

# Selective Randomization Inference for Adaptive Experiments

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# Collaborators



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- P-value:

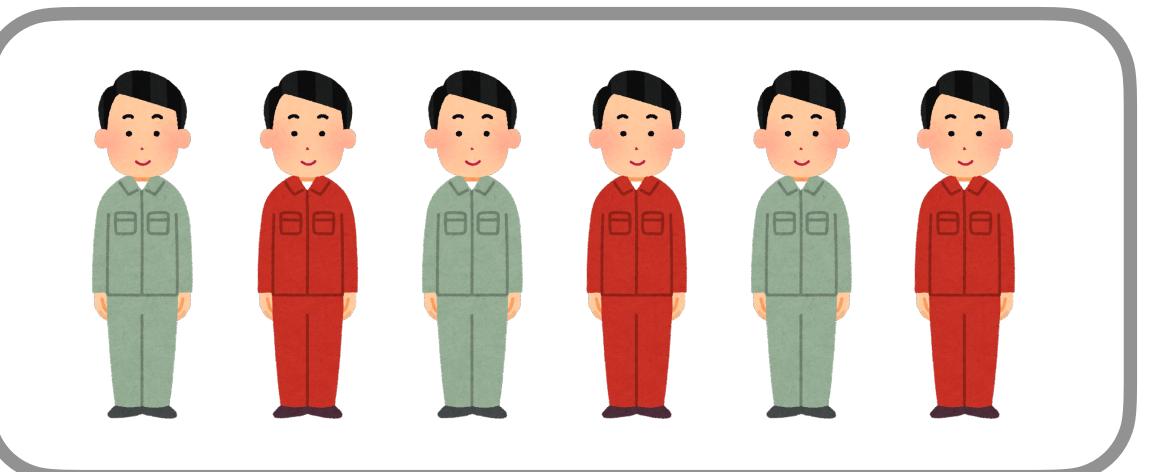
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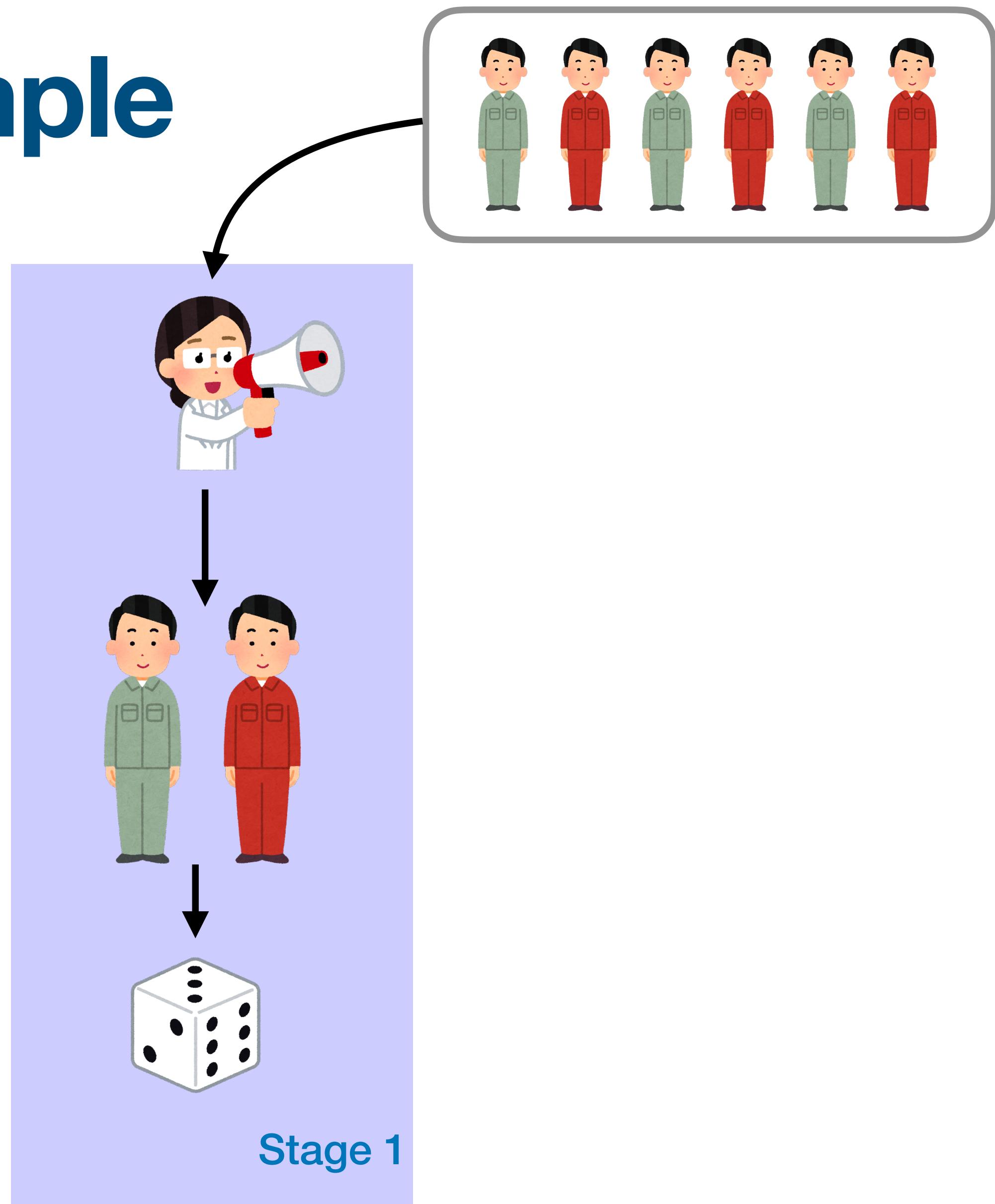
High genetic risk



Low genetic risk

FOURIER trial: Marston et al. (2020)

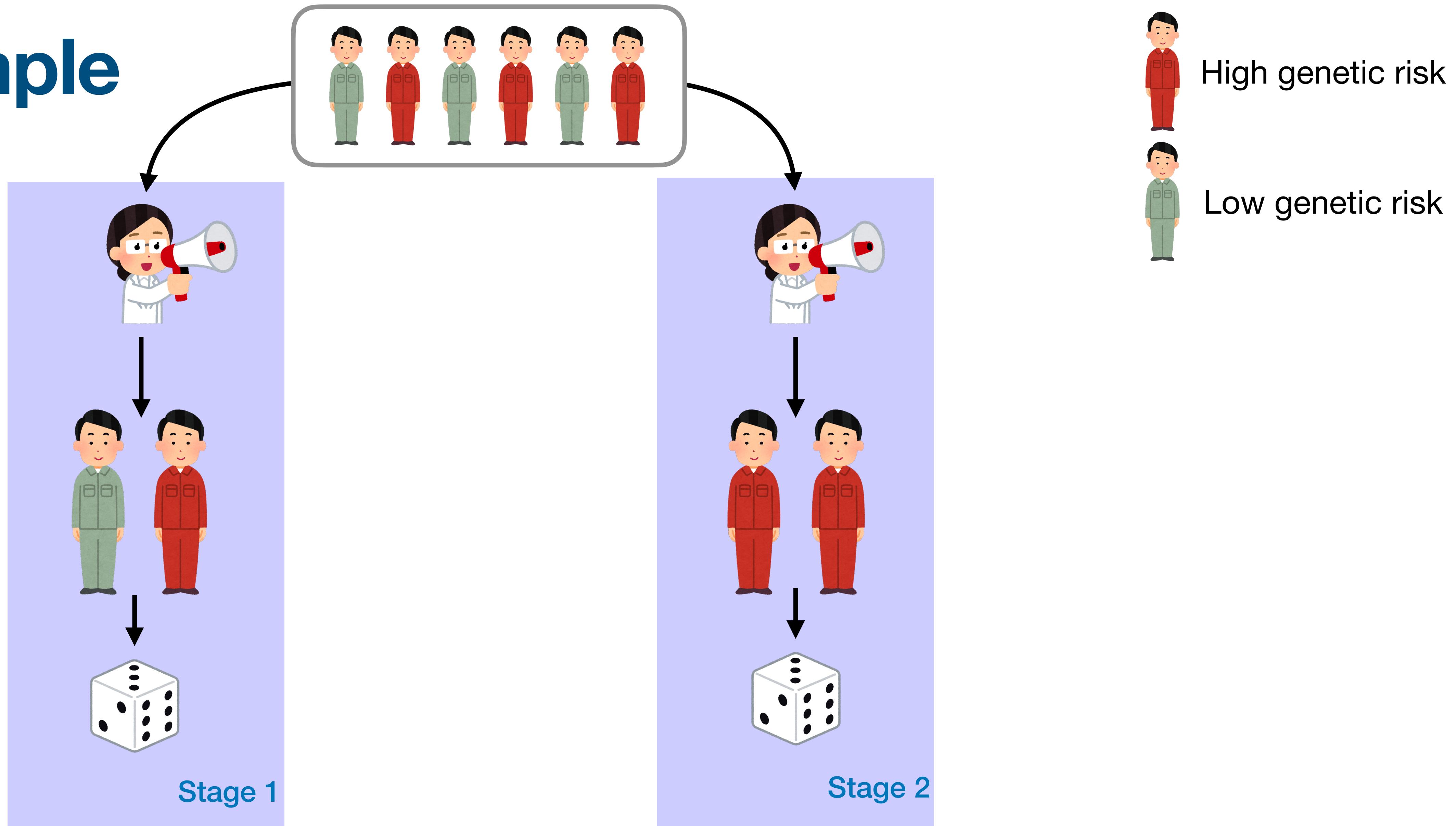
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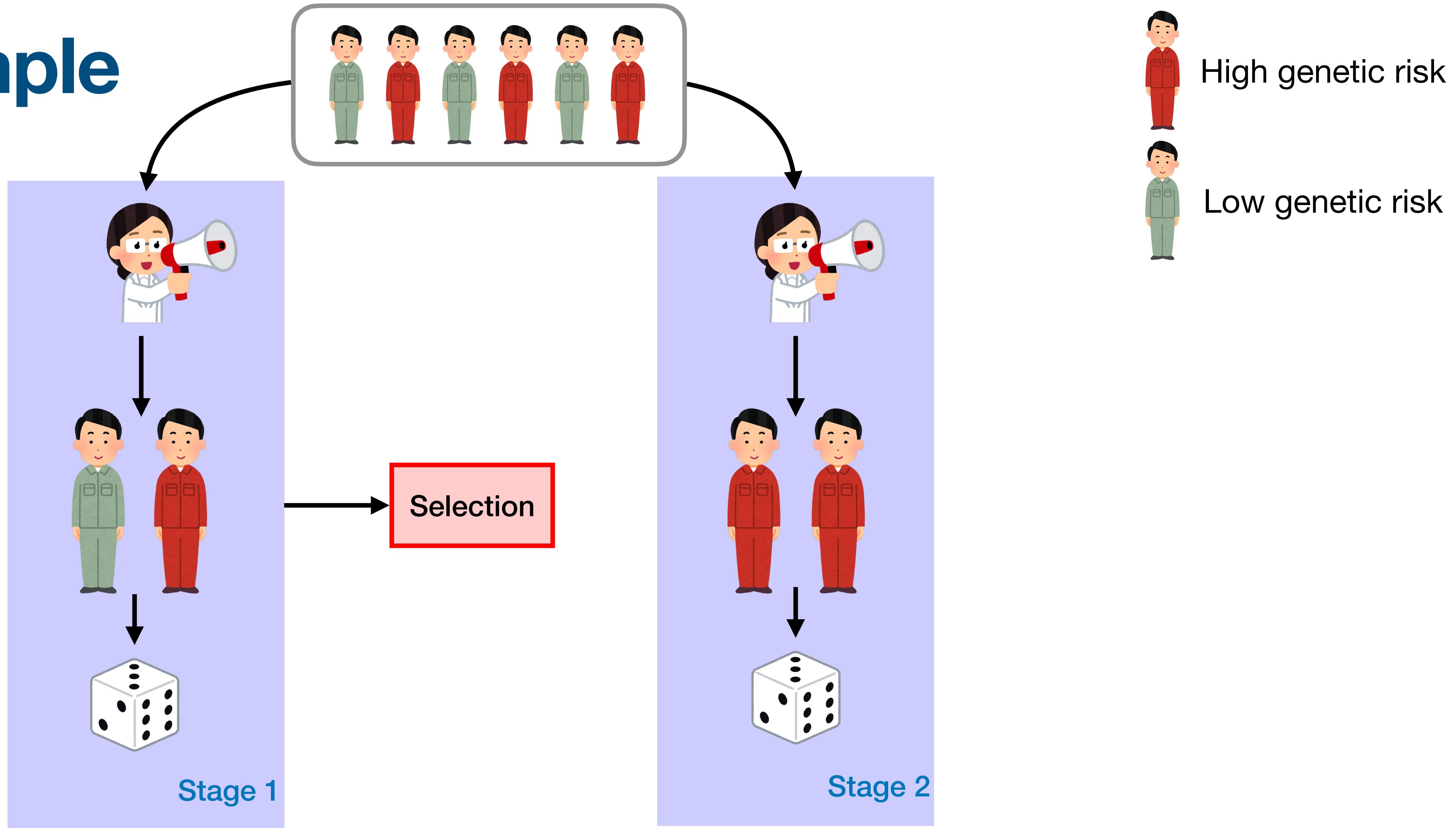
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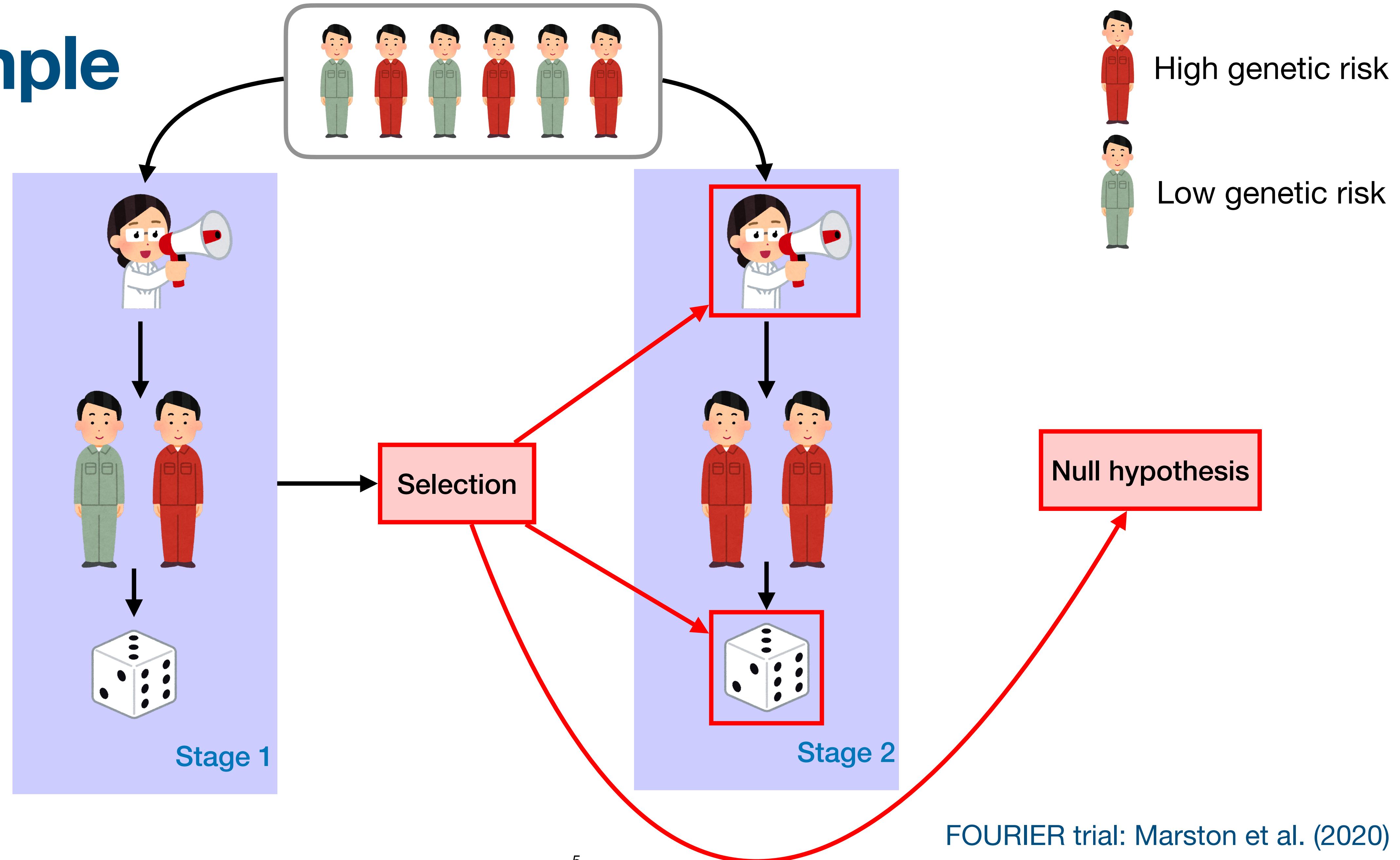
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# Graphical Model

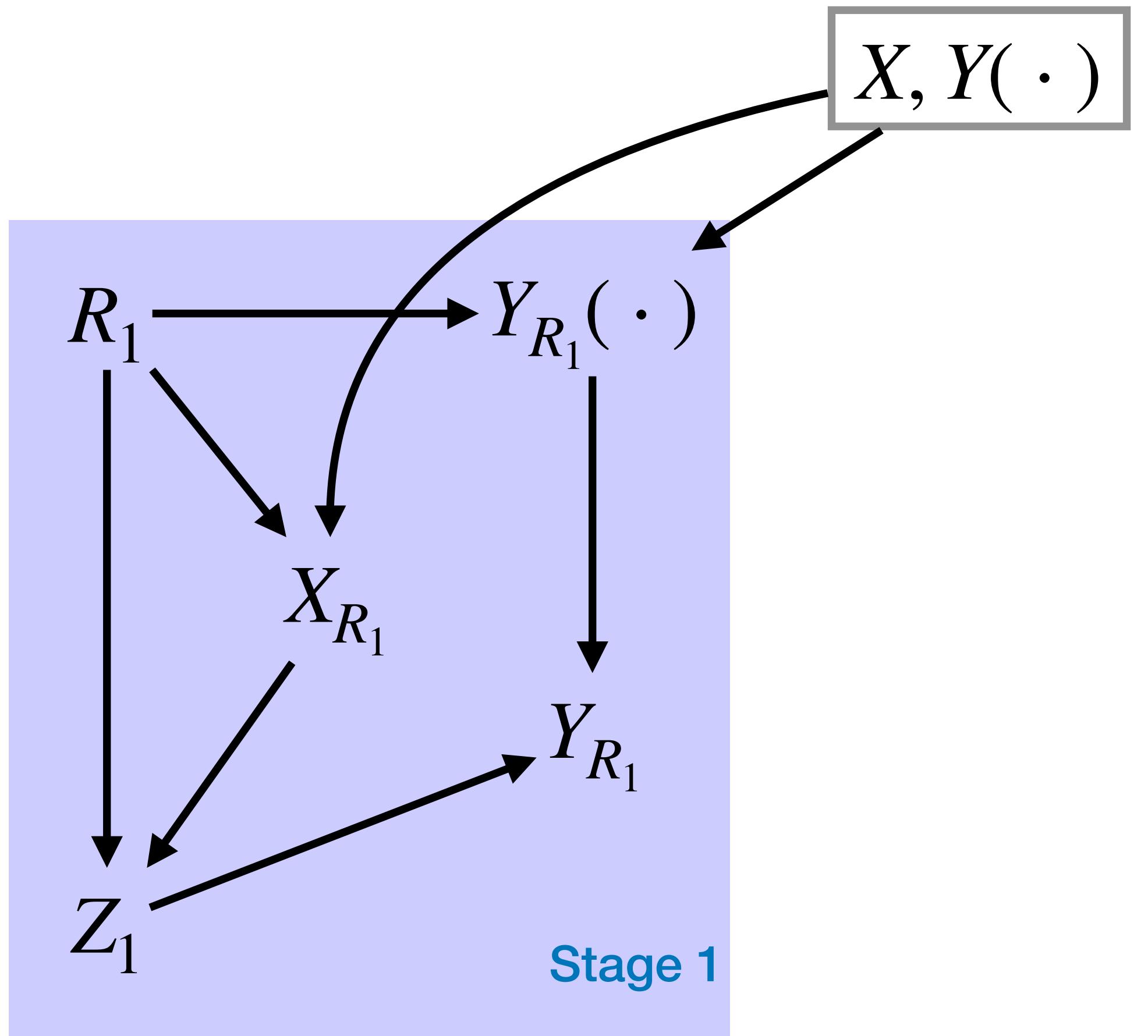
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- Covariates:  $X$
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$$X, Y(\cdot)$$

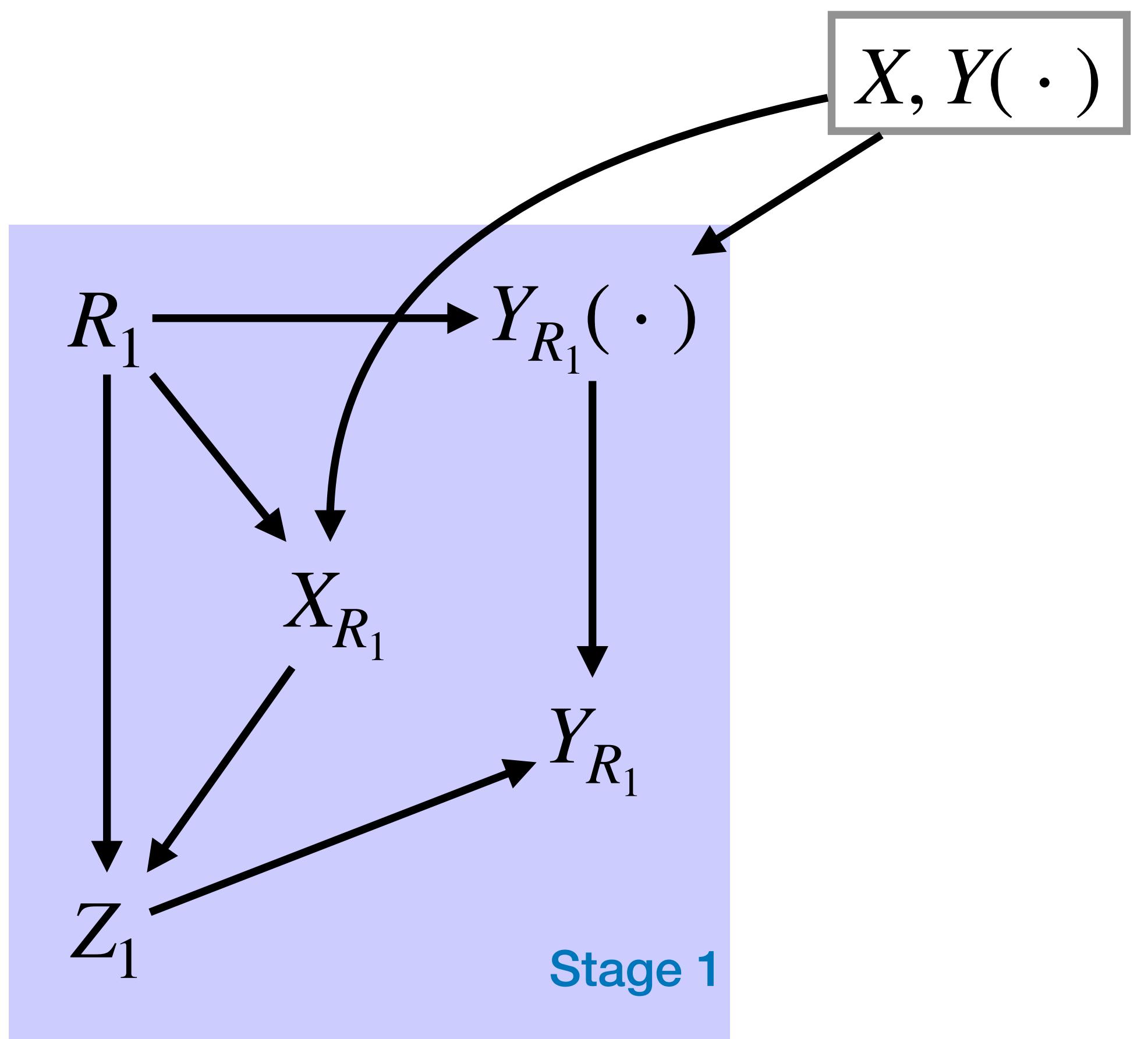
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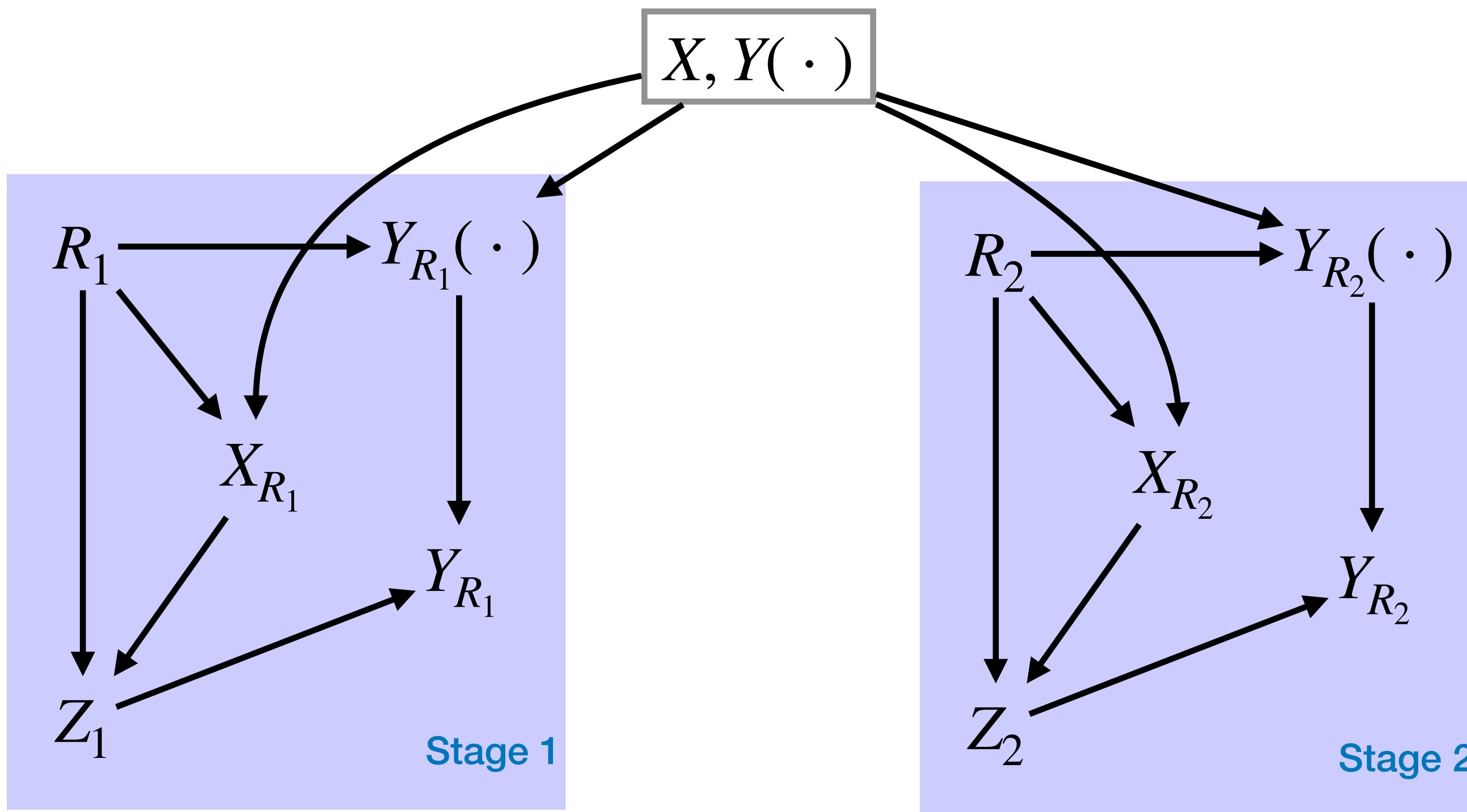


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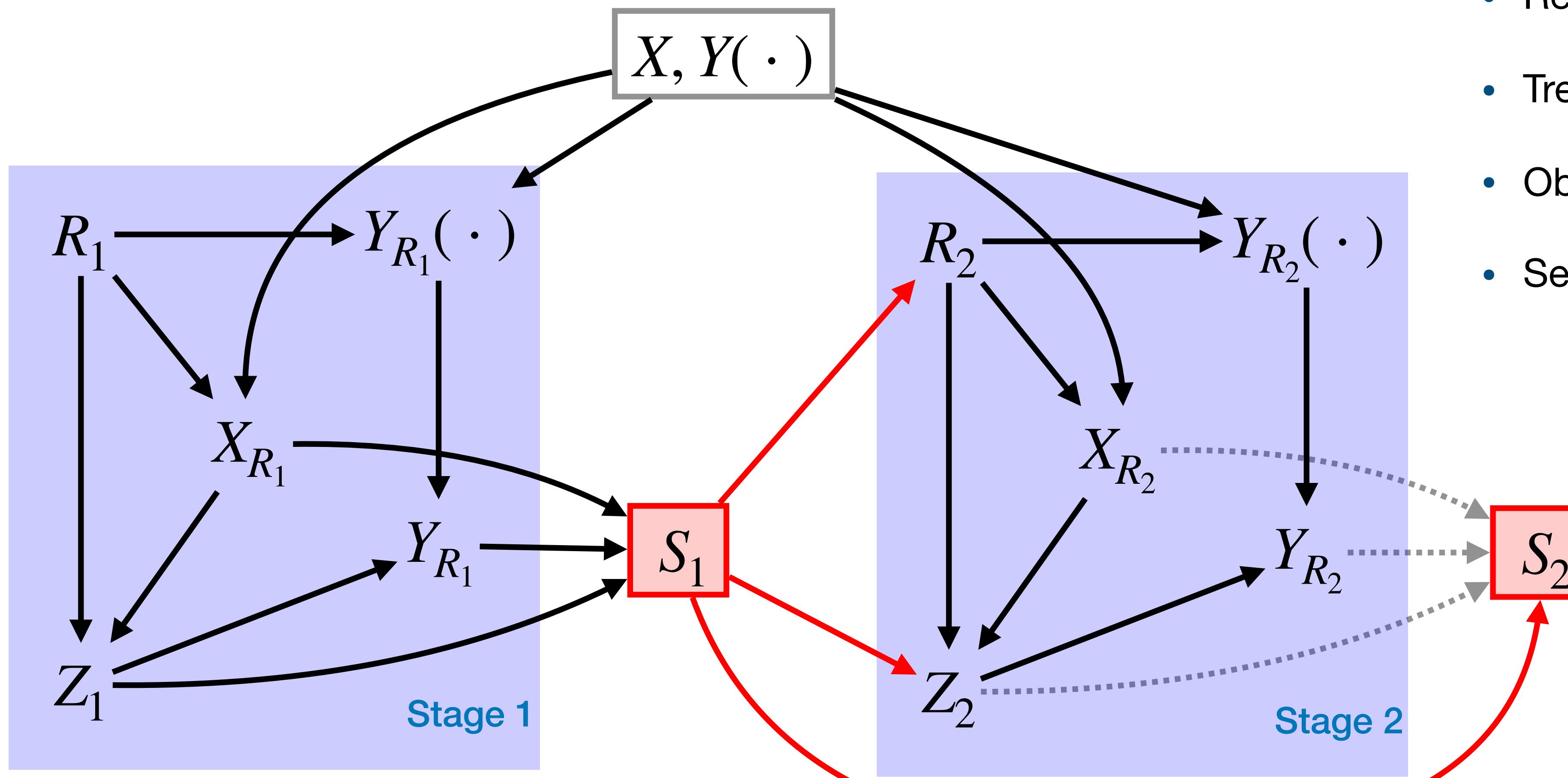
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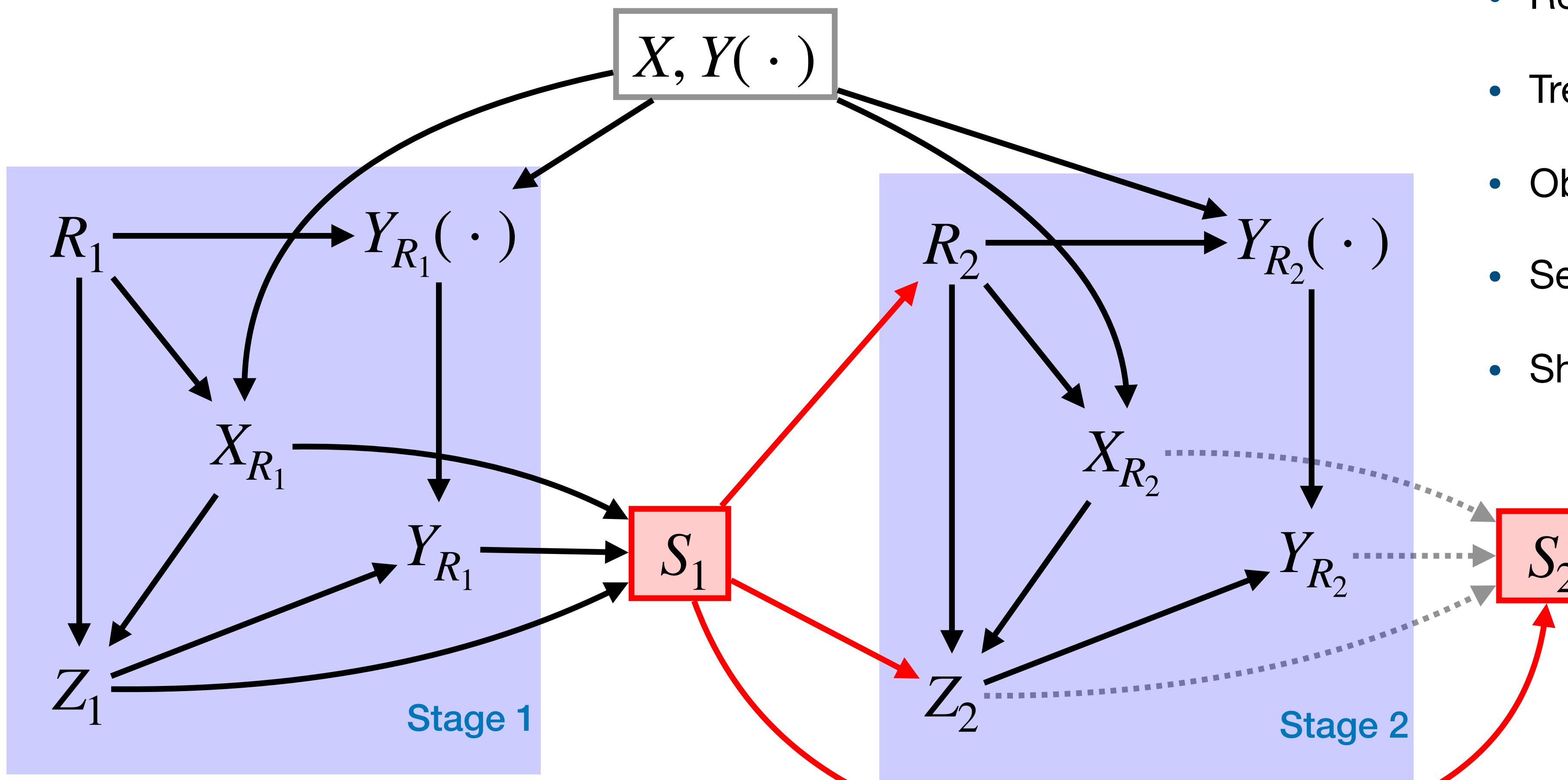
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- Short-hand:  $W = (R, X_R, Y_R(\cdot))$

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  - Bandit algorithms: exploration and exploitation
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- Analysing data from adaptive experiments despite the **dependence** between different data points

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- Is there a problem when the experiment is adaptive?

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  - Selective randomization inference:

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- Rejection sampling, Markov Chain Monte Carlo (MCMC)

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- Estimation:  $\tau$  such that  $P_{sel}(\tau) = 0.5$

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- 2 stages, 2 treatments  $Z_i \in \{0,1\}$ , 2 groups  $X_i \in \{\text{low}, \text{high}\}$
- Potential outcomes:  $Y_i(0) = Y_i(1) \sim N(0,1)$  i.i.d.
- First stage: 100 patients, Second stage: 40 patients

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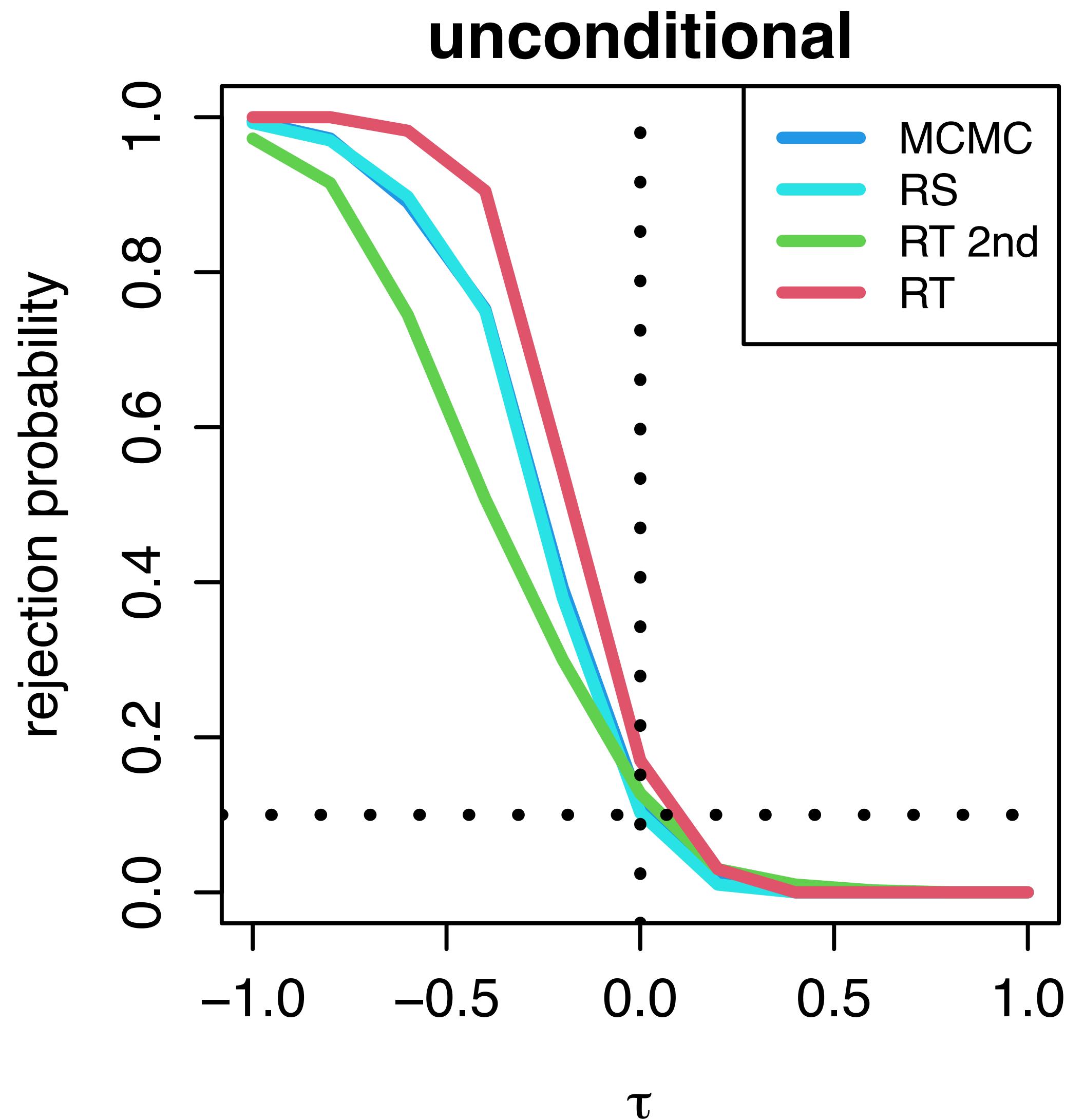
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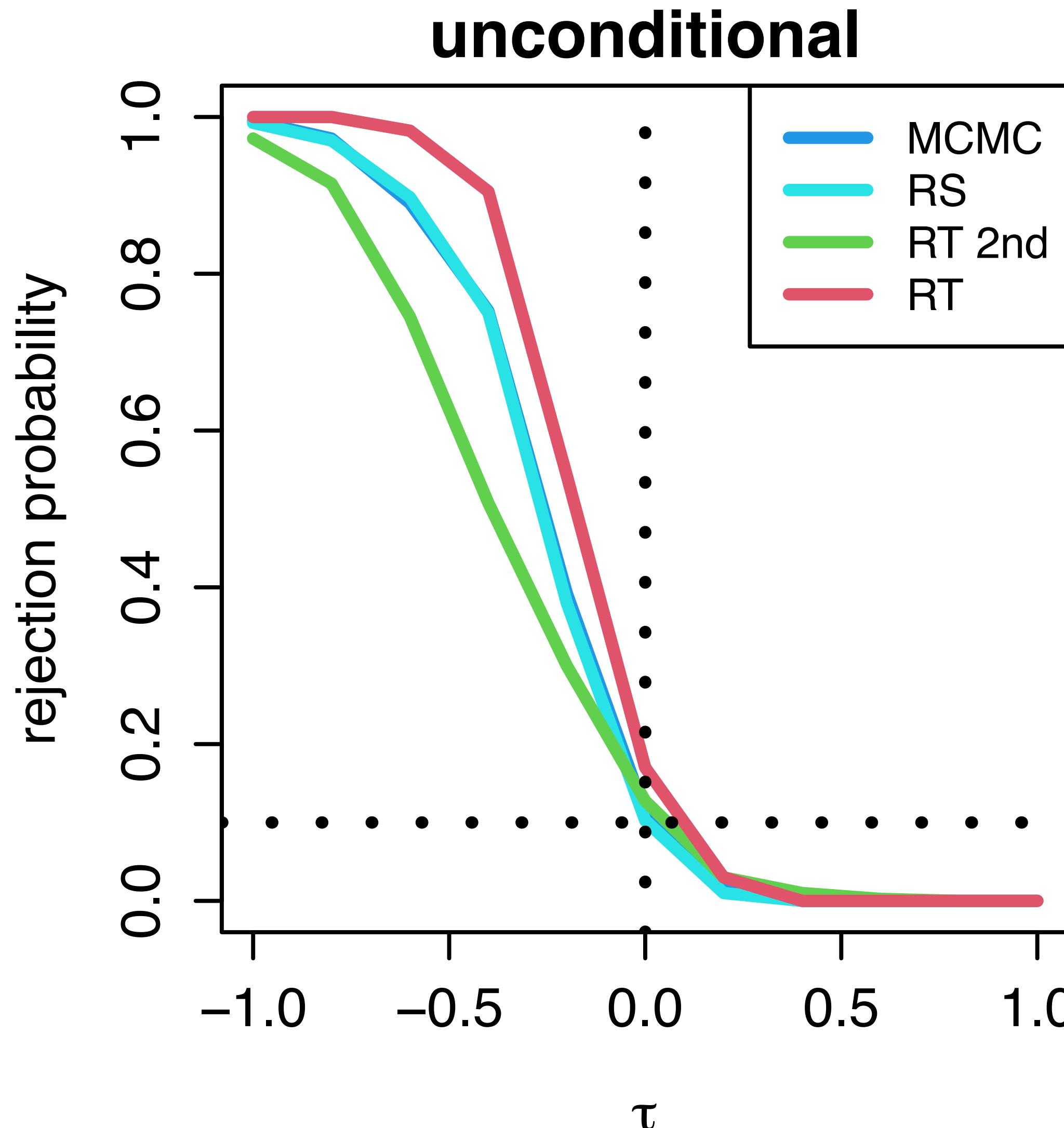
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- First stage: 100 patients, Second stage: 40 patients
- $\Delta = \text{standardized difference in SATEs between groups}$
- Selection variable:

$$S = \begin{cases} \text{only low,} & \Delta < \Phi^{-1}(0.2), \\ \text{only high,} & \Delta > \Phi^{-1}(0.8), \\ \text{both,} & \text{otherwise,} \end{cases} \quad \begin{array}{l} \text{recruit 40 from group } X_i = \text{low} \\ \text{recruit 40 from group } X_i = \text{high} \\ \text{recruit 20 from each group} \end{array}$$

# Power Analysis

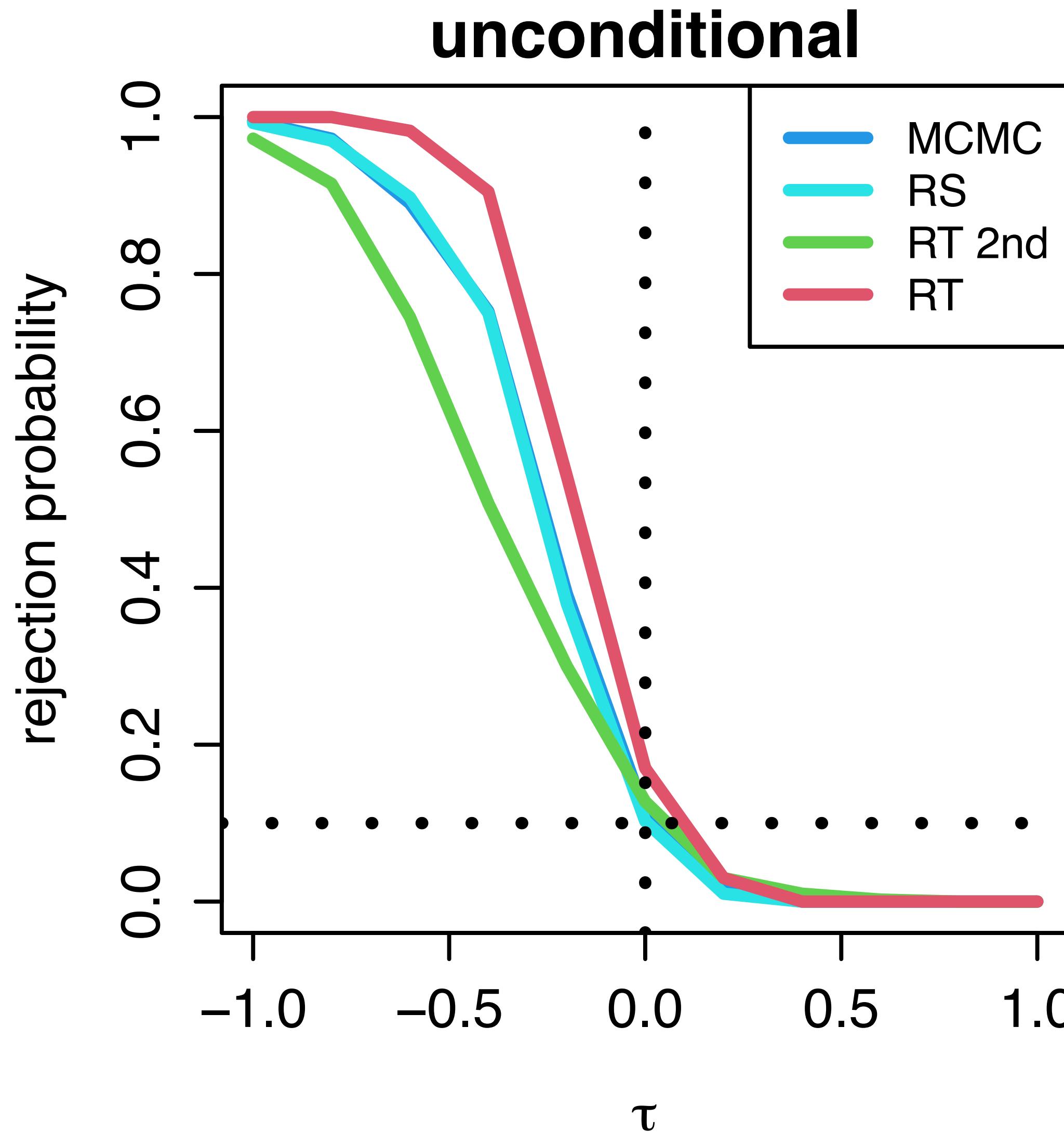


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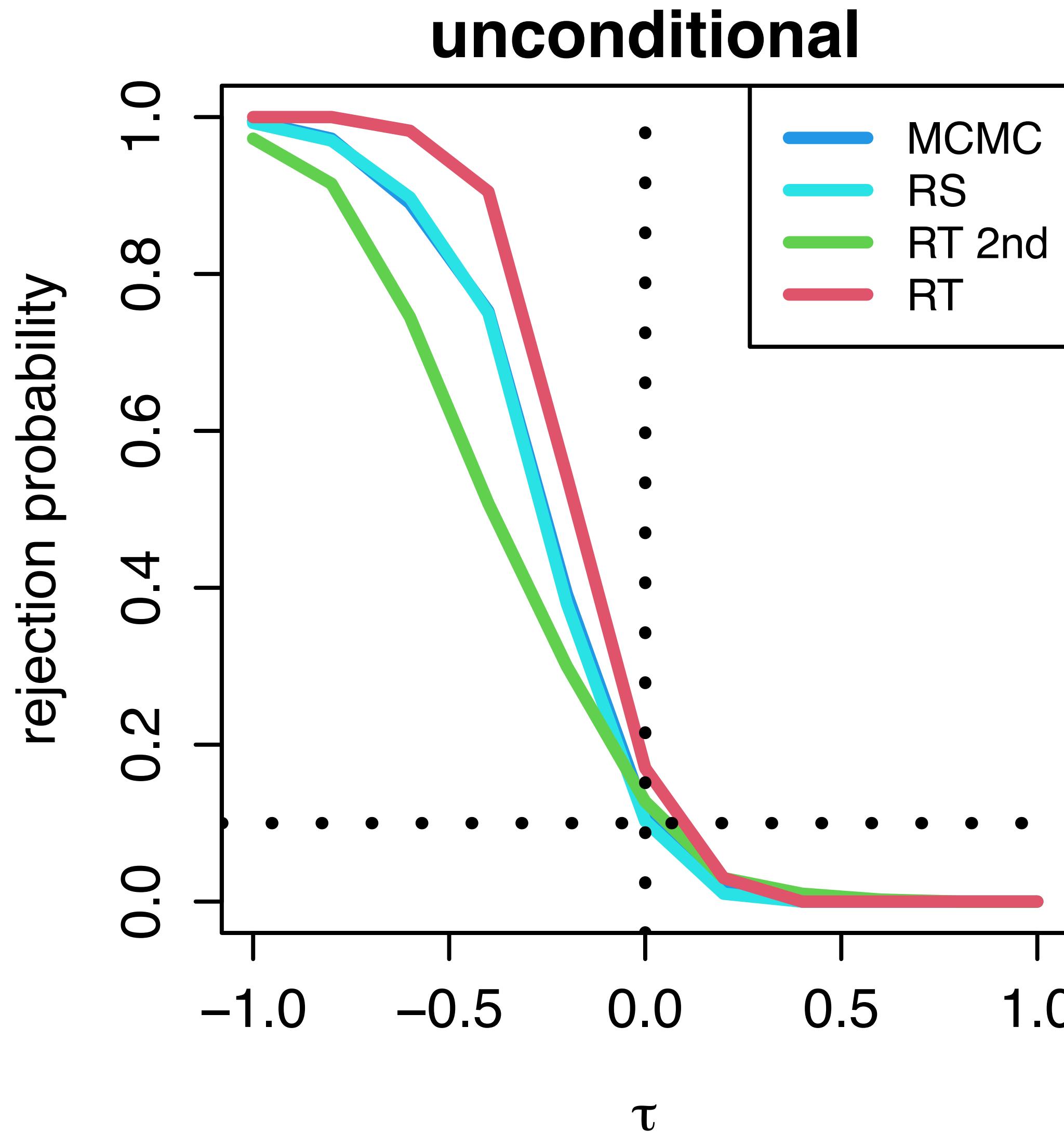
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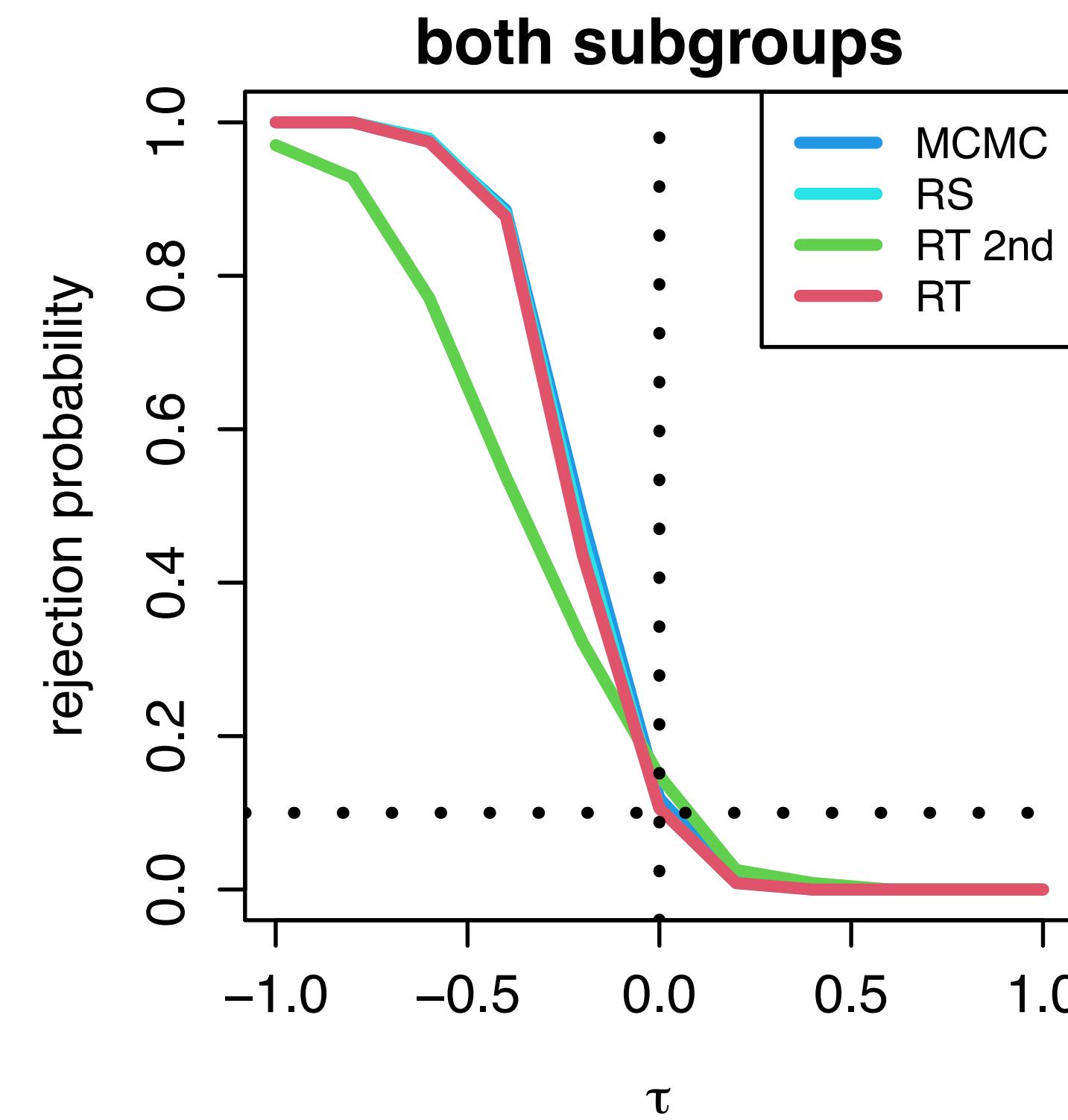
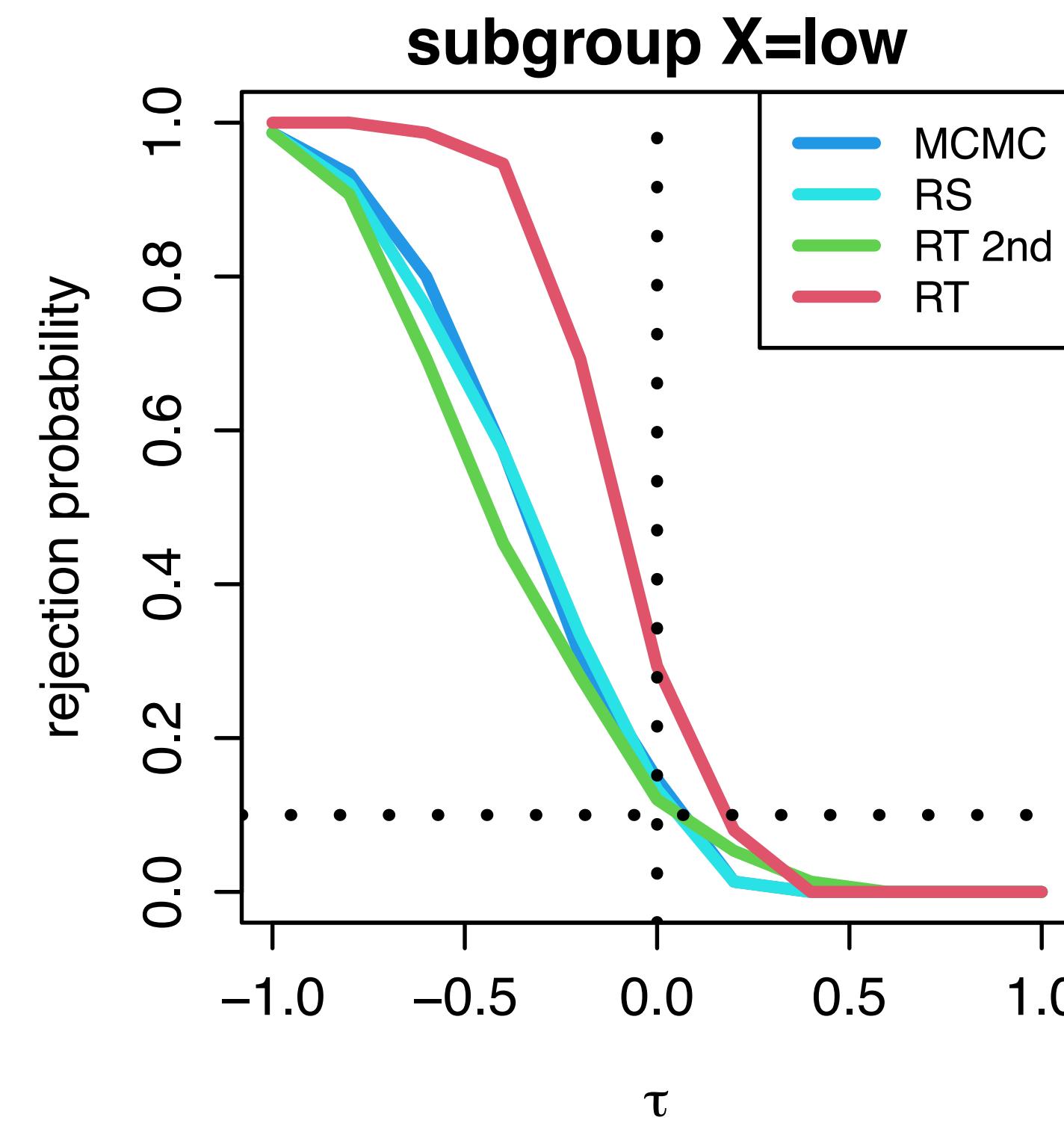
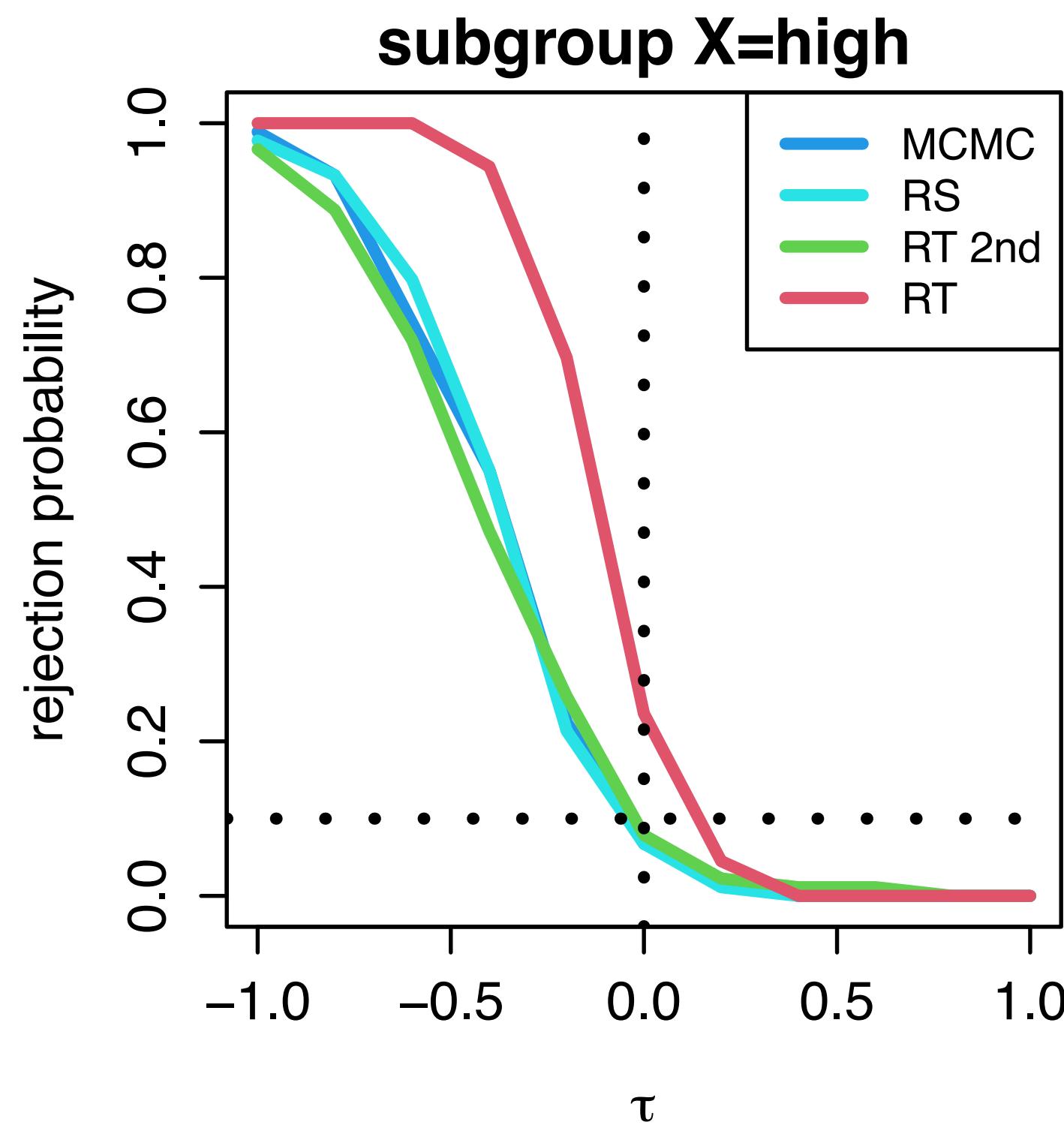
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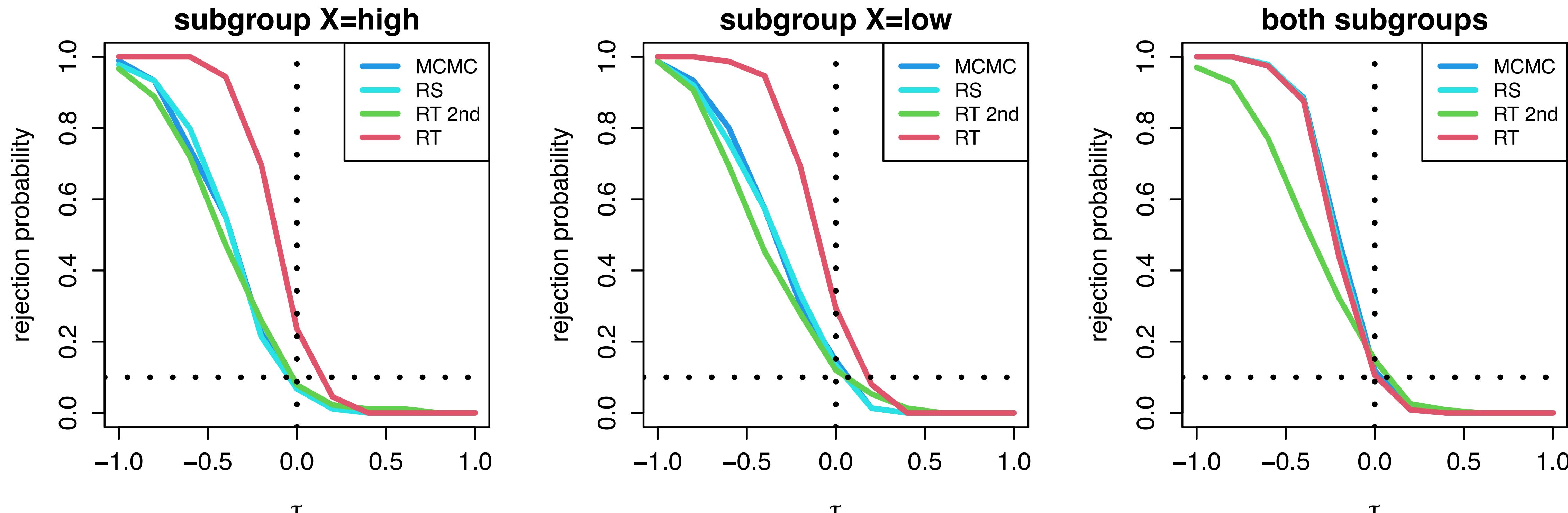


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- Selective RT: **valid and more powerful**
- Rejection sampling and MCMC lead to very similar approximations.

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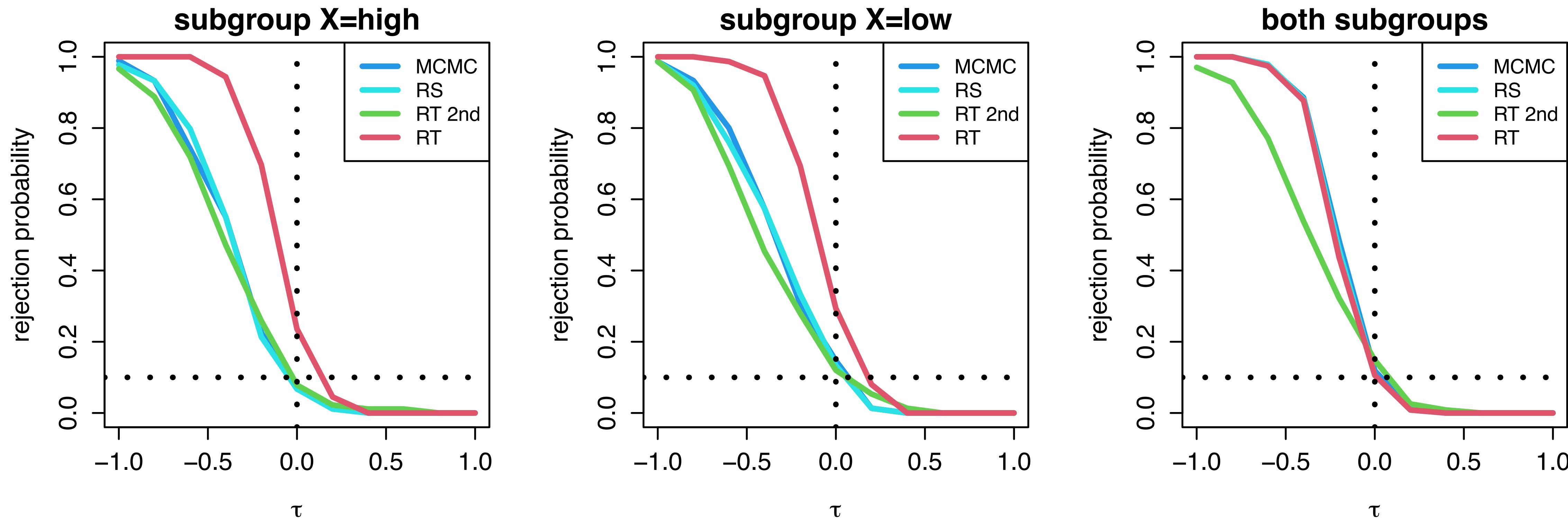


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- Type-I error control in every subgroup

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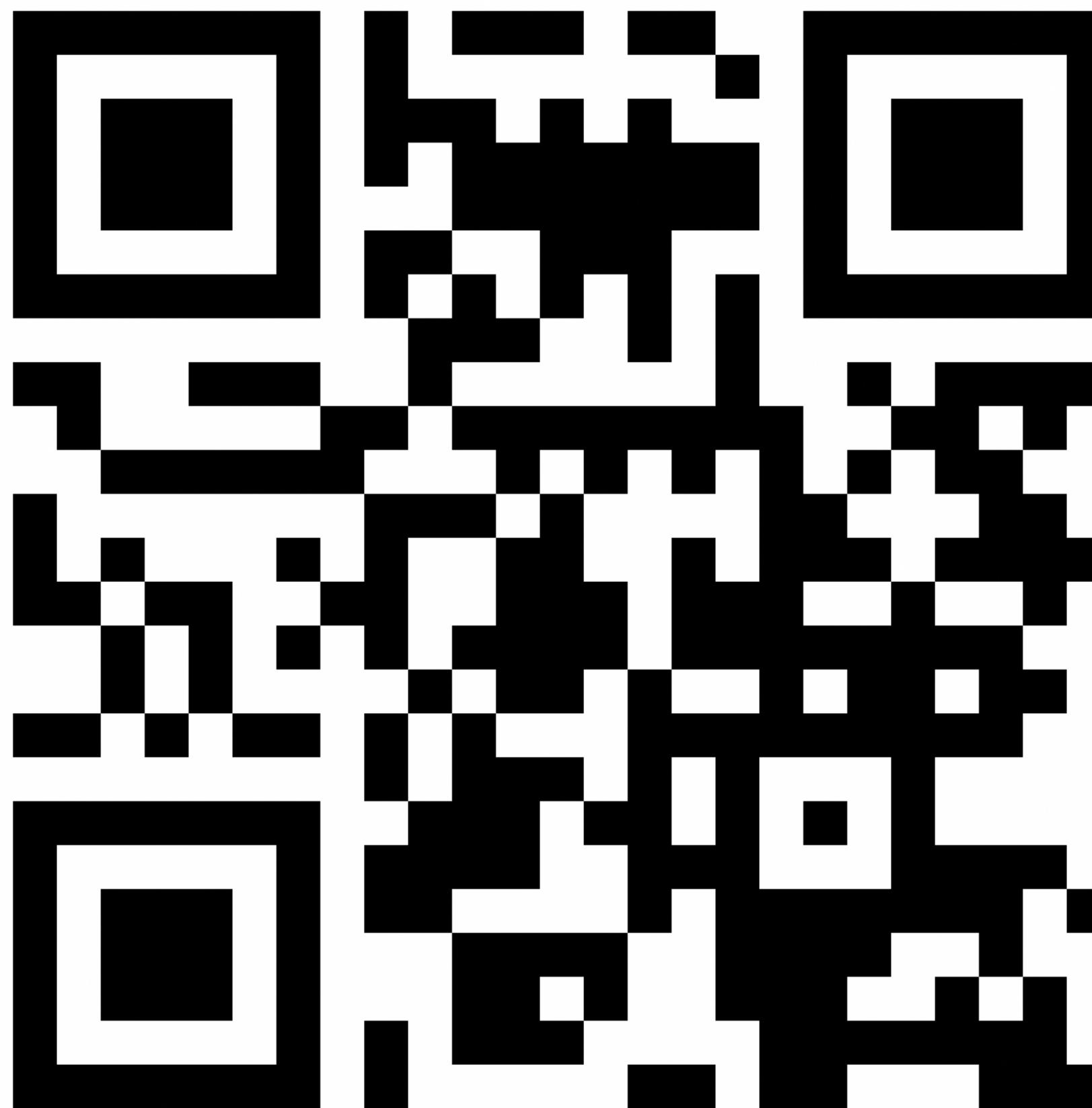


- Type-I error control in every subgroup
- Gain in power when there is a lot of “randomness left”

# Conclusion

- Experiments with adaptive treatments, recruitment and null hypothesis
- Visualization via DAGs
- **Key idea: Conditioning randomization p-value on the selection information**
- Computability under general assumptions
- Approximation via rejection sampling or MCMC

# Thanks for your attention!



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# References

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# Hold-out Units

