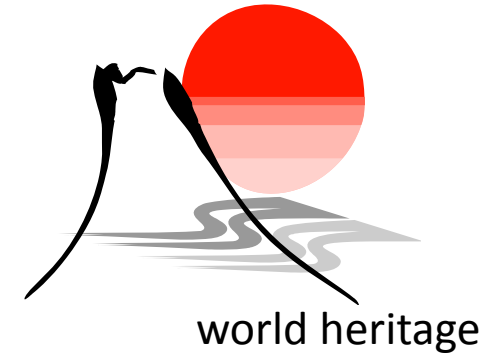
- If it is preferable to reject the hypotheses in prior families, we think that GA is appropriate
  - GA can flow the significant level flexibly in prior families
- On the other hand, if it is preferable to reject more hypotheses, we think that BTG is appropriate
  - BTG has  $\alpha$ -exhaustive property

### Future research

- GA can be modified to have  $\alpha$ -exhaustive property
- BTG and GA are Bonferroni-based methods. So, these methods can be expanded to those based on other procedures (ex. Holm, Hochberg etc). Performance of the expanded BTG and GA will be evaluated.

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**THANK YOU FOR YOUR  
ATTENTION !!!**

