Another View of the Classical Problem of Comparing Two Probabilities

Herman Chernoff

Department of Statistics, Harvard University, Cambridge, MA 02138, USA. (chernoff@hstat.harvard.edu)

Abstract. The usual calculation of the P-value for the classical problem of comparing probabilities is not always accurate. This issue arose in the context of a legal dispute which depended on when some written material was written in a diary. The problem raises some issues on the foundations of statistical inference.

1 Background

A legal case concerned the question of when a certain handwritten entry was made in a diary. One party contended that two paragraphs on a page were written before the entries on the reverse side of that page. It was admitted that the writing was all made under similar conditions by one writer. The other party claimed that the short first paragraph was written after the writing on the reverse side of that paragraph.

The writing was subjected to forensic examination by Erich Speckin of Speckin Forensic Laboratories. The underlying theory behind his

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examination is that the earlier writing leaves a convex impression on the page where the pen moves. The later writing on the reverse side introduces a number of intersections where the ink trails from the two sides cross. The convexity of the first impression often causes the later impression on the reverse side to leave a short skipping, a gap in the ink trail, which is frequently detectable by examination of enlarged photographs, when the intersection is at an appropriate angle.

The examination is not error free. Thus an intersection where a skipping should be detected does not always yield an observable skipping, whereas an intersection where a skipping should not be observed might lead to what the forensic examiner regards as an observable skipping. If the two error probabilities are \( \epsilon_f \) and \( q \) for the first and later sections written, then the probabilities of observing a skipping at an intersection are \( q_f = \epsilon_f \) and \( q_l = 1 - q \) respectively. If the two error probabilities are small enough, \( q_l \) is substantially greater than \( q_f \). Then the observed skippings at a reasonable number of intersections should provide evidence to determine whether a paragraph was written earlier or later than the writing on the reverse side.

As we shall see in the following section, the issue involves a one-sided test of the hypothesis that two unknown probabilities are equal. This, in turn, raises the question of how to calculate a conservative version of the P-value for this test and the inferential meaning of such a calculation.

2 Several relevant hypotheses

Both parties are willing to grant that the writings on both sides of the page were made by one person under similar conditions. One party claims that the two paragraphs on the first side were written before the material on the reverse side. The contending party claims that the first short paragraph on the first side was written later than the material on the reverse side. It accepts the claim that the second paragraph on the first side was written earlier than the material with which it intersects on the reverse side.

To test the reliability of the claim of the first party we should test the hypothesis \( H_1 : p_1 = p_2 = q_f \) vs. \( H_2 : p_1 = q > p_2 = q_f \), where \( p_1 \) and \( p_2 \) are the probabilities of observing a skipping for the first and second paragraphs. Mr. Speckin had observed \( X_1 = 7 \) skippings
out of $n_1 = 16$ intersections for the first paragraph, and $X_2 = 24$
skippings out of $n_2 = 261$ intersections for the second paragraph.
These data lead to the estimates $\hat{p}_1 = 7/16 = 0.4375$ and $\hat{p}_2 = 24/261 = 0.092$ for $p_1$ and $p_2$. From an informal point of view it
appears unlikely that the probability of observing a skipping at an
intersection is the same for both paragraphs.

The skippings due to the writing on the reverse side provide addi-
tional relevant data. It seemed to me that these data should not
be ignored. Indeed, the legal case for claiming a difference between
$p_1$ and $p_2$ might be weakened if the judge were suspicious about why
these data were not presented. At my request, Speckin examined
skippings due to writing on the reverse side. He had previously found
$X'_1 = 0$ opposite the first paragraph, and then he obtained $X'_2 = 126$
opposite the second paragraph.

Considering the new data, we are led to introduce additional hypo-
theses. These are $H_2 : p'_1 = p'_2$, $H_3 : p_1 = p_2 < p'_1 = p'_2$,
$K_2 : p'_2 > p'_1$, and $K_3 : p_1 = p'_2 > p_2 = p'_1$ where $p'_1$ and $p'_2$ are
the probabilities of an observed skipping for the reverse side of para-
graphs 1 and 2 respectively.

According to the claim that the material on the first side was
written before that of the reverse side, we should expect to find that
the data are consistent with $H_1$, $H_2$, and $H_3$ and we should test these
against the corresponding alternatives $K_1$, $K_2$, and $K_3$.

Since $q_1$ and $q_2$ are unknown characteristics of the writer, the
parameters $p_1, p_2, p'_1$ and $p'_2$ are unknown.

## 3 Three standard one-sided tests for the equal-
ity of two probabilities

The problem of testing $H_1$ vs. $K_1$ is one of the most studied problems
in the statistical literature. When I declared to two colleagues that
I had a new insight on the problem, each of them referred me to
relatively recent papers that they had published on this subject Little
(1989) and Rubin and Stern (1994).

For most purposes, the standard approaches are pretty reason-
able. In my case, a superficial view of the data is compelling. It was
my desire to be more conservative than necessary that led me to look
further. I shall list three of the standard approaches first.

One of the standard approaches, which we will call the Pearson
method, is to use as the test statistic

\[
Z_P = \frac{X_1/n_1 - X_2/n_2}{\sqrt{\hat{p}(1 - \hat{p})(n_1^{-1} + n_2^{-1})}}
\]

(1)

where \( n = n_1 + n_2 \) and \( \hat{p} = (X_1 + X_2)/n \) is the maximum likelihood estimate of the common value of \( p_1 \) and \( p_2 \) under the hypothesis \( H_1 \). If \( p_1 > p_2 \), \( Z_P \) will tend to be positive and the one-tail test of \( H_1 \) consists of rejecting \( H_1 \) if \( Z_P \) is sufficiently large. Under \( H_1 \) the asymptotic distribution of \( Z_P \) is \( N(0, 1) \), the normal distribution with mean 0 and variance 1. The P-value for this test is taken to be \( P_P = p_P(Z_P) \) where

\[
p_P(z) = P(Z_P \geq z|H_1) = 1 - \Phi(z)
\]

(2)

and \( \Phi \) is the cumulative distribution function for the standard normal distribution \( N(0, 1) \).

In the early history, lack of sufficient computer power made it important to develop an asymptotic theory, as a result of which some of the claims are approximations which become poor when the sample sizes are not large. Some of the recent publications, Little (1989), seek to determine how conservative some of these approximations are. In our application, \( n_2 \) is reasonably large, but \( n_1 \) is quite moderate in size.

A second approach, which we shall label the Yates approach, applies a continuity correction which improves the asymptotic approximation. This method can be explained by constructing a \( 2 \times 2 \) table with entries \( a = X_1 \), and \( b = X_2 \) in the first row, and \( c = n_1 - X_1 \) and \( d = n_2 - X_2 \) in the second row. The table has margins \( n_1 \) and \( n_2 \) at the bottom and \( m_1 = X_1 + X_2 \) and \( m_2 = n - m_1 \) on the side. See Table 1.

<table>
<thead>
<tr>
<th></th>
<th>( a = X_1 )</th>
<th>( b = X_2 )</th>
<th>( m_1 = X_1 + X_2 )</th>
</tr>
</thead>
<tbody>
<tr>
<td>( c = n_1 - X_1 )</td>
<td>( d = n_2 - X_2 )</td>
<td>( m_2 = n - m_1 )</td>
<td></td>
</tr>
<tr>
<td>( n_1 )</td>
<td>( n_2 )</td>
<td>( n )</td>
<td></td>
</tr>
</tbody>
</table>

The Yates test statistic is

\[
Z_Y = \sqrt{\frac{n}{n_1 n_2 m_1 m_2}}(ad - bc - n/2)
\]

(3)
which corresponds to the chi-square statistic and $Z_F$, except that $a$ and $d$ are decreased by 1/2 and $b$ and $c$ are increased by 1/2. Here, the P-value is given by $P_f = 1 - \Phi(Z_f)$.

A third approach is that of the Fisher Exact Test. This test capitalizes on the fact that the conditional probability of observing the $2 \times 2$ table, given the margins $m_1, m_2, n_1$ and $n_2$, does not depend on the unknown common value of $p_1$ and $p_2$ under the hypothesis. This conditional probability is:

$$z_F(a) = \frac{n_1!n_2!m_1!m_2!}{n!}$$

Note that for specified values of the margins the value of $a$ determines all the other entries. Thus $z_F$ may be regarded as a function of $a$, and we may treat $Z_F = z_F(X_1)$ as the test statistic (conditional on $X_1 + X_2$), with conditional P-value

$$P_F = \sum_{a=X_1}^{a^*} z_F(a)$$

where $a^* = \min(n_1, m_1)$ and $m_1 = X_1 + X_2$.

For the data given, the P-values for the three approaches on the test of $H_1$ vs. $K_1$ are $P_{f} = 1.0(-5), P_{\gamma} = 6.0(-5)$, and $P_{F} = 6.3(-4)$. In these representations the integer in parentheses represents the exponent of 10. These results bear out the general claim that both $P_{f}$ and $P_{\gamma}$ tend to be more conservative than $P_{F}$ in the sense that they do not argue as strongly as $P_{F}$ against the hypothesis being tested. The fact that $P_{\gamma}$ is so much greater than $P_{F}$ suggests that the asymptotic approximation needs the continuity correction in this case where the data strongly suggest that the hypothesis is false and at least one of the sample sizes is not large.

With the use of modern computers it is no longer necessary to rely on asymptotic approximations, and while $P_{f}$ is exact, both of the other P-values can be also computed exactly, assuming that the common value of $p$ under $H_1$ is $\tilde{p}$, to yield $P_{P1} = 2.9(-4)$ and $P_{P2} = 2.9(-4)$. The relatively large discrepancy between the exact and asymptotic results is apparently due, in part, to the large deviation effect in the tails of the distribution of the test statistic, as well as to the modest sample size $n_1$.

However, we have a more serious problem due to the fact that the hypothesis $H_1$ is a composite hypothesis, and the distribution of the data depend on the unknown common value of $p_1$ and $p_2$ under the hypothesis. That problem will be attacked in the next section.
4 The effect of composite hypotheses

Technically speaking, the P-value for a composite hypothesis is defined differently than what we have apparently done. If we have a composite hypothesis $H : \theta \in \Theta$ and a test statistic $T$, then the P-value is $P = \sup_{\theta \in \Theta} P(T; \theta)$ where $p(t; \theta) = P(T \geq t|H, \theta)$. In using $P_F$ and $P_Y$, we implicitly assumed that the asymptotic normal distribution $N(0,1)$, which is independent of the nuisance parameter, the common value of $p_1$ and $p_2$ assumed under $H_1$, represented the true distribution of the test statistic. This asymptotic distribution is not a very good fit in the tails; the true distribution of $Z_F$ and $Z_Y$ depends on $p$, and a true P-value should take into account $P(p) = p(T; p)$ for all values of $p \in (0,1)$.

Although the Fisher approach provides an exact test, that test is exact only if we condition on the margins. That procedure is appropriate in some tests of independence for $2 \times 2$ tables, but it is not exactly so here. In the case of testing for independence with specified margins, $X_1 + X_2$ would be an ancillary statistic, but it is only approximately ancillary in our problem Little (1989). Thus even the Fisher test yields a range of P-values depending on the nuisance parameter, and we should consider the maximum of these P-values. Note that to define a P-value, one must have a test statistic. For the case where the marginals are given, we could use $X_1$ as the test statistic. But if the marginals are not specified, a more natural test statistic would be the P-value using the Fisher exact test, assuming, without proper justification, that the marginals are specified.

Unfortunately, the range of P-values corresponding to various values of $p$ is uncomfortably large for $P_F$ and $P_Y$. Our calculations indicate that $P_F(p), P_Y(p)$ and $P_F(p)$ attain their maximum values of 3.29(-3), 1.03(-3), and 2.93(-4), at $p = 0.013, 0.0092$ and 0.15 respectively. On the other hand, if we do exact calculations, assuming that $\hat{p}$ is the only value of $p$ to consider, we would have values of 2.91(-4), 2.89(-4), and 2.86(-4) respectively. These compare with our original values of 1.0(-5), 6.0(-5) and 6.3(-4) respectively.

One of our difficulties comes from the poor quality of the asymptotic approximations in this example where one of the sizes is moderate, and large deviation effects are relevant. Another major problem comes from the fact that using the proper definition of the P-value provides another large effect. The range of values from 1.0(-5) to 3.3(-3) is enormous, although not enough to make one believe in $H_1$. Note that $P_F(p)$ is very stable. Somewhat surprisingly $P_F(p)$ tends
to be about $2.9(4)$ over a large range of values of $p$, but different from $P_F = 6.3(−4)$. Figure 1 shows how these $P$-values depend on $p$.

![Graph showing $p(p)$ for different $P$-values](image)

Figure 1: $P$-values for Pearson, Yates, Fisher, and LR

5 Modified $P$-values

While $P$-values have played a useful role in statistical inference, their use is subject to considerable criticism by Bayesians. One major limitation in the general case is that the $P$-value does not put enough emphasis on the role of the alternative hypotheses. What good is it to use the data to reject a hypothesis as unreasonable, if the same data
would show that the potential alternative is also unreasonable? This
latter criticism does not apply in the current problem. Granted that
P-values have limited inferential value, does it pay to use a precise
definition in this example? Here the maximum value of \( P_p(p) \) is
attained at \( p = 0.012 \) which is very inconsistent with the estimate
\( \hat{p} = 37/277 = 0.111 \) assuming \( H_1 \). A 95% confidence interval for \( p \)
would be \((0.077, 0.155)\). Over this range the maximum of the three
measures \( P_p(p), P_Y(p), \) and \( P_F(p) \) are 4.5(−4), 3.8(−4), and 2.9(−4)
respectively.

However, there is a mismatch between the confidence and P-
values. It seems strange to say that “I am 95% confident that the
P-value is 4.6(−4)”. With my remaining confidence, the P-value could
conceivably be be very large. It would seem more sensible to say
something like “I am 99.9% confident that the P-value is 0.001”! In-
creasing the confidence leads to increasing the range of potential val-
ues of \( p \), and consequently increasing the maximum P-value over that
range. With such matching we achieve approximately 99.95% confi-
dence for \( P_p(p) \leq 7.1(−4), P_Y(p) \leq 4.6(−4) \) and \( P_F(p) \leq 2.9(−4) \).
Thus a substantial increase in confidence leads to a relatively small
increase in P-values. Table 2 presents various level confidence in-
tervals of confidence \( \gamma \) for \( p_1 \) based on \( x \) successes out of \( n \) trials. Under
\( H_1 \) the intervals for \((x, n) = (31, 277)\) are the confidence intervals for
\( p \) while \((x, n) = (126, 277)\) is relevant for \( H_2 \).

Table 2. Intervals with confidence \( \gamma \) for a probability based on \( x \)
successes out of \( n \) trials.

\[
\begin{array}{cccc}
\text{x/n} & 24/277 & 31/277 & 126/277 & 133/277 \\
1-\gamma & \hline
5(-2) & 0.0630 & 0.12617 & 0.07732 & 0.15508 & 0.360519 & 0.51533 & 0.394999 & 0.54072 \\
1(-2) & 0.04872 & 0.13940 & 0.06836 & 0.16933 & 0.3773 & 0.53396 & 0.410193 & 0.55960 \\
5(-3) & 0.04606 & 0.14454 & 0.06518 & 0.17484 & 0.37071 & 0.54089 & 0.39519 & 0.56510 \\
1(-3) & 0.04092 & 0.1562 & 0.05884 & 0.18666 & 0.32695 & 0.55529 & 0.38120 & 0.58018 \\
5(-4) & 0.03888 & 0.16009 & 0.05664 & 0.19141 & 0.315158 & 0.56093 & 0.37574 & 0.58577 \\
1(-4) & 0.03490 & 0.16995 & 0.05116 & 0.20183 & 0.3101 & 0.57305 & 0.36405 & 0.59774 \\
5(-5) & 0.03330 & 0.17399 & 0.04967 & 0.20630 & 0.3155 & 0.57797 & 0.35937 & 0.60254 \\
1(-5) & 0.03023 & 0.18302 & 0.04570 & 0.21560 & 0.32556 & 0.58852 & 0.34920 & 0.61390 \\
5(-6) & 0.02900 & 0.18677 & 0.04414 & 0.21953 & 0.32152 & 0.59283 & 0.34507 & 0.61725 \\
1(-6) & 0.02641 & 0.19162 & 0.04083 & 0.22834 & 0.31266 & 0.60234 & 0.33600 & 0.62660 \\
5(-7) & 0.02530 & 0.19872 & 0.03951 & 0.23202 & 0.30904 & 0.60624 & 0.33228 & 0.63044 \\
1(-7) & 0.02422 & 0.20168 & 0.03669 & 0.23409 & 0.30102 & 0.61490 & 0.32406 & 0.63894 \\
5(-8) & 0.02326 & 0.20802 & 0.03576 & 0.23477 & 0.29772 & 0.61848 & 0.32067 & 0.64245 \\
1(-8) & 0.02502 & 0.21759 & 0.03312 & 0.25162 & 0.29037 & 0.62646 & 0.33313 & 0.65028 \\
5(-9) & 0.01979 & 0.22078 & 0.03214 & 0.25493 & 0.28734 & 0.62977 & 0.33000 & 0.65352 \\
1(-9) & 0.01821 & 0.22803 & 0.03000 & 0.26242 & 0.28055 & 0.63719 & 0.30302 & 0.66079 \\
\end{array}
\]
The comparison of these three approaches seems to favor the Fisher approach, where the maximum P-value of $P_F(p) = 2.93(-4)$ is relatively small, and takes place at $p = 0.145$ which is not far from $\hat{p}$. The large values of $P_T(p)$ and $P_Y(p)$ for small values of $p$ reflects the fact that the denominator of $Z_P$, which is a normalizing factor when $p$ is close to $\hat{p}$, no longer serves that purpose well when $p$ is very small.

6 Other approaches

Given the stability of the Fisher approach, it seems unlikely that there will be much room for improvement in this special problem of testing for the equality of two probabilities. For more complicated problems, e.g. testing $H_3$ vs. $K_3$, we may not have the benefit of the advice of Fisher to select such an effective test statistic. We propose to introduce two new methods, one using a Bayesian approach and both depending on the likelihood-ratio test statistic. We would hope that the use of the likelihood-ratio would free us of the need to depend on some clever choice of the test statistic.

We elaborate on the likelihood-ratio approach for which we use the abbreviation LR. The likelihood for this model is $L(p_1, p_2) = l(X_1, X_2; p_1, p_2)$ where

$$l(x_1, x_2; p_1, p_2) = p_1^{x_1} (1 - p_1)^{n_1 - x_1} p_2^{x_2} (1 - p_2)^{n_2 - x_2}.$$ 

Let $Z_L$ be the log of the likelihood-ratio for the two hypotheses $H_1$ and $K_1$ calculated at the maximum likelihood estimates under the two hypotheses.

$$Z_L = ((n_1 + n_2)H(\hat{p}) - n_1H(\hat{p}_1) - n_2H(\hat{p}_2))(\text{sgn}(\hat{p}_1 - \hat{p}_2))$$

where $H$ is the entropy given by

$$H(p) = -[p \log(p) + (1 - p) \log(1 - p)]$$

and $\hat{p}_1 = X_1/n_1$, $\hat{p}_2 = X_2/n_2$, and $\hat{p} = (X_1 + X_2)/(n_1 + n_2)$.

Then $p_L(z; p) = P(Z_L \geq z|H_1, p)$, $P_L(p) = p_L(Z_L; p)$, $P_{L1} = P_L(\hat{p})$, and

$$P_L(p) = \sum_{x_1 \in A} \left(\begin{array}{c} n_1 \\ x_1 \end{array}\right) \left(\begin{array}{c} n_2 \\ x_2 \end{array}\right) p^{x_1 + x_2} (1 - p)^{n_1 + n_2 - x_1 - x_2}$$

(8)
where \( x^* = (x_1^*, x_2^*) \) and \( A = \{ x^* : Z^*_L \geq Z_L \} \).

Finally we introduce \( P_B \) which is the posterior probability of the hypothesis \( H_1 \) based on a somewhat artificial and arbitrary prior distribution, for which the theoretical calculations are relatively simple. We assume that the prior distribution of \( p_1 \) and \( p_2 \) is a mixture of two distributions. One of these distributions is a beta distribution in \( p_0 \) on the line where \( p_1 = p_2 = p_0 \). The other distribution takes \( p_1 \) and \( p_2 \) to have independent beta distributions. With a slight abuse of notation we may write, for the distribution law \( L \) of \((p_1, p_2)\),

\[
L(p_1, p_2) = wB(a_0, b_0) + (1 - w)B(a_1, b_1) * B(a_2, b_2) \quad (9)
\]

to indicate that with probability \( w \), \( p_1 = p_2 = p_0 \) which has a beta distribution with parameters \((a_0, b_0)\) and with probability \( 1 - w \), \( p_1 \) has the beta distribution with parameters \((a_1, b_1)\) and \( p_2 \) has the beta distribution with parameters \((a_2, b_2)\) and \( p_1 \) and \( p_2 \) are independent.

Then the posterior distribution, given the data, \( X = (X_1, X_2) \), has the same form and can be written

\[
L(p_1, p_2 | X) = w^*B(e_0, b_0^*) + (1 - w^*)B(e_1^*, b_1^*) * B(e_2^*, b_2^*) \quad (10)
\]

where \( P_B = w^* \) is the posterior probability of the hypothesis \( H_1 \). The necessary calculations are described as follows. If the data yield \( X_1 \) successes out of \( n_1 \) trials with probability \( p_1 \) and \( X_2 \) successes out of \( n_2 \) trials with probability \( p_2 \), then

\[
a_1^* = a_1 + X_1, \quad b_1^* = b_1 + n_1 - X_1, \quad a_2^* = a_2 + X_2, \quad b_2^* = b_2 + n_2 - X_2, \quad a_0^* = a_0 + X_1 + X_2, \quad b_0^* = b_0 + n_1 + n_2 - X_1 - X_2,
\]

and

\[
\frac{w^*}{1 - w^*} = \frac{w}{1 - w} \frac{B_1 B_2 B_0^*}{B_1^* B_2^* B_0}
\]

(11)

where \( B_i = B(a_i, b_i) \), \( B_i^* = B(a_i^*, b_i^*) \), \( B \) is the beta function

\[
B(c_1, c_2) = \Gamma(c_1) \Gamma(c_2) / \Gamma(c_1 + c_2)
\]

and \( \Gamma \) is the gamma function.

Low values of \( a_i \) and \( b_i \) correspond to a prior beta distribution which, in some sense, assumes very little about the distribution of \( p_i \). For example \( a_1 = b_1 = 1 \) yields a uniform distribution on the interval from 0 to 1 for \( p_1 \). On the other hand \( a_1 = b_1 = 20 \) would correspond to a prior distribution which assumes that \( p_1 \) tends to be rather close to 0.5. There is a good deal of subjectivity in the process of applying
the Bayesian philosophy in our current problem, and I recommend it with reservations.

At this point let us summarize our results and notation on P-values. In general we have a test statistic \( T \) for testing a hypothesis \( H \), against an alternative \( K \), a conventional P-value, and a nuisance parameter \( \theta \in \Theta \). Let \( p(t; \theta) = P(T \geq t | H, \theta) \), \( P_1 = p(T; \theta) \), where \( \hat{\theta} \) is the maximum likelihood estimate of \( \theta \) under \( H \), and \( P_2 = \sup_{\theta \in \Theta} p(T; \theta) \). Given a 95\% confidence interval \( P_3 \) is the supremum of \( p(T; \theta) \) for \( \theta \) in that interval. Similarly, \( P_4 \) corresponds to the maximum P-value over an approximately matching confidence interval Table 3 summarizes the various P-values for testing \( H_1 \). (The last two digits of the tabled values of \( P_4 \) are not reliable).

| Table 3. P-values for testing \( H_1 \) vs. \( K_1 \). |
|------------------|------------------|------------------|------------------|
| Pearson | Yates | Fisher | LR |
| \( P \) | 1.04(-5) | 5.97(-5) | 6.28(-4) |
| \( P_1 \) | 2.91(-4) | 2.89(-4) | 2.86(-4) | 2.79(-4) |
| \( P_2 \) | 3.29(-3) | 1.03(-3) | 2.93(-4) | 1.04(-3) |
| \( P_3 \) | 4.62(-4) | 3.80(-4) | 2.93(-4) | 2.80(-4) |
| \( P_4 \) | 7.09(-4) | 4.56(-4) | 2.93(-4) | 2.80(-4) |

The posterior probability of \( H_1 \) for the parameters \( w = 0.5, a_0 = b_0 = a_1 = b_1 = a_2 = b_2 = 1 \) is \( w^t = 8.94(-3) \). With this prior distribution, for which \( P(p_1 < p_2) = 0.25 \), there is a small posterior probability, 4.59(-4), that \( p_1 < p_2 \). When using a Bayesian approach it often makes sense to use information based on previous experience. Suppose that we started with \( a_0 = b_0 = a_1 = b_1 = 5 \) and \( a_2 = 2, b_2 = 18 \). I chose these numbers without any real background experience, and without reference to Mr. Speckin who may have some relevant suggestions, merely to represent a possibility. The \( Be(5, 5) \) and \( Be(2, 18) \) distributions have means 0.5 and 0.1, and standard deviations 0.151 and 0.065. With these parameters to represent the prior distribution, we obtain the posterior probability of \( H_1 \) to be \( w^t = 5.76(-5) \).

7 The reverse side

Up to now, we have concentrated almost exclusively on the observed skippings on one side of the page. We have additional data from the reverse side where \( X'_1 = 0 \) and \( X'_2 = 126 \). These additional data could
be used to test $H_2$ vs. $K_2$ with the same methods used to test $H_1$ vs. $K_1$. Corresponding P-values are listed in Table 4.

<table>
<thead>
<tr>
<th></th>
<th>Pearson</th>
<th>Yates</th>
<th>Fisher</th>
<th>LR</th>
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</thead>
<tbody>
<tr>
<td>$P$</td>
<td>3.85(-5)</td>
<td>2.28(-4)</td>
<td>4.15(-5)</td>
<td></td>
</tr>
<tr>
<td>$P_1$</td>
<td>1.34(-5)</td>
<td>1.34(-5)</td>
<td>1.22(-5)</td>
<td>1.21(-5)</td>
</tr>
<tr>
<td>$P_2$</td>
<td>2.29(-2)</td>
<td>1.21(-3)</td>
<td>1.67(-5)</td>
<td>1.42(-5)</td>
</tr>
<tr>
<td>$P_3$</td>
<td>3.64(-5)</td>
<td>3.64(-5)</td>
<td>1.44(-5)</td>
<td>1.42(-5)</td>
</tr>
<tr>
<td>$P_4$</td>
<td>7.28(-5)</td>
<td>7.28(-5)</td>
<td>1.47(-5)</td>
<td>1.42(-5)</td>
</tr>
</tbody>
</table>

The posterior probability of $H_2$ for the parameters $w = 0.5, a_0 = b_0 = a_1 = b_1 = a_2 = b_2 = 1$ is $w^* = 6.65(-5)$. The posterior probability that $p_1 > p_2$ is 2.13(-5).

### 7.1 Combining P-values

One standard approach for combining the results of testing the two hypotheses $H_1$ and $H_2$ uses the fact that under the hypothesis to be tested the P-value has a uniform distribution and its negative logarithm has an exponential distribution, which may also be regarded as the gamma distribution with scale parameter 1 and shape parameter 1, i.e. $\Gamma(1,1)$. Then the sum of two such independent random variables has the $\Gamma(1,2)$ distribution, the tail probabilities of which can be used to provide a combined P-value. For example if the two P-values are 2.80(-4) and 1.42(-5) the sum of the two negative logarithms is 19.343 for which the tail probability is $P = 8.09(-8)$.

There are been three implicit assumptions in the analysis above. These are that the P-values are independent, that they have continuous distributions, and that the hypothesis being tested is *simple*, i.e. not composite. These assumptions are not satisfied here, but that probably has little influence here. The data are discrete and the P-values are not quite continuously distributed. The observed skippings, at an intersection on both sides of the page are probably not quite independent for all the candidate intersections. I believe that for a given value of $p, P(p)$ is reasonably well approximated by the combining analysis in spite of the failure of the assumptions. On the other hand this may not hold so well if we try to combine the maxima in this way.
7.2 Likelihood-ratio

An alternative to combining the P-values of the two tests is to test $H_3$ vs. $K_3$. Such a test introduces an interesting variation on our problem. It made little sense to get P-values for testing $K_1$ vs. $H_1$ or $K_2$ vs. $H_2$. As long as the observed proportions favored $K_1$ and $K_2$, such P-values could not be very small. One could not show that both $H_1$ and $K_1$ were inconsistent with the data. That is no longer the case for testing $H_3$ vs. $K_3$. Both of these alternative hypotheses are specific enough that each can be tested to see how consistent they are with the data. Is it the case that the skipping counts are not only inconsistent with $H_3$, but also inconsistent with the assumptions that (i) the first paragraph and the reverse of the second paragraph show similar probabilities of observed skippings, (ii) the same can be said for the second paragraph and the reverse of the first, and (iii) the probability for the first pair above exceeds that of the second pair?

The likelihood-ratio and Bayesian approaches are applicable for testing $H_3$ vs. $K_3$. These methods may be applied in a routine fashion and do not require the use of deep insight to develop a clever test statistic. On the other hand the resulting method does not directly confront assumptions (i) and (ii). To do so requires a modified test.

In this subsection we shall describe the likelihood-ratio method. We introduce the variables $R_{12} = (X_1 + X_2)/n$, $R_{34} = (X'_1 + X'_2)/n$, $R_{14} = (X_1 + X'_2)/n$, $R_{23} = (X_2 + X'_1)/n$, $R_0 = (X_1 + X_2 + X'_1 + X'_2)/2n$, and $X = (X_1, X_2, X'_1, X'_2)$. Under $H_3$, the maximum of the logarithm of the likelihood is given by

\[
l(X|H_3) = \begin{cases} 
-n[H(R_{12}) + H(R_{34})] & \text{if } R_{12} < R_{34} \\
-2nH(R_0) & \text{if } R_{12} \geq R_{34}
\end{cases}
\]

Similarly, the logarithm of the likelihood under $K_3$ attains its maximum of

\[
l(X|K_3) = \begin{cases} 
-n[H(R_{23}) + H(R_{14})] & \text{if } R_{23} < R_{14} \\
-2nH(R_0) & \text{if } R_{23} \geq R_{14}
\end{cases}
\]

and we may use

\[
Z_T = l(X|K_3) - l(X|H_3)
\]

as our test statistic.

The P-value for testing $H_3$ is $P_H = p_H(Z_T)$ where

\[
p_H(z) = \sup_{p_1 \leq p_1'} P(Z_T \geq z|H_3, p_1, p_1')
\]

(13)
Similarly, the P-value for testing $K_3$ is $P_K = p_K(Z_T)$ where

$$p_K(z) = \sup_{p_1 > p_1'} P(Z_T \leq z | K_3, p_1, p_1')$$

(14)

Table 5 presents the results for testing $H_3$ and $K_3$ in analogy with the notation of Table 3.

Table 5. P-values for testing $H_3$ vs. $K_3$ and $K_3$ vs. $H_3$

<table>
<thead>
<tr>
<th>P</th>
<th>$P_H$</th>
<th>$P_K$</th>
</tr>
</thead>
<tbody>
<tr>
<td>$P_1$</td>
<td>3.33(-8)</td>
<td>6.02(-1)</td>
</tr>
<tr>
<td>$P_2$</td>
<td>5.05(-8)</td>
<td>1.00(-0)</td>
</tr>
<tr>
<td>$P_3$</td>
<td>3.49(-8)</td>
<td>9.73(-1)</td>
</tr>
<tr>
<td>$P_4$</td>
<td>5.91(-5)</td>
<td>9.73(-1)</td>
</tr>
</tbody>
</table>

7.3 Bayesian approach

If we assign prior probabilities $w_3$ and $1 - w_3$ to $H_3$ and $K_3$, a natural version of the prior distribution of $(p_1, p_1')$ may be written as

$$L(p_1, p_1') = w_3 Be(a_1, b_1) * Be(a_2, b_2) + (1 - w_3) Be(a_2, b_2) * Be(a_1, b)$$

(15)

Letting $X$ represent the complete data, we have

$$L(X|H_3, p_1, p_1') = p_1^{X_1 + X_2} (1 - p_1)^{n - X_1 - X_2} (p_1')^{X_1'} + X_2' (1 - p_1')^{n - X_1' - X_2'}$$

(16)

and

$$L(X|K_3, p_1, p_1') = p_1^{X_1 + X_2} (1 - p_1)^{n - X_1 - X_2} (p_1')^{X_1' + X_2'} (1 - p_1')^{n - X_1' - X_2'}$$

(17)

It follows that the posterior distribution is given by

$$L(p_1, p_1'|X) = w_3 B_1 * B_2 + (1 - w_3) B_3 * B_4$$

(18)

where $B_i = Be(a_i, b_i^*)$ for $1 \leq i \leq 4$, $a_1^* = a_1 + X_1 + X_2$, $b_1^* = b_1 + n - X_1 - X_2$, $a_2^* = a_2 + X_1' + X_2'$, $b_2^* = b_2 + n - X_1' - X_2'$, $a_3^* = \ldots$
\[ a_2 + X_1 + X_2', \quad b_3' = b_2 + n - X_1 - X_2', \quad a_4' = a_1 + X_1' + X_2, \quad b_4' = b_1 + n - X_1' - X_2, \]
and
\[
\frac{w^*_i}{1 - w^*_i} = \frac{w_3}{1 - w_3} \frac{B_{31}B_{32}}{B_{33}B_{34}}
\]

where \( B_{3i} = B(a_i', b_i') \) for \( 1 \leq i \leq 4 \).

Applying the prior distribution with \( w = 0.5, a_1 = b_1 = a_2 = b_2 = 1 \), we obtain the posterior probability of \( H_3 \) to be \( w^* = 5.38(-7) \).

### 8 Historical Remarks

Much of this work was done under the pressure of responding to a request to provide a document for a legal case. It should not be surprising that a subsequent search of the literature reveals that for this topic which has been studied extensively, much of what has been written above has been presented elsewhere. In particular, papers by G. Barnard (1947) and E.S. Pearson (1947) have anticipated many of the ideas presented here. Barnard introduced the idea that we can select the test statistic so as to minimize the maximum of the P-values over the range \((0,1)\) of \( p \). This led to the CSM test, which he subsequently renounced in favor of the conditionality of the Fisher Exact test. The CSM test was developed in the context of relatively small sample sizes at a time when computers were rather limited. For small sample sizes, the possibility that the maximum is attained for unlikely values of \( p \) does not seem very pressing as it is in our case. Pearson was interested in good asymptotic approximations, and incidentally indicated that one should expect our method of using \( P_F \), the P-value of the Fisher Exact test, as the test statistic should yield \( P_F(p) \) to be almost independent of \( p \). It is slightly inappropriate to use an abbreviation of Pearson to describe one of the standard tests because of the contributions of Karl Pearson and an abbreviation of Fisher to describe the test statistic pioneered by Karl's son Egon Pearson. Tocher (1950) demonstrated that the test which uses \( P_F \) as the test statistic is optimal among similar tests of the hypothesis of equality. This implies that the Fisher Exact test is optimal, but does not endorse \( P_F \) as the proper P-value to use in interpreting the data.

As computing power grew, various investigators, including Berkson (1978a) criticized the use of the Fisher Exact Test and \( P_F \), suggesting that \( P_F \) was too conservative, that the test was not as powerful as it should be, and that a major justification for the use of
conditioning on the margins, the ancillarity of the margins, was not justified Berksen (1978b). Yates (1984) responded, defending the application of conditioning on the margins, which he regarded as implicit in the Yates continuity correction as well as the Fisher Exact Test, and claimed that it was obvious to both Fisher and him that there was very little, if any, relevant information in the margins. Yates claimed that the application of Neyman-Pearson theory and power considerations has served to confuse the real underlying issues.

The paper by Yates and the discussion indicates some lack of consensus about the inferential role and definition of the P-value. While the composite nature of our hypothesis $H_1$ complicates matters a bit and led me to introduce the four measures $P_1, P_2, P_3,$ and $P_4,$ the issue applies even in the case of a simple hypothesis. The formal definition of a P-value involves the concept of a test and the use of a test statistic $T$. Presumably, some decision, terminal or temporary, is to be made depending on whether or not $T$ exceeds some value and the P-value is less than a corresponding value $a$, in which case the data are regarded as significant. That corresponding value is so often taken to be 0.05 or 0.01, for the sake of discussion or to use known tables, that these numbers have assumed undue importance among users of statistical methods. In principle, the appropriate critical P-value should depend in part on the alternative hypotheses and the costs of wrong decisions, and possibly on some background information.

In those cases where the data are not supposed to lead to a terminal decision, the P-value is regarded as a measure of the inferential impact of the data in support of the hypothesis. As such a measure it is incomplete. It says little about the alternative and about the model. In fact the example Fisher presented Barnard, discussed by Barnard in Yates (1984), which led Barnard to accept the conditioning argument and reject CSM, demonstrated exactly this weakness of the inferential interpretation of the P-value. While the conditioning argument is relevant enough so that both S.S. Pearson and Tukey argue in favor of $P_T$ as the appropriate test statistic, one must be careful not to give $P_T$ undue influence as a measure of belief. The fact, that $P_T$ is conditional and technically not the P-value corresponding to a test statistic, is relevant for those who depend on the formal definition of a P-value, but is not ordinarily a major factor in deciding how to interpret it.
Rubin and Stern (1994) present a Bayesian approach alternative to our measures $P_3$ and $P_4$. Assuming $H_3$ and a prior distribution on the unknown common value of $p$, they calculate a posterior distribution using the data $(X_1, X_2)$. This yields a corresponding marginal predictive distribution of future values $(X'_1, X'_2)$ of the data, in terms of which they calculate $P^* = P(T(X'_1, X'_2) \geq T(X_1, X_2) | T(X_1, X_2))$ to serve as a posterior predictive $P$-value.

At a late date, I was informed of a paper by Berger and Boos (1994) which introduces the idea of using as a $P$-value the following relative of $P_3$ and $P_4$, $P = \sup_{\theta \in \Gamma} p(T; \theta) + (1 - \gamma)$ where $\Gamma$ is a confidence region of confidence $\gamma$ for a large value of $\gamma$, e.g., 0.999.

9 Conclusions

The standard Yates and Pearson methods for testing for the equality of two probabilities do not provide reliable estimates for $P$-values when one or both sample sizes are moderate, the evidence is strong, and conservative claims are desired. The sources of difficulty are poor asymptotic approximations in the tails of the distributions and the composite nature of the hypothesis to be tested.

One way to deal with this situation is to use exact calculations and to maximize the $P$-value over a suitable range of values of the nuisance parameter. In this special problem, the nuisance parameter is the supposedly common value of the two probabilities being compared. The appropriate range would be a confidence interval with confidence matched to the $P$-value attained.

While the Fisher Exact Test yields a conditional $P$-value, and not a true $P$-value, using this conditional $P$-value as a test statistic provides excellent results, since the resulting $P$-values are almost constant over a wide range of values of the nuisance parameter, and the maximum is relatively small and it is attained at a reasonable value of $p$.

For more general problems, the use of the likelihood-ratio as a test statistic relieves the statistician of the need to find clever test statistics. This approach works about as well as using the Fisher $P$-values for the test statistic in our special problem. It applies easily to the problem of testing $H_3$.

Because the $P$-value pays insufficient attention to the alternative hypothesis, and the composite nature of the hypothesis to be tested, it is good policy to supplement calculations of the $P$-value with alter-
native considerations. Although, the conditional $P_F$ is not a formal P-value, it recommends itself for two reasons. In my case it provides a value which can be presented to a judge as a conservative value. Also it has the advantage of not being influenced by the margins which carry little relevant information.

On the other hand, neither the true P-value nor the conditional value is adequate without considering, formally or informally, the decision theoretic background of the problem, involving the costs of decisions and prior information from past experience or supplementary data.

Perhaps the result that I found, and probably should not have found, most surprising is the difference between the conventional P-value for the Fisher Exact Test and the one obtained by using it as the test statistic.

Suppose that in my legal problem I had to present one number to reflect the information in the case of the limited data on the first side. In that case I would choose the conditional P-value to present to the judge. Suppose that I had to make a terminal decision in some research effort based on those data. Then I would not depend on the P-value, but would use a decision theoretic approach, based mainly on the likelihood-ratio, and considering costs and some informal prior knowledge. Using a Bayesian approach might be sensible. If I were involved in an ongoing research project where I was required to provide one number as a P-value, I would insist on supplementing one of the $P_1$ values with the likelihood-ratio, $\exp(-Z_L)$, which in this case is 2.51(-3). Finally, if I were not constrained to present a single number, I like $P_F(p)$, supplemented by $p^*_F(Z_F)$, which is 5.11(-1) in our case, where $p^*_F(z) = P(Z_F \geq z|p_1 = \hat{p}_1, p_2 = \hat{p}_2)$. A similar analysis could be done with the likelihood-ratio, yielding 5.09(-1).

In short, there is no simple solution to the problem of inference, and there is no one number that is adequate to describe the inferential content of the data.

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