

Hypothesis Tests With Separation

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Separation commonly occurs in political science, usually when the presence (or absence) of a binary explanatory variable perfectly predicts the presence or absence of a binary outcome (e.g., Bell and Miller 2015; Mares 2015; Vining, Wilhelm, and Collens 2015). For example, Barrilleaux and Rainey (2014) find that being a Democrat perfectly predicts a governor accepting Medicaid funds under the Affordable Care Act. Under separation, the usual maximum likelihood estimates are unreasonably large and the Wald p -values are highly misleading.

As a solution, some methodologists propose using a Bayesian prior distribution to regularize the estimates, which we can alternatively consider as *penalized* maximum likelihood estimator. Zorn (2005; see also Heinze and Schemper 2002) points political scientists toward the penalized maximum likelihood estimator proposed by Firth (1993), which is equivalent to Jeffreys' prior distribution (Jeffreys 1946). As an alternative, Gelman et al. (2008) recommend a Cauchy prior distribution. Both of these methods ensure finite

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estimates in theory and usually produce reasonably-sized estimates in practice.

But Rainey (2016) points out that the parameter estimates (and especially the confidence intervals) depend largely on the chosen prior distribution or penalty. Indeed, many priors that guarantee finite estimates can lead to meaningfully different conclusions. He argues that the set of *a priori* “reasonable” and “implausible” parameters depends on the substantive application, so context-free defaults (like Jeffreys’ and Cauchy priors) might not produce reasonable results. Rainey (2016) concludes that “[w]hen facing separation, researchers must *carefully* choose a prior distribution to nearly rule out implausibly large effects” (p. 354).

But it’s not always easy to include prior information, and some scholars prefer to avoid injecting prior information into their model. How can researchers proceed in these situations? In particular, can they obtain useful p -values to test hypotheses in the usual frequentist framework without using prior information?

Below, I show that while the popular Wald test produces misleading (even nonsensical) p -values under separation, likelihood ratio tests and score tests behave in the usual manner. As such, researchers can produce meaningful p -values to test hypotheses with standard frequentist tools under separation *without the use of prior information*.

Statistical Theory

Point Estimates

Maximum likelihood provides a general and powerful framework for obtaining estimates of model parameters. In our case of logistic regression, we write the probability π_i that an event occurs for observation i (or that the outcome variable $y_i = 1$) as

$$\pi_i = \text{logit}^{-1}(X_i\beta) \text{ for } i = 1, 2, \dots, n, \quad (1)$$

where X represents a matrix of explanatory variables and β represents a vector of coefficients. Then we can derive the likelihood function $L(\beta|y)$

$$L(\beta|y) = \pi_i^{y_i} (1 - \pi_i)^{(1-y_i)}, \text{ where } \pi_i = \text{logit}^{-1}(X_i\beta) \quad (2)$$

and the log-likelihood function

$$\ell(\beta|y) = \log L(\beta|y) = y_i \log(\pi_i) + (1 - y_i) \log(1 - \pi_i). \quad (3)$$

For logistic regression (and many other models), researchers typically use numerical algorithms to locate the value $\hat{\beta}^{ML}$ of β that maximizes ℓ . For examples, Stata's `logit` command used a modified Newton-Raphson algorithm (`link`) and R's `glm()` function uses iteratively reweighted least squares. Then, to test hypotheses, we rely on certain features of ℓ , depending on the test.

Hypothesis Tests

With the point estimates in hand, researchers typically compare their research hypothesis H_R to a null hypothesis H_0 . For a logistic regression, the researcher might compose a null hypothesis as $H_0 : \beta \in B_0 \subset R^n$, which leaves the research hypothesis as $H_R : \beta \in B_0^C$. Depending on the data, the researcher may then choose to reject H_0 in favor of H_R or fail to distinguish between the two.

To fix ideas, I focus on the simple point null hypothesis $H_0 : \beta_s = 0$. However, the intuition and conclusions generalize to more complex hypotheses.

The literature offers three common methods to assess the null hypothesis—the “holy trinity” of hypothesis tests: the Wald test, the likelihood ratio test, and the score test. For practical reasons, most regression tables in political science include p -values and stars based on the Wald test. While usually the most convenient of the three tests, the Wald test is uniquely ill-suited for testing hypotheses under separation. The likelihood

ratio and score tests, on the other hand, work as expected. Below I briefly describe each test, explain why the Wald test works poorly under separation, and describe why the likelihood ratio and score tests perform better.

Wald Test

Of the three methods, researchers usually report the Wald test because it requires fitting only the full model. The Wald procedure uses the shape of the log-likelihood function around the maximum to estimate the precision of the point estimate. If small changes in the parameter near the maximum lead to large changes in the log-likelihood function, then we can treat the maximum likelihood estimate as precise. However, if large changes in the parameter lead to small changes in the likelihood function, then we must treat the maximum likelihood estimate as imprecise.

More formally, the Wald test uses the second derivative to quantify the curvature of ℓ at $\hat{\beta}^{ML}$. Further, we can estimate the standard error $\widehat{SE}(\hat{\beta}_i^{ML})$ as

$$\widehat{SE}(\hat{\beta}_i^{ML}) = \left(-\frac{\partial^2 \ell(\hat{\beta}_i^{ML}|y)}{\partial^2 \hat{\beta}_i^{ML}} \right)^{-\frac{1}{2}}. \quad (4)$$

For large samples, the maximum likelihood estimate approximately follows a normal distribution centered at the true value of β with a standard deviation of $\widehat{SE}(\hat{\beta}_i^{ML})$. Wald (1943) advises us how compare the estimate with the standard error: the statistic $Z_w = \frac{\hat{\beta}_i^{ML}}{\widehat{SE}(\hat{\beta}_i^{ML})}$ follows a standard normal distribution approximate follows a standard normal distribution. (Casella and Berger 2003, pp. ???, Greene 2012, pp. 527-529).

However, *this approach works poorly when dealing with separation*. Under separation, the log-likelihood function at the found maximum is numerically flat or nearly so. The flatness produces very large standard error estimates, which has a troubling dependence on the error tolerance of the algorithm.

For a logistic regression model with a separating variable, we can state a precise result

that relates the size of the coefficient of the separating variable to the Wald estimate of its standard error.

Theorem 1 *Suppose that a binary explanatory variable s with coefficient β_s perfectly predicts the outcome y_i such that when $s_i = 1$ then $y_i = 1$. Then the log-likelihood function increases in β_s and the quantity $\lim_{\beta_s \rightarrow \infty} \left[\left(-\frac{\partial^2 \ell(\beta_s | y)}{\partial^2 \beta_s} \right)^{-\frac{1}{2}} - \beta_s \right] = \infty$. Thus, under separation, the estimated standard error will be much larger than the coefficient for the separating variable for any algorithm that obtains a sufficiently large coefficient.*

Theorem 1 shows that, so long as the researcher uses a sufficiently precise algorithm, the Wald test will *never* reject the null hypothesis.

Theorem 1 has an important practical implication, given in Corollary 1.

Corollary 1 *Suppose a data-generating process that results in separation with probability p . Then the power of the Wald test cannot exceed $1 - p$.*

This has a counter-intuitive implication that highlights the poor behavior of the Wald test under separation. Increasing the coefficient of a potentially separating variable increases the chance of separation. But Corollary 1 established that this must *decrease* the power of the test. This leads to a pathological result, moving the coefficient further from zero must *decrease* the power of the test.

Figure 1 shows this intuition for a typical, non-monotonic log-likelihood function (i.e., without separation) and for a monotonic log-likelihood function (i.e., with separation). In the absence of separation, the curvature of the log-likelihood function around the maximum speaks to the evidence against the null hypothesis. But under separation, the likelihood function, *by definition*, is flat at the maximum, regardless of the relative likelihood of the data under the null hypothesis.

To see the importance of this result, suppose an absurd example in which a binary treatment perfectly predicts 500 successes and 500 failures (i.e., $y = x$ always). Of course, these data are *extremely* unlikely under the null hypothesis that the coefficient for the treatment indicator equals zero. The exact p -value for the null hypothesis that

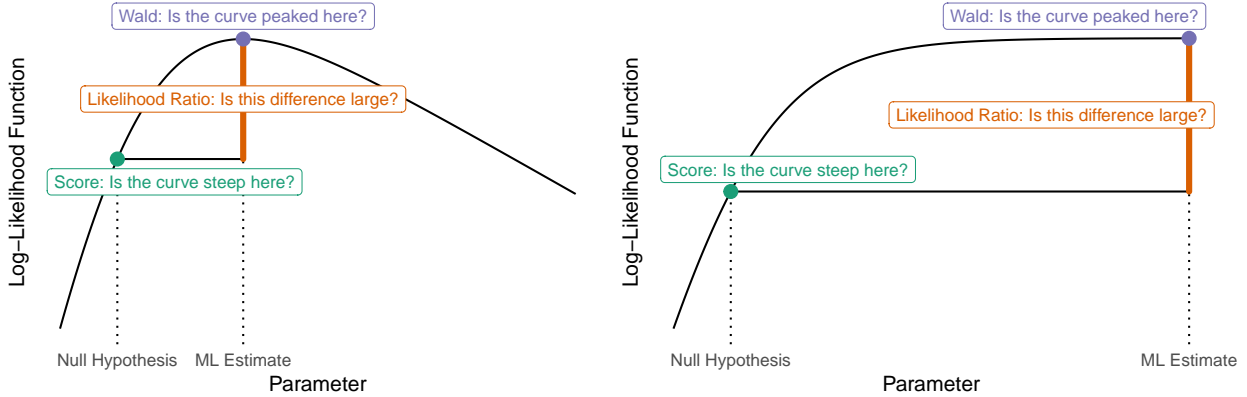


Figure 1: A figure summarizing logic of the "holy trinity" of hypothesis tests. The Wald test relies on the curvature around the maximum of the log-likelihood function, which breaks down under separation. But the likelihood ratio and score test rely on *other* features of the log-likelihood function, that are not meaningfully impacted by separation.

successes and failures are equally likely under both treatment and control equals $2 \times \left(\frac{1}{2}\right)^{500} \times \left(\frac{1}{2}\right)^{500} = \frac{2}{2^{1000}} \approx \frac{2}{10^{301}}$. (For comparison, there are about 10^{80} atoms in the known universe.) Yet, the default `glm()` routine in R calculates a Wald p -value of 0.998 with the default precision and 1.000 with the maximum precision.

When dealing with separation, the approach of using curvature around the maximum to estimate the relative likelihood of the restricted model breaks down; researchers cannot use the Wald test to obtain reasonable p -values for the coefficient of a separating variable.

Likelihood Ratio Test

The likelihood ratio test resolves the problem of the flat log-likelihood by comparing the maximum log-likelihood of two models: an "unrestricted" model ML that imposes no bounds on the estimates and a "restricted" model ML_0 that constrains the estimates to the region suggested by the null hypothesis. If the data are much more likely under the unrestricted estimate than under the restricted estimate, then the researcher can reject the null hypothesis.

More formally, the likelihood ratio test compares the value of the unrestricted log-likelihood $\ell(\hat{\beta}^{ML}|y)$ to the restricted log-likelihood $\ell(\hat{\beta}^{ML_0}|y)$. For the simple null hypoth-

esis $H_0 : \beta_s = 0$, we can understand ML_0 as a separate model fit without the explanatory variable s .

Wilks' theorem (1938) advises us how to compare the two likelihoods: $D = 2 \times [\ell(\hat{\beta}^{ML}|y) - \ell(\hat{\beta}^{ML_0}|y)]$ follows a χ^2 distribution with degrees of freedom equal to the number of constrained dimensions (Casella and Berger 2003, pp. 488-492, Greene 2003, pp. 484-486).

Figure 1 also shows the intuition of the likelihood ratio test. The figure highlights that the gap between the unrestricted and restricted maximum summarizes the evidence against the null hypothesis. Further, the logic does not break down under separation. Unlike the Wald test, the likelihood ratio test can reject the null hypothesis under separation.

Score Test

The score test resolves the problem of the flat log-likelihood by focusing on gradient of the log-likelihood function at the null hypothesis. If the null hypothesis is correct, then the log-likelihood should not be increasing much around that point. But if the log-likelihood function is increasing rapidly at the null hypothesis, this casts doubt on the null hypothesis. The score test uses two quantities: the score function $S(\beta) = \frac{\partial \ell(\beta|y)}{\partial \beta}$ and the Fisher information $I(\beta) = -E_{\beta} \left(\frac{\partial^2 \ell(\beta|y)}{\partial^2 \beta} \right)$. When evaluated at the null hypothesis, the score function quantifies the slope and the Fisher information quantifies the variance of that slope in repeated samples.

If the score at the null hypothesis is large relative to its standard deviation, then researchers can reject the null hypothesis. Rao (1948) advises us how evaluate the assess the statistic: $Z_s = \frac{S(\beta_s^0)}{\sqrt{I(\beta_s^0)}}$ follows a standard normal distribution. (Rao 1948, Casella and Berger 2003, pp. 494-495, Greene 2003, pp. 489-490).

Figure 1 finally shows the intuition of the score test. The figure highlights that the slope of the log-likelihood function under the null hypothesis summarizes the evidence

against the null hypothesis. As with the likelihood ratio test, the logic works even under separation, and the score test can reject the null hypothesis under separation.

Table 1 summarizes the three tests. Most importantly, the likelihood ratio and score test rely on features of the log-likelihood function that are not meaningfully affected by a monotonic log-likelihood function created by separation. The Wald test, on the other hand, cannot provide a reasonable tests under separation (see Theorem 1). For more details on the three tests, see Silvey (1959) or Greene (2012, pp. 524–530).

Table 1: A table summarizing the "holy trinity" of hypothesis tests.

Test	Feature	Statistic and Distribution
Wald	Curvature of the log-likelihood function around the maximum.	$Z_w = \frac{\hat{\beta}_i^{ML}}{\widehat{SE}(\hat{\beta}_i^{ML})}$ follows a standard normal distribution.
Likelihood Ratio	Relative log-likelihoods of the unrestricted and restricted models.	$D = 2 \times [\ell(\hat{\beta}^{ML} y) - \ell(\hat{\beta}^{ML_0} y)]$ follows a χ^2 distribution with degrees of freedom equal to the number of constrained dimensions.
Score	Slope of the log-likelihood function at the null hypothesis.	$Z_s = \frac{S(\beta_s^0)}{\sqrt{I(\beta_s^0)}}$ follows a standard normal distribution.

Simulations

To evaluate the performance of the various methods for testing hypothesis under separation, I use a diverse collection of data-generating processes (DGPs) that feature separation in more than 10% of repeated samples. Importantly, I cannot focus on data sets *with separation* because separation is a feature of a particular sample. Instead, I focus on DGPs that *sometimes* feature separation (e.g., in 15% of repeated samples, in 50% of repeated samples, etc.).

To create the collection of DGPs, I imagine the logistic regression model $\Pr(y = 1) = \text{logit}^{-1}(\beta_{\text{cons}} + \beta_s s + \beta_{z_1} z_1 + \dots + \beta_{z_k} z_k)$ and a researcher testing the null hypothesis that the binary explanatory variable s (that might produce separation) has no effect on a binary outcome variable y . I vary the frequency that $s = 1$, the value of β_{cons} , the number of control variables (k), and the total number of observations. For each DGP, I use Monte

Carlo simulation to compute the power function as β_s varies for each of the three tests: Wald test, likelihood ratio test, and score test. For completeness, I also compute the power function for Wald tests using the two Firth and Cauchy PML estimates.

I use all 944 (logically possible) combinations of the values below. When the null hypothesis true (i.e., $\beta_s = 0$), the percent of repeated samples with separation ranges from 98% (ctk) to 14% (ctk) in this collection.

A Close Look at a Single DGP

To clarify the simulation results, I provide the details for a single DGP. Following the structure discussed above, Table 2 shows the parameters for this single DGP.

Table 2

Parameter	Value
Total Number of Observations	50
Frequency that $s = 1$	5
The Value of the Intercept β_{cons}	0
The Number of Control Variables k	2

Table 3 shows the power function for each of the three tests, as well as the ideal power, and the percent of the data sets that featured separation for the parameter combination and the value of β_s . For this particular DGP, separation is relatively rare when β_s —the coefficient for the potentially separating variable—is near zero. But for $\beta_s = \pm 2$, about 50% of the data sets feature separation. And for β_s larger/smaller than ± 4 , more than 90% of the data sets feature separation.

This DGP clearly shows the implication of Theorem 1. Even though the data sets with separation should allow the researcher to reject the null hypothesis, at occasionally, the power of the Wald test is near 0%, even for very large effects. The likelihood ratio and score tests, on the other hand, perform as expected. For both alternatives, the power of the test when $\beta_s = 1$ is about 5%, as designed, and the power approaches 100% relatively quickly as β_s moves away from zero. For this particular DGP, the likelihood ratio is a

more powerful test, than the score test, but the likelihood ratio and score tests work reasonably well, unlike the Wald test.

Table 3

β_s	Ideal Power	Percent with Separation	Wald Test Power	Likelihood Ratio Test Power	Score Test Power
-10	As high as possible.	100%	0%	99%	85%
-7		99%	0%	99%	85%
-4		89%	0%	90%	77%
-2		45%	2%	49%	42%
-1		13%	2%	18%	15%
0	5%	4%	1%	6%	5%
1	As high as possible.	19%	0%	20%	16%
2		50%	0%	51%	41%
4		90%	0%	89%	70%
7		99%	0%	97%	76%
10		100%	0%	98%	78%

This table shows the power for the Wald, likelihood ratio, and score tests for a DGP that often features separation.

A Broad Look at Many DGPs

Table ?? shows the broad collection of parameters I used in the simulations. All combinations of the values below yield 150 unique combinations. I excluded combinations in which the frequency that $s = 1$ exceeded the number of observations, which leaves 110 combinations. For each of these 110 combinations, I use Monte Carlo simulations to estimate the probability that each method rejects the null hypothesis as the coefficient β_s varies from -10 to 10 (across the particular values shown in Table 3). I also exclude any scenario in which the parameter and the coefficient β_s would sometimes produce data sets with no variation in the outcome variable. As such, one might consider this a diverse collection of DGPs that sometimes generate separation but almost always have variation in the outcome.

Figure 2 shows the statistical power of each of the three tests as the chance of separation varies. Most starkly, the power of the Wald test is bounded above my the chance of no-separation. Intuitively, as the coefficient of the potential separating variable

Table 4

Parameter	Value
Total Number of Observations	50, 250, 500
Frequency that $s = 1$	5, 25, 50, 100, 250
The Value of the Intercept β_{cons}	-8, -5, -2.5, -1, 0
The Number of Control Variables k	2, 6

increases, the power of the test should *increase*. But because a large coefficient makes separation more likely, a large coefficient *decreases* the power of the test. The likelihood ratio and score test behave as expected.

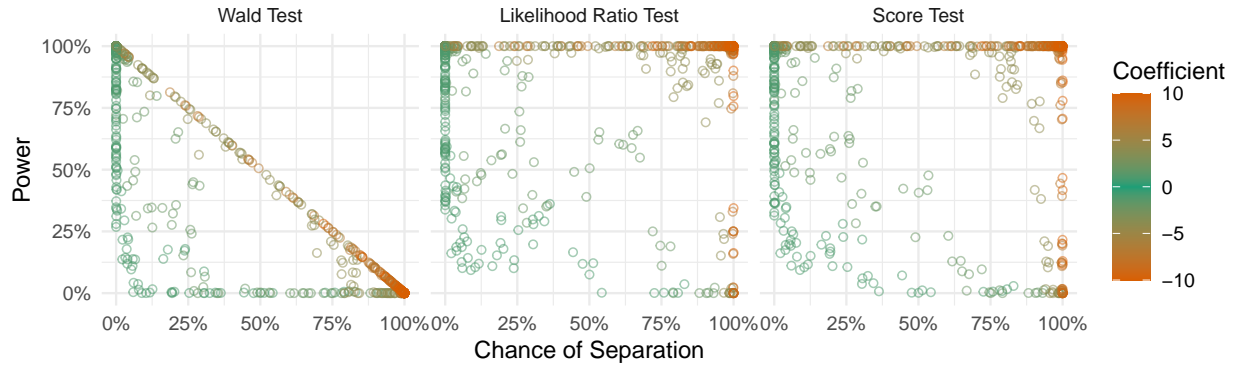


Figure 2: caption here

Figure 3 plots and summarized the power function for each of the 110 DGPs in the diverse collection.¹ The fine lines show the power function for each DGP and the color of the points indicates the chance of separation at that point. The heavy, solid lines show the average of the collection of power functions. The heavy, dashed lines show the power functions for the other two tests, for comparison. First, the Wald test power functions show it poor properties. For most of the power functions, as the true coefficients grows larger in magnitude from about three, the test becomes less powerful. This occurs because separation becomes more likely and the test cannot reject the null hypothesis when separation occurs. Second, the likelihood ratio and score test behave reasonably well.

¹I exclude parameter combinations that might not produce variation in the outcome variable, so I do not compute the power function across the entire range of the coefficient for every power function. For 47 of the 110 DGPs, I exclude a portion of the range. For 63 of the 110, I compute the power function across the entire range from -10 to 10.

Most importantly, the size of the likelihood ratio and score tests is about 5% when the coefficient equals zero and grows as the coefficient moves away from zero. The likelihood ratio test is slightly preferred, at least of this collection of DGPs, because it is slightly more powerful when the coefficient does not equal zero, especially for large, negative coefficients.²

This simulations illustrate Theorem 1 and Corollary 1. Figure 2 shows that the power of the Wald test never exceeds the chance of no-separation. For data sets with large effects, separation might be very likely and—counter-intuitively—the chance of rejecting the null hypothesis might be very low. The simulations also, show, though, that the likelihood ratio and score tests behave as expected.

Re-Analysis of Barrilleaux and Rainey (2014)

To illustrate frequentist hypothesis testing under separation, I reanalyze data from Barrilleaux and Rainey (2014) (that Rainey (2016) considers in great detail). Barrilleaux and Rainey (2014) examine U.S. state governors decisions to support or oppose the Medicaid expansion under the 2010 Affordable Care Act. But because all Democratic governors supported the expansion, separation occurs—being a Democratic governor perfectly predicts support for Medicaid expansion.

I focus on their first hypothesis: *Republican governors are more likely to oppose the Medicaid expansion funds than Democratic governors*. Barrilleaux and Rainey adopt a fully Bayesian approach, modeling the probability that a state's governor opposes the Medicaid expansion as a function of the governor's partisanship and several other covariates. Here, I re-estimate their logistic regression model using several frequentist procedures. The appendix provides the full details. Table 5 presents the estimates and p -values for the coefficient of the binary Democratic governor variable.

Because no Democratic governors oppose the expansion, being a Democrat perfectly

²This asymmetry occurs because I only consider negative intercepts.

Table 5

Estimator	Coef. Est.	SE Est.	Wald <i>p</i> -Value	LR <i>p</i> -Value	Score <i>p</i> -Value
ML with Default Precision	-20.35	3,224	0.99	0.00	0.01
ML with Maximum Precision	-35.22	15 million	1.00	0.00	0.01
PML with Jeffreys Penalty	-2.68	1.42	0.06		
PML with Cauchy Penalty	-3.38	1.63	0.04		

This table shows the *p*-values from several procedures that researchers might use when dealing with separation in logistic regression models. The Wald test for maximum likelihood estimates relies on unreasonable standard errors that depend heavily on the precision of the algorithm and, as a consequence, produces unrealistic *p*-values. However, the *p*-values from the likelihood ratio test seem reasonable and resemble the *p*-values from the more conservative penalized maximum likelihood approaches.

predicts non-opposition. Therefore, the coefficient estimates are implausibly large. To address this, Barrilleaux and Rainey use a Bayesian approach. Other authors suggest penalized maximum likelihood, but Rainey (2016) shows that the choice of penalty can impact the results. This example shows that the likelihood ratio and score tests offer reasonable *p*-values without using prior information or a identifying a suitable penalty.

Conclusion

Separation commonly occurs in political science. When this happens, I shows that the usual Wald *p*-values are highly misleading. But researchers cannot always use suitable prior information to address a monotonic likelihood function. Even without a suitable prior or penalty, I show that the standard likelihood ratio and score tests behave in the usual way. As such, researchers can use the likelihood ratio and score tests to produce meaningful *p*-values under separation *without the use of prior information*.

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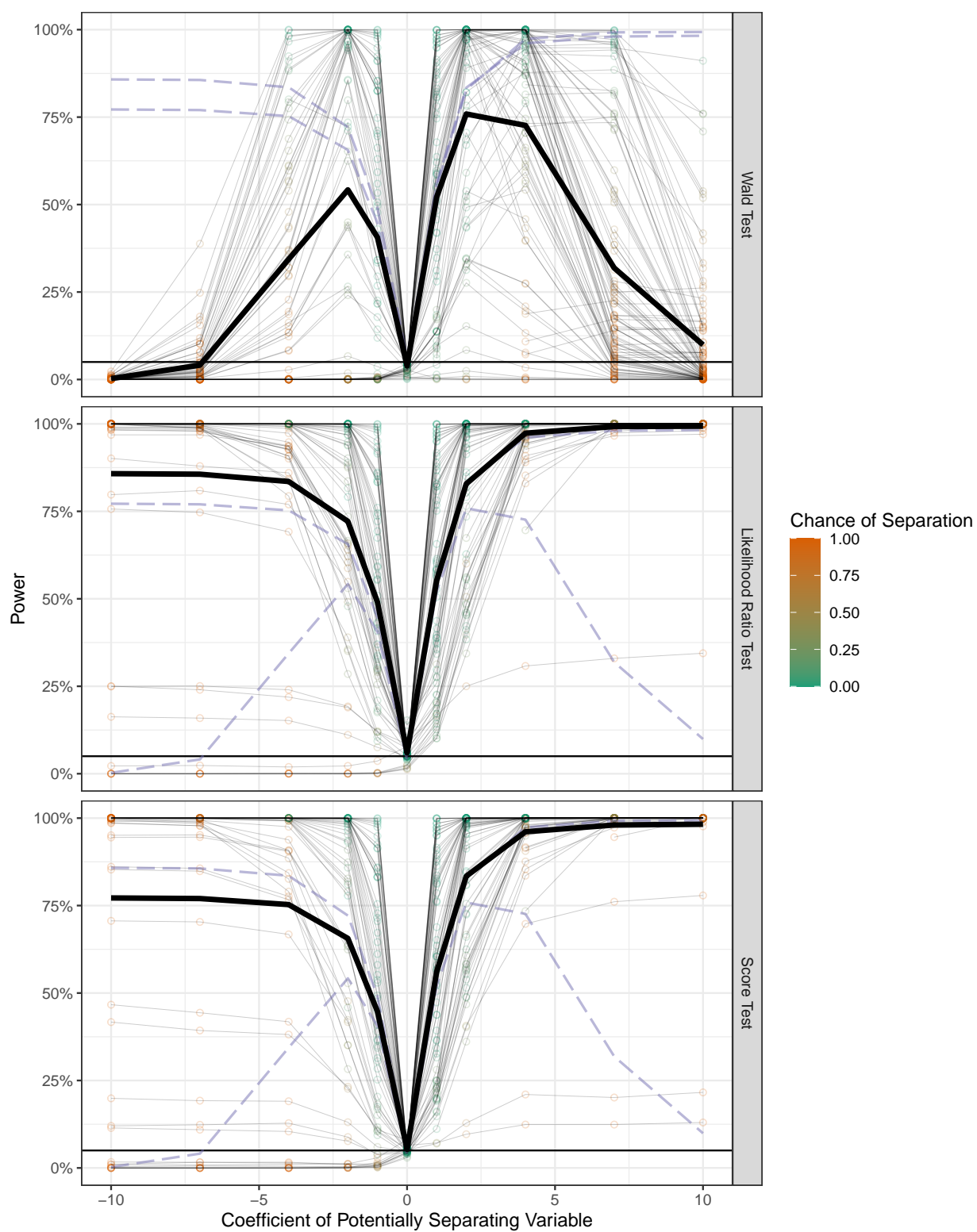


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