

Analysis of 2x2 contingency tables: Hypothesis tests and confidence intervals

Different versions of the Pearson chi squared tests, the Fisher exact test, and Wald confidence intervals are widely used for contingency tables. Unfortunately, some of these methods are also commonly used in situations when they perform poorly, and better alternatives exist. I will present recommended methods for 2x2 tables in different situations.

Stian Lydersen

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TUTORIAL IN BIostatistics
Recommended tests for association in 2 x 2 tables

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SUMMARY
 The asymptotic Pearson chi-squared test and Fisher's exact test have long been the most used for testing association in 2 x 2 tables. Unconditional tests preserve the significance level and generally are more powerful than Fisher's exact test for the majority of small samples, but previously were disadvantaged by being computationally demanding. The disadvantage is now gone as software to facilitate unconditional tests has been available for years. However, Fisher's exact test still may outperform other tests when the sample size is small and the effect size is small. This article describes and compares asymptotic and exact confidence intervals that are available in common software packages. We illustrate the performance of the intervals and make recommendations for both small and moderate sample sizes.

Keywords:
 2 x 2 table, NNT, odds ratio, relative risk, risk difference

1 Introduction and notation
 In a comparison of two independent binomial proportions, it is often of interest to know whether a dichotomous variable is associated with a dichotomous variable. The dichotomous variable is often the occurrence of an event (e.g. death) and the other variable is the presence of certain disease, and the proportions of diseased patients are compared between exposed and unexposed groups. Another example is unmatched case-control studies where the proportions of exposed subjects are compared between cases and controls.

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Data from Essential Science Indicators

Article

Recommended confidence intervals for two independent binomial proportions

Morten W Fagerland¹, Stian Lydersen² and Petter Laake³

Abstract
 The relationship between two independent binomial proportions is commonly estimated and presented using the difference between proportions, the number needed to treat, the ratio of proportions or the odds ratio. Several different confidence intervals are available, but they can produce markedly different results. Some of the traditional approaches, such as the Wald interval for the difference between proportions and the exact logit interval for the ratio of proportions, do not perform well when the sample size is large. Better intervals are available. This article describes and compares asymptotic and exact confidence intervals that are available in common software packages. We illustrate the performance of the intervals and make recommendations for both small and moderate sample sizes.

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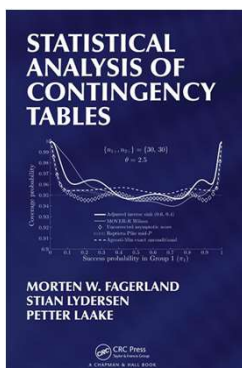
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Fagerland, M. W., Lydersen, S., & Laake, P.:
 "Statistical Analysis of Contingency Tables."
 Chapman and Hall/CRC, 2017.

www.contingencytables.com

Norsk:
 Krysstall (kontingenstall)



Hypothesis tests for associations in 2x2 tables



TABLE 4.2

A possible result of Muriel Bristol's blind taste test

	Guessed		Total
	Milk first	Tea first	
Poured			
Milk first	3	1	4*
Tea first	1	3	4*
Total	4*	4*	8*

*Fixed by design



TABLE 4.3

Treatment of epinephrine in children with cardiac arrest (Perondi et al., 2004)

Treatment	Survival at 24h		Total
	Yes	No	
Standard dose	7 (21%)	27 (79%)	34*
High dose	1 (2.9%)	33 (97%)	34*
Total	8 (12%)	60 (88%)	68*

*Fixed by design

The number of successes n_{i1} in row number i is assumed $\text{bin}(n_{i+}, \pi_i)$



Expected counts m_{ij} under H_0 :**TABLE 4.4**

Exposure to GADA for children with IPEX versus IPEX-like syndromes (Lampasona et al., 2013)

GADA	Cases/controls		Total
	IPEX	IPEX-like	
Positive	6.26 (69%)	6.74 (29%)	13 (48%)
Negative	6.74 (31%)	7.26 (71%)	14 (52%)
Total	13*	14*	27*

*Fixed by design

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TABLE 4.1The observed counts of a 2×2 table

	Success	Failure	Total
Group 1	n_{11}	n_{12}	n_{1+}
Group 2	n_{21}	n_{22}	n_{2+}
Total	n_{+1}	n_{+2}	N

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The estimated expected cell counts of the 2×2 table under the null hypothesis are

$$m_{ij} = n_{i+}n_{+j}/N, \quad i, j = 1, 2, \quad (4.5)$$

and the Pearson chi-squared test statistic is

$$T_{\text{Pearson}}(\mathbf{n}) = \sum_{i,j} \frac{(n_{ij} - m_{ij})^2}{m_{ij}} = \frac{N(n_{11}n_{22} - n_{12}n_{21})^2}{n_{1+}n_{2+}n_{+1}n_{+2}}. \quad (4.6)$$

Asymptotically chi squared distributed with 1 d.f. under H_0 .Cochran's criterion: OK if all $m_{ij} \geq 5$.

Example Table 4.4:

T = 4.464, p=0.035

The asymptotic Pearson chi-squared test does not necessarily preserve the significance level, and may perform poorly in small samples.

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Yates (1934) proposed the following modification of Equation 4.6:

$$T_{\text{YatesCC}}(\mathbf{n}) = \sum_{i,j} \frac{(|n_{ij} - m_{ij}| - 1/2)^2}{m_{ij}} = \frac{N(|n_{11}n_{22} - n_{12}n_{21}| - N/2)^2}{n_{1+}n_{2+}n_{+1}n_{+2}}. \quad (4.7)$$

The modification in Equation 4.7 is called a continuity correction, and the purpose of the correction is to obtain asymptotic P -values closer to exact P -values. The use of Yates's continuity correction—and other continuity corrections suggested in the literature—has been widely debated; see, for instance, the historical review in Hitchcock (2009). Many now consider such corrections to be no more than “interesting historic curiosities” (Hirji, 2006, p. 149).

Example Table 4.4:

T Yates CC = 2.984, p=0.084

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Possible counts x_{ij} if the marginals are fixed (or conditioned on)**TABLE 4.1**The observed counts of a 2×2 table

	Success	Failure	Total
Group 1	x_{11}		n_{1+}
Group 2			n_{2+}
Total	n_{+1}	n_{+2}	N

Additional notation:

 $\mathbf{n} = \{n_{11}, n_{12}, n_{21}, n_{22}\}$: the observed table $\mathbf{x} = \{x_{11}, x_{12}, x_{21}, x_{22}\}$: any possible table

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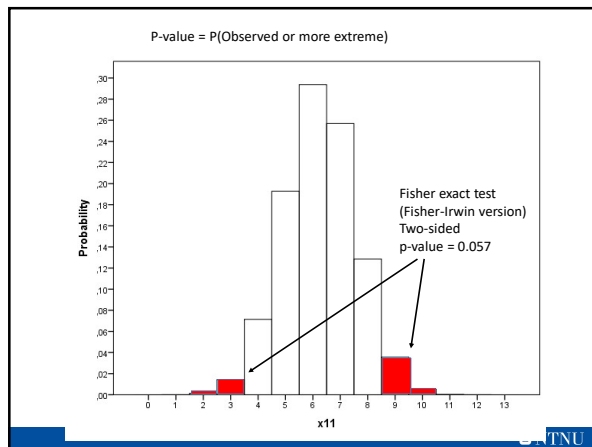
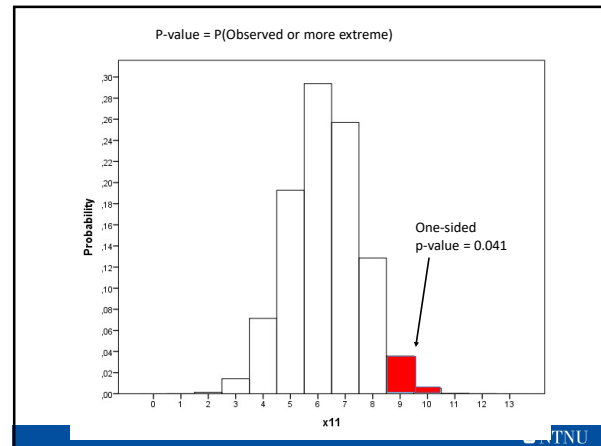
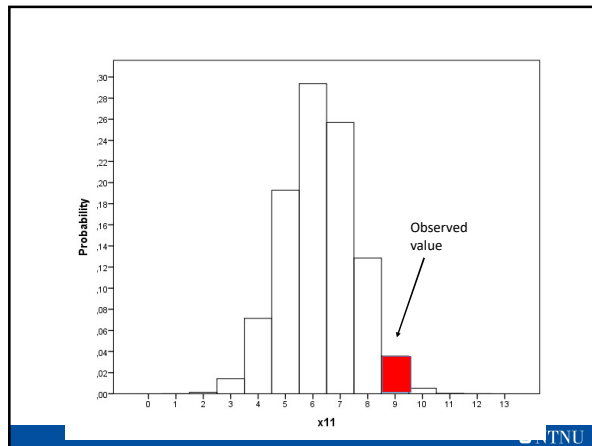
Fisher's exact test uses the following fact:

Conditional on the marginals, x_{11} is hypergeometrically distributed under H_0

$$f(x_{11} | n_{1+}, n_{2+}, n_{+1}, n_{+2}) = \frac{\binom{n_{1+}}{x_{11}} \binom{n_{2+}}{n_{+1} - x_{11}}}{\binom{N}{n_{+1}}}. \quad (4.2)$$

This distribution does not depend on the unknown common success probability (nuisance parameter) under H_0 !

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What is more extreme (than the observed value)? Different test statistics (Norsk: testobservator) may be used to measure «distance» from the null hypothesis:

- Use the conditional point probability as in the Fisher-Irwin version of the Fisher exact test
- The Pearson chi squared statistic
- Etc
- In a one-sided test in a 2x2 table, the choice of (monotonely increasing) statistics does not matter.
- Also the case for twice the smallest tail two sided tests
- The choice of statistic matters for other two sided tests in 2x2 tables, and for rxc tables with $r > 2$ or $c > 2$.
- In 2x2 tables, the conditional probability (Fisher) and the Pearson statistics generally perform well.

Example output from SPSS

Operation type * Severe nausea Crosstabulation

		Severe nausea		Total	
		No	Yes		
Operation type	CABG	Count 39	12	51	
	% within Operation type	76.5%	23.5%	100.0%	
Other	Count 3	5	8		
	% within Operation type	37.5%	62.5%	100.0%	
Total		Count 42	17	59	
		% within Operation type	71.2%	28.8%	100.0%

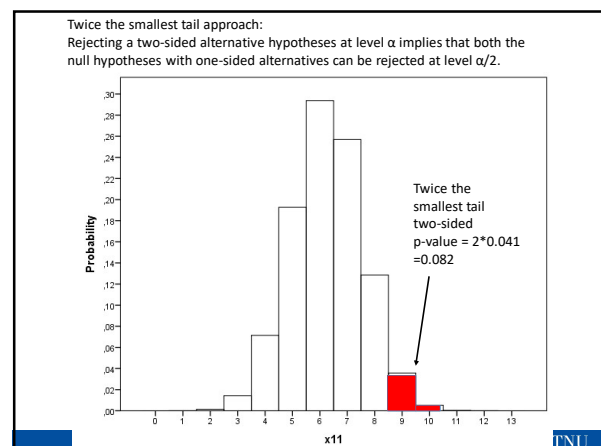
Chi-Square Tests

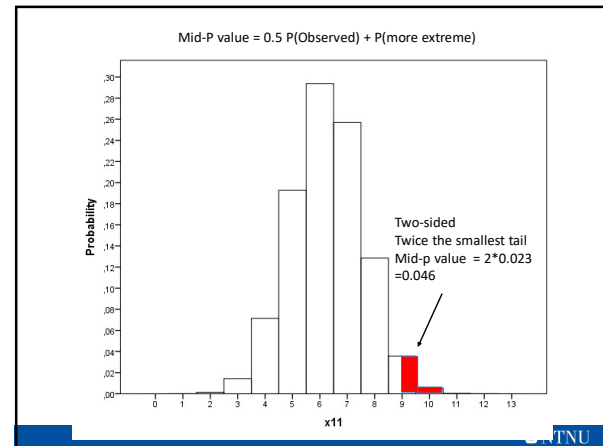
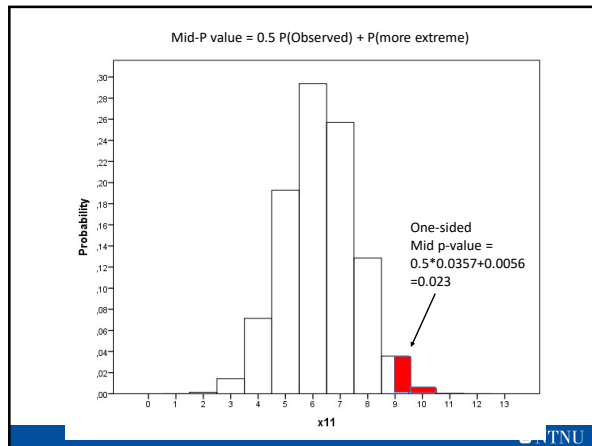
	Value	df	Asymp. Sig. (2-sided)	Exact Sig. (2-sided)	Exact Sig. (1-sided)
Pearson Chi-Square	5,120 ^a	1	,024	,037	,037
Continuity Correction ^b	3,397	1	,065		
Likelihood Ratio	4,620	1	,032	,090	,037
Fisher's Exact Test				,037	,037
N of Valid Cases	59				

a. Computed only for a 2x2 table

b. 1 cells (25.0%) have expected count less than 5. The minimum expected count is 2.31.

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Mid-p tests and Mid p confidence intervals:

- Solid theoretical justifications (Fagerland, Lydersen, Laake, 2017, page 28-29, and references therein)
- Reduces the conservatism of exact methods
- Do not preserve nominal significance level (tests) and nominal coverage (confidence intervals), but the violations are usually not serious
- An ideal p-value is $U(0,1)$ under H_0 . Exact p-values for categorical data are right skewed. Mid-p values have expectation 0.5 and is approximately $U(0,1)$
- In most cases the mid-p approach gives methods with better properties than those based on asymptotic normal theory. A notable exception is testing for equality of paired binomial distributions, where the McNemar asymptotic test is better than the McNemar mid-P test (Fagerland, Lydersen, Laake, 2017, page 29)

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4.4.7 Exact Unconditional Tests

The probability of observing an arbitrary table \mathbf{x} under H_0 is then given by Equation 4.3 with $\pi = \pi_1 = \pi_2$, which we rewrite slightly to

$$f(\mathbf{x} | \pi, \mathbf{n}_+) = \binom{n_{1+}}{x_{11}} \binom{n_{2+}}{x_{21}} \pi^{x_{11}+x_{21}} (1-\pi)^{N-x_{11}-x_{21}},$$

where $\mathbf{n}_+ = \{n_{1+}, n_{2+}\}$ denotes the fixed row sums. An explicit expression for the exact unconditional P -value is

$$P\text{-value} = \max_{0 \leq \pi \leq 1} \left\{ \sum_{\Omega(\mathbf{x} | \mathbf{n}_+)} I[T(\mathbf{x}) \geq T(\mathbf{n})] \cdot f(\mathbf{x} | \pi, \mathbf{n}_+) \right\}, \quad (4.14)$$

where $\Omega(\mathbf{x} | \mathbf{n}_+)$ denotes the set of all tables with row sums equal to \mathbf{n}_+ .

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TABLE 4.5

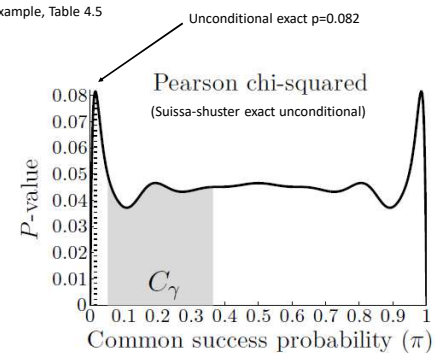
Genotype and presence of XFS in the eyes (Ritland et al., 2007)

Genotype	XFS		Total
	Yes	No	
CHRNA4-CC	0	16 (18%)	16 (18%)
CHRNA4-TC/TT	15 (17%)	57 (65%)	72 (82%)
Total	15 (17%)	73 (83%)	88* (100%)

*Fixed by design

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Example, Table 4.5



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The Berger and Boos Procedure

It may be argued that maximizing π over the entire nuisance parameter space, $0 \leq \pi \leq 1$, is unreasonable because the interval contains values that are highly unlikely in light of the observed data. This argument was the crux of Fisher's criticism (Fisher, 1945) of Barnard's proposition of the exact unconditional test. The Berger and Boos procedure is a remedy (Berger and Boos, 1994). It restricts the nuisance parameter space to C_γ : a $100(1-\gamma)\%$ exact confidence interval for π , where γ is taken to be very small. To make sure that the actual significance level is bounded by the nominal level, the value of γ is added to the P -value:

$$P\text{-value} = \max_{\pi \in C_\gamma} \left\{ \sum_{\Omega(\mathbf{x}|\mathbf{n}_+)} I[T(\mathbf{x}) \geq T(\mathbf{n})] \cdot f(\mathbf{x}|\pi, \mathbf{n}_+) \right\} + \gamma.$$

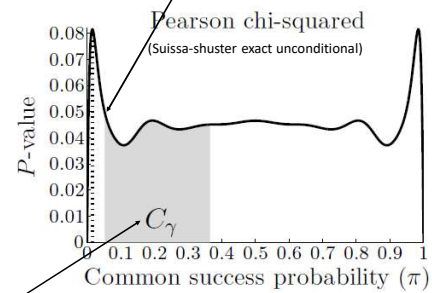
For the Suissa-Shuster exact unconditional test, Lydersen et al. (2012b) found $\gamma = 0.0001$ to be approximately optimal under rather general conditions.

In addition to avoiding computation over unrealistic values of the nuisance parameter, Berger (1996) states two other advantages of using the Berger and Boos procedure: (i) maximization over C_γ is computationally easier than over $0 \leq \pi \leq 1$; and (ii) the resulting test can have higher power than the ordinary exact unconditional test. The user manual of the software package StatXact also notes that using the Berger and Boos procedure provides greater computational stability (StatXact 11, 2015, p. 528). A common method to form the confidence interval C_γ is to use the Clopper-Pearson exact interval (see Section 2.4.7).

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Example, Table 4.5

Unconditional exact with BB ($\gamma=0.0001$) $p=0.0499$



99.99% CI for the common success probability

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Evaluation criteria for hypothesis tests:

1. The actual significance level (ASL) should ideally equal the nominal significance level (usually 5%). If the ASL level is lower, say 2% or 3%, the test is conservative. If the ASL is higher, the test is liberal.
2. Among tests with acceptable ASL, we prefer the one with highest power.

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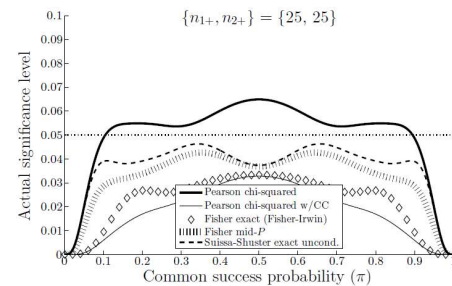


FIGURE 4.4

Actual significance levels of five commonly used tests

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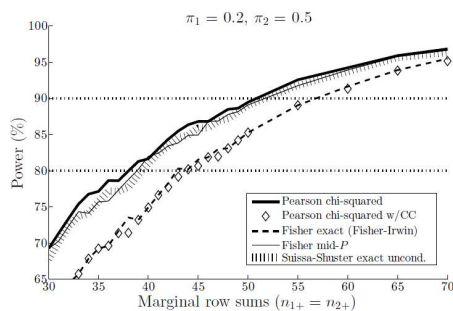


FIGURE 4.10

Power of five commonly used tests

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Fisher's exact test:

Advantage: Always preserves level

Disadvantage: Conservative (lower power than necessary)

Pearson's asymptotic test:

Advantage: Not conservative

Disadvantage: Does not always preserve level.

Pearson's asymptotic test with Yates' continuity correction:

Disadvantage: Does not always preserve level

Disadvantage: Conservative

Fisher (and other) Mid-p:

Advantage: Not conservative

Disadvantage: Does not always preserve level (but seldom large violations)

Exact unconditional tests:

Advantage: Not conservative

Advantage: Always preserves level

Disadvantage: Computer-intensive in moderate and large sample sizes

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Exact unconditional tests are available in

- StatXact (Cytel software)
- R package «Exact»
- <http://www4.stat.ncsu.edu/~boos/exact/>
- ...

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Exact Unconditional Homogeneity/Independence Tests for 2x2 Tables

This performs exact, unconditional tests of homogeneity (binomial model) or independence (multinomial model) for 2x2 tables. These tests are usually **uniformly more powerful than Fisher's exact test**. See references below.

Cell Counts:

1	33
7	27

Model: For binomial model, the row totals are the fixed binomial sample sizes.
☒ binomial ☐ multinomial

Hypothesis: The one-sided hypotheses are
 Ho: odds ratio ≥ 1 vs H_a: odds ratio < 1 [More explanation](#)
☐ one-sided ☒ two-sided

Test Statistic:
☐ Fisher's Exact-Boschloo ☒ z-pooled ☐ z-unpooled

Confidence Interval Method:
☒ yes ☐ no

Confidence Coefficient:
 0.9999

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Exact Unconditional Test Output

Model: BINOMIAL
 Hypothesis: TWO-SIDED
 Test statistic: Z POOL - SIGNED CHI SQUARE - SCORE TEST

The Input Table

1	33	34
7	27	34
8	60	68

99.9990% confidence interval used.
 confidence interval for common p: (0.0193, 0.3303)

Fisher's exact conditional p-value = 0.0544
 test statistic = -2.2583
 Unconditional p-value = 0.0262
 Maximum p-value occurred at common p = 0.1324

[Go Back to the input form](#)

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TABLE 4.24

Recommended tests and confidence intervals (CIs) for 2 × 2 tables

Analysis	Recommended methods	Sample sizes
Tests for association	Fisher mid-P*	all
	Suissa-Shuster exact unconditional†	small/medium
	Fisher-Boschloo exact uncond.†	small/medium
	Pearson chi-squared*	large

*These methods have closed-form expression

†Preferably with the Berger and Boos procedure ($\gamma = 0.0001$)

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Confidence intervals in 2x2 tables

Effect measures in 2x2 tables,
 comparing two binomial probabilities π_1 and π_2 :

Probability difference
 (risk difference, absolute risk reduction, attributable risk)

$$\Delta = \pi_1 - \pi_2$$

Ratio of probabilities (risk ratio, relative risk)

$$\phi = \pi_1 / \pi_2$$

Odds ratio

$$\theta = \frac{\pi_1 / (1 - \pi_1)}{\pi_2 / (1 - \pi_2)}$$

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Difference between probabilities:
Maximum likelihood estimate (difference between sample proportions):

$$\hat{\Delta} = \hat{\pi}_1 - \hat{\pi}_2 = \frac{n_{11}}{n_{1+}} - \frac{n_{21}}{n_{2+}}.$$

The traditional Wald confidence interval for Δ is based on the asymptotic normal distribution of $\hat{\Delta}$:

$$\hat{\Delta} \pm z_{\alpha/2} \sqrt{\frac{\hat{\pi}_1(1-\hat{\pi}_1)}{n_{1+}} + \frac{\hat{\pi}_2(1-\hat{\pi}_2)}{n_{2+}}}, \quad (4.19)$$

where $z_{\alpha/2}$ is the upper $\alpha/2$ percentile of the standard normal distribution.

A continuity corrected version of the Wald interval, due to Yates (1934), can be expressed as shown in Fleiss et al. (2003, p. 60):

$$\hat{\Delta} \pm \left[z_{\alpha/2} \sqrt{\frac{\hat{\pi}_1(1-\hat{\pi}_1)}{n_{1+}} + \frac{\hat{\pi}_2(1-\hat{\pi}_2)}{n_{2+}}} + \frac{1}{2} \left(\frac{1}{n_{1+}} + \frac{1}{n_{2+}} \right) \right]. \quad (4.20)$$

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Agresti-Caffo (2000) CI for the original table is simply the Wald CI for the table with one failure and one success added to each group:

TABLE 4.1

The observed counts of a 2×2 table

	Success	Failure	Total
Group 1	n_{11} +1	n_{12} +1	n_{1+} +2
Group 2	n_{21} +1	n_{22} +1	n_{2+} +2
Total	n_{+1} +2	n_{+2} +2	N +4

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Agresti and Caffo (2000) proposed a simple, yet effective procedure for computing a confidence interval: add one success and one failure (pseudo-frequencies, see Section 2.4.4) in each sample and calculate the Wald confidence interval on the resulting data:

$$\tilde{\pi}_1 - \tilde{\pi}_2 \pm z_{\alpha/2} \sqrt{\frac{\tilde{\pi}_1(1-\tilde{\pi}_1)}{\tilde{n}_{1+}} + \frac{\tilde{\pi}_2(1-\tilde{\pi}_2)}{\tilde{n}_{2+}}}, \quad (4.21)$$

where

$$\tilde{n}_{1+} = n_{1+} + 2, \quad \tilde{n}_{2+} = n_{2+} + 2, \quad \tilde{\pi}_1 = (n_{11} + 1)/\tilde{n}_{1+}, \quad \tilde{\pi}_2 = (n_{21} + 1)/\tilde{n}_{2+}.$$

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Newcombe (1998b) proposed a confidence interval for Δ formed as a combination of the Wilson score (Section 2.4.3) confidence limits for π_1 and the Wilson score confidence limits for π_2 . Denote the interval for π_1 by (l_1, u_1) and the interval for π_2 by (l_2, u_2) . The Newcombe hybrid score confidence interval (L, U) for Δ is given by

$$L = \hat{\Delta} - \sqrt{(\hat{\pi}_1 - l_1)^2 + (u_2 - \hat{\pi}_2)^2} \quad (4.22)$$

and

$$U = \hat{\Delta} + \sqrt{(\hat{\pi}_2 - l_2)^2 + (u_1 - \hat{\pi}_1)^2}. \quad (4.23)$$

Wilson score CI:

$$\frac{2n\hat{\pi} + z_{\alpha/2}^2 \pm z_{\alpha/2} \sqrt{z_{\alpha/2}^2 + 4n\hat{\pi}(1-\hat{\pi})}}{2(n + z_{\alpha/2}^2)},$$

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Example, two binomials:
Treatment of children with cardiac arrest.
(Perondi et al, NEJM, 2004)

Epinephrine treatment	survival at 24 hours		Total
	yes	no	
High dose	1	33	34
Standard dose	7	27	34
total	8	60	68

Fisher's exact test, two-sided $p=0.054$

Exact z-pooled (Suissa & Shuster) unconditional test: $p=0.028$

Estimated probability difference: $1/34 - 7/34 = -0.176$

95% CI: Wald: -0.324 to -0.029

Agresti-Caffo: -0.322 to -0.012

Newcombe hybrid score: -0.340 to -0.019

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Evaluation criteria for confidence intervals:

1. Coverage probability: Ought to not dip (much) below the nominal coverage (usually 95%). This is the primary criterion.
2. Interval width: Among intervals with similar coverage, we prefer the narrower.
3. Interval location: We say that an interval is located too distally if it is located too far out from the centre of symmetry for the effect measure (the midpoint). If the interval is located too close to the midpoint, we say that the interval is too mesially located. A $1-\alpha$ confidence interval has ideal location if both the left and right non-coverage are equal to $\alpha/2$.

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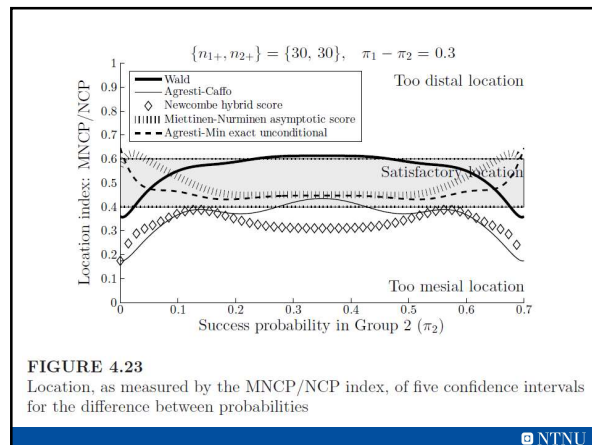
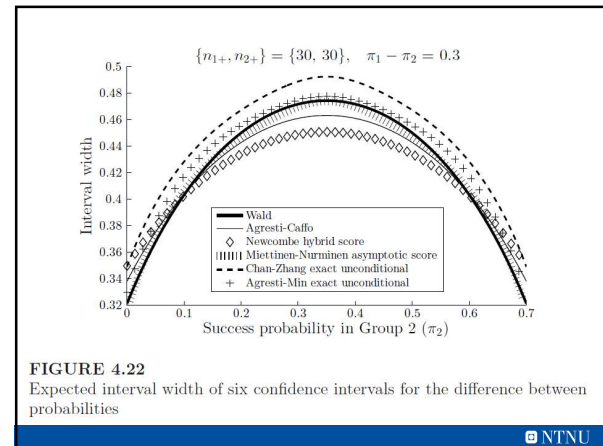
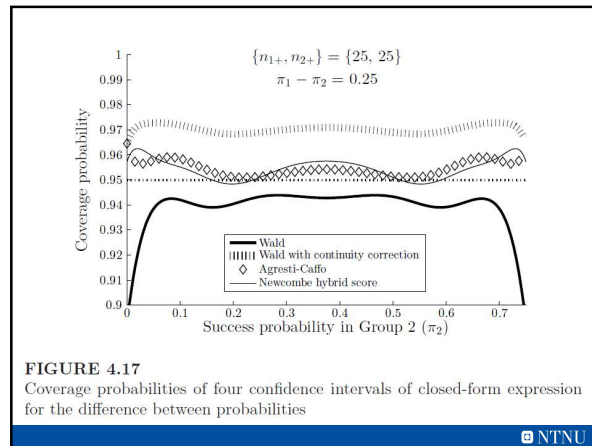
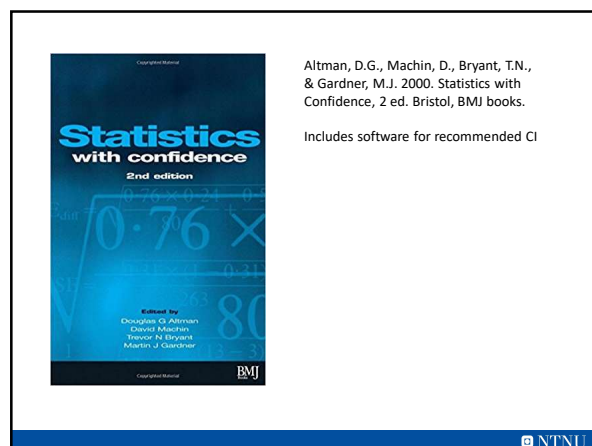


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	Fisher-Boschloo exact uncond. [†]	small/medium
	Pearson chi-squared [†]	large
CIs for difference between probabilities	Agresti-Min exact unconditional [†]	small/medium
	Agresti-Caffo [*]	medium/large
	Newcombe hybrid score [*]	medium/large
	Miettinen-Nurminen asympt. score	medium/large
	Wald [*]	large
CIs for number needed to treat	The reciprocals of the limits of the recommended intervals for the difference between probabilities	
CIs for ratio of probabilities	Adjusted inverse sinh [*]	all
	MOVER-R Wilson [*]	all
	Koopman asymptotic score	all
	Agresti-Min exact unconditional [†]	small/medium
	Katz log [*]	large
CIs for odds ratio	Adjusted inverse sinh [*]	all
	MOVER-R Wilson [*]	all
	Baptista-Pike mid- P	all
	Agresti-Min exact unconditional [†]	small/medium
	Woolf logit [*]	large

^{*}These methods have closed-form expression
[†]Preferably with the Berger and Boos procedure ($\gamma = 0.0001$)

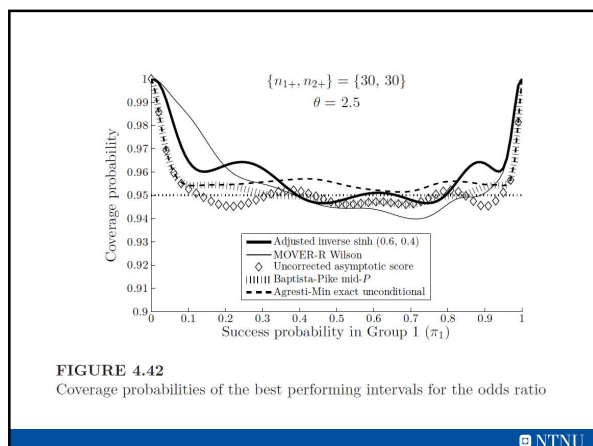


STATISTICAL ANALYSIS OF CONTINGENCY TABLES

MORTEN W. FAGERLAND
STIAN LYDERSEN
PETTER LAAKE

Tables with r rows and c columns:

- 1x2
- 1xc
- 2x2
- Ordered rx2 and ordered 2xc
- Unordered, singly ordered and doubly ordered rxc
- Paired 2x2 and paired cxc
- 2x2k and other stratified tables



Fagerland, M. W., Lydersen, S., & Laake, P.: "Statistical Analysis of Contingency Tables." Chapman and Hall/CRC, 2017.

- "This book should be a very useful reference for anyone who wants an overview of the relevant literature (much of it quite recent) or who routinely needs to analyze contingency tables." Alan Agresti.
- "I highly recommended it for masters and doctoral students in statistics ... and other fields requiring the analysis of discrete data." Karim F. Hirji
- "I strongly recommend the book both to statisticians and to researchers in health and social disciplines." Robert G. Newcombe
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