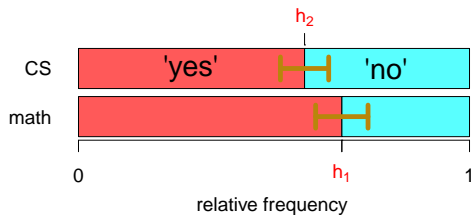


Proportions



All examples are fictitious. All data are simulated and the graphics were created with the statistical program package R.

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Sämtliche Beispiele sind frei erfunden. Alle Daten sind simuliert und die Grafiken wurden mit statistischen Programmpaket R erstellt.

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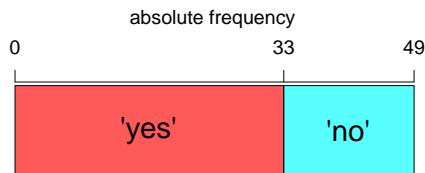
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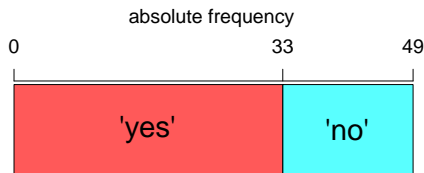
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- Thus, graphical representation, e.g., in a *barplot* (in R: `barplot()`)



Relative frequencies

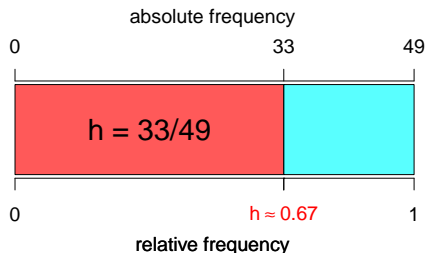
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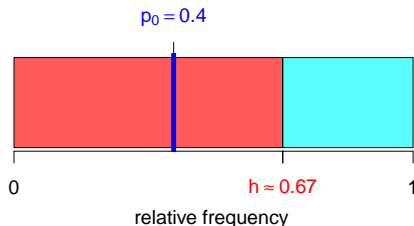
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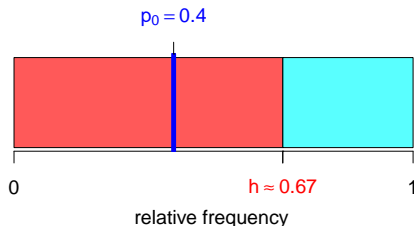
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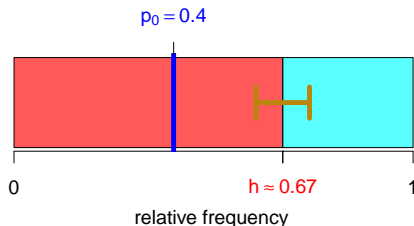
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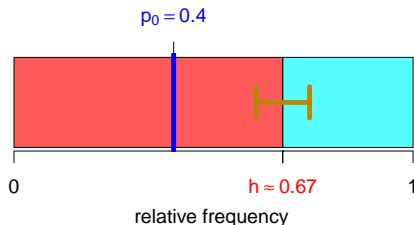
The discrepancy can be judged in the context of a statistical model...
...in which we can speak of the **variability** of the proportion



Relative frequency

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- n data in two categories (here: 'yes' and 'no')



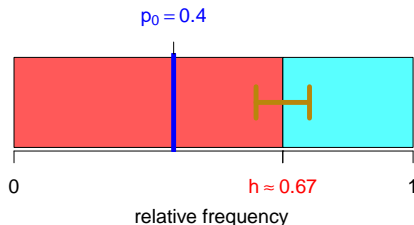
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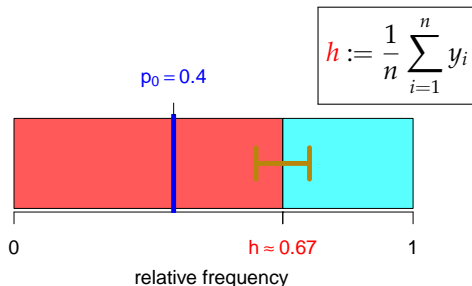
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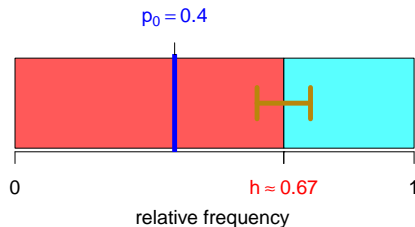


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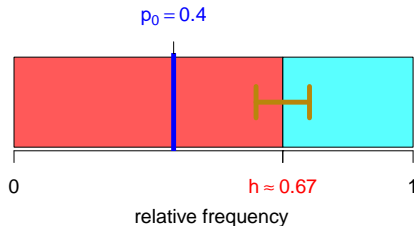
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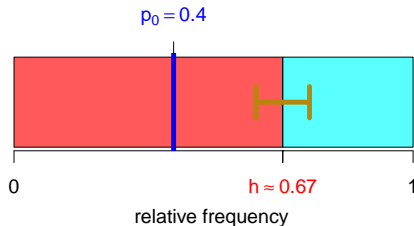
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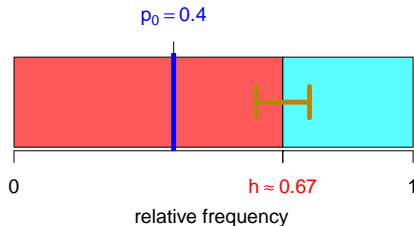
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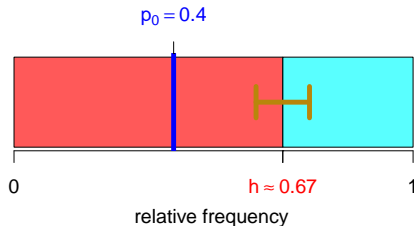
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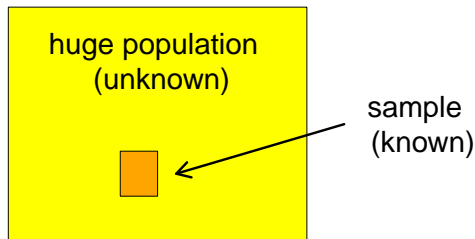
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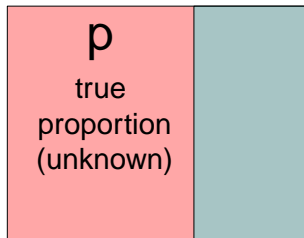
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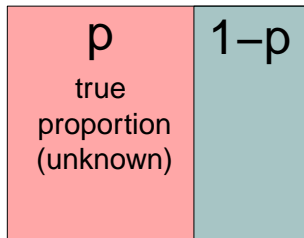
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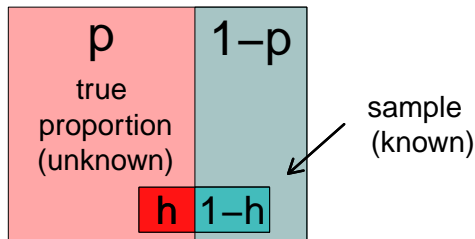
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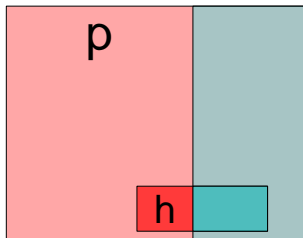
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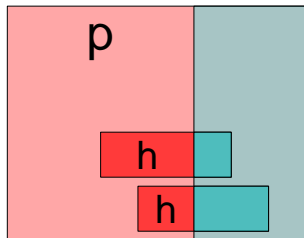
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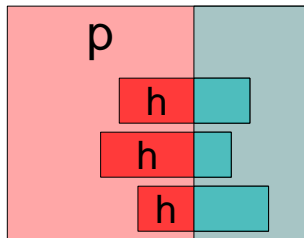
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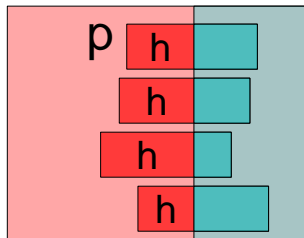
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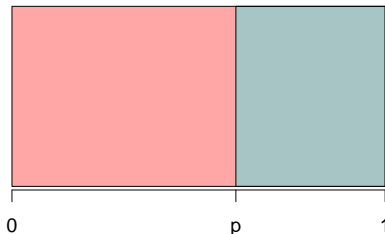
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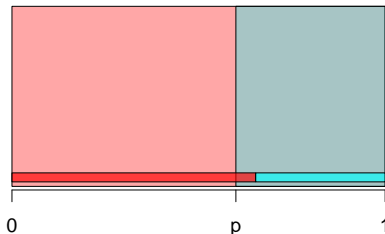
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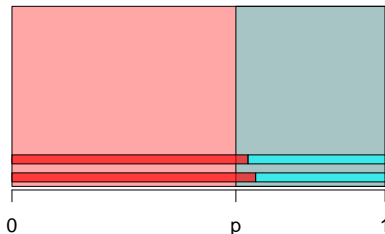
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- It holds for the expectation and the variance

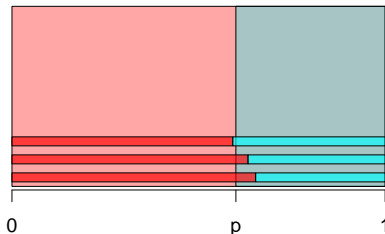
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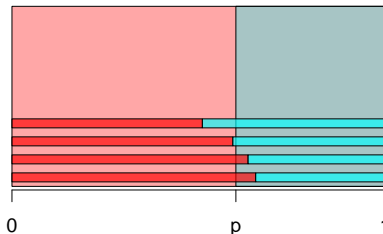
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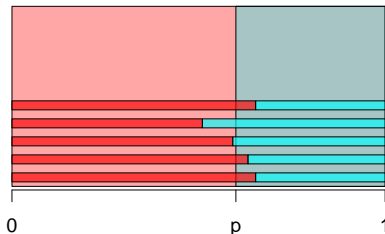
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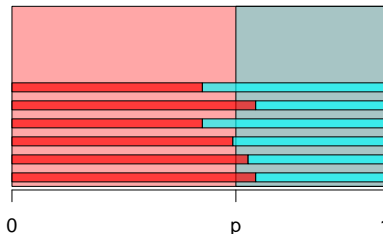
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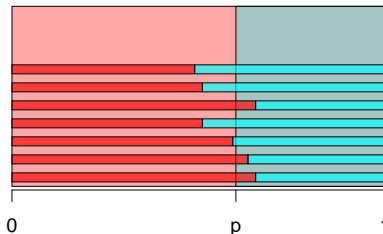
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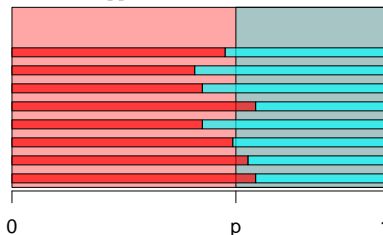
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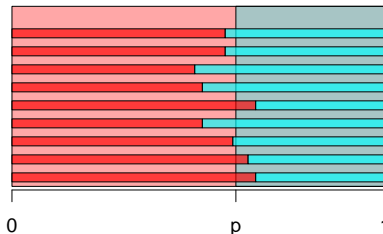
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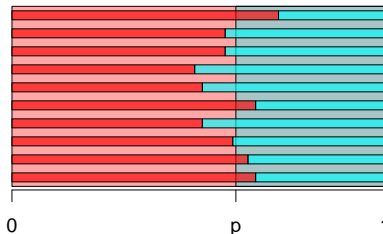
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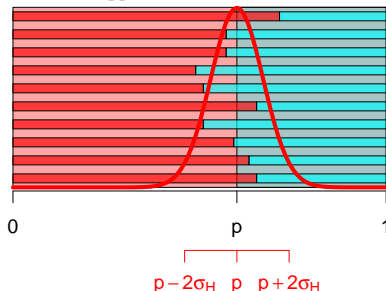
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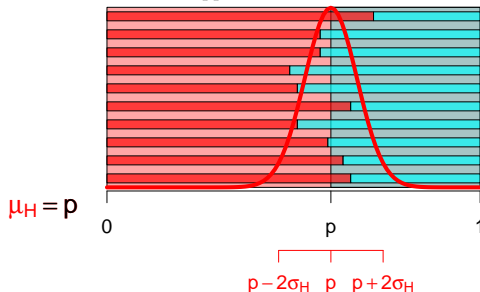
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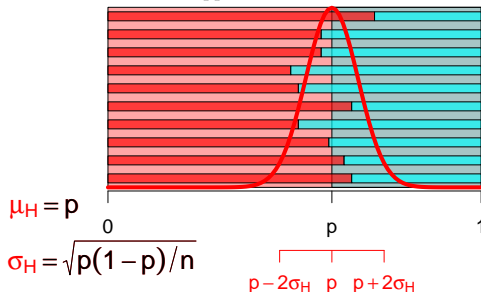
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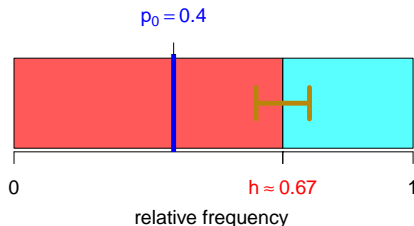
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 - but deviations of 'many' SE_H are unlikely

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- the estimation based on the data yields

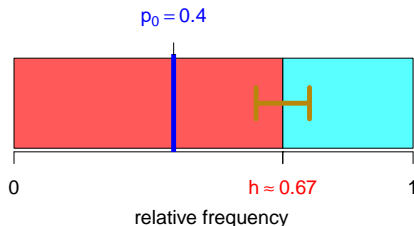
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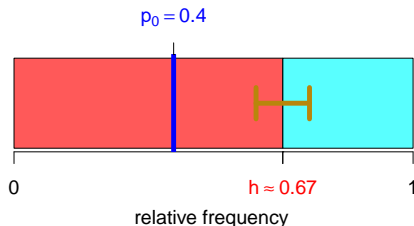
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- the discrepancy of the observed frequency $h = 0.67$ and the claimed proportion $p_0 = 0.4$ is

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- this is extremely far, given the typical deviation of H to be about $1 \cdot SE_H$

Asymptotic one-sample test for frequencies

- Let Y_1, \dots, Y_n be i.i.d. RVs with $Y_1 \sim \text{ber}(p)$ and $p \in (0, 1)$
- and let $q_{1-\alpha/2}$ be the $(1 - \alpha/2)$ -quantile of the $N(0, 1)$ -distribution

Under $H_0 : p = p_0$ it holds approximately for large n that

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$$I := (H - q_{1-\alpha/2} \cdot SE_H, H + q_{1-\alpha/2} \cdot SE_H)$$

overlaps the parameter p_0 with probability about $1 - \alpha$

- 'Structure' as in the t -test: $Z = (\spadesuit - \clubsuit) / \heartsuit$ and $I = (\spadesuit - q \cdot \heartsuit, \spadesuit + q \cdot \heartsuit)$

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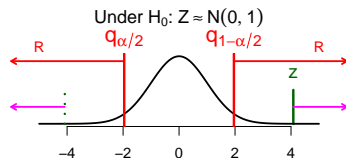
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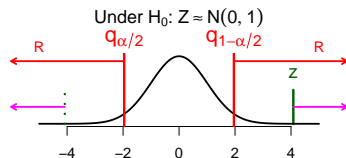
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- Thus again: Equivalence of test and confidence interval
 $\alpha = \mathbb{P}_{H_0}(Z \in R) = \dots = \mathbb{P}_{H_0}(I \not\ni p_0)$ (while R denotes the rejection area of the two-sided test)

Evaluation of the data



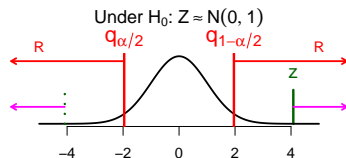
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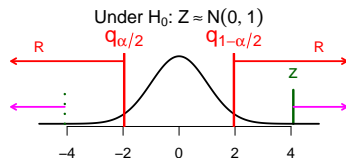
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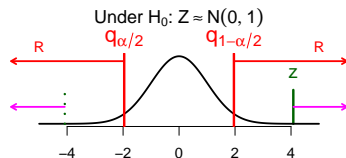
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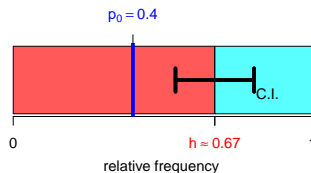
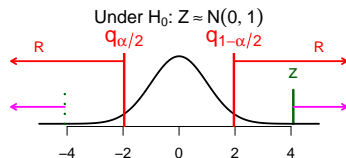
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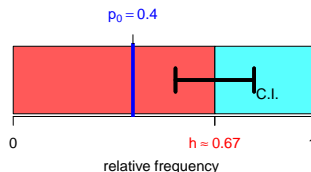
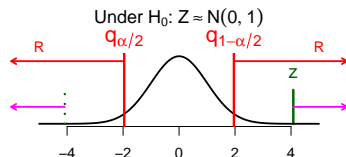
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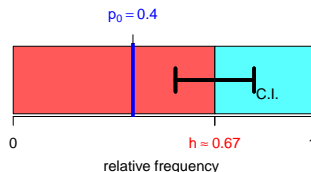
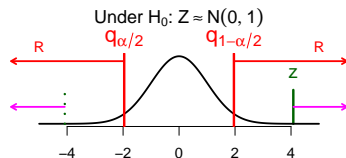
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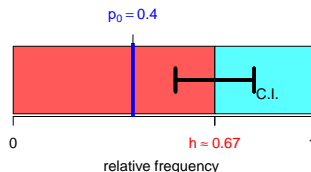
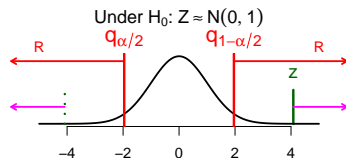
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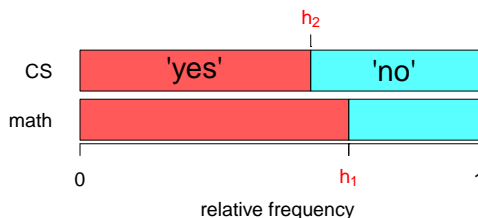
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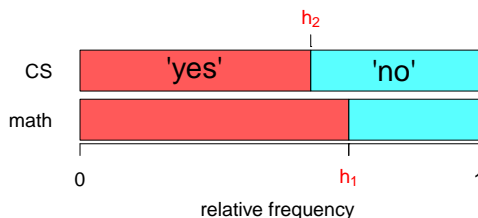
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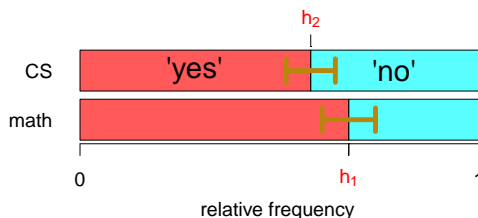
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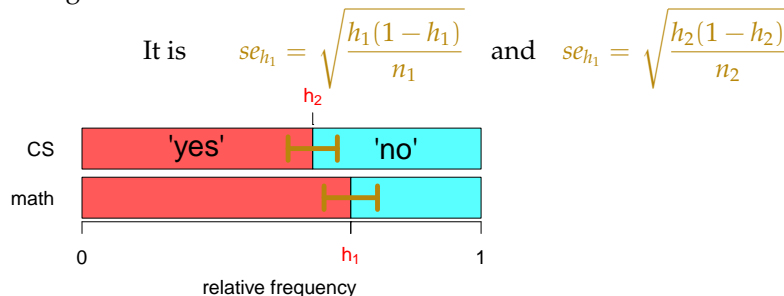
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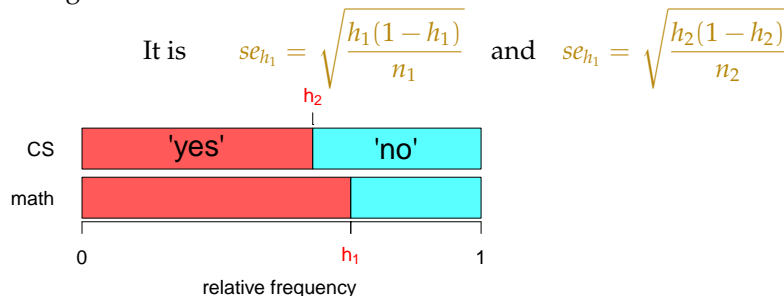
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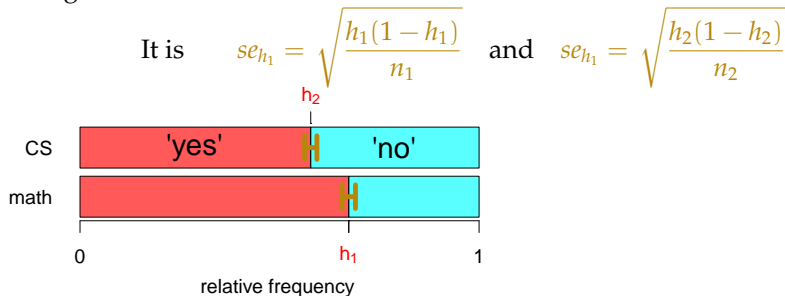


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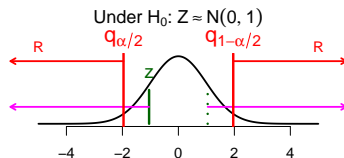
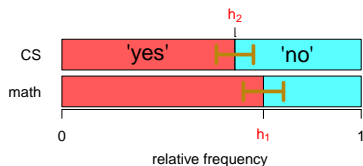
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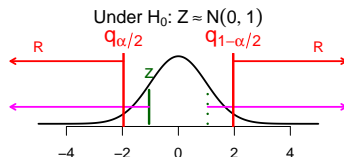
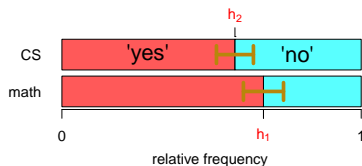
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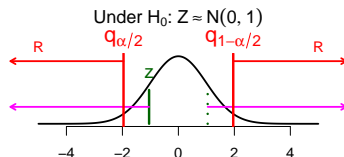
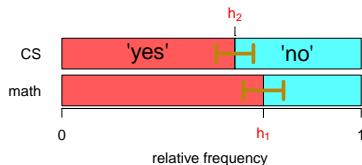
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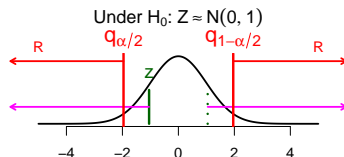
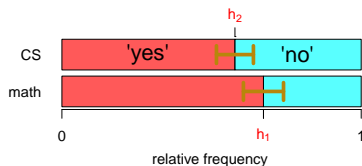
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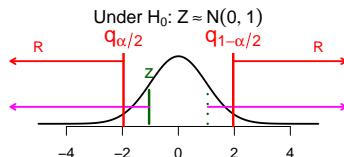
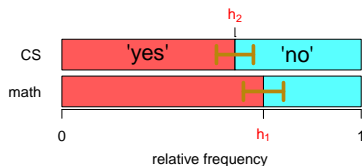


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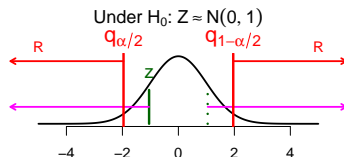
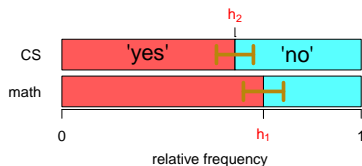
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- Because $z \notin R$ we cannot reject H_0 on the 5%-level
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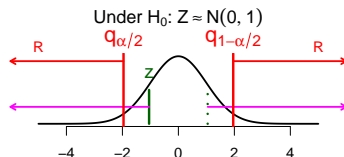
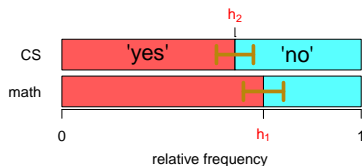
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- Interpretation: the discrepancy of the observed proportions barely gives us reason to doubt the null hypothesis. If H_0 holds true, then in about every third case we will observe a discrepancy, that is at least as large as in our data ($p \approx 1/3$)

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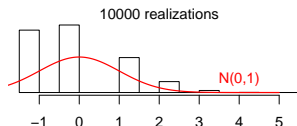
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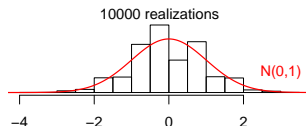
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H after rescaling, with $n=10$ and $p=0.1$



H after rescaling, with $n=40$ and $p=0.6$



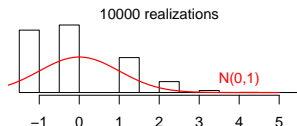
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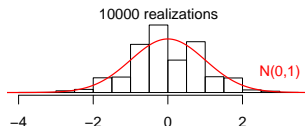
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- left: approximation unreasonable :- (right: quite plausible :-)

Thank you!