

ON DETERMINING THE POWER OF A TEST AFTER DATA COLLECTION

by

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ABSTRACT

The term retrospective power describes methods for estimating the true power of a test after data have been collected. These methods have been recommended by some authors when null hypothesis of a test cannot be rejected. This report uses simulations to study power as a construct of an observed effect, variance, sample size, and set level of significance under the balanced one-way analysis of variance model for normally distributed populations with constant variance.

Retrospective power, as a construct of sample data, is not recommended when the null hypothesis of a test cannot be rejected. When the p-value of the test is large, estimates for true power tend to fall below the 0.80 level and width-minimized confidence limits for true power tend to be wide.

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1 Introduction

For many researchers, failing to reject the null hypothesis of statistical test is disconcerting. In some cases, the research design is questioned: was the study powerful enough to detect a meaningful effect? While valid statistical methods exist for calculating power before data have been collected, determining power retrospectively is seen in the literature as controversial. Here we explore power, as described in Thomas (1997), as a construct of sample data using the balanced one-way ANOVA model. Section 2 discusses methods for calculating prospective power; methods for estimating retrospective power; and concerns associated with estimating retrospective power. Section 3 outlines the balanced one-way ANOVA model; a method for estimating power using the observed effect, variance, sample size, and set significance level; and a method for constructing a minimal $(1 - \alpha)100\%$ confidence interval for true power. In addition, Section 3 contains conditions for simulations and implementation methods used for simulations. Section 4 summarizes the simulation results, displaying the location of true power, median estimated power, average estimated power, average p-value of tests and average confidence limits produced from the simulations. Section 4 also discusses the effect of population conditions on power calculations. Flow charts, code, and complete output are given in appendices A-D. By examining power after data have been collected and statistical tests have been performed, the perspective shifts from calculating power to estimating power.

2 Literature Review

Calculating the power of a statistical test before data have been collected, denoted *prospective power*, is generally regarded by statisticians as a good practice for obtaining conclusive results (American Statistical Association 1999). Power calculations are useful in determining adequate sample sizes for an up coming study (Lenth 2001). The power of a statistical test can be computed using a set significance level, sample size, effect, and variance (Lenth 2001; Gerard et al. 1998). Lenth (2001) provides a list of discussion-techniques for eliciting meaningful effect and variance values for a potential study. Power for t-tests, multiple regression, one-way anova, and other common statistical methods can be computed using the SAS procedure POWER (SAS Institute Inc. 2004b). Power for linear models with fixed class effects and contrast statements can be calculated using the SAS procedure GLMPOWER (SAS Institute Inc. 2004a). For many researchers, prospective power calculations are within reach.

Calculating the power of a statistical test after data have been collected, i.e. *retrospective power*, is less accepted by many statisticians as compared to prospective power (Hoenig and Heisey 2001; Thomas 1997; Yuan and Maxwell 2005). Retrospective power has been proposed for interpreting results when the null hypothesis cannot be rejected (Taylor and Muller 1995; Hogarty and Kromrey 2001) and for determining the sample size needed for the observed data to show statistically significant results (Steidl et al. 1997; Thomas 1997). Some popular methods for calculating retrospective power included: an observed sample size and given significance level where effect and variance come from literature (Yuan and Maxwell 2005); an observed effect, observed sample size, and given significance level (Yuan and Maxwell 2005) where variance comes from literature; an observed variance, observed sample size, and given significance level (Thomas 1997; Yuan and Maxwell 2005; Steidl et al. 1997) where effect comes from literature; and an observed effect, an observed sample size, an

observed variance, and given significance level (Gerard et al. 1998; Thomas 1997; Hoenig and Heisey 2001). Of these four methods only the last method is reviewed here.

Calculating retrospective power using an observed effect, an observed variance, an observed sample size, and given significance level has been shown to produce biased estimates when using an observed noncentrality parameter (Gerard et al. 1998). Thomas (1997) suggests adjusting the observed noncentrality parameter as shown in Wright and O'Brien (1988) and Johnson et al. (1995) to remove the bias in calculating power. The adjusted observed noncentrality parameter can produce negative values, which are typically set to zero (Gerard et al. 1998). A median estimator for the noncentrality parameter has been proposed by Taylor and Muller (1996) as a means for correcting the retrospective power calculation. The median estimator for the noncentrality parameter under-estimates half the time and over-estimates half the time. Here, the adjusted observed noncentrality parameter is explored.

Concerns about using retrospective power procedures to represent true power exist in the literature. Zumbo and Hubley (1998) provide a Bayesian argument against retrospective power as a logical representative for true power. Hoening and Heisey (2001) note that retrospective power obtain through observed values is redundant with the p-value of the test.

Due to the bias associated with retrospective power, confidence intervals for effect have been proposed in lieu of retrospective power analyses when the null hypothesis cannot be rejected (Gerard et al. 1998; Thomas 1997; Steidl et al. 1997).

In this paper, we examine the method presented in Thomas (1997) for estimating the power of a test in the context of the balanced one-way ANOVA model.

3 Methods

3.1 Model

To investigate post-hoc power, the balanced one-way analysis of variance model is considered. The following briefly reviews this model and the usual analysis as described in Kuehl (2000) and Milliken and Johnson (2009). The means model for this situation is

$$y_{ij} = \mu_i + \varepsilon_{ij}, \text{ with } \varepsilon_{ij} \text{ 's i.i.d } N(0, \sigma^2) \text{ for } i=1, \dots, p, j=1, \dots, n. \quad (1.1)$$

Here y_{ij} represents the observed value of the j th observation from the i th treatment group and μ_i denotes the population mean of the i th treatment group. The term ε_{ij} represents the random error associated with the j th observation from the i th treatment group. The errors are assumed independent and identically distributed from a normal population with mean 0 and variance σ^2 .

The null and alternative hypotheses are that of no difference and that of some difference among the p population means, respectively:

$$\begin{aligned} H_0 &: \mu_1 = \dots = \mu_p \\ H_a &: \text{at least two } \mu_i \text{'s not equal} \end{aligned} \quad (1.2)$$

When the sample means are not “too different” from each other we deem the data consistent with the null hypothesis of equal means. We do not conclude that the population means are all the same but acknowledge that sample means as too similar to be called different. When at least two sample means exist which “strongly” differ we favor the alternative hypothesis and conclude that not all of the population means are equal to one another.

Two estimates of variance are used to measure the amount of evidence against the null hypothesis. The first estimate of variance is based only on the assumptions of the model. The second estimate of variance is based on the assumptions of the model and the assumption that

the null hypothesis is true. Of these two estimates of variance, the second estimate has a larger expected mean square than the first estimate when the null hypothesis is false.

The mean square for error is the first estimate of σ^2 and is calculated by taking the average of the sample variances $s_1^2, s_2^2, \dots, s_p^2$:

$$\begin{aligned}
MSE &= \frac{s_1^2 + s_2^2 + \dots + s_p^2}{p} \\
&= \frac{1}{p(n-1)} \sum_{i=1}^p \sum_{j=1}^n (y_{ij} - \bar{y}_{i\cdot})^2 \\
&= \frac{1}{p(n-1)} \left\{ \sum_{i=1}^p \sum_{j=1}^n y_{ij}^2 - n \sum_{i=1}^p \bar{y}_{i\cdot}^2 \right\} \\
&= \frac{1}{p(n-1)} \left\{ \sum_{i=1}^p \sum_{j=1}^n y_{ij}^2 - n \bar{Y}^T \bar{Y} \right\} \\
&= \frac{SSE}{DFE},
\end{aligned} \tag{1.3}$$

where $\bar{y}_{i\cdot} = \frac{1}{n} \sum_{j=1}^n y_{ij}$, $\bar{Y}^T = [\bar{y}_{1\cdot}, \bar{y}_{2\cdot}, \dots, \bar{y}_{p\cdot}]$, SSE is the error sum of squares, and DFE is the

error degrees of freedom. Under the assumptions of the model,

$$\frac{(DFE)(MSE)}{\sigma^2} \sim \chi^2_{(DFE)} \tag{1.4}$$

and

$$E(MSE) = \sigma^2. \tag{1.5}$$

The mean square for treatments (MST) is the second estimate of σ^2 and is calculated as the sample variance of the sample means $\bar{y}_{1\cdot}, \bar{y}_{2\cdot}, \dots, \bar{y}_{p\cdot}$. That is,

$$\begin{aligned}
MST &= \frac{n}{p-1} \sum_{i=1}^p (\bar{y}_{i\cdot} - \bar{y}_{\cdot\cdot})^2 \\
&= \frac{n}{p-1} \left\{ \sum_{i=1}^p \bar{y}_{i\cdot}^2 - p\bar{y}_{\cdot\cdot}^2 \right\} \\
&= \frac{1}{p-1} \left\{ \sum_{i=1}^p \frac{y_{i\cdot}^2}{n} - \frac{\bar{y}_{\cdot\cdot}^2}{np} \right\} \\
&= \frac{1}{p-1} \left\{ n\bar{Y}^T \bar{Y} - \frac{n}{p} (\bar{Y}^T J)^2 \right\} \\
&= \frac{SST}{DFT},
\end{aligned} \tag{1.6}$$

where $y_{i\cdot} = \sum_{j=1}^n y_{ij}$, $y_{\cdot\cdot} = \sum_{i=1}^p \sum_{j=1}^n y_{ij}$, $J^T = [1, 1, \dots, 1]$, SST is the treatment sum of squares, and

DFT is the treatment degrees of freedom. Under the assumptions of the model and the assumption that the null hypothesis is true

$$\frac{(DFT)(MST)}{\sigma^2} \sim \chi^2_{(DFT)} \tag{1.7}$$

and

$$E(MST) = \sigma^2. \tag{1.8}$$

Under the assumptions of the model and the assumption that the null hypothesis is not true

$$\frac{(DFT)(MST)}{\sigma^2} \sim \chi^2_{(DFT, \lambda)} \tag{1.9}$$

and

$$E(MST) = \sigma^2 + \frac{n}{p-1} \sum_{i=1}^p (\mu_i - \bar{\mu}_{\cdot\cdot})^2, \tag{1.10}$$

where

$$\lambda = \frac{n}{\sigma^2} \sum_{i=1}^p (\mu_i - \bar{\mu}_{\cdot\cdot})^2 \tag{1.11}$$

denotes the noncentrality parameter.

Under the assumptions of the model and assumption that the null hypothesis is true,

$$\frac{[(DFT)(MST) / \sigma^2] / DFT}{[(DFE)(MSE) / \sigma^2] / DFE} \sim F_{(DFT, DFE)}, \quad (1.12)$$

where (1.7) and (1.9) are independently distributed χ^2 random variables. Under the

assumptions of the model and the assumption that the alternative hypothesis is true,

$$\frac{[(DFT)(MST) / \sigma^2] / DFT}{[(DFE)(MSE) / \sigma^2] / DFE} \sim F_{(DFT, DFE, \lambda)}. \quad (1.13)$$

The observed F -statistic for the test is calculated as the ratio of the mean square for treatments over the mean square for error. This statistic is used to measure the amount of evidence in the sample data against the null hypothesis,

$$F = \frac{MST}{MSE} = \frac{[(DFT)(MST) / \sigma^2] / DFT}{[(DFE)(MSE) / \sigma^2] / DFE} \quad (1.14)$$

Given the assumptions of the means model and the assumption that the null hypothesis is true, the mean square for error should be of similar size to the mean square for treatments, practically speaking. However, given the null hypothesis is false, the mean square for error should be smaller than the mean square for treatments. An uncommonly large F -statistic leads to the conclusion that not all population means are equal. The p -value of the test gives the probability of obtaining an F -statistic greater than or equal to the one observed. The more evidence in the sample data against the null hypothesis of equal means the smaller the p -value.

3.2 Estimated Power

Power is estimated using the observed effect and the observed sampling variance. Let the cumulative distribution function of the noncentral F -distribution be denoted as

$$F(DFT, DFE, \lambda) . \quad (2.1)$$

The power of the F -test is defined as

$$\text{power} = 1 - F(F_{crit} | DFT, DFE, \lambda) , \quad (2.2)$$

where F_{crit} is the $100 \cdot (1 - \alpha)$ percentile from a central F -distribution with numerator degrees of freedom DFT , denominator degrees of freedom DFE , level of significance α , and noncentrality parameter λ (Thomas 1997). A positively-biased estimate for the power of an F -test is

$$\widehat{\text{power}} = 1 - F(F_{crit} | DFT, DFE, \hat{\lambda}) \quad (2.3)$$

where $\hat{\lambda} = SST / MSE$ (Thomas 1997). The estimated noncentrality parameter can be adjusted to remove the upward bias. Therefore, an unbiased estimate of power (Wright and O'Brien 1988, and Johnson et al. 1995, cited in Thomas 1997) is

$$\widehat{\text{power}}_{adj} = 1 - F(F_{crit} | DFT, DFE, \hat{\lambda}_{adj}) \quad (2.4)$$

where $\hat{\lambda}_{adj}$ is the adjusted estimated noncentrality parameter and is computed as

$$\hat{\lambda}_{adj} = [\hat{\lambda} \cdot (DFE - 2) / DFE] - DFT . \quad (2.5)$$

Note that for $\hat{\lambda} \cdot (DFE - 2) / DFE$ less than DFT , $\hat{\lambda}_{adj}$ can be negative.

Upper and lower confidence limits for the true power are therefore

$$\widehat{\text{power}}_U = 1 - F\left(F_{crit} \mid DFT, DFE, \hat{\lambda}_U\right) \quad (2.6)$$

and

$$\widehat{\text{power}}_L = 1 - F\left(F_{crit} \mid DFT, DFE, \hat{\lambda}_L\right), \quad (2.7)$$

where $\hat{\lambda}_U$ and $\hat{\lambda}_L$ satisfy $F\left(F_{obs} \mid DFT, DFE, \hat{\lambda}_U\right) = \alpha_U$ and $F\left(F_{obs} \mid DFT, DFE, \hat{\lambda}_L\right) = 1 - \alpha_L$ in

which α_U is the upper tail probability and α_L lower tail probability (Thomas 1997). The upper and lower tail probabilities α_U and α_L are those which define the $100 \cdot (1 - \alpha_U - \alpha_L)$ percent confidence interval for true power.

3.3 Confidence Interval Optimization

Confidence limits for true power were minimized to explore retrospective power under a “best” case scenario. Confidence limits set under $\alpha_L = \alpha_U = \alpha / 2$ are not generally optimal (i.e. narrowest confidence interval) because of the skewed nature of the noncentral F -distribution. Setting upper and lower probabilities equal may under-represent, or over-represent, true power.

For a user-specified level of significance α , we consider the $100(1 - \alpha)\%$ confidence interval for true power, where

$$\alpha = \alpha_L + \alpha_U. \quad (3.1)$$

We denote the inverse cumulative distribution function of the noncentral F -distribution by

$$F^{-1}\left(\alpha_c, DFT, DFE, \hat{\lambda}_{adj}\right), \quad (3.2)$$

where α_c is a lower tail probability, DFT the numerator degrees of freedom, DFE the denominator degrees of freedom, and $\hat{\lambda}_{adj}$ the noncentrality parameter. We denote the quantile associated with α_L as

$$F_L = F^{-1}(\alpha_L, DFT, DFE, \hat{\lambda}_{adj}), \quad (3.3)$$

and the quantile associated with α_U as

$$F_U = F^{-1}(1 - \alpha_U, DFT, DFE, \hat{\lambda}_{adj}). \quad (3.4)$$

The width of the $100(1 - \alpha)\%$ confidence interval for true power is minimized by finding the distance between the lower and upper quantiles such that

$$W = F_U - F_L \quad (3.5)$$

is minimized. Note, for fixed α we have the constraint that $\alpha = \alpha_L + \alpha_U$ (3.1) which gives

$$\alpha_U = \alpha - \alpha_L. \quad (3.6)$$

As such, we wish to find the value of α_L which minimize the distance

$$W(\alpha_L) = F^{-1}(1 - \alpha + \alpha_L, DFT, DFE, \hat{\lambda}_{adj}) - F^{-1}(\alpha_L, DFT, DFE, \hat{\lambda}_{adj}), \quad (3.7)$$

for

$$\alpha_L \in (0, \alpha). \quad (3.8)$$

A method for finding α_L such that $W(x)$ is minimal is by the Golden Section Search (GSS) as described in Press et al. (2007). For a unimodal function $W(x)$ defined for $x \in [a, b]$, the Golden Section Search method seeks to bracket the value k in $[a, b]$ such that $W(k) < W(x)$ for all other $x \in [a, b]$. The value k is considered found when brackets are

reached whose distance is within a user-specified tolerance level. The GSS procedure is named for its use of the mathematical constant

$$\varphi = \frac{-1 + \sqrt{5}}{2} \approx 0.618034 . \quad (3.9)$$

The ends points a and b are used to construct two new possible bracket points. We calculate the first new point, which we call c , by

$$c = a + \varphi(b - a) , \quad (3.10)$$

and the second new point, which we call d , by

$$d = a + \varphi^2(b - a) . \quad (3.11)$$

These new points are then evaluated with the function $W(x)$ such that

$$u = W(c) \quad (3.12)$$

and

$$v = W(d) . \quad (3.13)$$

One of two cases may arise:

$$(1) u > v$$

$$(2) u \leq v$$

For the first case (Figure 3.1.a) the value k must be in $[a,c]$ since the function $W(x)$ is unimodal. The interval $[a,c]$ will then have width $c - a = \varphi(b - a)$ (3.10). For the second case (Figure 3.1.b) the value k must be in $[d,b]$, again since the function $W(x)$ is unimodal. Using the equation $d = a + \varphi^2(b - a)$ (3.11) and the fact that $\varphi^2 = 1 - \varphi$, it can be shown that $b - d = \varphi(b - a)$. Therefore,

Figure 3.1.a: Update Bracket Interval for Minimand for v less than u

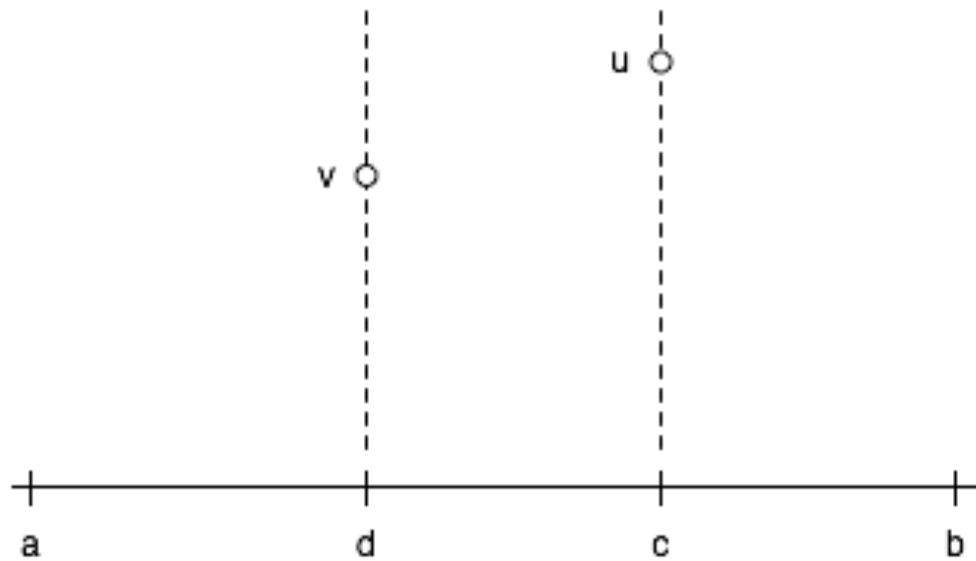
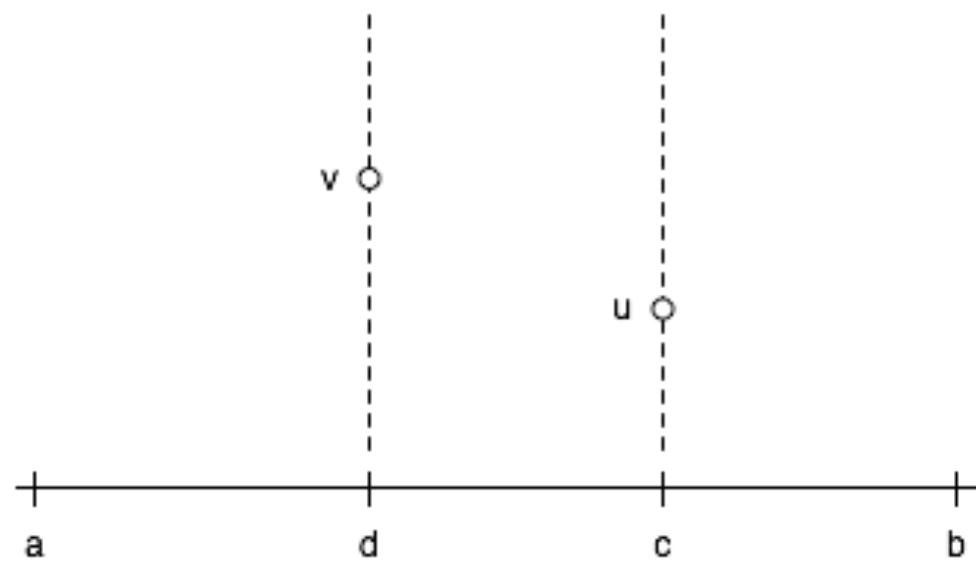


Figure 3.1.b: Update Bracket Interval for Minimand for v greater than u



$$c - a = b - d , \quad (3.14)$$

that is, the interval width for case (1) equals the interval width for case (2). That is, we get a consistent reduction factor. Additionally,

$$\varphi = \frac{\varphi^2(b-a)}{\varphi(b-a)} = \frac{d-a}{b-d} \quad (3.15)$$

and

$$\varphi = -1 + \frac{1}{\varphi} = -1 + \frac{\varphi(b-a)}{\varphi^2(b-a)} = -\frac{d-a}{d-a} + \frac{\varphi(b-a)}{d-a} = \frac{-(d-a) + \varphi(b-a)}{d-a} = \frac{c-d}{d-a} \quad (3.16)$$

so that

$$\frac{c-d}{d-a} = \frac{d-a}{b-d} \quad (3.16)$$

that is, using φ to pick c and d maintains the spacing among points as the brackets are updated. Therefore, we only need to calculate one new point and make one new function evaluation per interval reduction.

3.4 Parameter Settings

Based on the balanced one-way ANOVA model, true power and estimated power were obtained for a range of population conditions and simulation outcomes, respectively, for three sizes of number of treatments, $p = 2, 3$, and 4 .

The two-treatment class has a single symmetric mean arrangement. The three-treatment class has one symmetric mean arrangement and one asymmetric mean arrangement. The four-treatment class has two symmetric mean arrangements and one asymmetric mean arrangement. For each mean arrangement a range of population standard deviations and sample sizes are considered. These population conditions are summarized in Table 3.1. Table 3.1 also includes under each mean arrangement the sum of squared differences between population means and their grand population mean which is the numerator of the noncentrality parameter λ (1.10). These values were chosen so that treatment populations should be discernibly different at a 95% empirical interval when the population standard deviation is “small” and/or sample size is “large,” and indiscernibly different when the population standard deviation is “large” and/or sample size is “small.”

Table 3.1: Summary of Parameter Settings for Simulations

Case	Mean = μ_i	$\sum(\mu_i - \bar{\mu}_*)^2$
Two-Sample Symmetric	30, 40	50.0000
Three-Sample Symmetric	30, 35, 40	50.0000
Three-Sample Asymmetric	30, 40, 40	66.6667
Four-Sample Symmetric A	30, 35, 35, 40	50.0000
Four-Sample Symmetric B	30, 30, 40, 40	100.0000
Four-Sample Asymmetric	30, 40, 40, 40	75.0000
	Standard Deviation = σ	2.5 to 10.0 by 0.5 and 20.0
	Sample Size = n	5 to 14 by 1 and 15 to 50 by 5
	Significance Level = α	0.05

3.5 Iteration

For each combination of mean arrangement, population standard deviation, and sample size, 100 simulations were produced. Each simulation produced p random samples where p denotes the number of treatment populations. The p samples were produced to have equal samples size n . Each random sample was produced from a random seed value, an integer between 1 and 1,000,000,000, obtained through the website www.random.org. Random samples were constructed to reflect a normally distributed population with mean μ_i and equal population standard deviation where $i = 1, \dots, p$. Note that random samples were initially generated in part by the SAS® software function RANNOR. RANNOR was seen to produce dependent samples when called multiple times within the IML procedure. Therefore, RANNOR was encapsulated in a macro which resulted in independent random number generation.

For each simulation, the following calculations were made from the generated data. An ANOVA table was calculated as outlined in 3.1.1. True power and estimated power were calculated as outlined in 3.1.2. Confidence limits for true power were calculated and optimized as outlined in 3.1.2 and 3.1.3, respectively. Tolerance for bracket intervals in 3.1.3 was set to 1E-4 as suggested by the procedure FMINBND in MATLAB (2009) and the package OPTIMIZE in R (R Development Core Team 2009).

Several problems were noted with the SAS software function PROBF which was recommended by Thomas (1997) for calculating power (SAS Institute Inc. 2006a). First, the SAS function PROBF reports an invalid argument when the supplied noncentrality parameter is large with respect to numerator degrees of freedom, denominator degrees of freedom, and F critical value. This problem is noted by O'Brien (1988). Taylor and Muller (1996) suggest setting the noncentrality large enough such that PROBF evaluates to zero or one where

appropriate. The SAS software function CDF (SAS Institute Inc. 2006b) offers some relief to the PROBF issue though invalid arguments can still be had for extreme parameter values.

Second, PROBF reports an error when the supplied noncentrality parameter is negative. Gerard et al. (1998) suggest setting the noncentrality parameter to zero. For some random random seed values the data produced result with a negative adjusted estimated noncentrality parameter. The negative value occurs in some instances when adjusting the estimated noncentrality parameter (2.5), as noted previously.

Finally, FNONCT reports an error when either of the probabilities $1 - \alpha_L$ or α_U are greater than the probability associated with the observed F statistic from a central F -distribution with degrees of freedom DFT and DFE. Thomas (1997) suggests setting the noncentrality parameter equal to zero. In this case the error associated with FNONCT is noted in the SAS software documentation.

For each combination of mean arrangement, population standard deviation, and sample size, calculations for the 100 simulations are averaged. Notable averages include average estimated power and average upper and lower confidence limits about true power. The median for estimated power was also determined. Missing values were deleted.

The SAS program (Version 9.1.3) used can be found in Appendix B. Flow charts for the SAS program can be found in Appendix A (arranged through OmniGraffle Pro® version 5). This document was processed using the iWork '09® application Pages. Mathematical expression were set using MathType® version 6.

4 Results

Output for simulation summaries is plotted as power versus sample size for population standard deviations 2.5 to 10.0 by 0.5 and 20.0. Power ranges from 0.0 to 1.0 and sample size ranges from 5 to 14 by 1 and 15 to 50 by 5. The average p-value of the test is presented in each plot. The complete display of simulation plots can be found in Appendix C. In the Results, four plots are presented from each treatment class for each treatment mean arrangement, power plotted against sample size, where sample size ranges from 1 to 35 at population standard deviations of 2.5, 5.0, 10.0, and 20.0. A maximum sample size of 35 is presented since power approached 1.0 for sample sizes over 35 for the simulation configuration used here. Population standard deviations 2.5 and 20.0 represent two extreme cases for variability while population standard deviations 5.0 and 10.0 represent two medium cases for variability. Plots for each treatment arrangement appear as a moving-window like landscape from a moving vehicle.

4.1 Two-Sample Symmetric Case

Plots for population standard deviations 2.5, 5.0, 10.0, and 20.0 with a maximum sample size of 35 are seen in Figure 4.1.a, 4.1.b, 4.1.c, and 4.1.d, respectively. The p-value for the test decreases as the sample size increases, such that the null hypothesis can be rejected when $n \geq 5$, $n \geq 6$, $n \geq 14$, and $n > 35$ for respective values of $\sigma = 2.5$, $\sigma = 5.0$, $\sigma = 10.0$, and $\sigma = 20.0$. True power is always between the average lower and upper confidence limits for true power. Power and confidence limits are higher and closer together at larger sample sizes, as expected. For $\sigma = 2.5$, true power is very close to the average upper limit. For $\sigma = 5.0$, true power is closer to the average upper limit than the lower limit for $n < 30$; is half-way between the average upper and lower limits when both limits and true power approach 1.0 for $n \geq 30$. For $\sigma = 10.0$, true power starts (at $n = 5$) around the average lower limit, and is located half-way between average confidence limits at $n = 9$. For $n > 9$, true power is closer to the average upper limit than the average lower limit. When $\sigma = 20.0$, true power is located near the average lower limit (for $n = 5$), and obtains an approximate location half-way between average limits for $n = 30$. True power exhibits an increasing positive trend toward the average upper limit for each plot, though the incline of the curve decreases for larger values of σ .

Power curves appear generally ordered as,

$$\widehat{power}_{true} > \widehat{power}_{med} > \widehat{power}_{avg}. \quad (4.1)$$

The general ordering of power curves is strictly seen for σ of 2.5 and 5.0. For $\sigma = 10.0$, average estimated power starts (at $n = 5$) above median estimated power but adopts the general ordering for sample sizes 10 and greater. When σ equals 20.0, average estimated power starts (at $n = 5$) above true power, and median estimated power starts below true power. Average estimated power appears (for $\sigma = 20.0$) below true power at sample sizes 13 and greater, and below median estimated power at sample sizes 35 and greater. For samples sizes greater than 35, the general ordering appears to hold for any value of σ presented here.

Two-Sample Symmetric Case: Power vs Sample size

Figure 4.1.a: Sigma 2.5

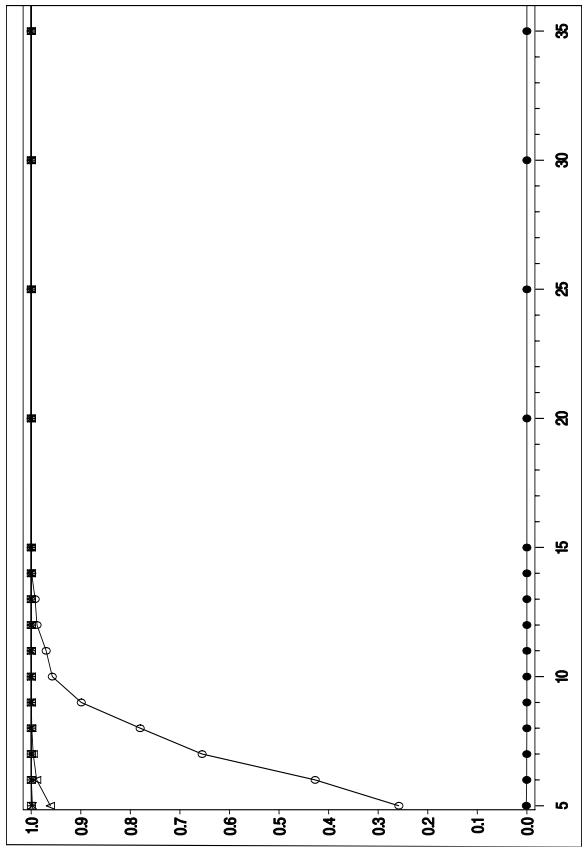


Figure 4.1.b: Sigma 5.0

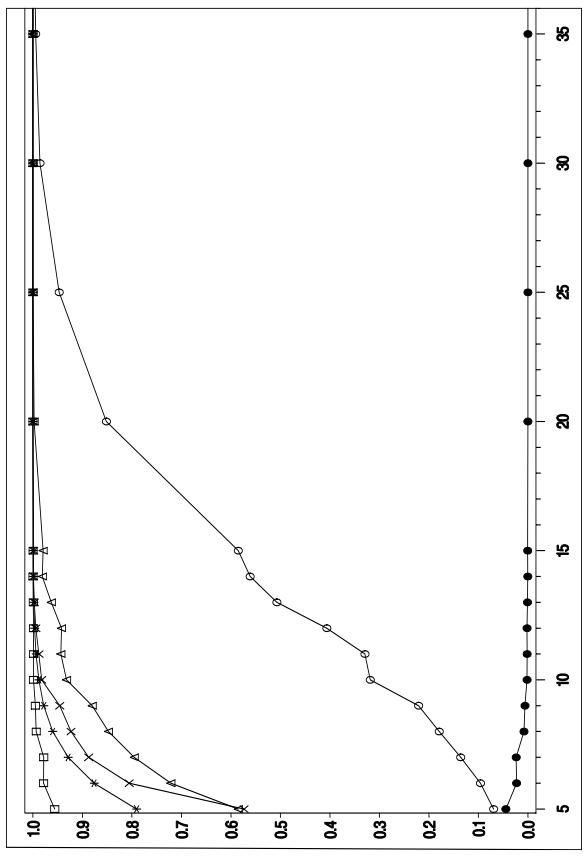


Figure 4.1.c: Sigma 10.0

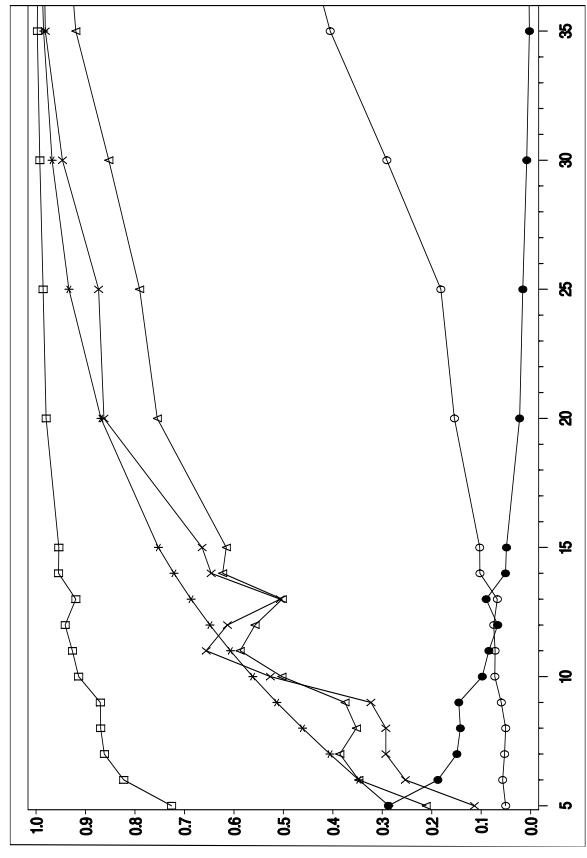
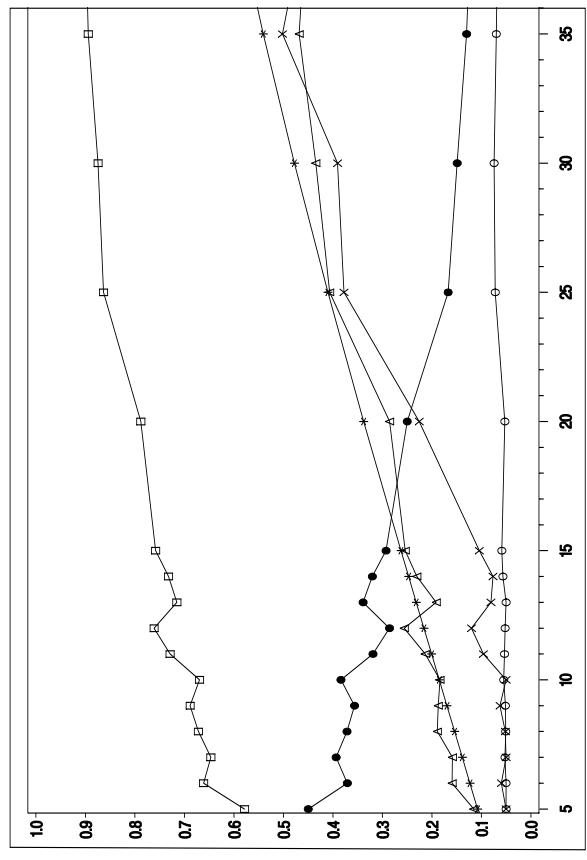


Figure 4.1.d: Sigma 20.0



□ ◻ Avg Upper 95% Estimated Power	*** True Power
△ ▲ Avg Estimated Power	× × Median Estimated Power
○ ○ Avg P-Value	● ● Avg Power

4.2 Three-Sample Symmetric Case

Plots for population standard deviations 2.5, 5.0, 10.0, and 20.0 with a maximum sample size of 35 are seen in Figure 4.2.a, 4.2.b, 4.2.c, and 4.2.d, respectively. Similarly to the two-sample case, the p-value for the test decreases as the sample size increases, such that the null hypothesis can be rejected when $n \geq 5$, $n \geq 8$, $n \geq 16$, and $n > 35$ for values of $\sigma = 2.5$, $\sigma = 5.0$, $\sigma = 10.0$, and $\sigma = 20.0$, respectively. True power is always contained within the average limits. As with the two-sample case, power and confidence limits tend to be higher and closer together for larger sample sizes. True power is very close to the average upper limit for $\sigma = 2.5$. For $\sigma = 5.0$, true power is located near the upper average limit and is approximately half-way between the average limits for $n \geq 35$. When $\sigma = 10.0$, true power is close to the average lower limit (at $n = 5$), and is approximately half-way between the confidence limits at $n = 12$. For $\sigma = 20.0$, true power is nearer to the average lower limit than the average upper limit for sample sizes up to 30, but for $n = 35$ true power is approximately half-way between the confidence limits. At all values σ considered here, true power tend toward the average upper limit.

As with the two-sample case, the general ordering (4.1) tends to hold. The general ordering of power is seen for $\sigma = 2.5$ and 5.0. For $\sigma = 10.0$, average estimated power starts (at $n = 5$) above the median estimated power, but takes on the general ordering at sample sizes 13 and greater. For $\sigma = 20.0$, true power begins (at $n = 5$) below average estimated power, and above median estimated power. Average estimated power (for $\sigma = 20.0$) is below true power at sample sizes 30 and greater, and below median estimated power at sample sizes greater than 35. The general position of power curves (4.1) tends to hold when samples sizes are large.

Three-Sample Symmetric Case: Power vs Sample size

Figure 4.2.a: Sigma 2.5

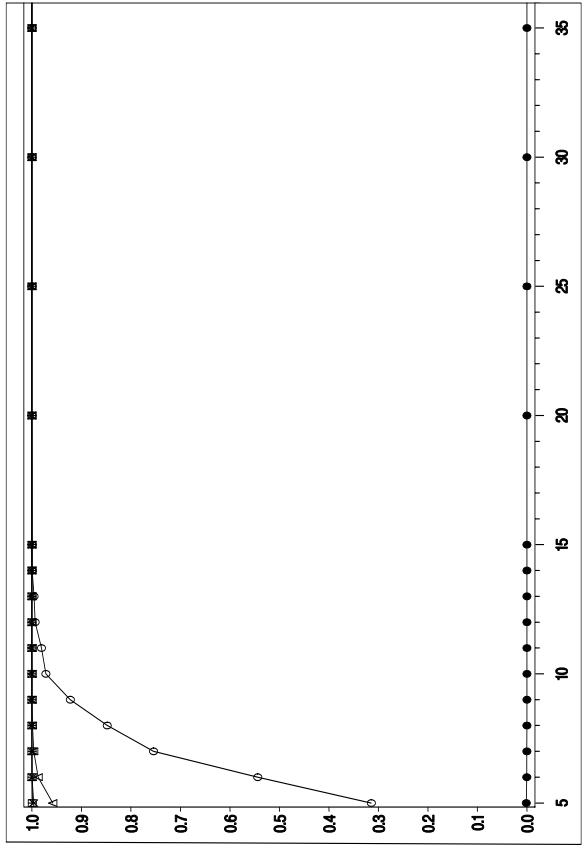


Figure 4.2.b: Sigma 5.0

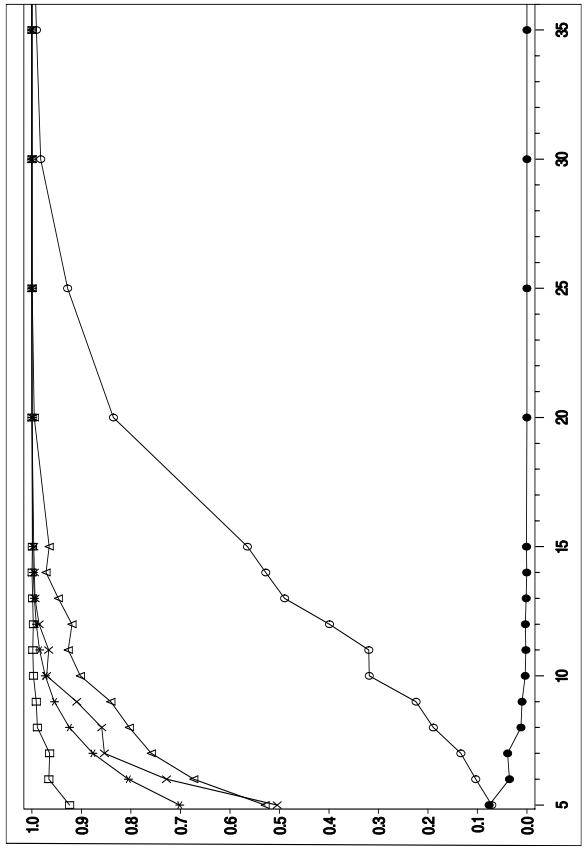


Figure 4.2.c: Sigma 10.0

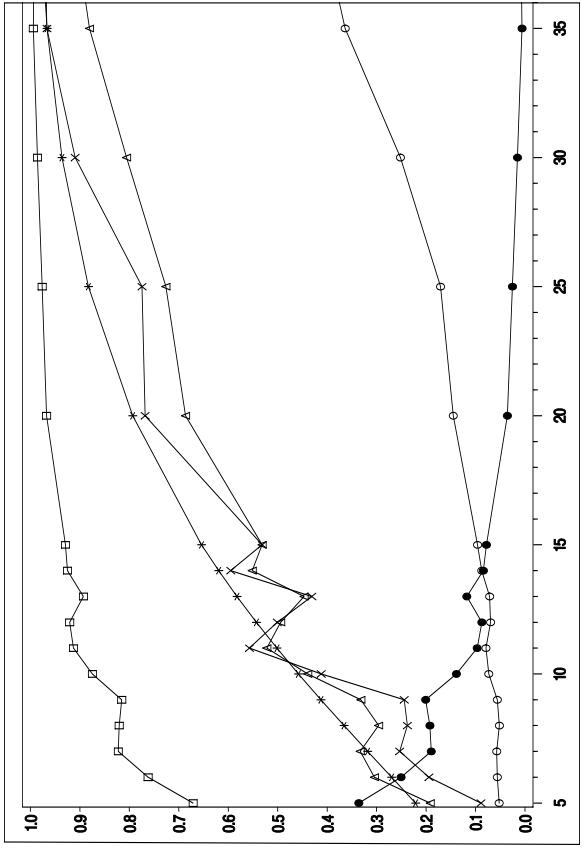
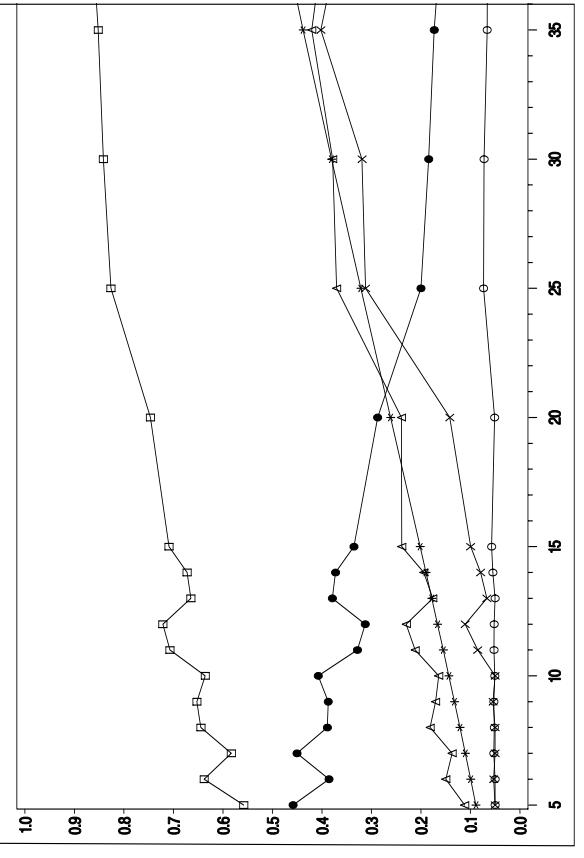


Figure 4.2.d: Sigma 20.0



$\square \square \square$	Avg Upper 95% Estimated Power	$\ast \ast \ast$	True Power
$\triangle \triangle \triangle$	Avg Estimated Power	$\times \times \times$	Median Estimated Power
$\bullet \bullet \bullet$	Avg P-Value		

4.3 Three-Sample Asymmetric Case

Plots for population standard deviations 2.5, 5.0, 10.0, and 20.0 with a maximum sample size of 35 are seen in Figure 4.3.a, 4.3.b, 4.3.c, and 4.3.d, respectively. Again, the p-value for the test decreases as the sample size increases, such that the null hypothesis can be rejected when $n \geq 5$, $n \geq 6$, $n \geq 14$, and $n > 35$ for respective values of $\sigma = 2.5$, $\sigma = 5.0$, $\sigma = 10.0$, and $\sigma = 20.0$. True power is between the average upper and lower limits for all values of the population standard deviation considered here. As with the previous cases, power and confidence limits are higher and closer together for larger sample sizes. For $\sigma = 2.5$, true power is indistinguishable from the average upper limit. For $\sigma = 5.0$, true power is closer to the average upper limit than the average lower limit, and true power is approximately half-way between the average upper and lower limits when both limits approach 1.0 ($n \geq 30$). For $\sigma = 10.0$, the average upper and lower limits are centered around true power for $n = 9$. When $\sigma = 20.0$, true power is located near the average lower limit (for $n = 5$), and is approximately half-way between the average upper and lower limits at $n = 30$. As with previous cases, true power shows an upward trend toward the average upper limit for larger sample sizes.

As with prior cases, the general ordering of true power, median estimated power, and average estimated power (4.1) tends to hold. For $\sigma = 2.5$ and 5.0, the general ordering of power curves can be seen in Figure 4.3.a. For $\sigma = 10.0$, average estimated power starts (at $n = 5$) above the median estimated power, but adopts the general ordering for sample sizes larger than 10. When $\sigma = 20.0$, average estimated power starts (at $n = 5$) above true power, and median estimated power starts (at $n = 5$) below true power. Average estimated power appears below true power for sample sizes 15 and greater, and below median estimated power for sample sizes greater than 35. As with the previous cases, the general order of the power curves (4.1) is maintained for large sample sizes.

Three-Sample Asymmetric Case: Power vs Sample size

Figure 4.3.a: Sigma 2.5

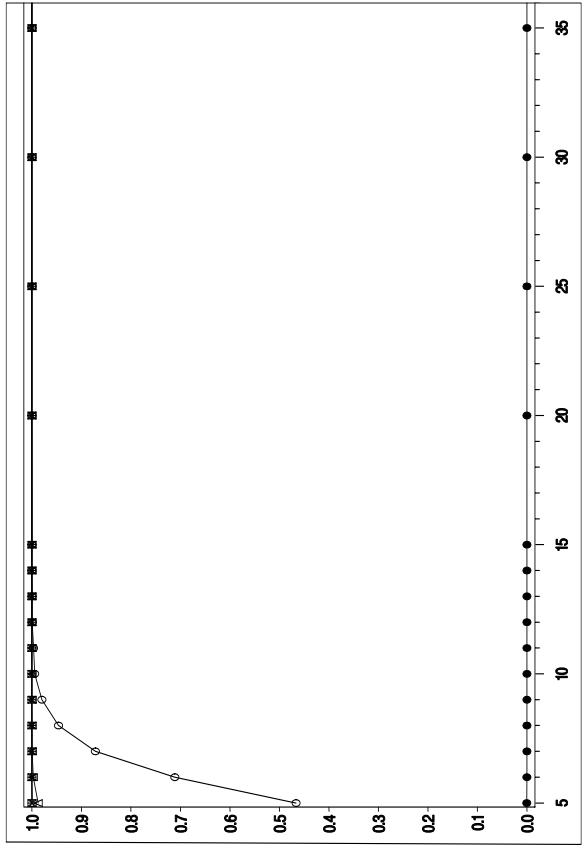


Figure 4.3.c: Sigma 10.0

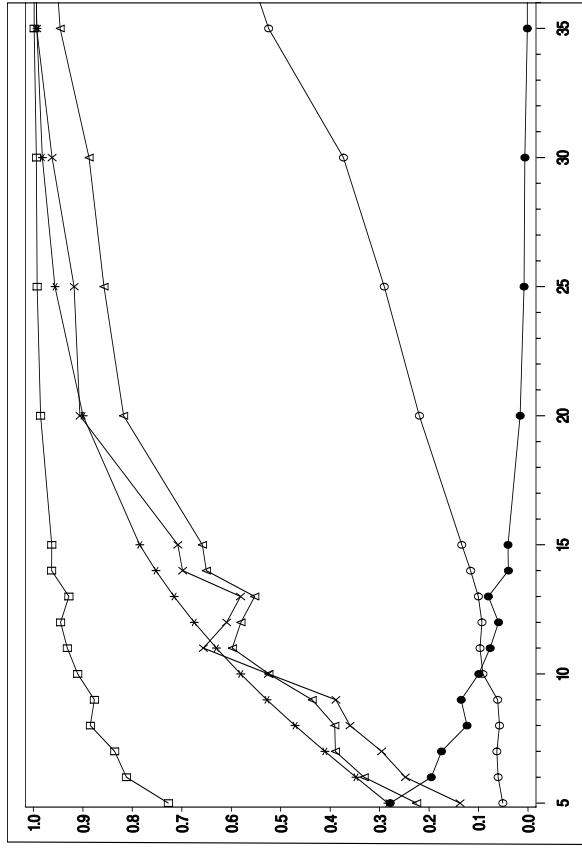


Figure 4.3.b: Sigma 5.0

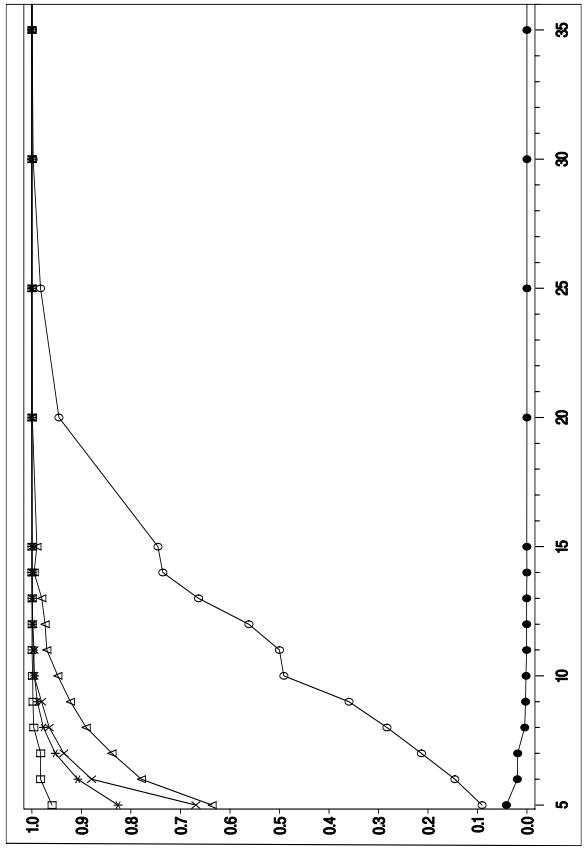
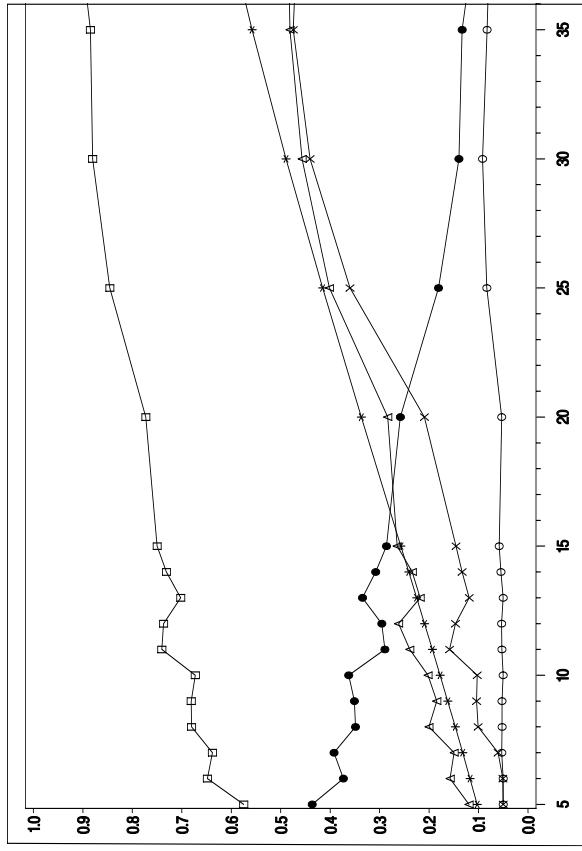


Figure 4.3.d: Sigma 20.0



Avg Lower 95% Estimated Power
● ● Avg P-Value

4.4 Four-Sample Symmetric Case A

Plots for population standard deviations 2.5, 5.0, 10.0, and 20.0 with a maximum sample size of 35 are seen in Figure 4.4.a, 4.4.b, 4.4.c, and 4.4.d, respectively. For larger sample sizes, the p-value for the test decreases such that the null hypothesis can be rejected when $n \geq 5$, $n \geq 8$, $n \geq 17$, and $n > 35$ for respective values of $\sigma = 2.5$, $\sigma = 5.0$, $\sigma = 10.0$, and $\sigma = 20.0$. True power, as seen in previous cases, is always between the average upper and lower limits for true power. Further, true power and average confidence limits tend toward 1.0 for large sample sizes. For $\sigma = 2.5$, true power is identical with the average upper confidence limit. For $\sigma = 5.0$, average confidence limits center are symmetric about true power when both limits equal 1.0. For $\sigma = 10.0$, average upper and lower confidence limits are centered around true power when $n = 14$. For $\sigma = 20.0$, true power is close to the average lower limit. True power displays a positive increasing trend toward the average upper limit, where the steepness of the curve reduces for larger values of σ .

The general order of the power curves (4.1) again tends to occur. The typical ordering always occurs for $\sigma = 2.5$ and 5.0. For $\sigma = 10.0$, median estimated power starts (at $n = 5$) below the median estimated power, but maintains the typical ordering at sample sizes 14 and greater. For $\sigma = 20.0$ and $n = 5$, power curves are ordered as average estimated power, true power, and median estimated power. The average estimated power (for $\sigma = 20.0$) appears below true power at sample sizes 15 and greater, and below median estimated power at sample sizes larger than 35. As with previous cases, the typical order of power curves (4.1) is seen for larger sample sizes.

Four-Sample Symmetric Case A: Power vs Sample size

Figure 4.4.a: Sigma 2.5

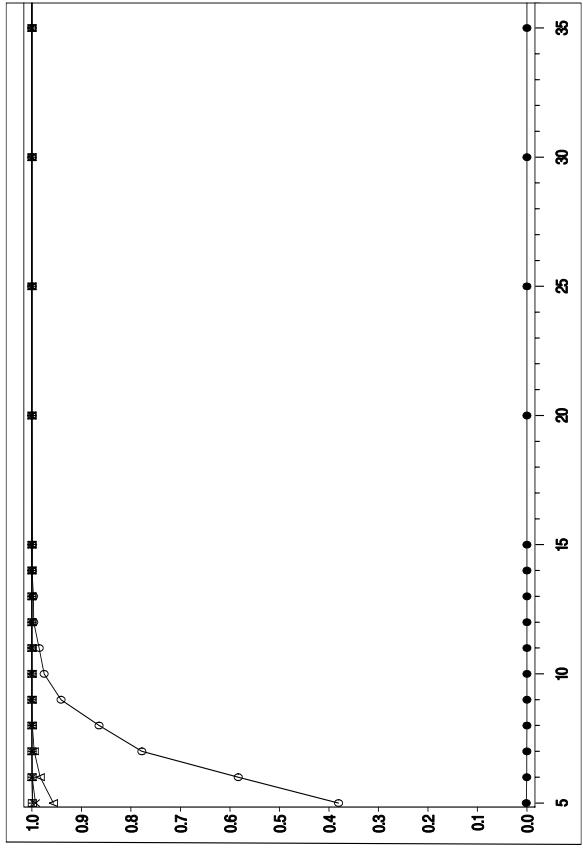


Figure 4.4.b: Sigma 5.0

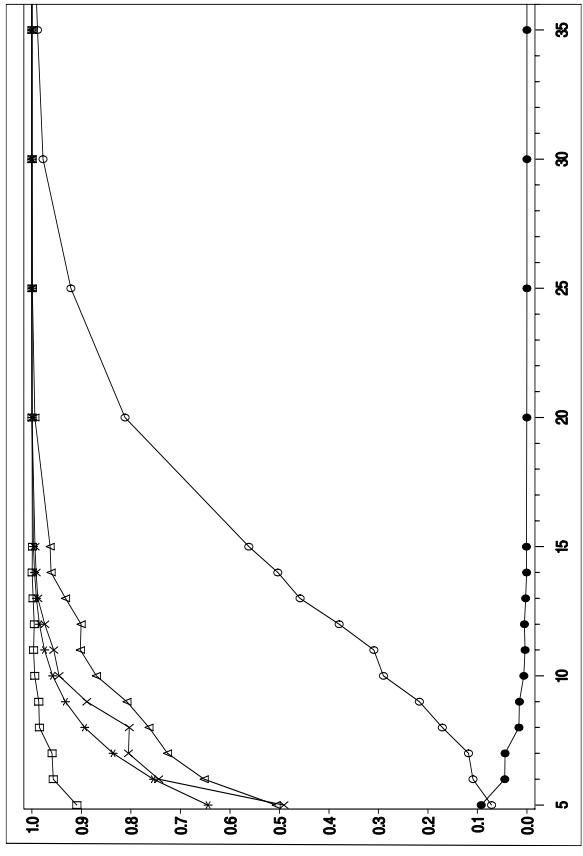


Figure 4.4.c: Sigma 10.0

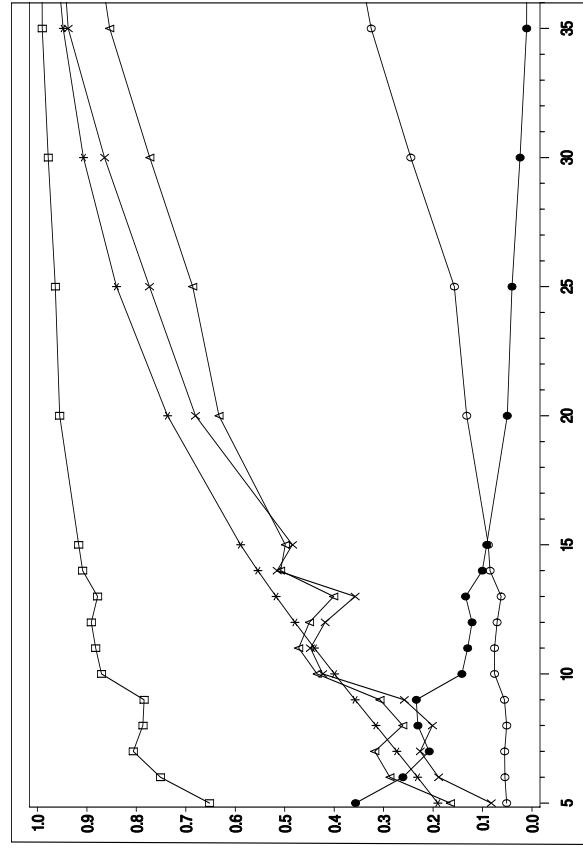
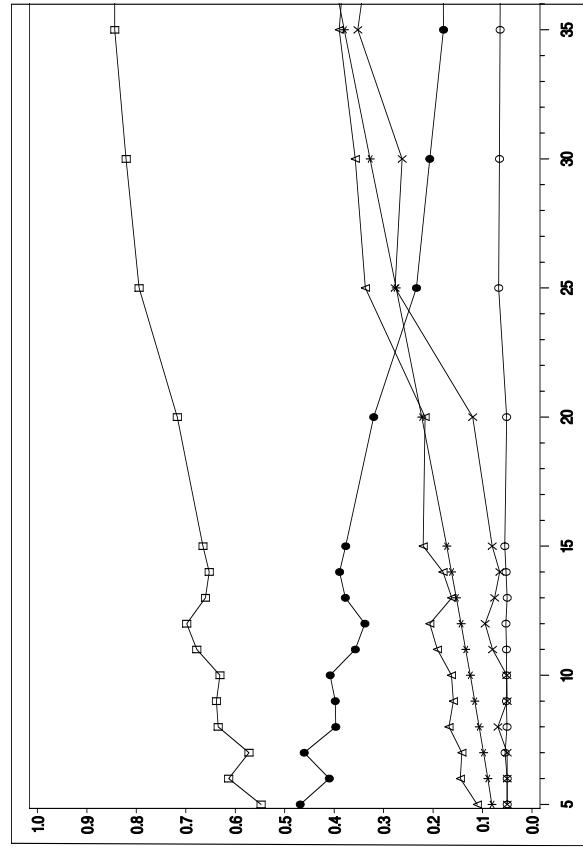


Figure 4.4.d: Sigma 20.0



$\square \circ \square$ Avg Upper 95% Estimated Power	$\ast \ast \ast$ True Power
$\triangle \triangle \triangle$ Avg Estimated Power	$\times \times \times$ Median Estimated Power

4.5 Four-Sample Symmetric Case B

Plots for population standard deviations 2.5, 5.0, 10.0, and 20.0 with a maximum sample size of 35 are seen in Figure 4.5.a, 4.5.b, 4.5.c, and 4.5.d, respectively. As the sample size increases, the p-value for the test decreases such that the null hypothesis can be rejected when $n \geq 5$, $n \geq 5$, $n \geq 10$, and $n > 35$ for $\sigma = 2.5$, $\sigma = 5.0$, $\sigma = 10.0$, and $\sigma = 20.0$, respectively. As with previous cases, average lower and upper confidence limits always contain true power. True power and confidence limits become narrower and approach 1.0 for larger sample sizes. For $\sigma = 2.5$, true power is identical to the average upper limit. For $\sigma = 5.0$, true power is just below the average upper limit and is centered between average limits when both limits approach 1.0. For $\sigma = 10.0$, true power begins (at $n = 5$) around the average lower limit, and centers between the average confidence limits for $n = 8$. When $\sigma = 20.0$, average confidence limits are symmetric around true power for $n = 25$. As with prior cases, true power has an increasing positive trend toward the average upper limit for each plot. Notably, the rate of increase of the true power curve decreases for larger values of σ .

As seen previously, average estimated power, median estimated power, and true power are consistently between the lower and upper confidence limits for true power. Power curves have the typical order (4.1) as in previous cases. The usual ordering of power is seen for $\sigma = 2.5$ and 5.0. For $\sigma = 10.0$, average estimated power starts (at $n = 5$) above the median estimated power, yet takes on the general ordering at sample sizes 10 and greater. For $\sigma = 20.0$, true power is below average estimated power and above median estimated power for $n = 5$. Further, for $\sigma = 20.0$, average estimated power appears below true power at sample sizes 13 and greater, and below median estimated power at sample sizes 30 and greater. As seen earlier, the general ordering (4.1) holds for large sample sizes.

Four-Sample Symmetric Case B: Power vs Sample size

Figure 4.5.a: Sigma 2.5

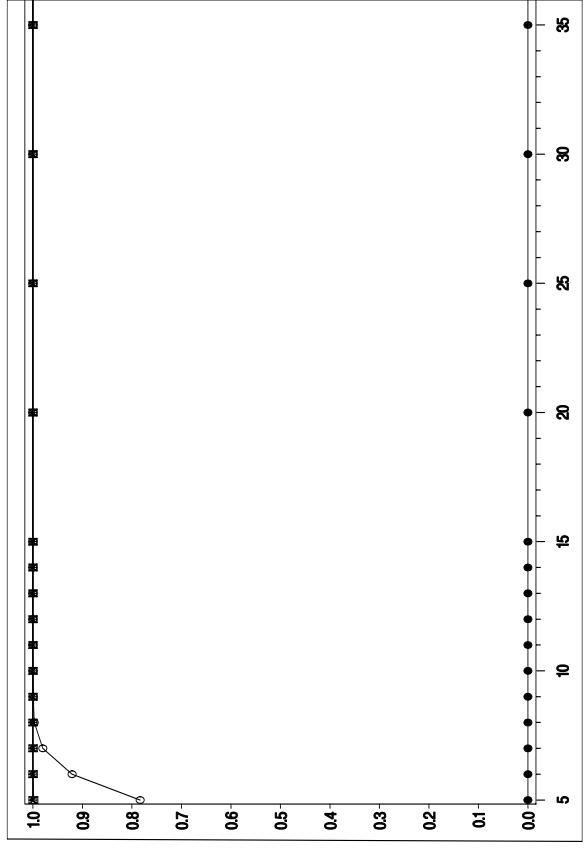


Figure 4.5.b: Sigma 5.0

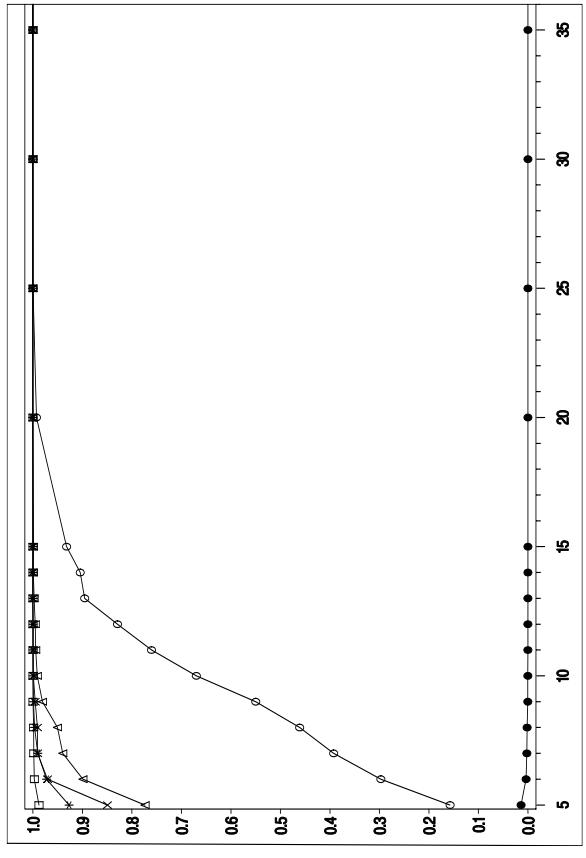


Figure 4.5.c: Sigma 10.0

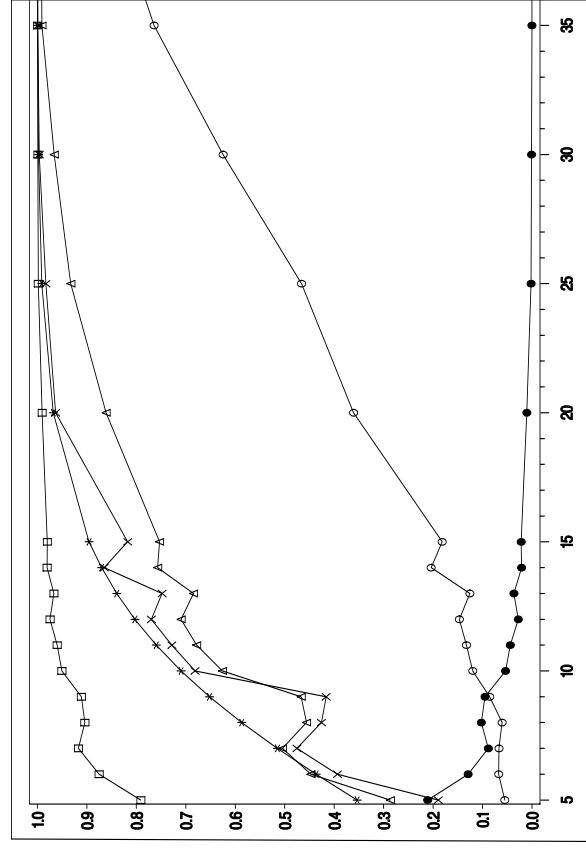
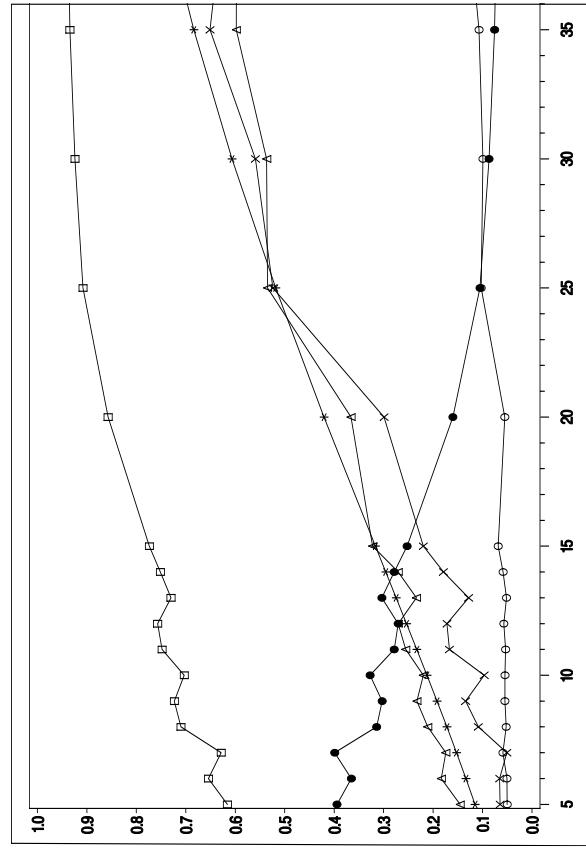


Figure 4.5.d: Sigma 20.0



□○□ AVG Upper 95% Estimated Power	*** True Power
△△△ AVG Lower 95% Estimated Power	××× Median Estimated Power
□●□ AVG Estimated Power	●●● AVG P-Value

4.6 Four-Sample Asymmetric Case

Plots for population standard deviations 2.5, 5.0, 10.0, and 20.0 with a maximum sample size of 35 are seen in Figure 4.6.a, 4.6.b, 4.6.c, and 4.6.d, respectively. The p-value for the test decreases as the sample size increases, as expected, such that the null hypothesis can be rejected when $n \geq 5$, $n \geq 6$, $n \geq 14$, and $n > 35$ for respective values of $\sigma = 2.5$, $\sigma = 5.0$, $\sigma = 10.0$, and $\sigma = 20.0$. True power lies between the average lower and upper limits for all values of σ . As seen in prior cases, true power and average confidence limits are higher and closer together for large sample sizes. For $\sigma = 2.5$, true power is indistinct from the average upper limit. For $\sigma = 5.0$, true power is close to the average upper limit and is approximately very close (at power approximately 1.0) with the average upper limit at $n = 10$. For $\sigma = 10.0$, average confidence limits are equally spaced about true power when $n = 9$. When $\sigma = 20.0$, average upper and lower limits are centered around true power for $n = 30$. As usual, true power shows an increasing upward trend toward the average upper confidence limit for each plot. Notably, the incline of the curve for true power climbs with less rapidity for larger values of σ .

The power curves are hold the usual ordering (4.1) as seen previously. The general ranking of power curves (4.1) is exemplified for $\sigma = 2.5$ and 5.0. For $\sigma = 10.0$, average estimated power starts (at $n = 5$) above the median estimated power but takes on the general ordering at sample sizes 9 and greater. For $\sigma = 20.0$, average estimated power starts above true power, while median estimated power starts below true power. Average estimated power (for $\sigma = 20.0$) appears below true power at sample sizes 20 and greater, and below median estimated power at sample sizes greater than 35. When sample sizes are large, the power curves follows the general ordering (4.1).

Four-Sample Asymmetric Case: Power vs Sample size

Figure 4.6.a: Sigma 2.5

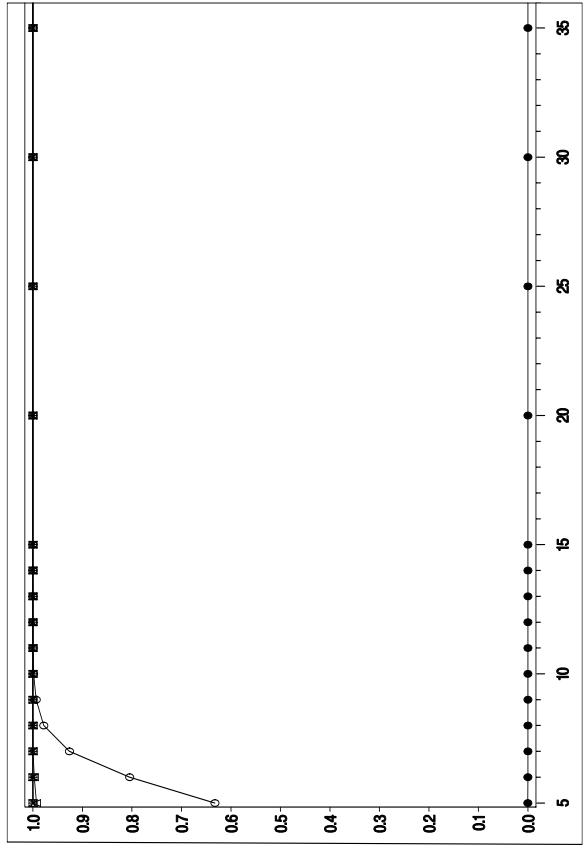


Figure 4.6.b: Sigma 5.0

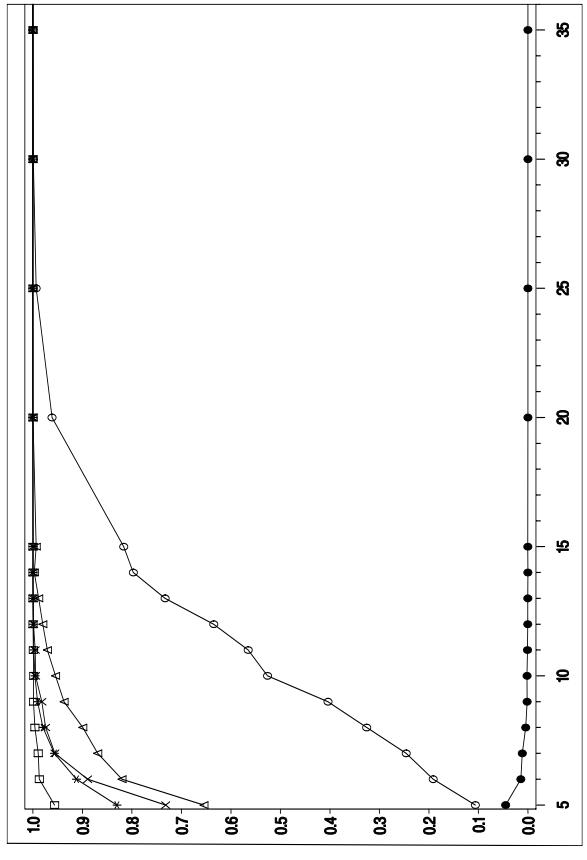


Figure 4.6.c: Sigma 10.0

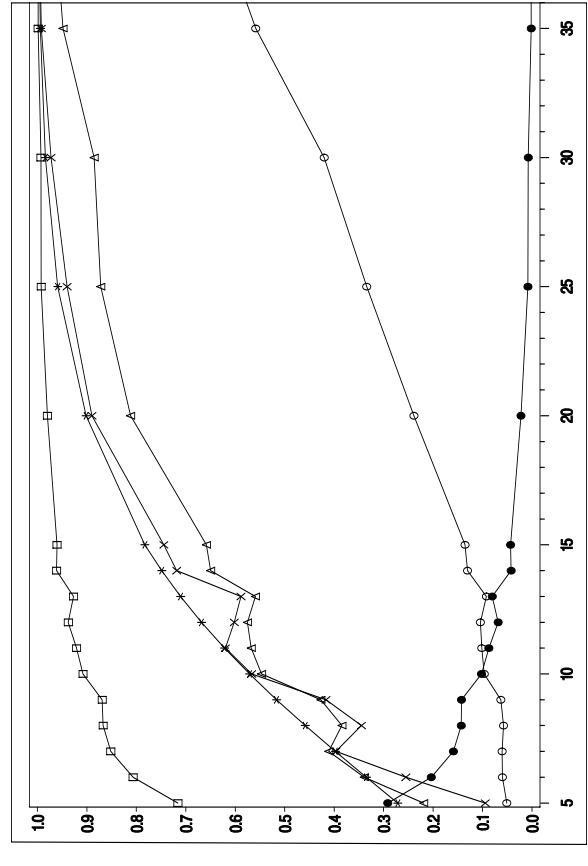
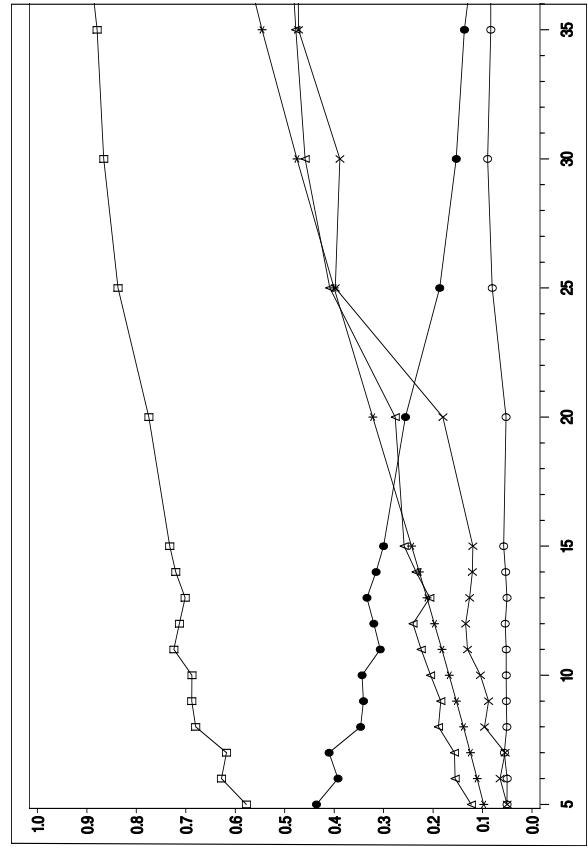


Figure 4.6.d: Sigma 20.0



□—○—○ AVG Upper 95% Estimated Power	***—*—* True Power
△—△—△ AVG Estimated Power	×—×—× Median Estimated Power
■—●—● AVG P-Value	

4.7 Symmetric Cases

For symmetric treatment arrangements, confidence limits for the two-sample case, three-sample case, and four-sample case A do not appear substantially different. As an exception, confidence limits for the four-sample case B are narrower than the other symmetric cases. Average lower and upper confidence limit widths are ordered, from least to greatest, as four-sample case B, four-sample case A, three-sample case, and two-sample case. Average lower and upper confidence limits always contain true power.

For all symmetric cases, true power shows a positive increasing trend toward the average upper limit. For $\sigma = 2.5$ and 5.0 , all symmetric cases show true power near the average upper limit. For $\sigma = 10.0$, true power starts (at $n = 5$) close to the average lower limit, reaches a half-way distance between average limits for $n = 5$ to 15 , and then proceeds toward the average upper limit. For $\sigma = 20.0$, true power is generally close to the average lower limit for all symmetric cases, though the two-sample case and four-sample case B reach a half-way distance between average limits at $n = 30$ and $n = 25$, respectively.

4.8 Asymmetric Cases

For the two asymmetric cases, confidence interval widths are noticeably different from one another. The four-sample case has narrower average lower and upper confidence limits for true power than the three-sample case. True power is always between the average lower and upper confidence limits.

For both asymmetric cases, true power exhibits an upward trend toward the average upper limit. For $\sigma = 2.5$ and 5.0 , true power is never half-way between the average upper and lower limits. For $\sigma = 10.0$, true power starts (at $n = 5$) around the average lower limit, reaches the half-way distance between average limits around $n = 8$ to 9 , and then moves closer to the

average upper limit. For $\sigma = 20.0$, true power is close to the average lower limit, but reaches the half-way distance between average limits around $n = 30$ to 35.

4.9 Symmetric Cases versus Asymmetric Cases

In both symmetric and asymmetric cases, true power is always between the average lower and upper limits. Generally, symmetric cases appear to have wider average confidence limits than the asymmetric cases, with the exception of the four-sample case B, which has the narrowest confidence limits among all cases considered. Symmetric and asymmetric cases both show that true power is closer to the average upper limit than the lower limit for large sample sizes. In terms of average limit widths, cases are ordered from least to greatest as follows: four-sample case B, four-sample asymmetric case, three-sample asymmetric case, four-sample case A, three-sample symmetric case, then the two-sample case.

Both symmetric cases and asymmetric cases have power curves typically ordered, from highest to lowest, as true power, median estimated power, then average estimated power. The typically ordering of power curves is seen for $\sigma = 2.5$ and 5.0. For $\sigma = 10.0$, average estimated power starts (at $n = 5$) above median estimated power, but crosses below median estimated power for samples sizes between 10 to 15. For $\sigma = 20.0$, average estimated power is above true power, and median estimated power is below true power. Further (for $\sigma = 20.0$), average estimated power drops below true power, then proceeds below median estimated power as sample sizes increase. All power curves are within the average lower and upper limits.

5 Conclusion

Simulations were constructed for a range of population conditions under the balanced one-way ANOVA model. Each simulation case was iterated 100 times and plots for average estimated power, median estimated power, true power, average lower 95% limit, and average upper 95% limit were produced. An average p-value for F -tests was also plotted for each simulation setup.

When the population standard deviation was small (from $\sigma = 2.5$ to 5.0), confidence limits for true power were wider for small sample sizes and narrower for medium-to-large sample sizes. Under all sample sizes considered here, average limits contained true power. Notably, true power was never half-way between the average upper and lower confidence limits except when sample sizes were large and both limits were very close to 1.0. Additionally, true power was always close to the average upper limit and above the average estimate for power. When significant differences exist among the population treatment means, the average p-value is small, true power is large, and the average retrospective estimate for power provides a higher lower limit for true power than the average lower limit.

When the population standard deviation was large (from $\sigma = 10.0$ to 20.0), average confidence limits for true power were generally wide for small and large sample sizes. True power was contained within the average limits: close to the average lower limit for small sample sizes and progressive toward the average upper limit for large sample sizes. In all cases, the average estimate for power was present above true power (when sample sizes were small) and below true power (when sample sizes were large). When the null hypothesis of equal population means was difficult to reject, i.e. when the p-value of the test was large, true power was low and confidence limits for true power were wide.

When the null hypothesis cannot be rejected, the test performed is seen as under powered. The results here show that true power is hard to determine retrospectively as laid out

in Thomas (1997) when the null hypothesis cannot be rejected, since the average 95% confidence limits were wide and the average estimate for power fails to act as a lower bound for true power under all values of σ .

The presence of nonsignificant results can lead researchers to question the power of their test. When power has not been calculated *a priori*, researchers may see *retrospective* methods as a viable technique for “figuring out what power was.” As an example of Thomas’ technique to calculate power *post hoc*, two possible consulting situations are described below. Each uses an observed F -statistic, numerator and denominator degrees of freedom, and the desired level of significance. The p-value of the test is also provided.

For the first example, consider an observed F -statistic of 1.5 with 1 numerator degree of freedom and 40 denominator degrees of freedom with a p-value of 0.2278. An estimate for true power is

$$\widehat{\text{power}} = 1 - F(4.085 | 1, 40, 0.425) = 0.0975,$$

where $F_{0.05,1,40} = 4.085$ is the upper tail critical value from a central F -distribution with 1 numerator degree of freedom and 40 denominator degrees of freedom, and $\hat{\lambda}_{adj} = ((1.5)(1)(40 - 2)) / 40 - 1 = 0.425$ is the adjusted estimated noncentrality parameter. The minimized 95% confidence limits for true power are

$$\widehat{\text{power}}_L = 1 - F(4.085 | 1, 40, 0.00000) = 0.05 \text{ and}$$

$$\widehat{\text{power}}_U = 1 - F(4.085 | 1, 40, 8.2801704) = 0.8017,$$

where the estimated lower and upper noncentrality parameters $\hat{\lambda}_L = 0.000000$ and $\hat{\lambda}_U = 8.2801704$ are determined by the lower and upper tail probabilities $\alpha_L = 0.000048$ and

$\alpha_U = 0.049952$ through use of the Golden Section Search method (Section 3.3). Using the estimate for power, a researcher would conclude that the test was under powered since this estimated value is below 0.8 (Cohen 1988). Notably, using the point estimate to represent true power is uninformative since the confidence limits have true power anywhere from 0.05 to 0.8017. The confidence interval here is similar to the confidence limits in Figure 4.1.d for the symmetric two population mean arrangement with $\sigma = 20.0$ at a sample size of 22. Figure 4.1.d likens the clients' observations to an extreme case where discerning treatment groups as different is difficult.

For the second example, consider an observed F -statistic of 4.05 with 1 numerator degree of freedom and 40 denominator degrees of freedom with a p-value of 0.0509. Power is estimated as

$$\widehat{\text{power}} = 1 - F(4.085 | 1, 40, 2.8475) = 0.3773,$$

where $F_{0.05,1,40} = 4.085$ is the upper tail critical value from a central F -distribution with 1 numerator degree of freedom and 40 denominator degrees of freedom, and $\hat{\lambda}_{adj} = ((4.05)(1)(40 - 2)) / 40 - 1 = 2.8475$ is the adjusted estimated noncentrality parameter.

Through the Golden Section Search (Section 3.3), the minimized lower and upper tail probabilities are found to be $\alpha_L = 0.000048$ and $\alpha_U = 0.049952$ which yield lower and upper estimated noncentrality parameters $\hat{\lambda}_L = 0.000000$ and $\hat{\lambda}_U = 13.59003$. The estimated 95% lower and upper confidence limits for true power are given by

$$\widehat{\text{power}}_L = 1 - F(4.085 | 1, 40, 0.000000) = 0.05 \text{ and}$$

$$\widehat{\text{power}}_U = 1 - F(4.085 | 1, 40, 13.59003) = 0.9491.$$

Using the estimate for power, a researcher would conclude that the test was under-powered since this estimated value is below 0.8 (Cohen 1988). Notably, using the point estimate to represent true power is uninformative since the confidence limits have true power anywhere from 0.05 to 0.9491. The confidence interval here is similar to the confidence limits in Figure 4.1.d for the two-sample symmetric case with $\sigma = 20.0$ at a sample size of 8. Figure 4.5.c likens the clients observations to the a case where discerning treatment groups as different is difficult.

When determining the power of a test *post hoc*, power is not calculated but estimated. For researchers looking to resolve power after observations have been collected, confidence limits for true power can help quantify the uncertainty inherent in a *post hoc* power calculation. Examples 1 and 2, along with the plots in the results section, show a high degree of variability when the null hypothesis cannot be rejected. Hence, researchers are recommended to incorporate power analyses in the *planning* stages of their studies, as opposed to waiting until data have been collected.

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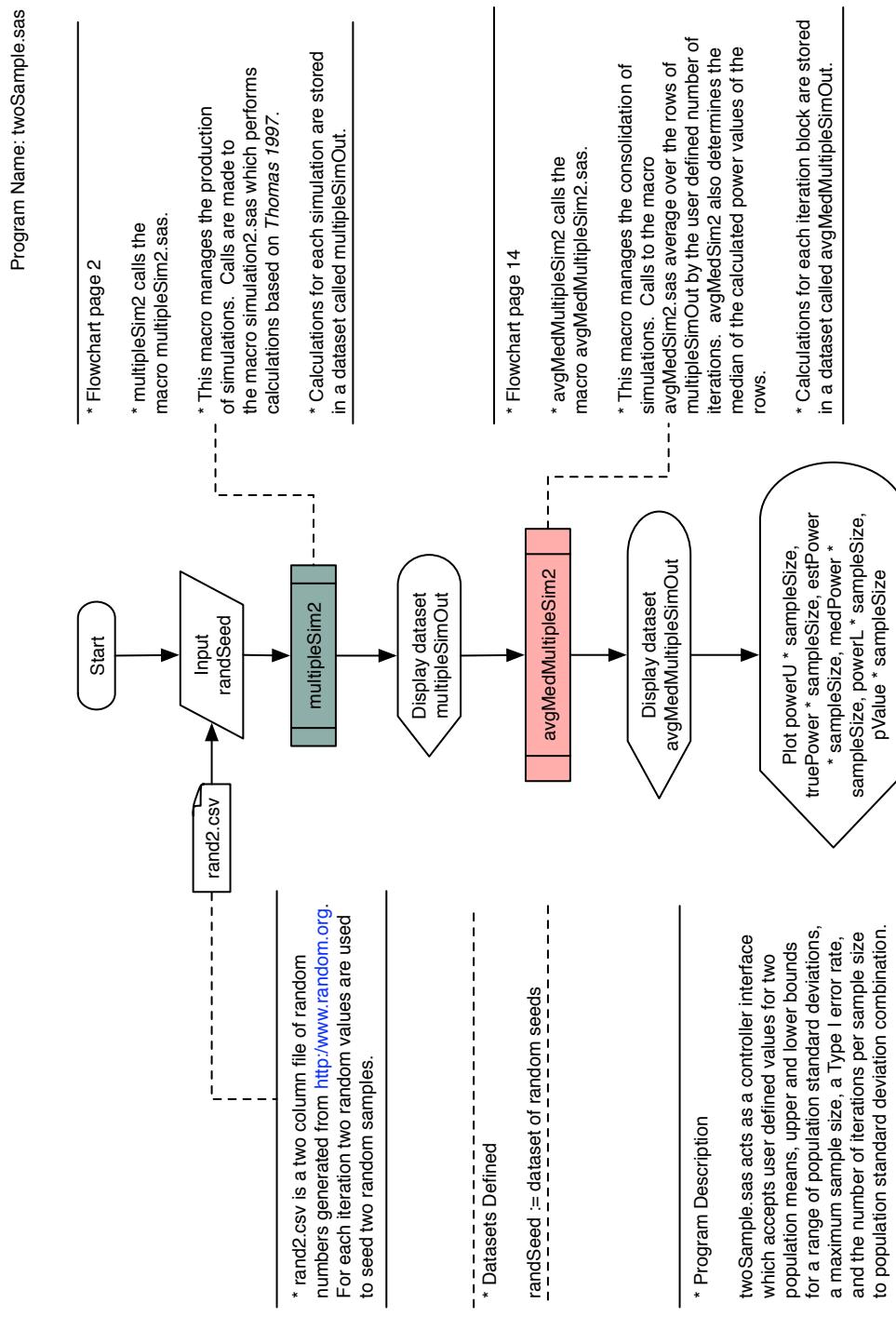
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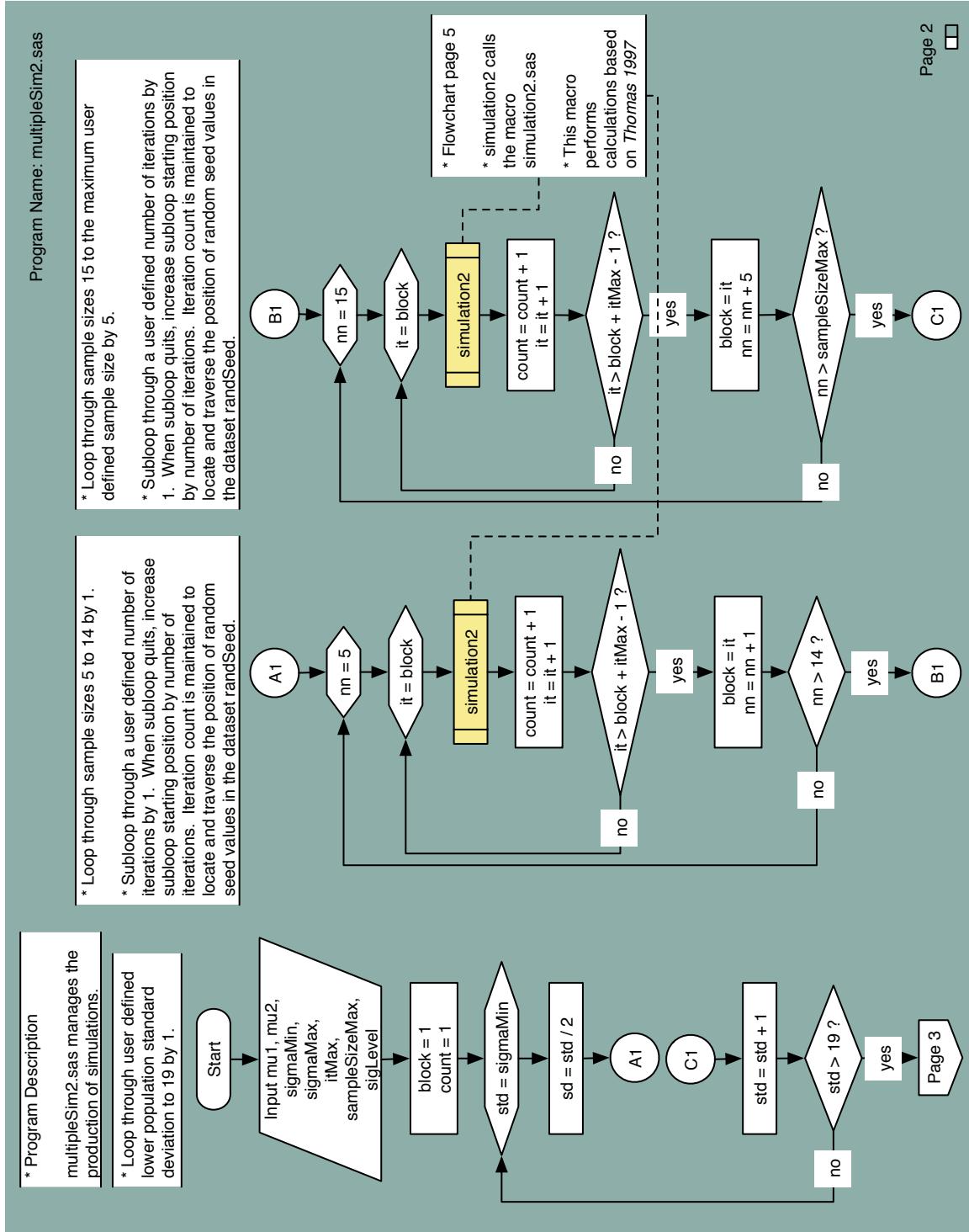
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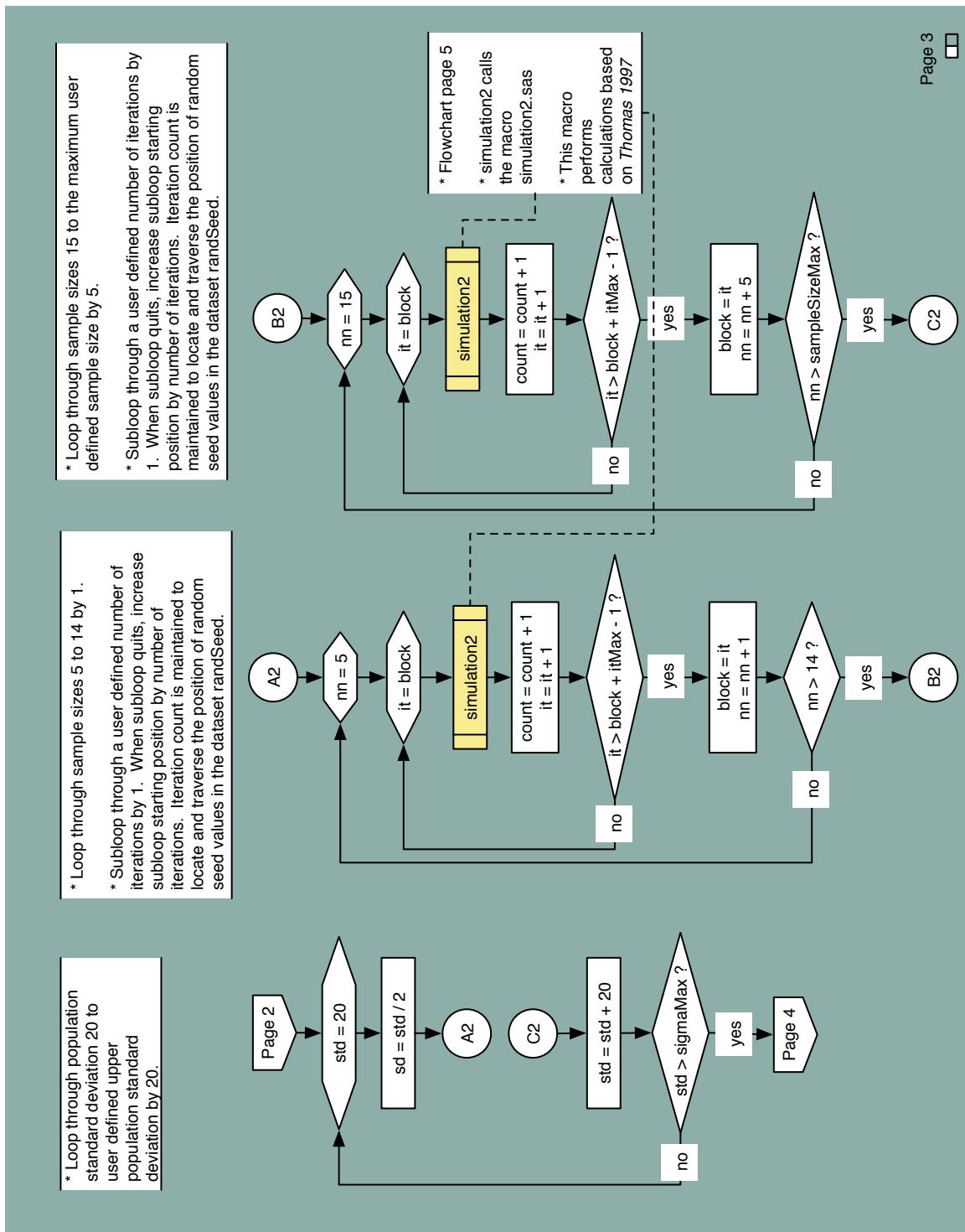
Appendix A: Two-Sample Flow Chart of Power Simulations



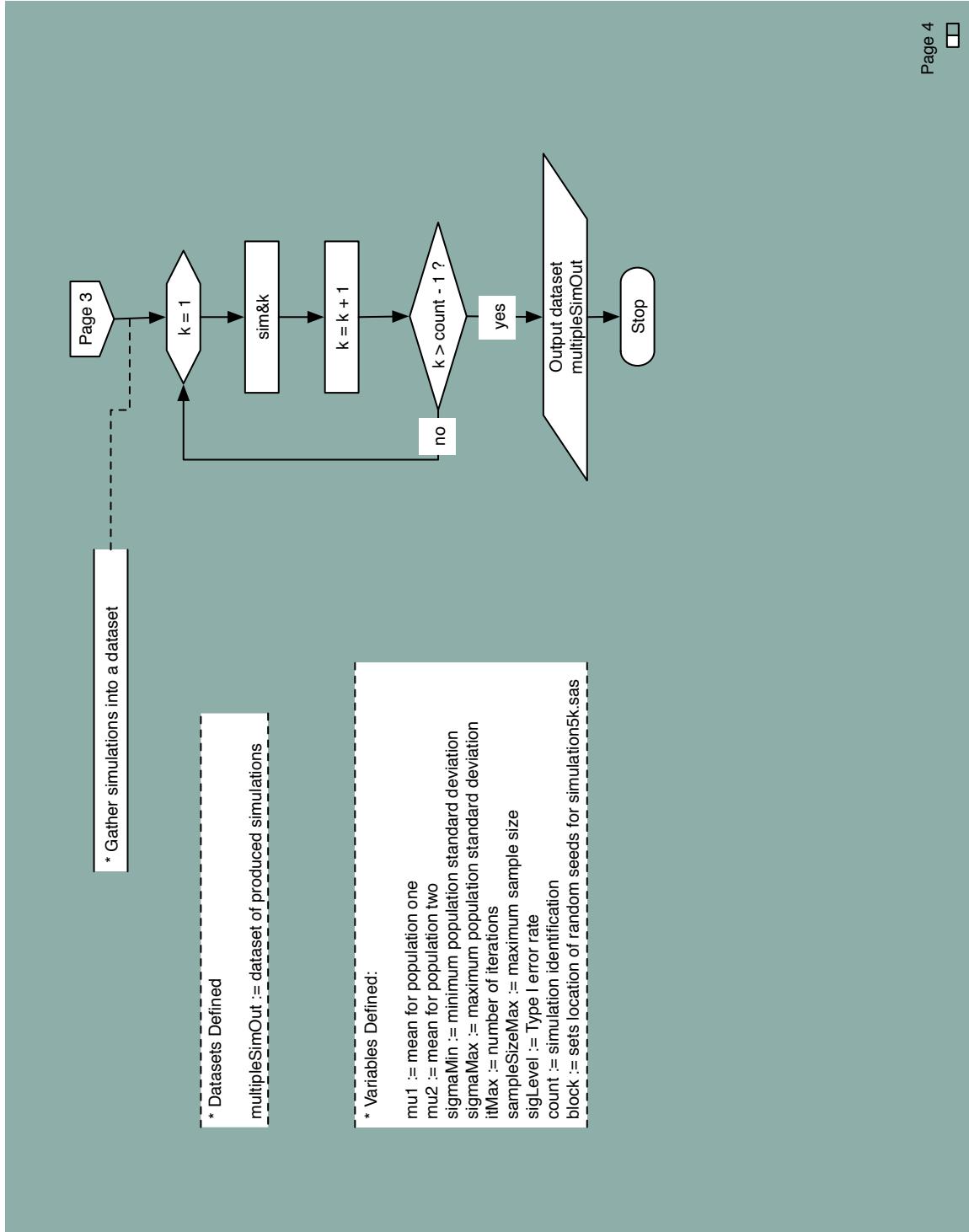
Appendix A: Two-Sample Flow Chart of Power Simulations



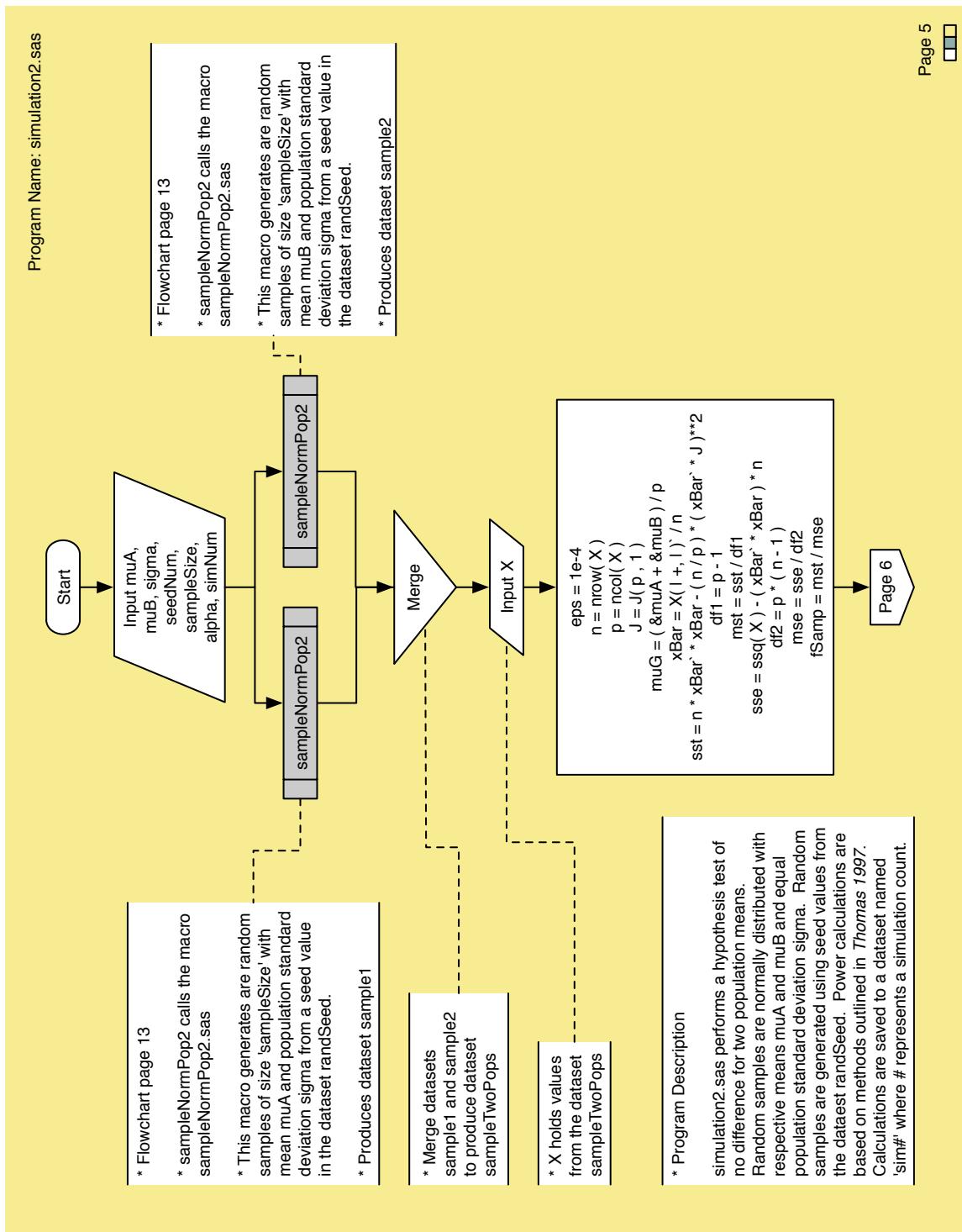
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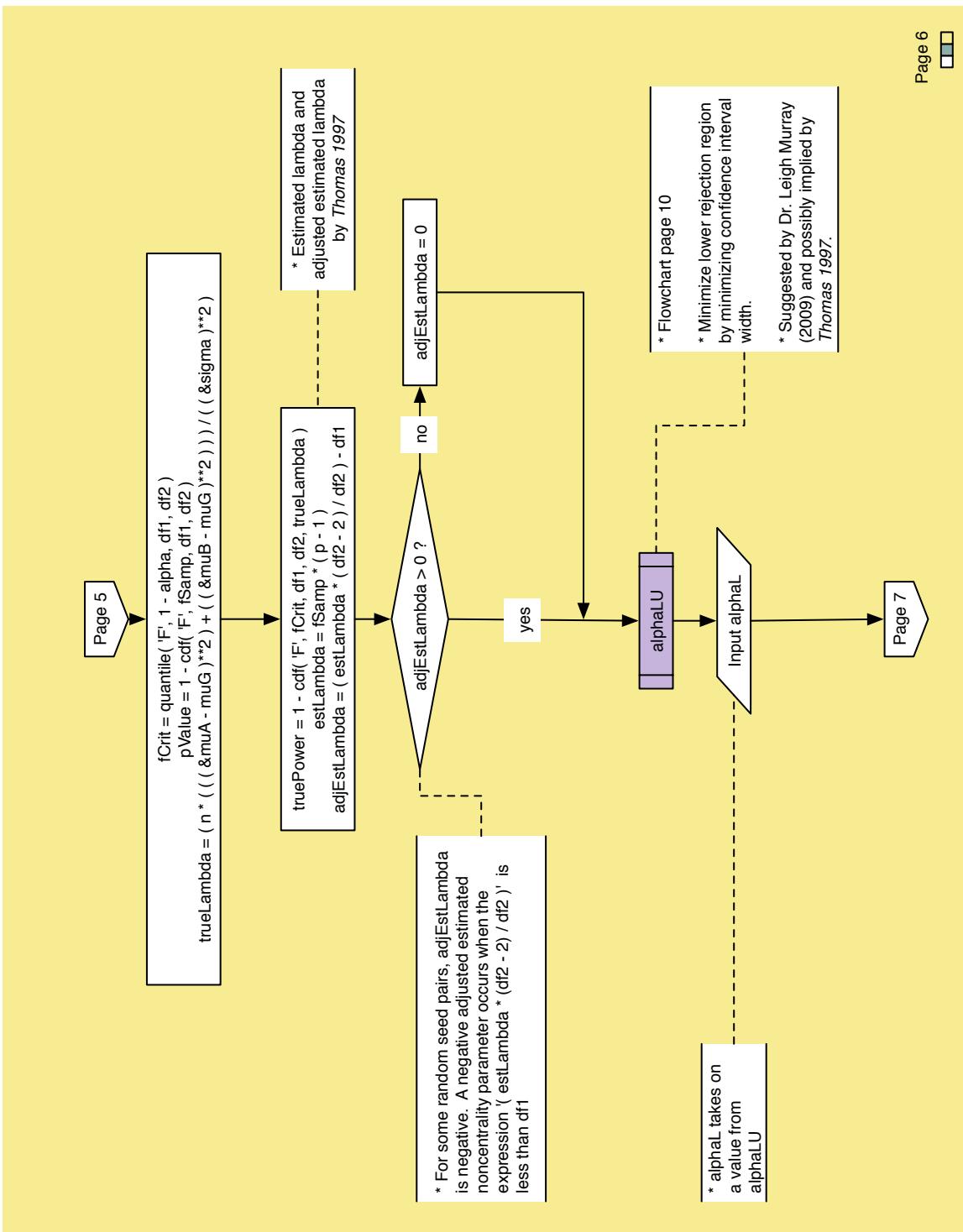
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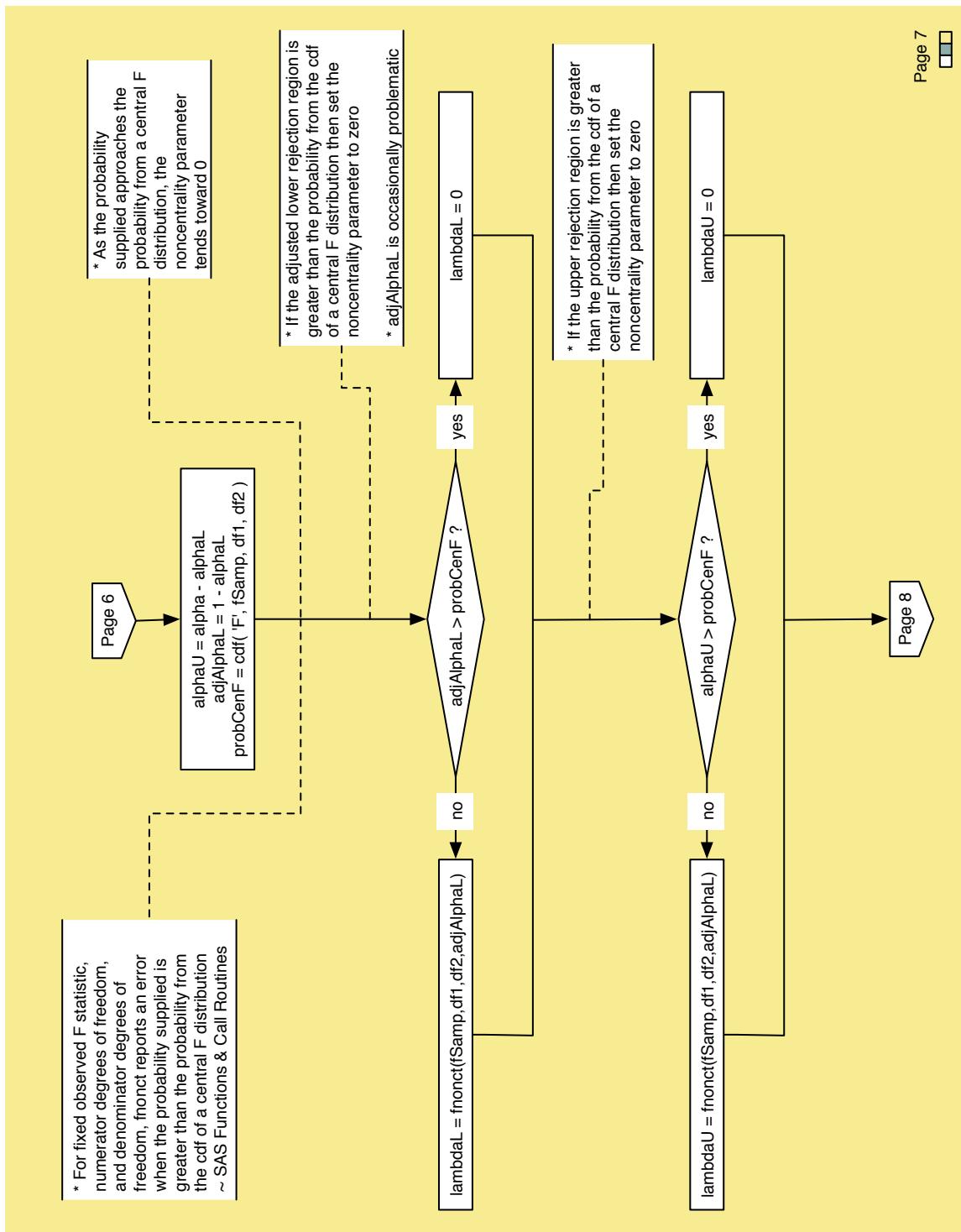
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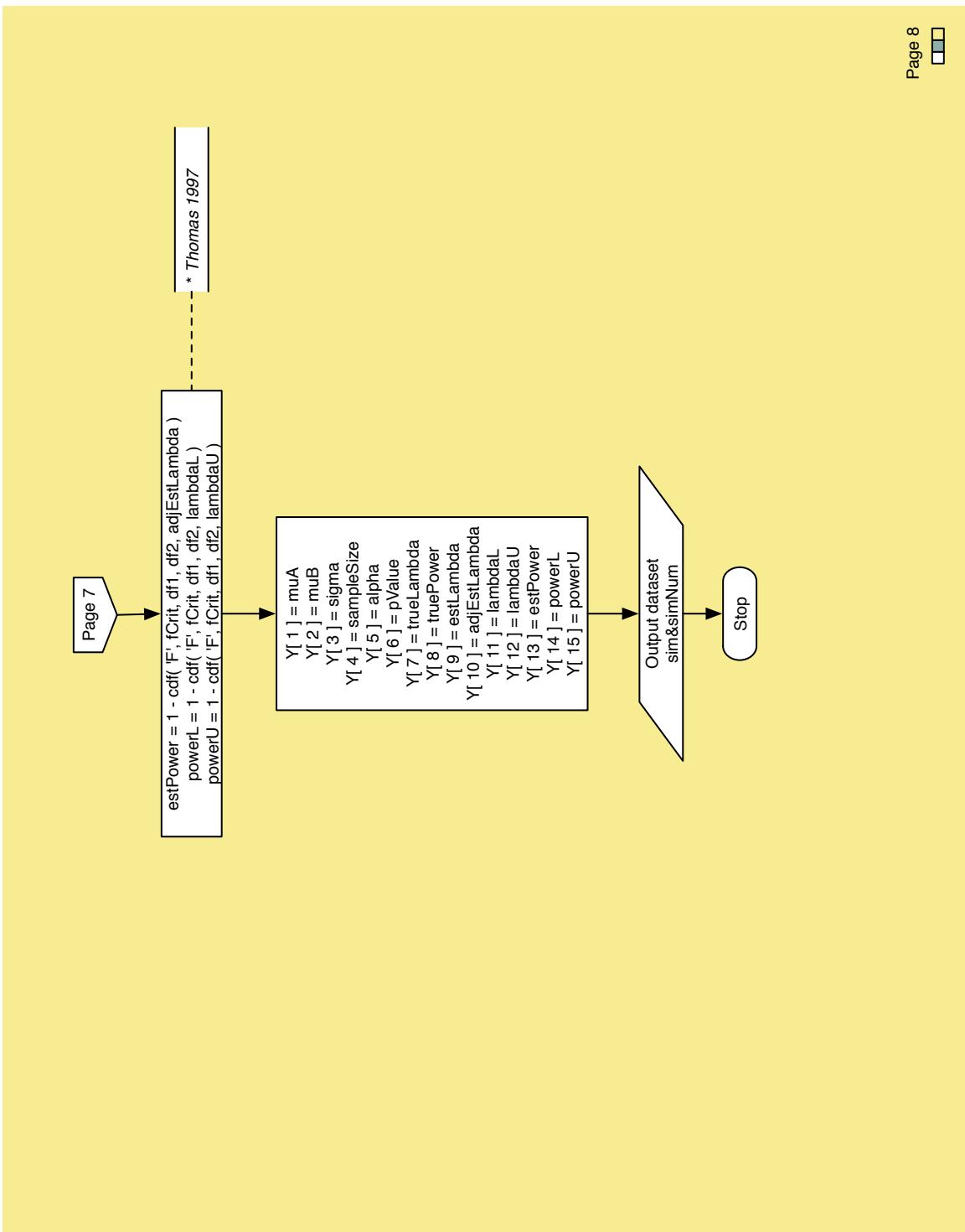
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Appendix A: Two-Sample Flow Chart of Power Simulations

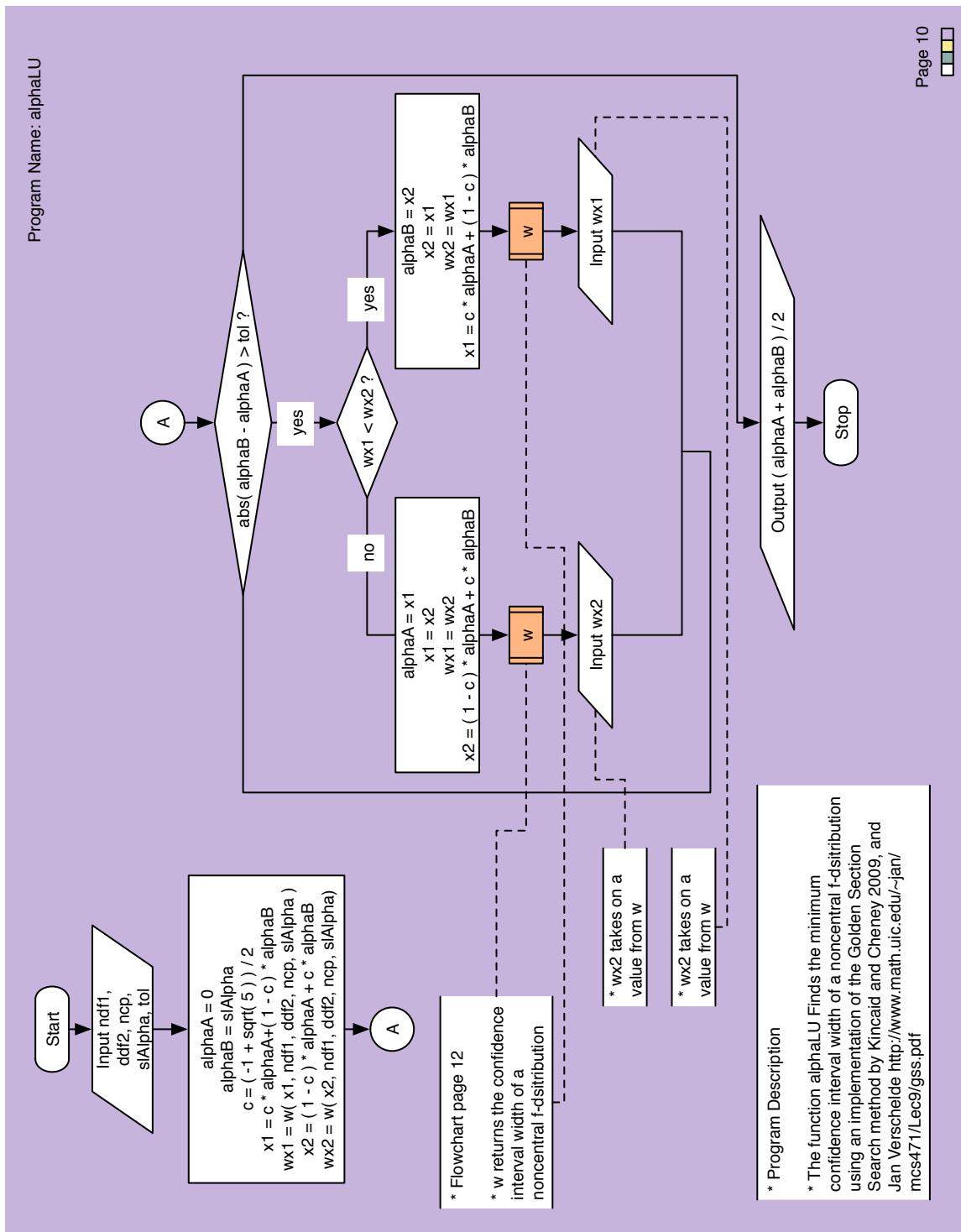


Appendix A: Two-Sample Flow Chart of Power Simulations

```
* Variables Defined
muA := mean for population one
muB := mean for population two
sigma := population standard deviation
seedNum := location of random seeds
sampleSize := sample size
alpha := level of significance
simNum := simulation identification
X := samples from normal populations
eps := tolerance for Golden Section Search methods
n := number rows
p := number columns
xBar := sample mean
sst := sum of squares for treatment
df1 := numerator degrees of freedom
mst := mean square for treatment
sse := sum of squares for error
df2 := denominator degrees of freedom
mse := mean square for error
fSamp := observed F statistic
fCrit := critical value from a central F distribution
pValue := p-value for the test
trueLambda := noncentrality parameter
truePower := power for the test
estLambda := calculated noncentrality parameter
adjEstLambda := adjusted calculated noncentrality parameter
alphaL := optimized lower critical region
alphaU := optimized upper critical region
probCenF := probability of observed F statistic of the central F distribution
estPower := calculated power
powerL := lower confidence interval for calculated power
powerU := upper confidence interval for calculated power

* Datasets Defined
sampleTwoPops := random samples dataset
sim&simNum := simulation dataset
```

Appendix A: Two-Sample Flow Chart of Power Simulations

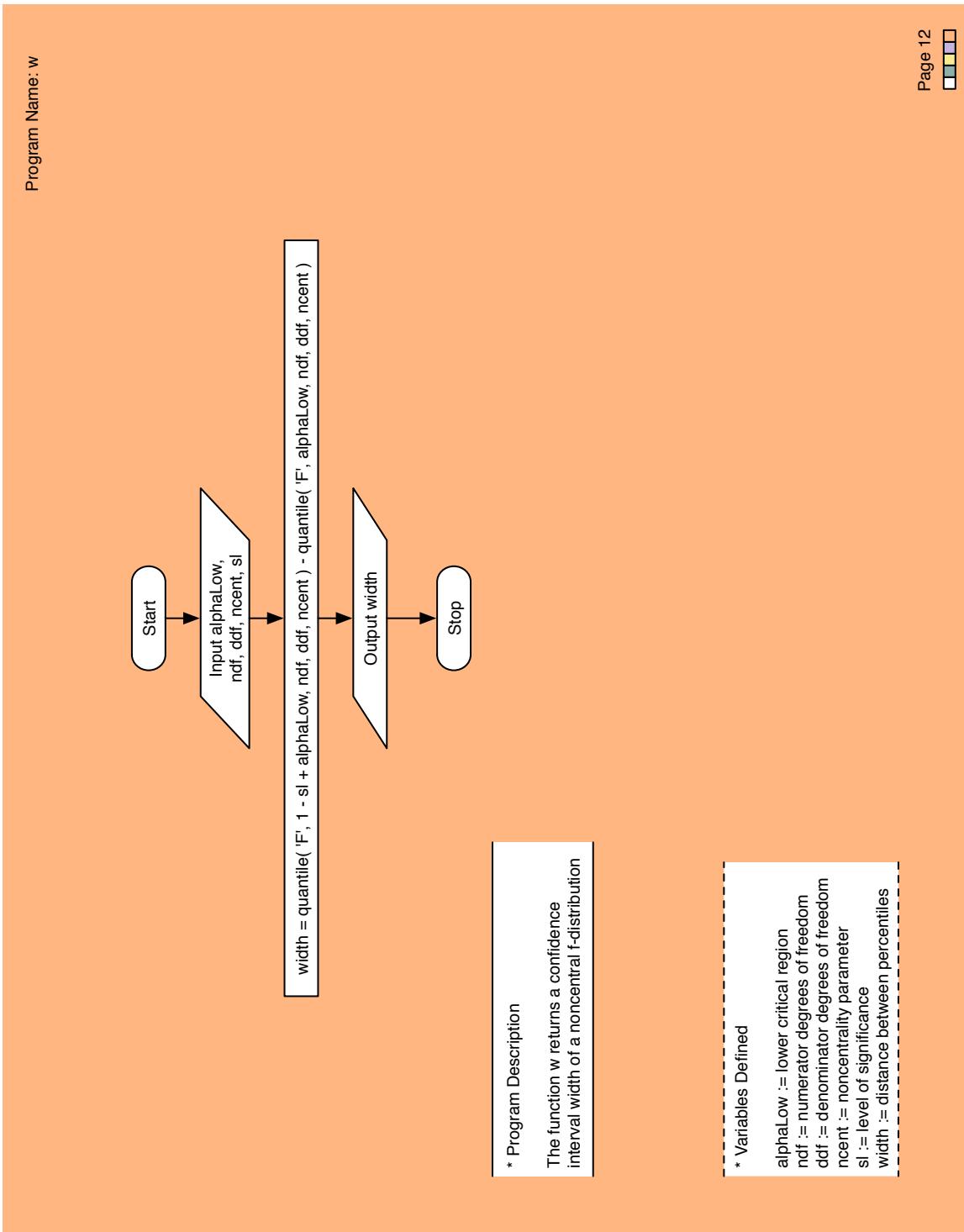


Appendix A: Two-Sample Flow Chart of Power Simulations

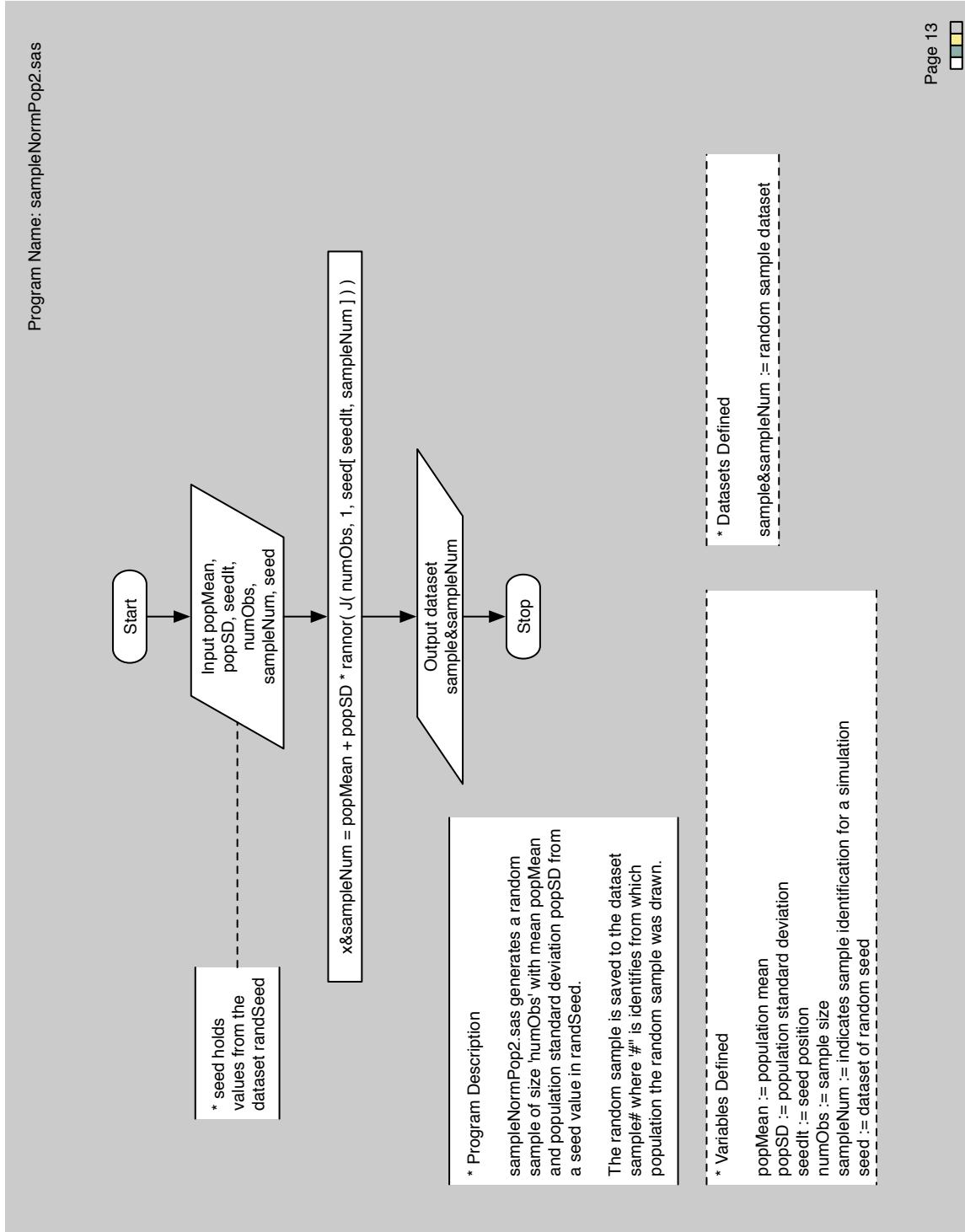
* Variables Defined

```
ndf1 := numerator degrees of freedom  
ddf2 := denominator degrees of freedom  
ncp := noncentrality parameter  
slAlpha := level of significance  
tol := tolerance for Golden Section Search method  
alphaA := lower bound for optimized lower alpha  
alphaB := upper bound for optimized lower alpha  
c := Golden ratio constant reduction factor  
x1 := percentile associated with alphaA  
wx1 := height associated with x1  
x2 := percentile associated with alphaB  
wx2 := height associated with x2
```

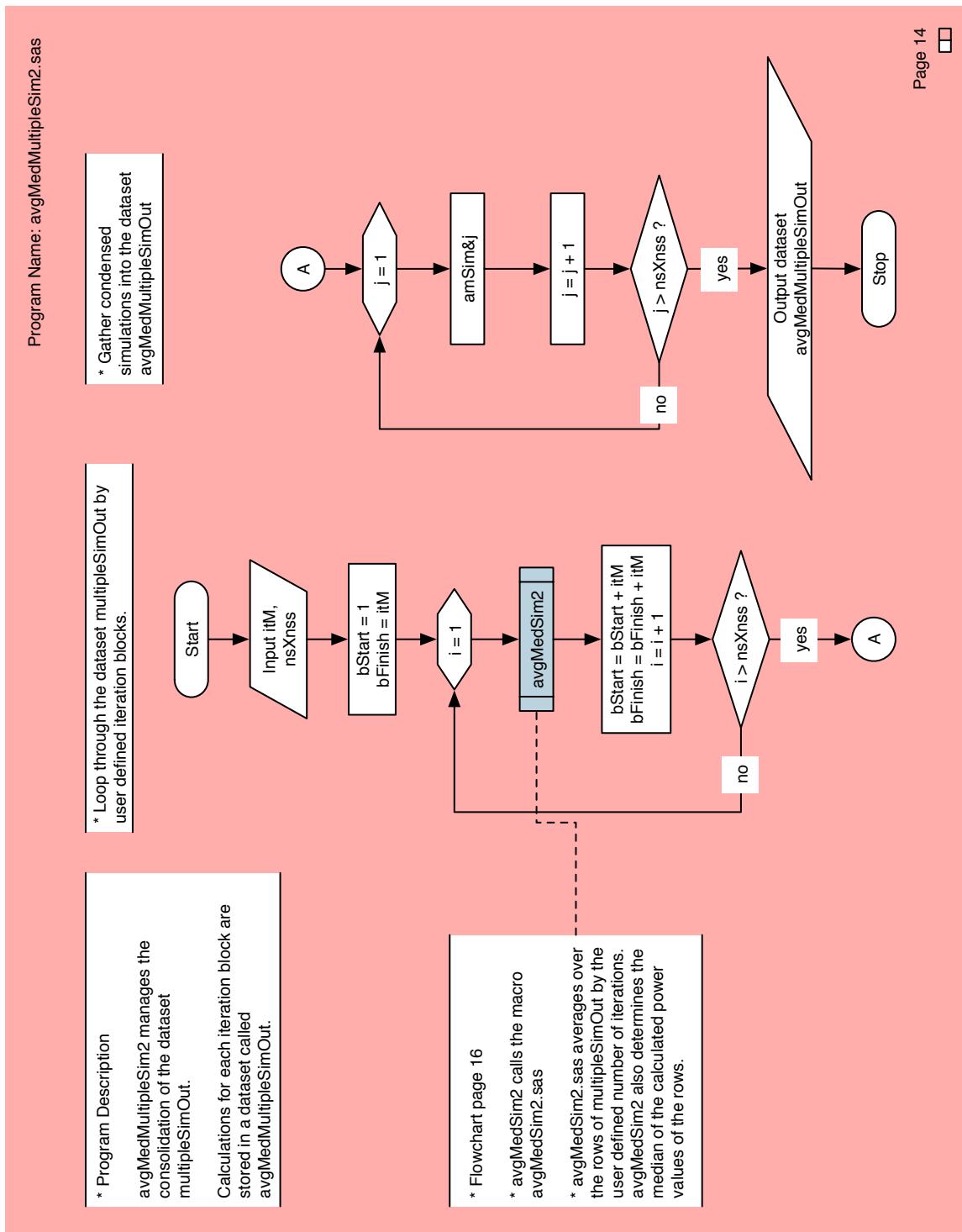
Appendix A: Two-Sample Flow Chart of Power Simulations



Appendix A: Two-Sample Flow Chart of Power Simulations



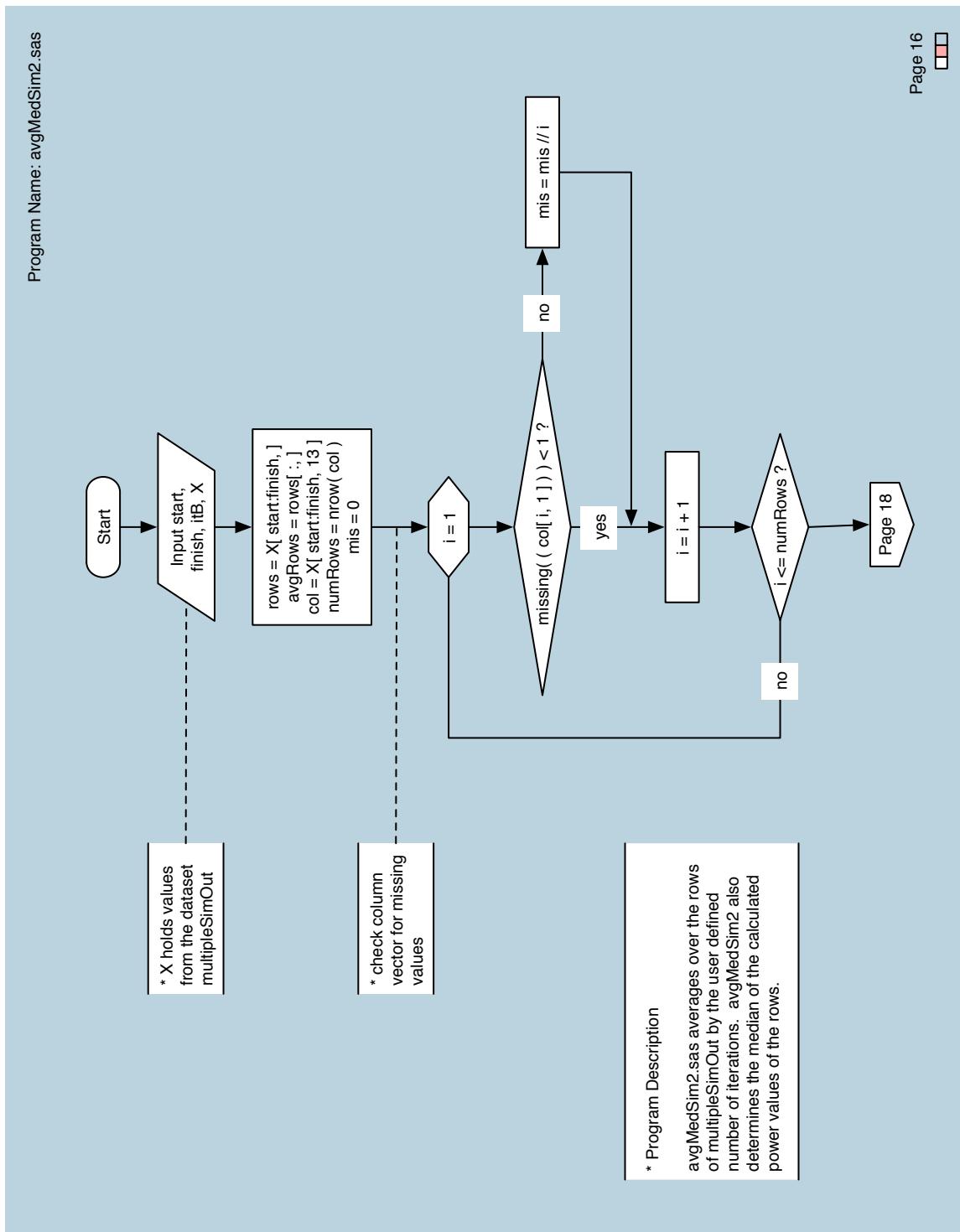
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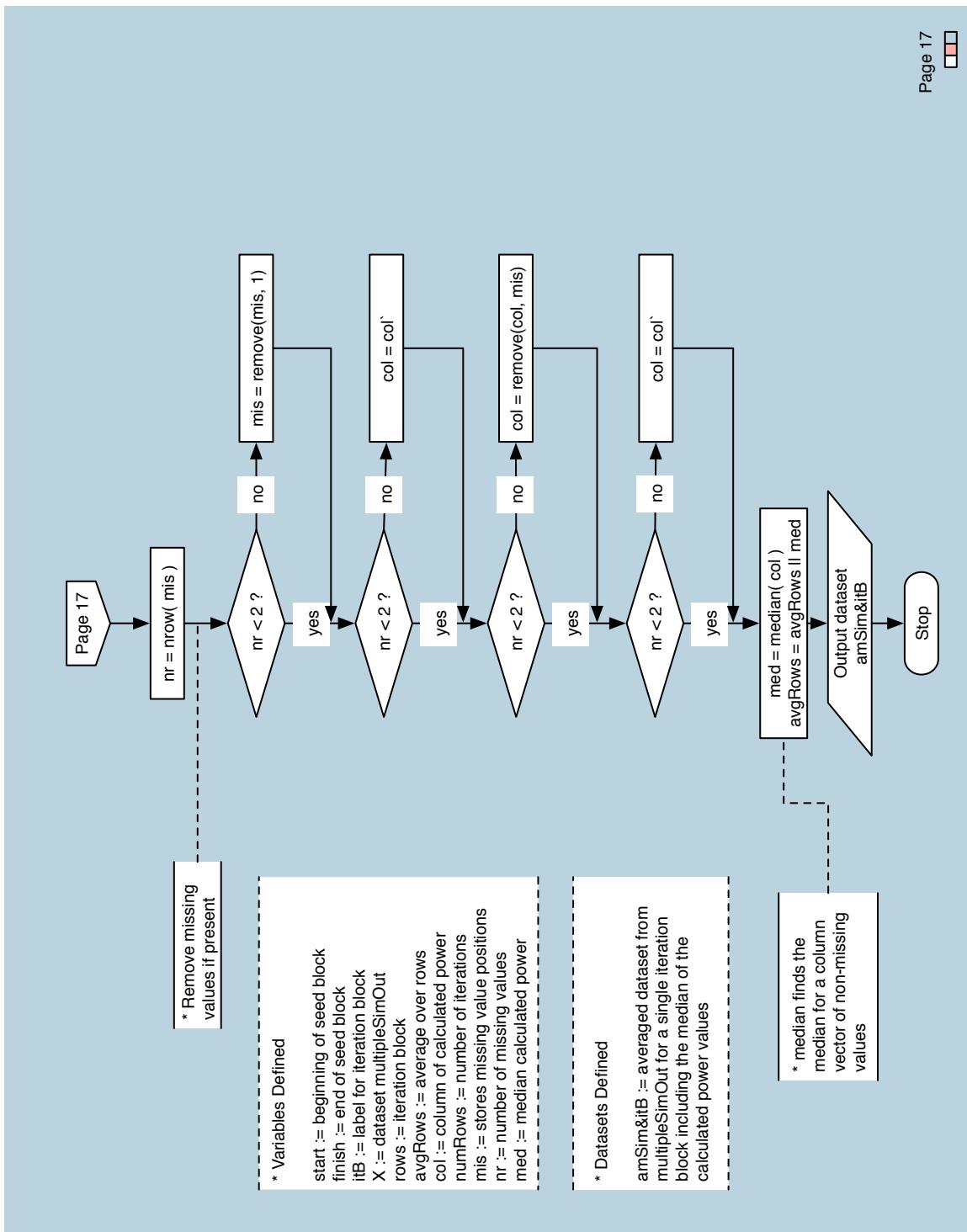
Appendix A: Two-Sample Flow Chart of Power Simulations

```
* Variables Defined  
  
itM := maximum number of iterations  
nsXns := number of population standard deviations considered multiplied by  
the number of sample sizes considered  
bStart := start of seed block  
bFinish := end of seed block  
  
-----  
* Datasets Defined  
  
avgMedMultipleSimOut := averaged dataset of multipleSimOut by iteration  
blocks including the median for the calculated power values in each iteration  
block
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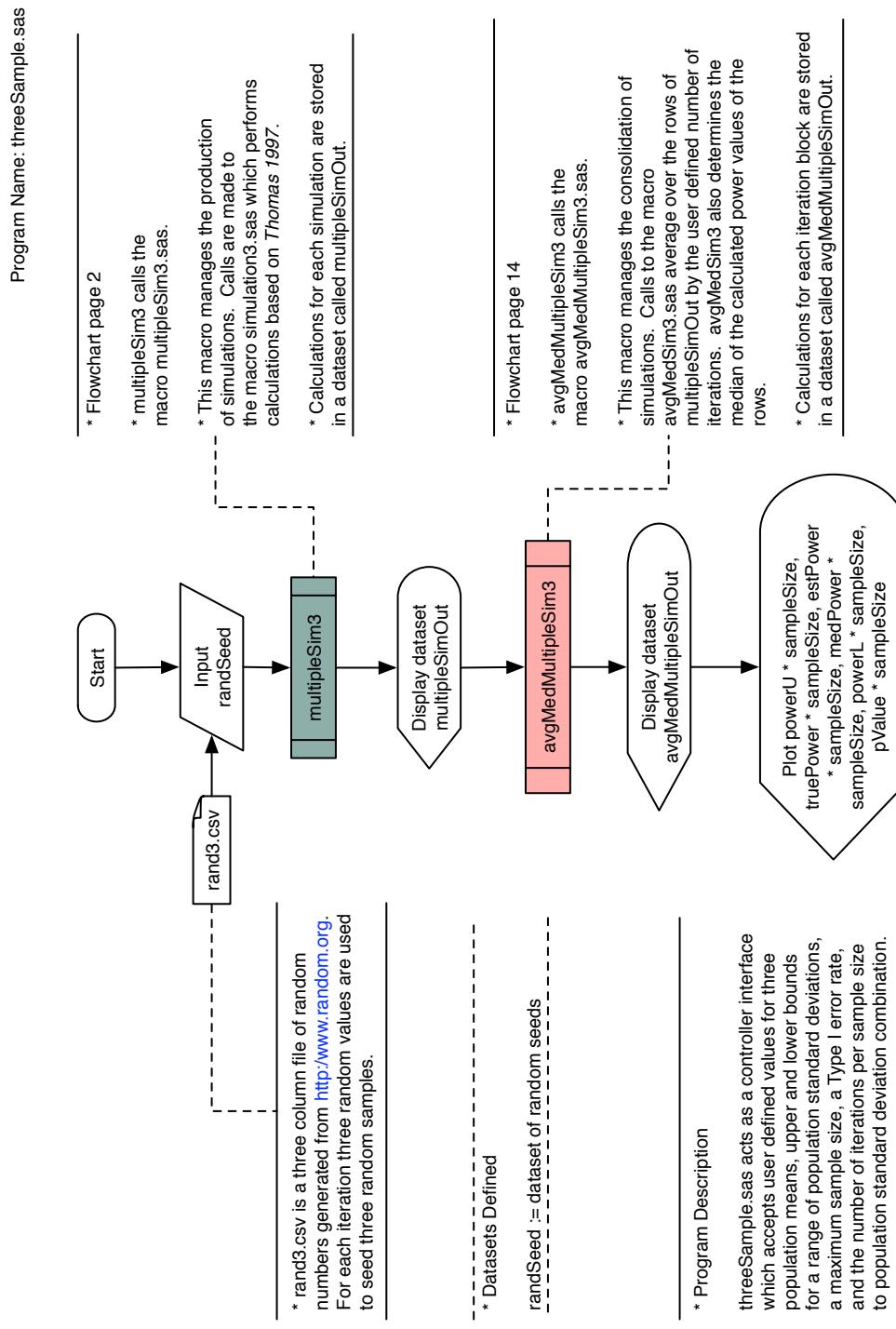
Appendix A: Two-Sample Flow Chart of Power Simulations



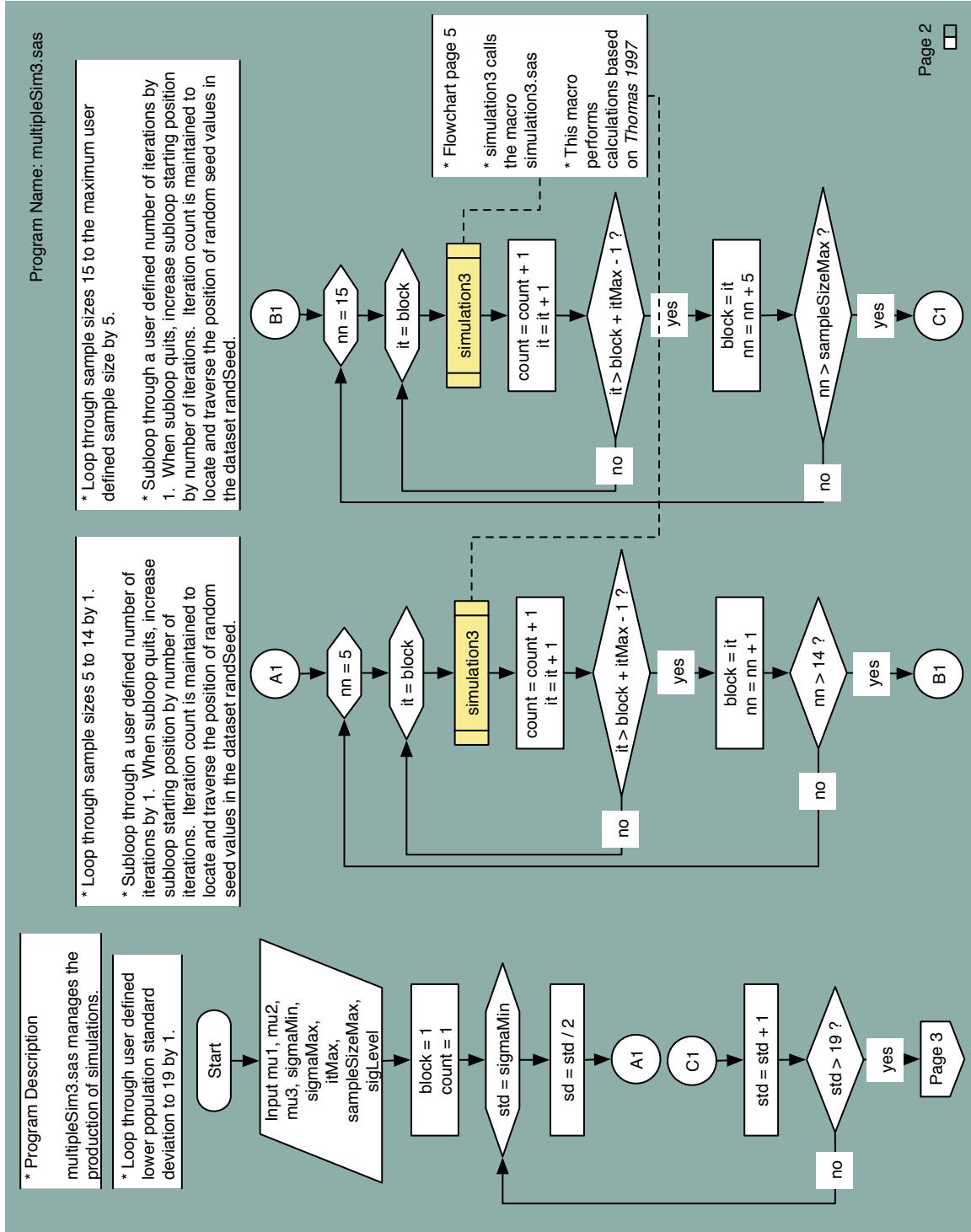
Appendix A: Two-Sample Flow Chart of Power Simulations



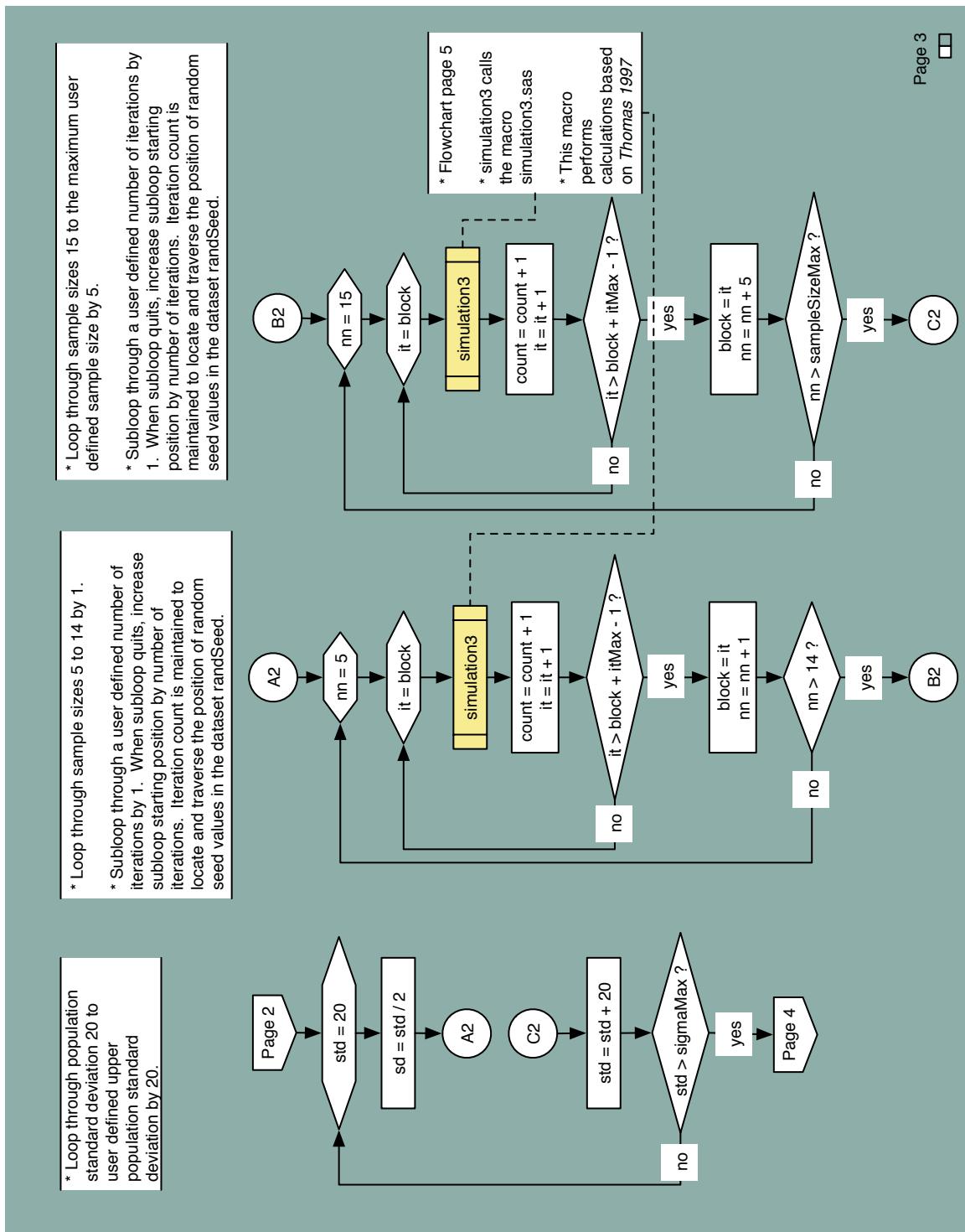
Appendix A: Three-Sample Flow Chart of Power Simulations



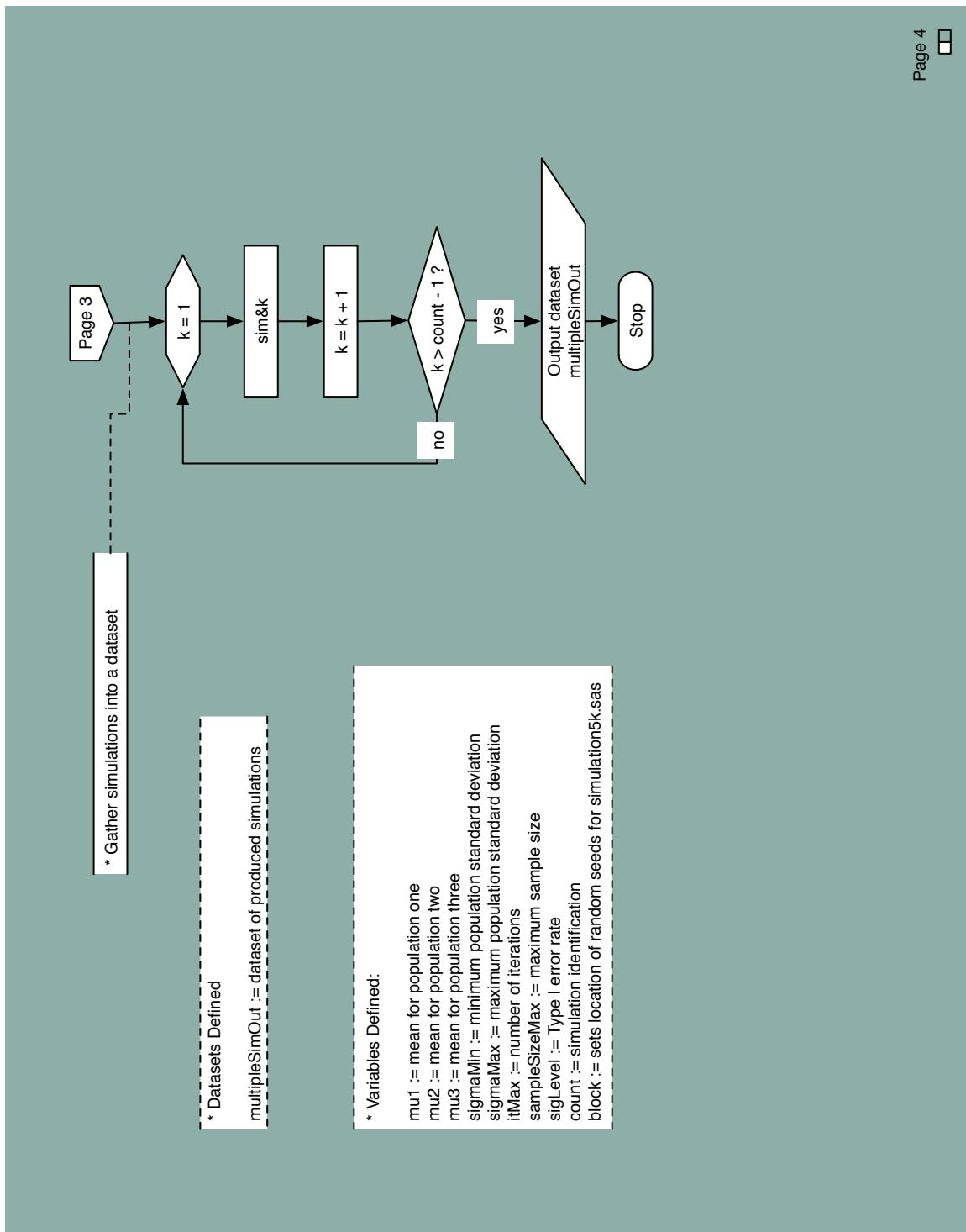
Appendix A: Three-Sample Flow Chart of Power Simulations



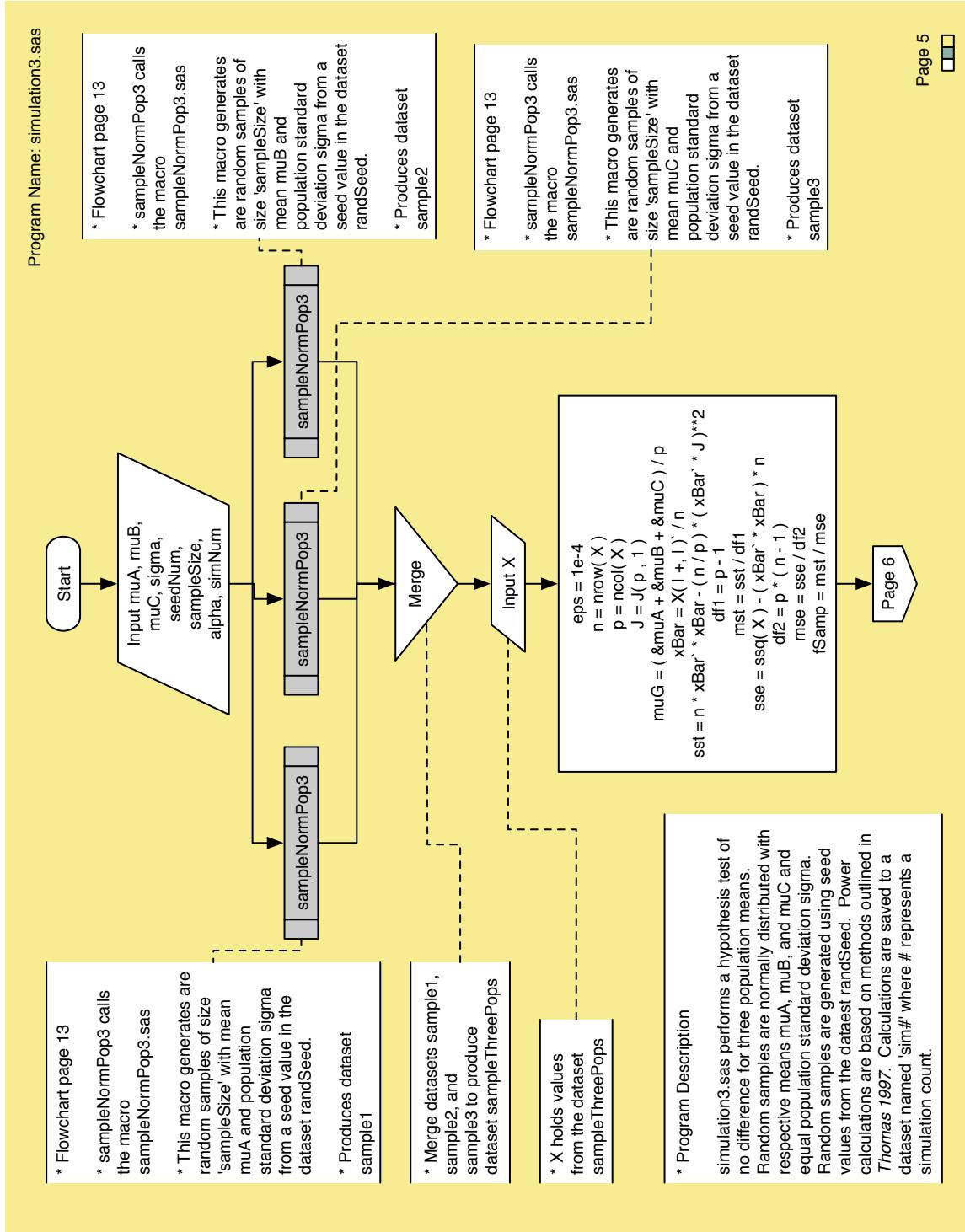
Appendix A: Three-Sample Flow Chart of Power Simulations



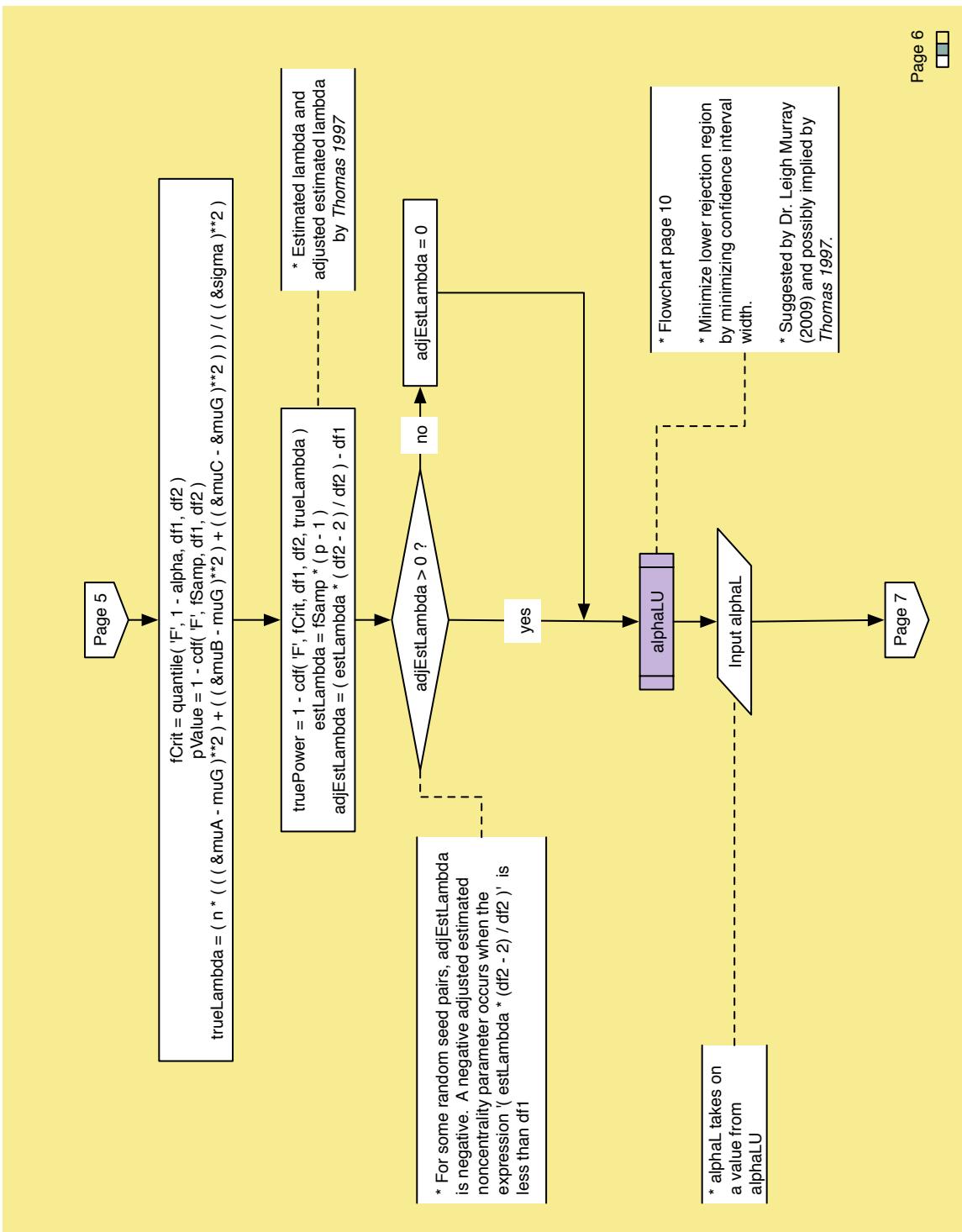
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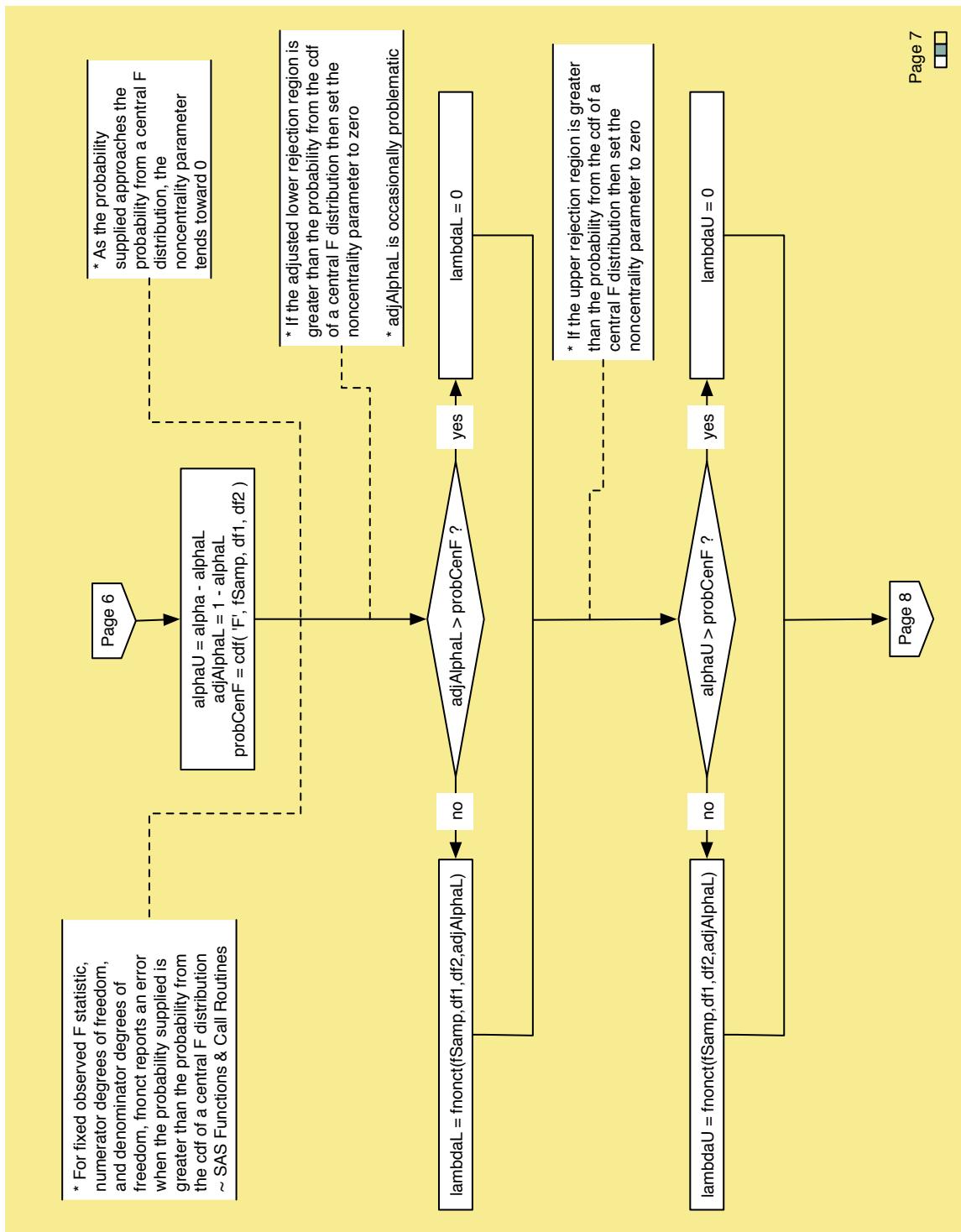
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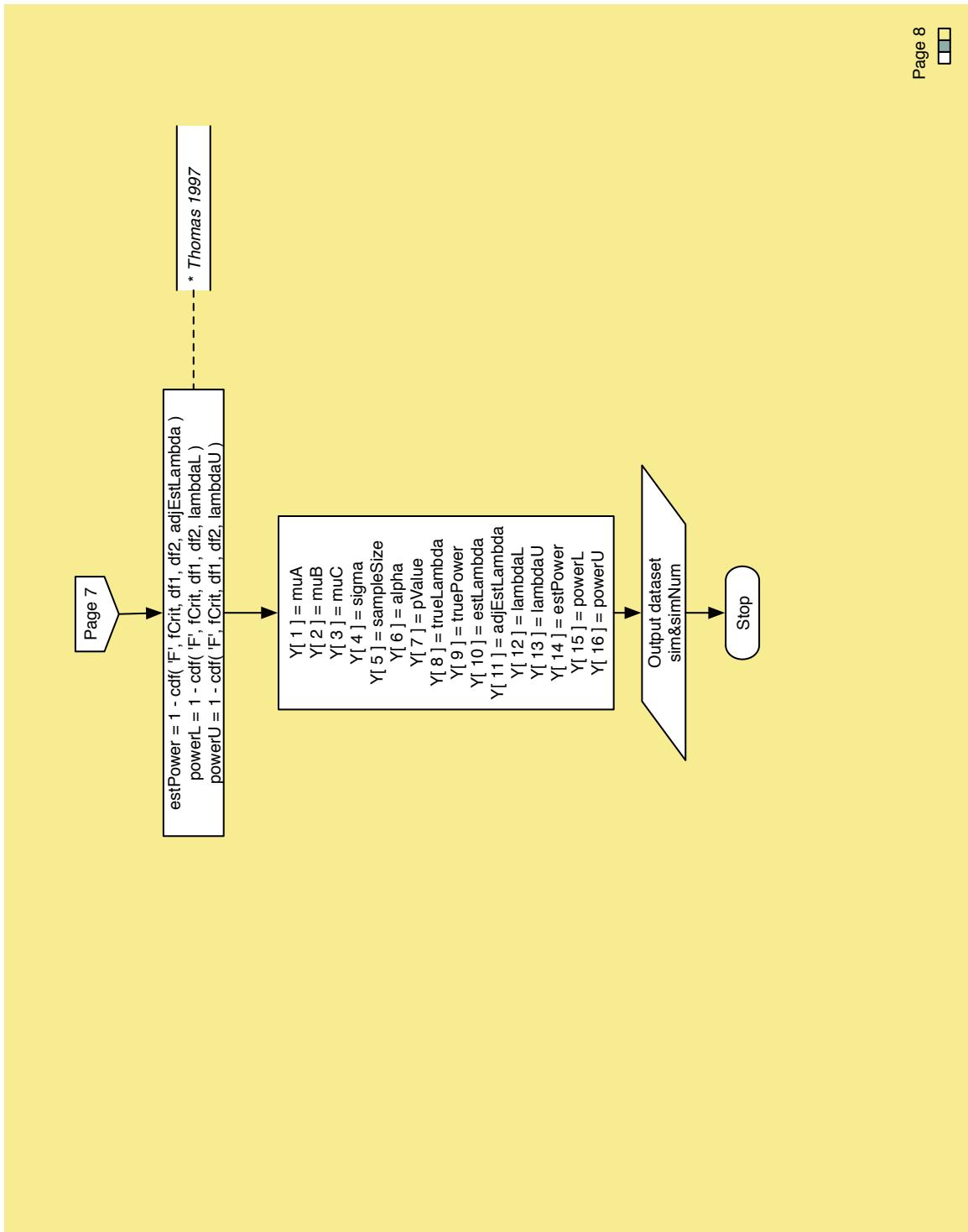
Appendix A: Three-Sample Flow Chart of Power Simulations



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Appendix A: Three-Sample Flow Chart of Power Simulations



Appendix A: Three-Sample Flow Chart of Power Simulations

* Variables Defined

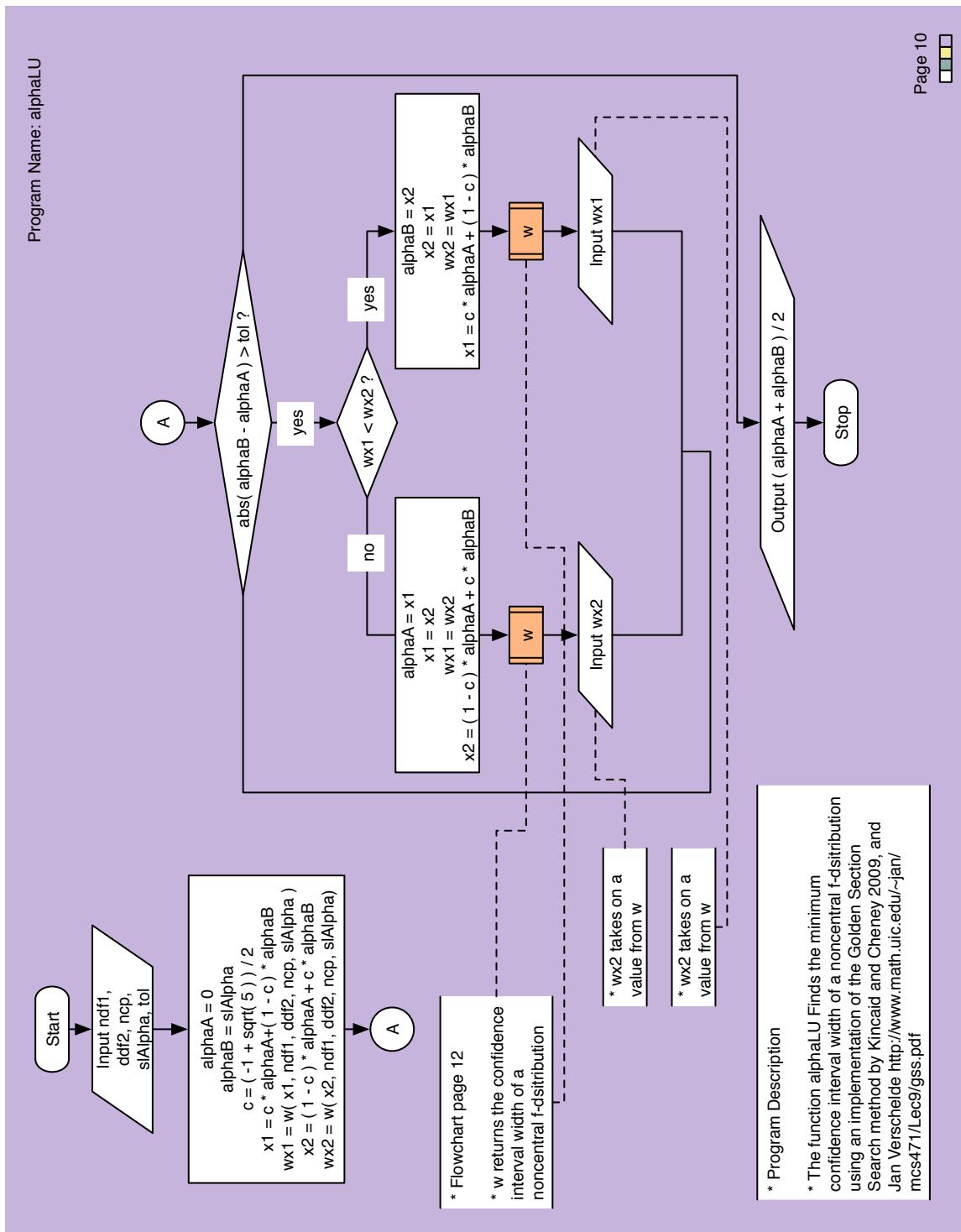
```
muA := mean for population one
muB := mean for population two
muC := mean for population three
muG := average of the three population means
sigma := population standard deviation
seedNum := location of random seeds
sampleSize := sample size
alpha := level of significance
simNum := simulation identification
X := samples from normal populations
eps := tolerance for Golden Section Search methods
n := number rows
p := number columns
xBar := sample mean
sst := sum of squares for treatment
df1 := numerator degrees of freedom
mst := mean square for treatment
sse := sum of squares for error
df2 := denominator degrees of freedom
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powerL := lower confidence interval for calculated power
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```

* Datasets Defined

```
sampleThreePops := random samples dataset
sim&simNum := simulation dataset
```



Appendix A: Three-Sample Flow Chart of Power Simulations

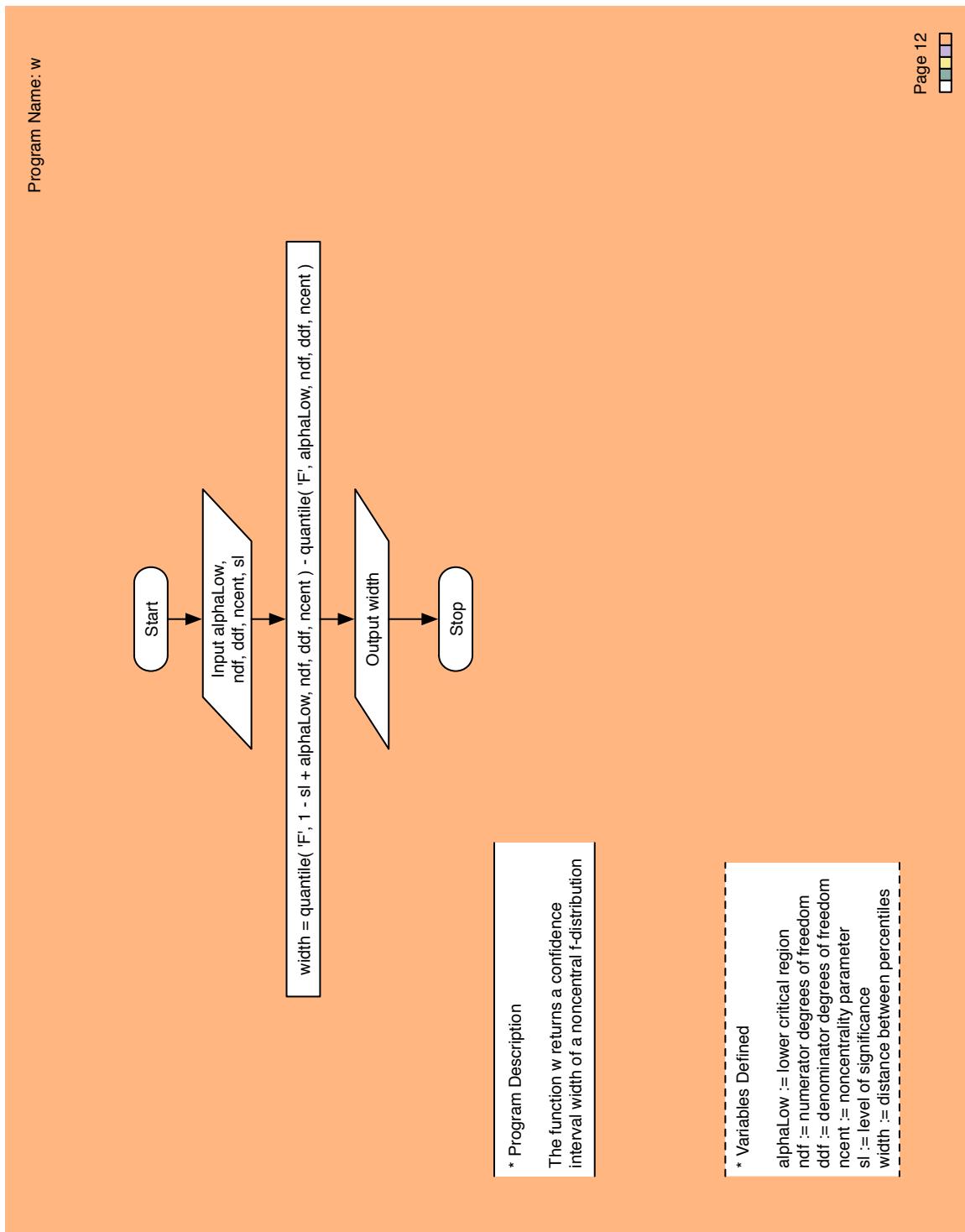


Appendix A: Three-Sample Flow Chart of Power Simulations

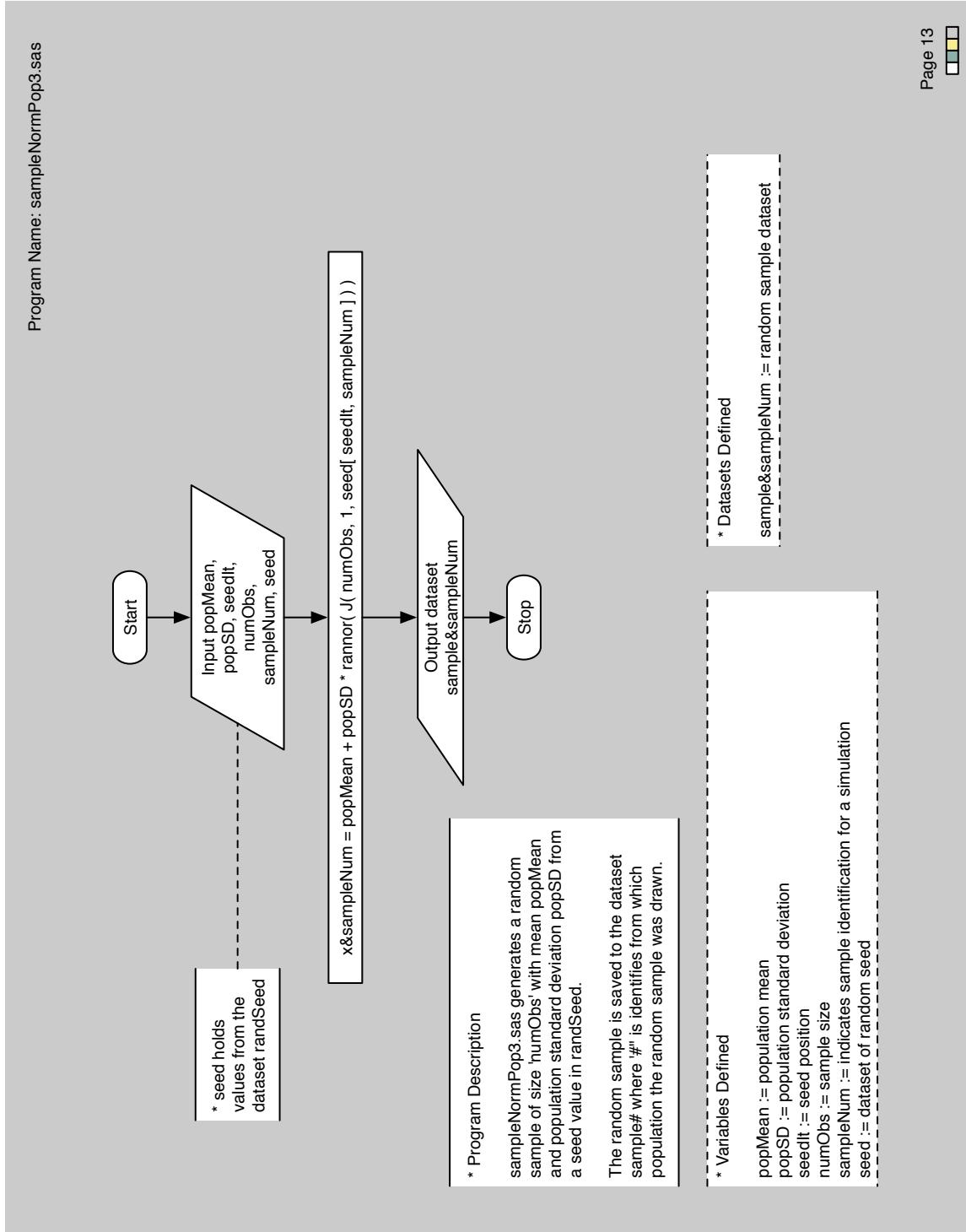
* Variables Defined

```
ndf1 := numerator degrees of freedom  
ddf2 := denominator degrees of freedom  
ncp := noncentrality parameter  
slAlpha := level of significance  
tol := tolerance for Golden Section Search method  
alphaA := lower bound for optimized lower alpha  
alphaB := upper bound for optimized lower alpha  
c := Golden ratio constant reduction factor  
x1 := percentile associated with alphaA  
wx1 := height associated with x1  
x2 := percentile associated with alphaB  
wx2 := height associated with x2
```

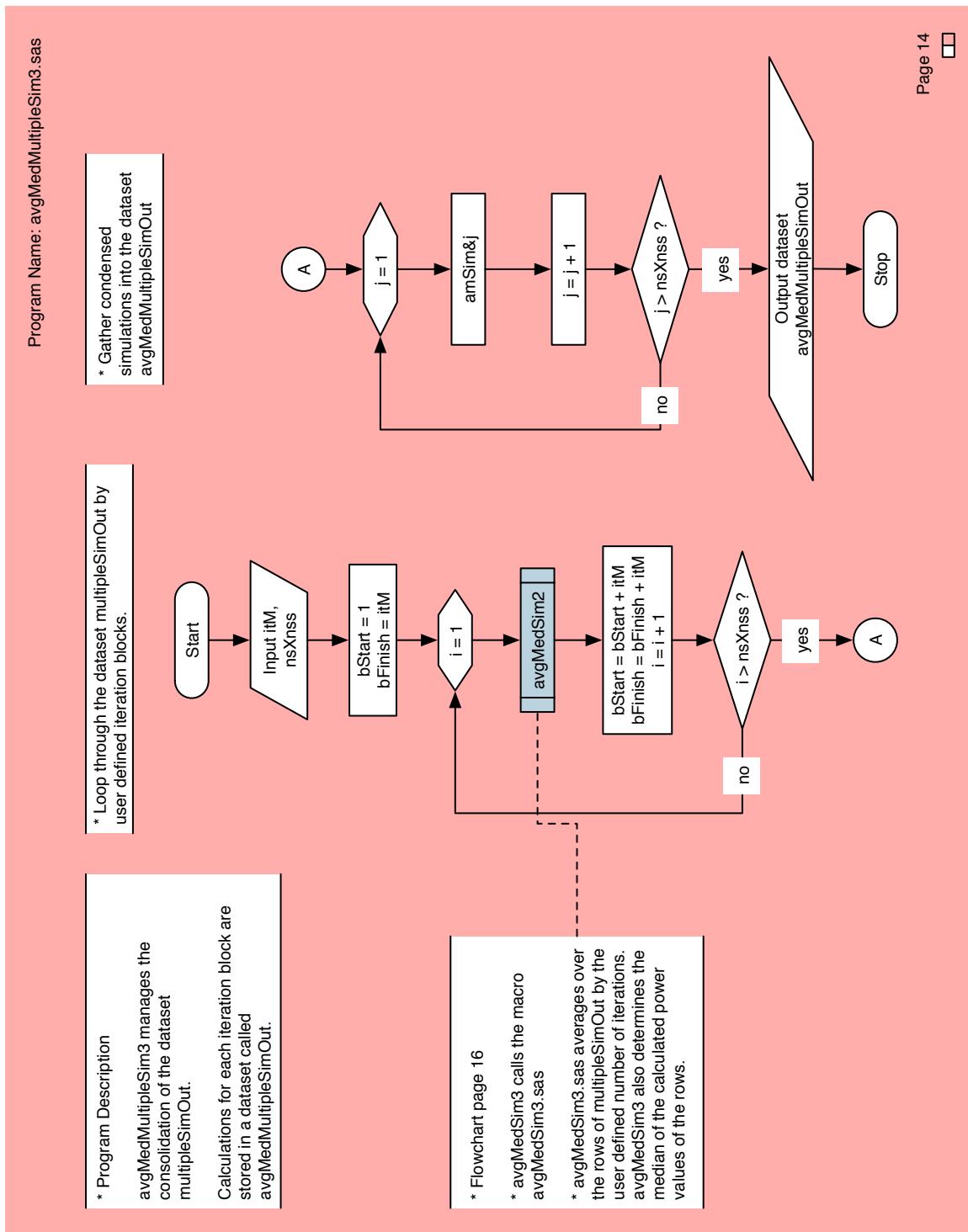
Appendix A: Three-Sample Flow Chart of Power Simulations



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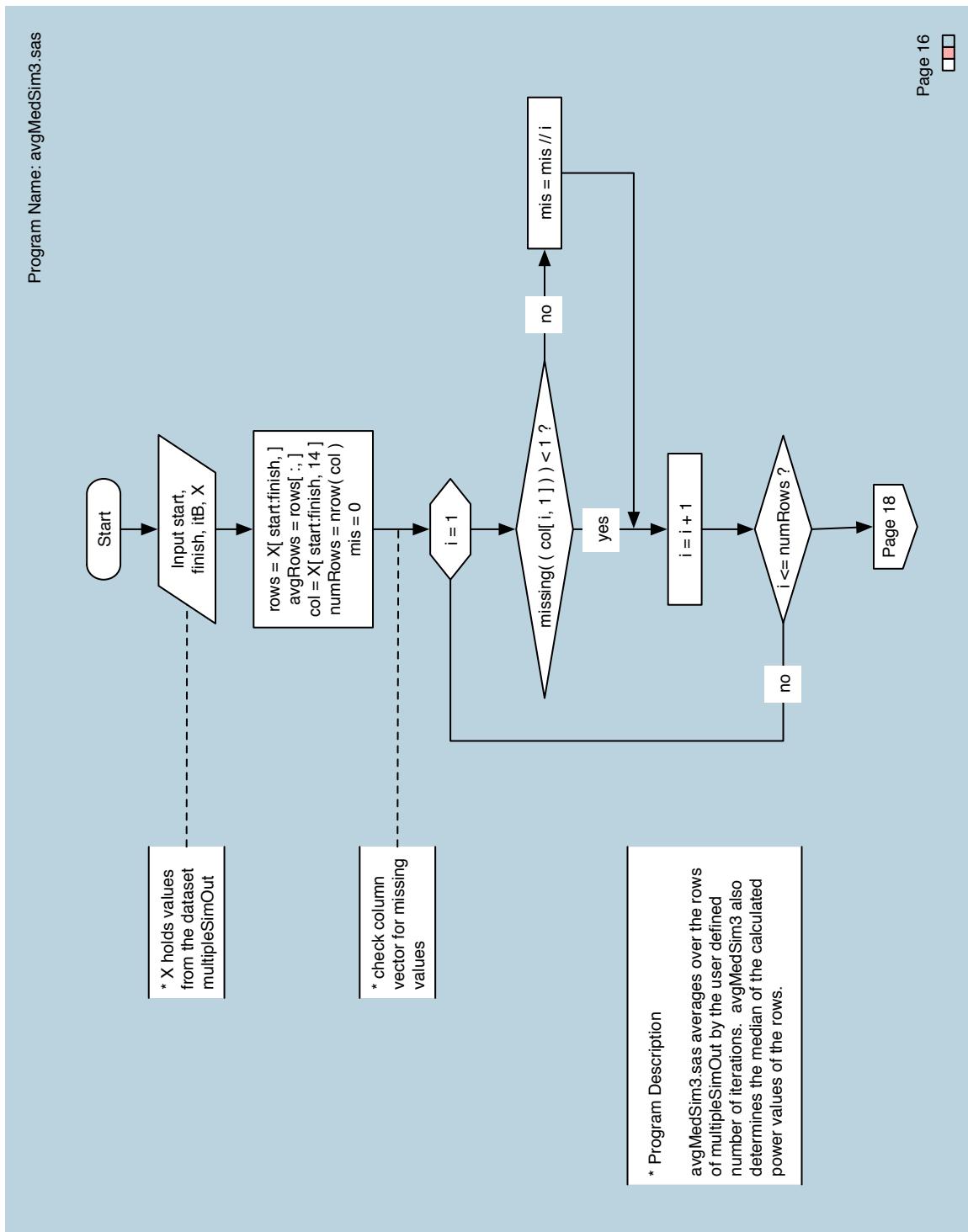
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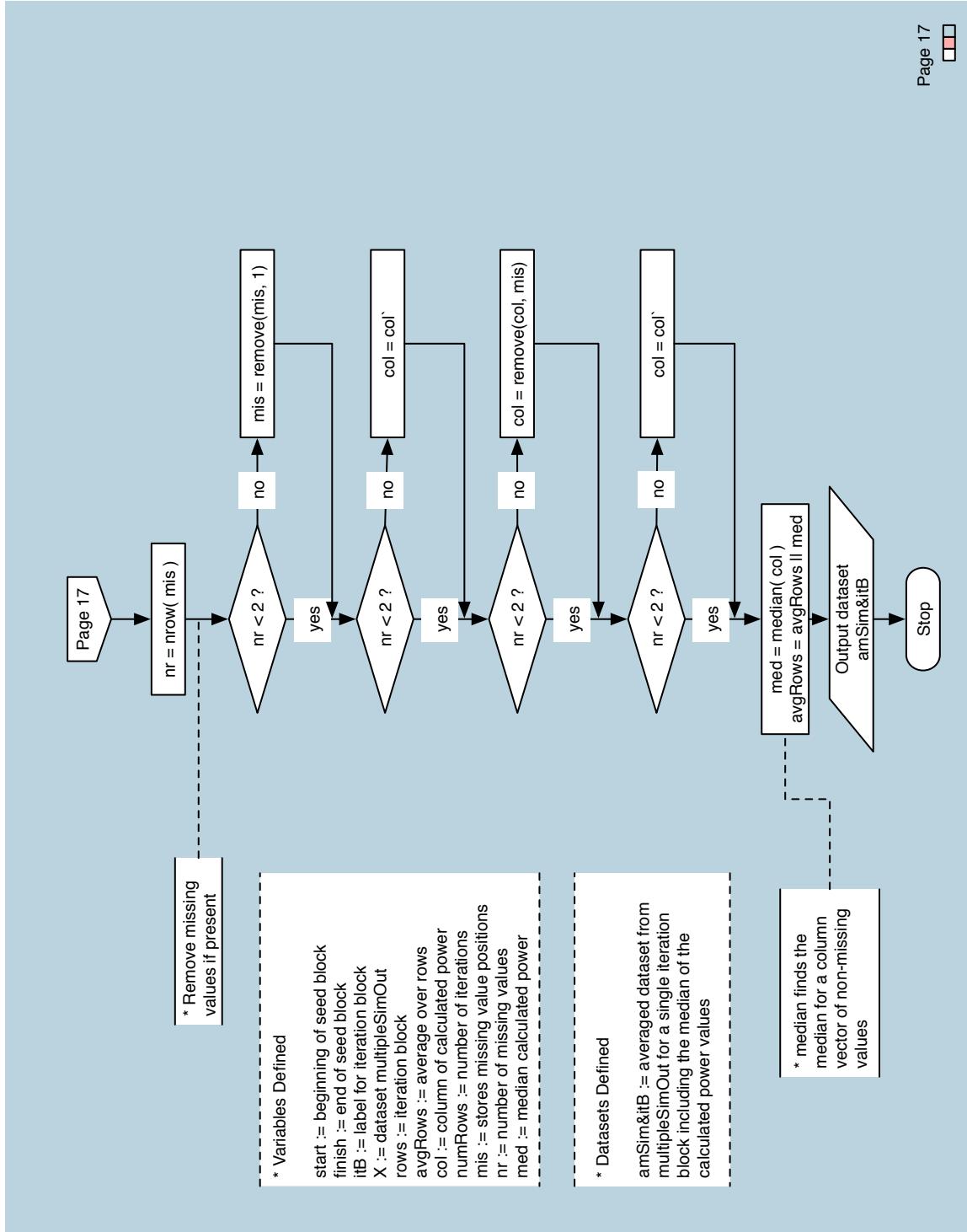
Appendix A: Three-Sample Flow Chart of Power Simulations

```
* Variables Defined  
  
itM := maximum number of iterations  
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the number of sample sizes considered  
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* Datasets Defined  
  
avgMedMultipleSimOut := averaged dataset of multipleSimOut by iteration  
blocks including the median for the calculated power values in each iteration  
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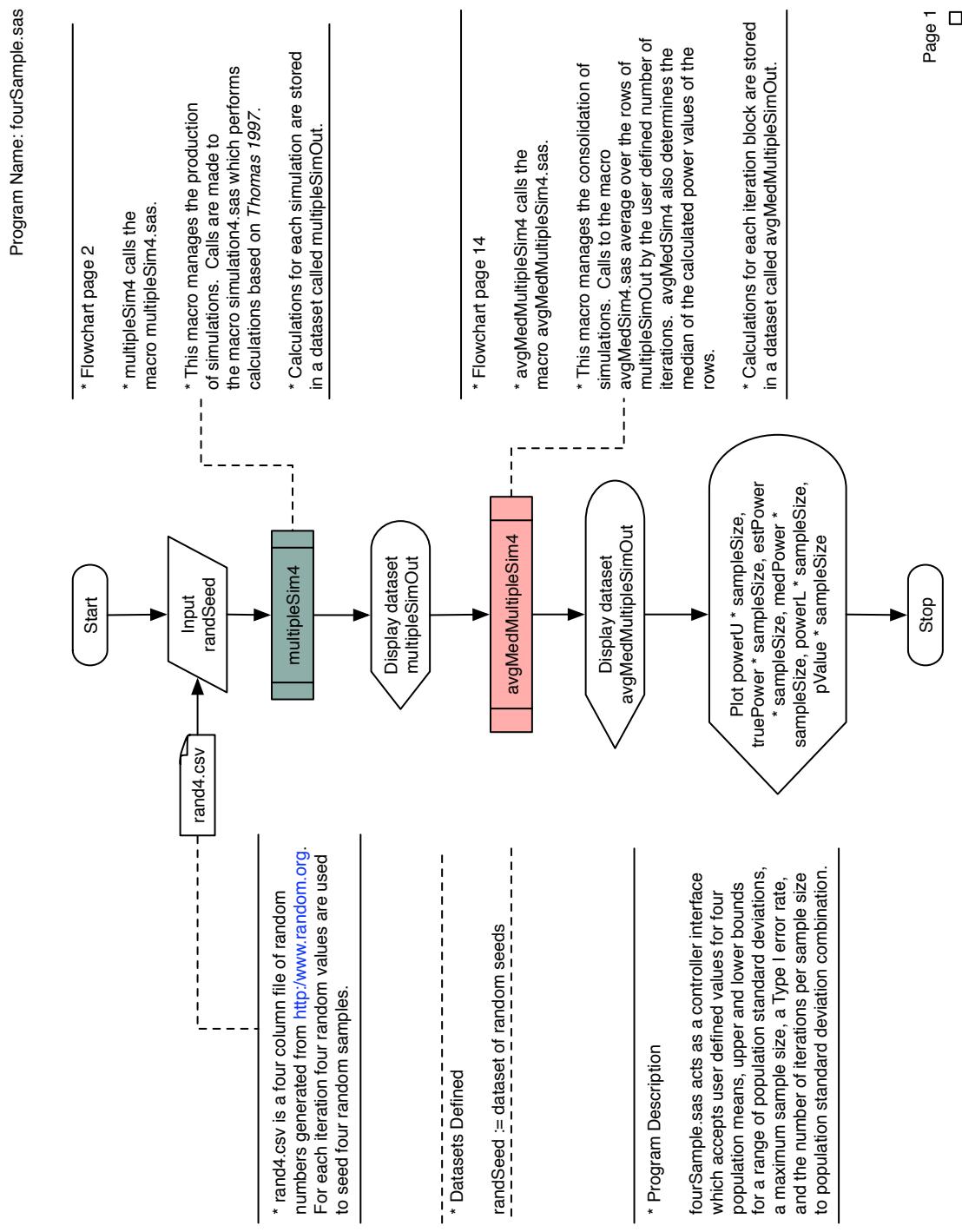
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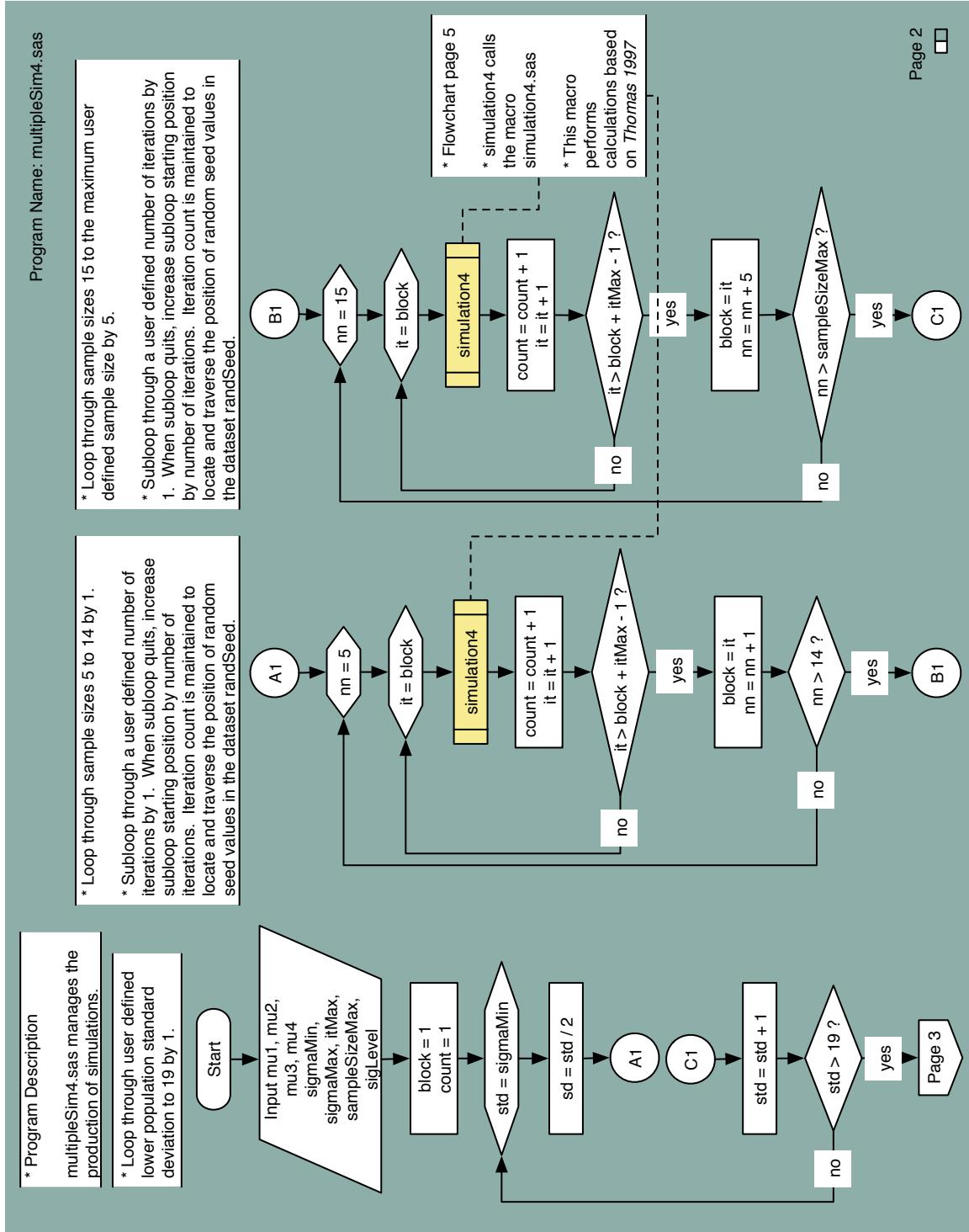
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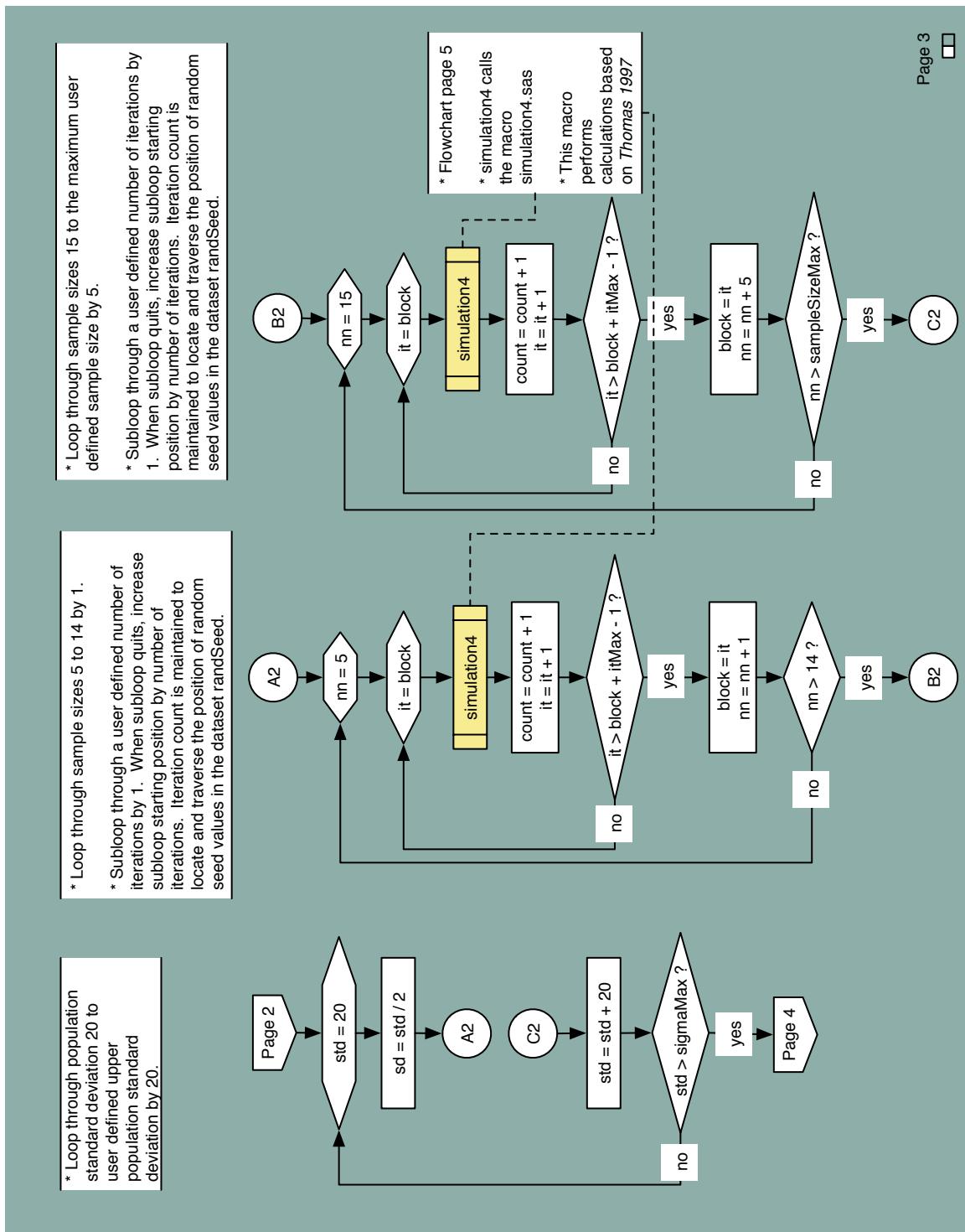
Appendix A: Four-Sample Flow Chart of Power Simulations



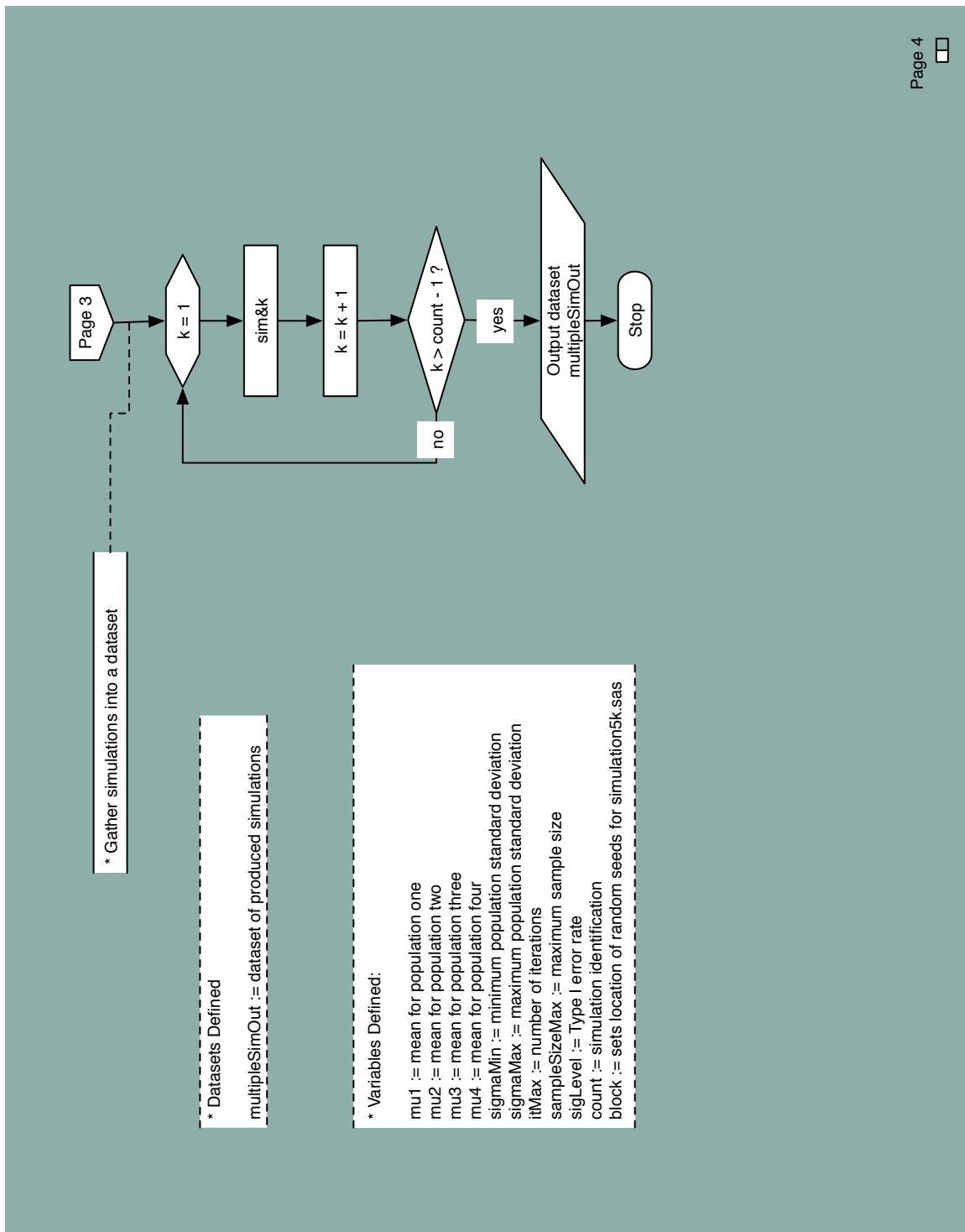
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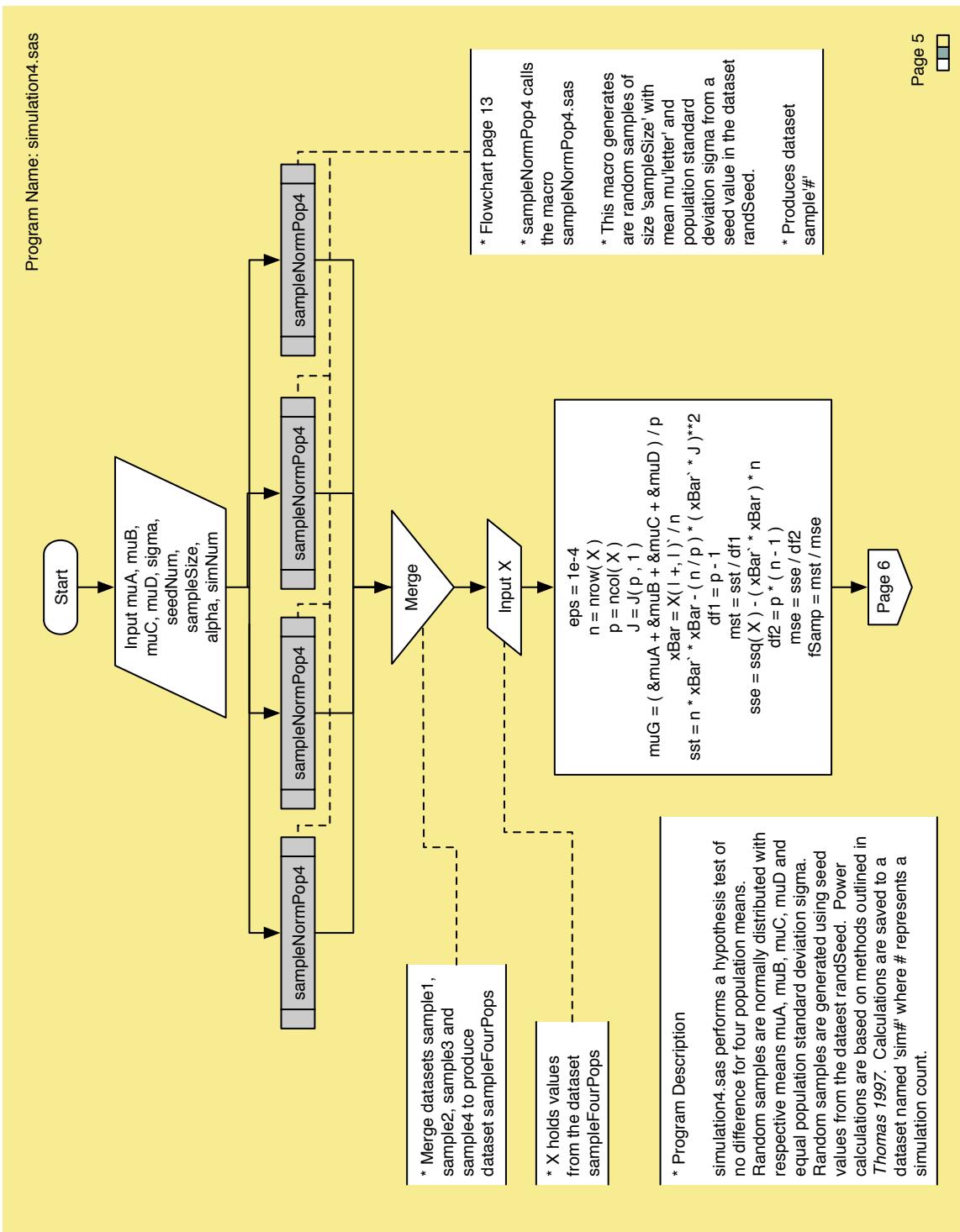
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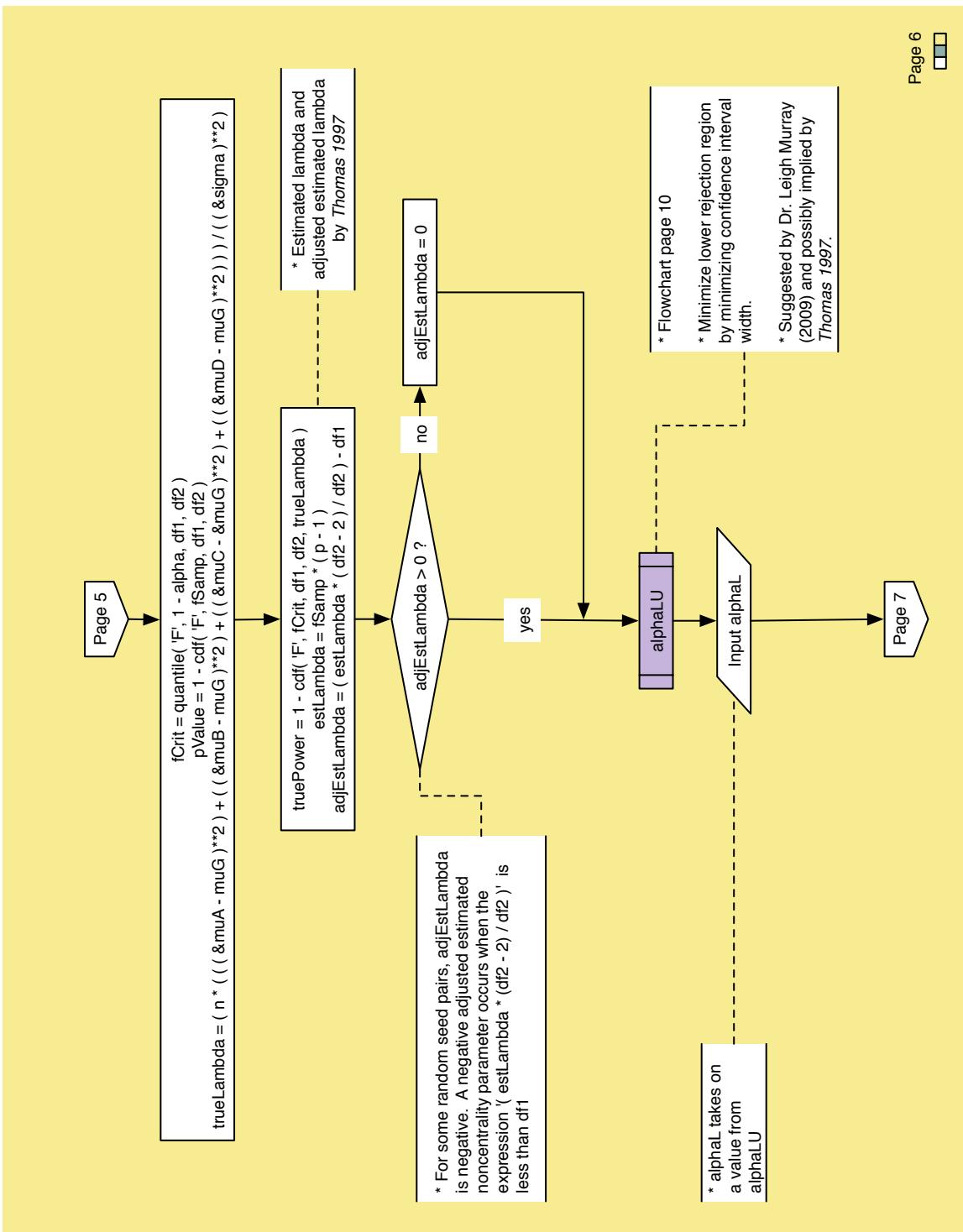
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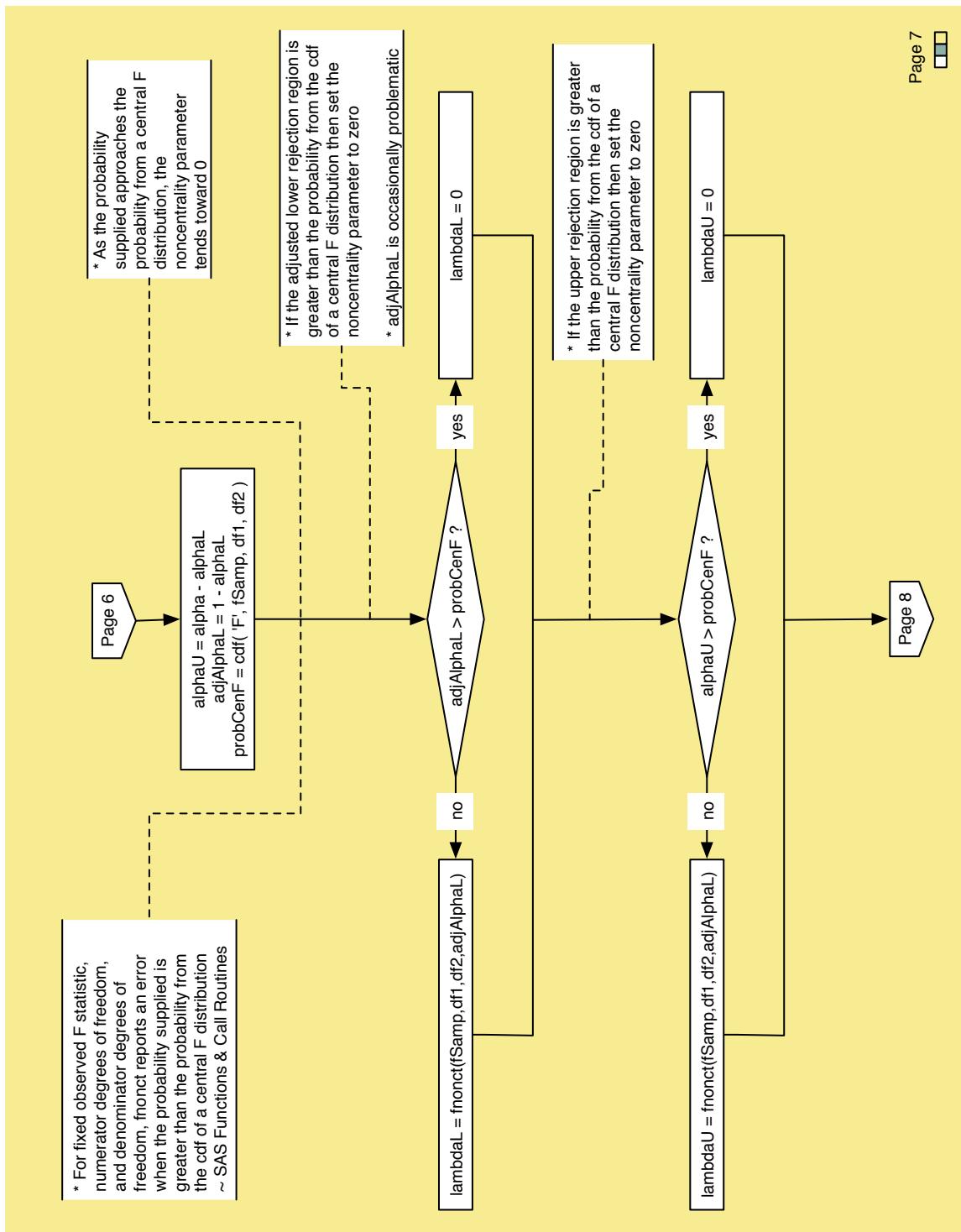
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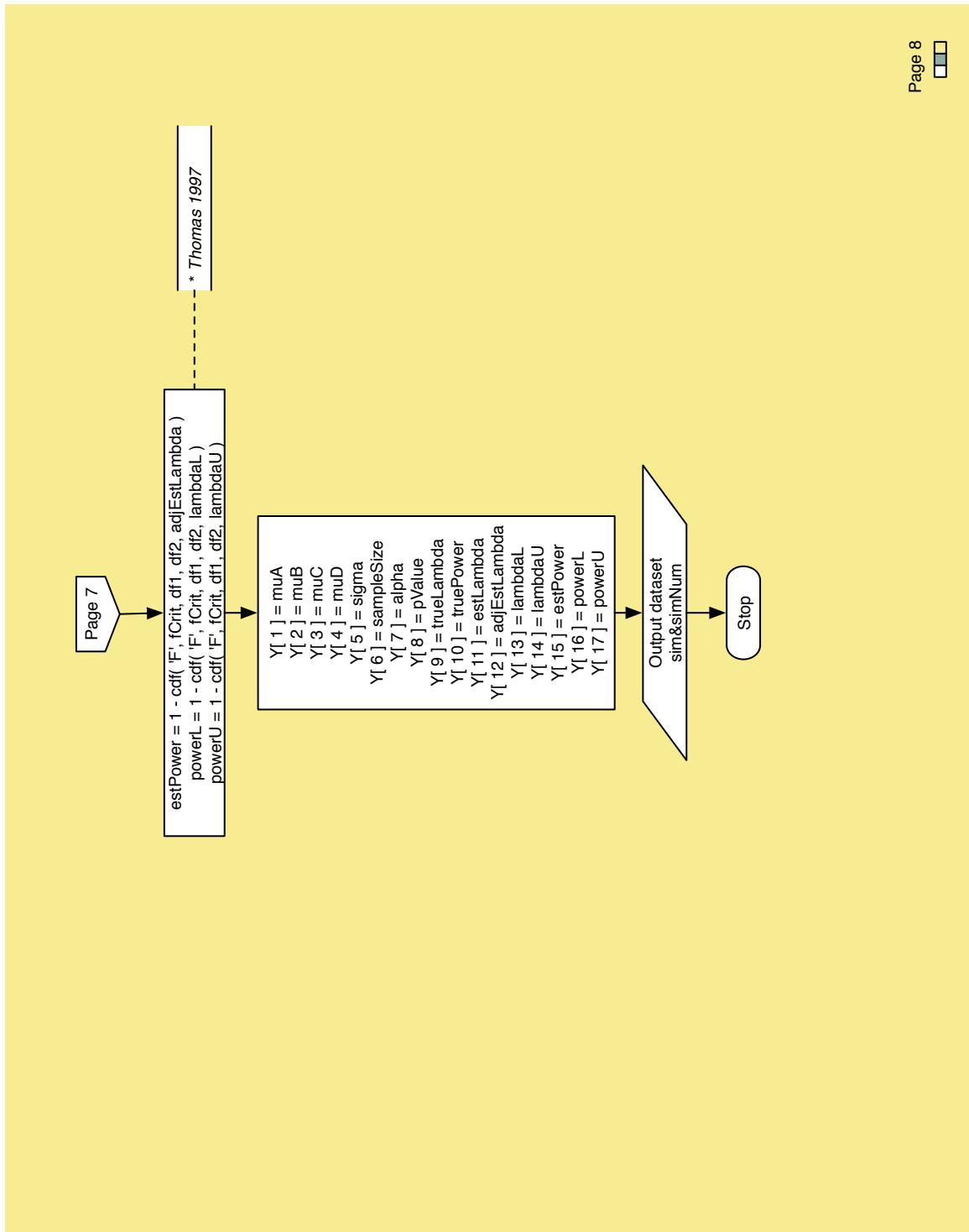
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Appendix A: Four-Sample Flow Chart of Power Simulations



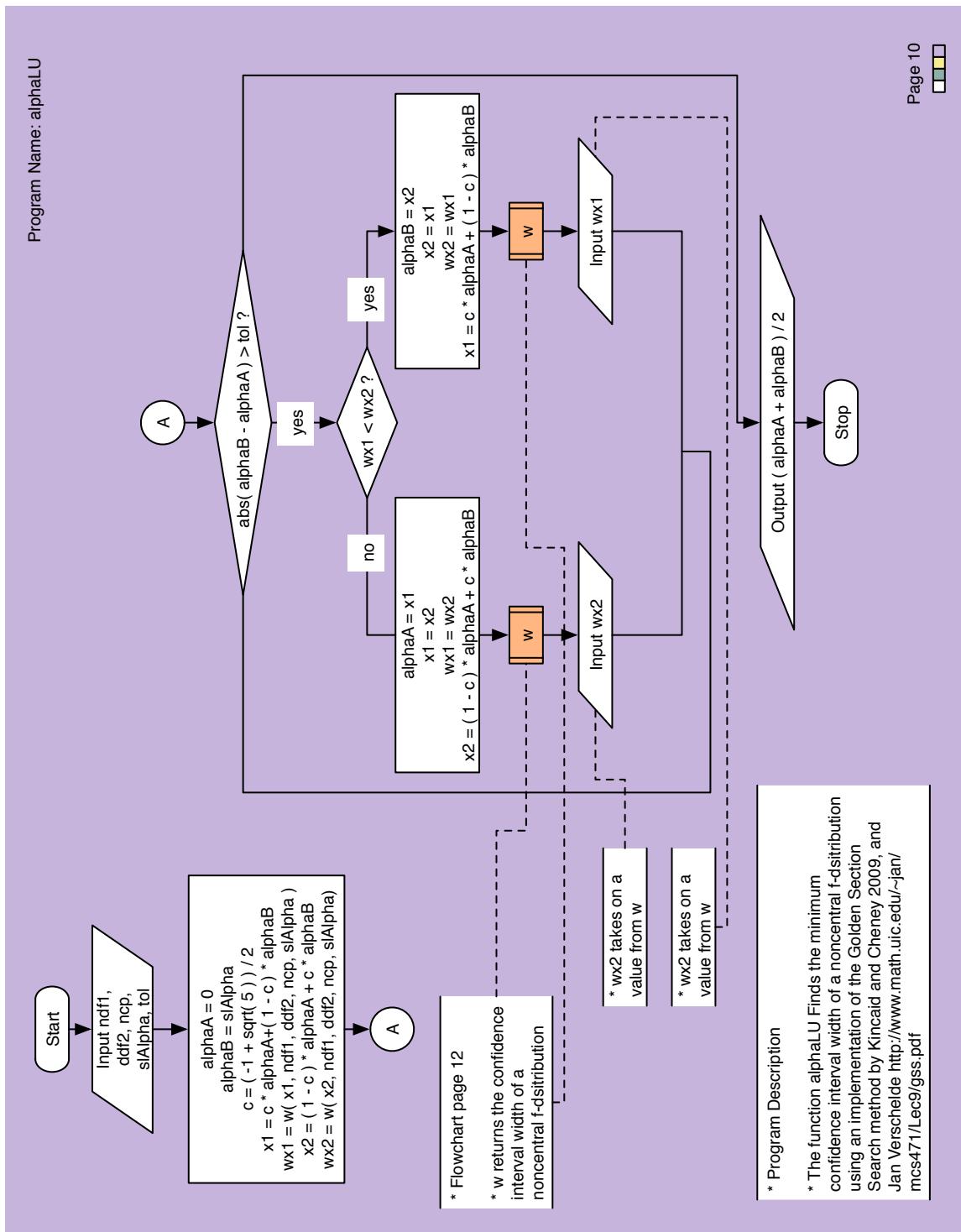
Appendix A: Four-Sample Flow Chart of Power Simulations

```
* Variables Defined
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muB := mean for population two
muC := mean for population three
muD := mean for population four
muG := average of the tree population means
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pValue := p-value for the test
trueLambda := noncentrality parameter
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alphaL := optimized lower critical region
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probCenF := probability of observed F statistic of the central F distribution
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powerL := lower confidence interval for calculated power
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* Datasets Defined
sampleFourPops := random samples dataset
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```



Appendix A: Four-Sample Flow Chart of Power Simulations

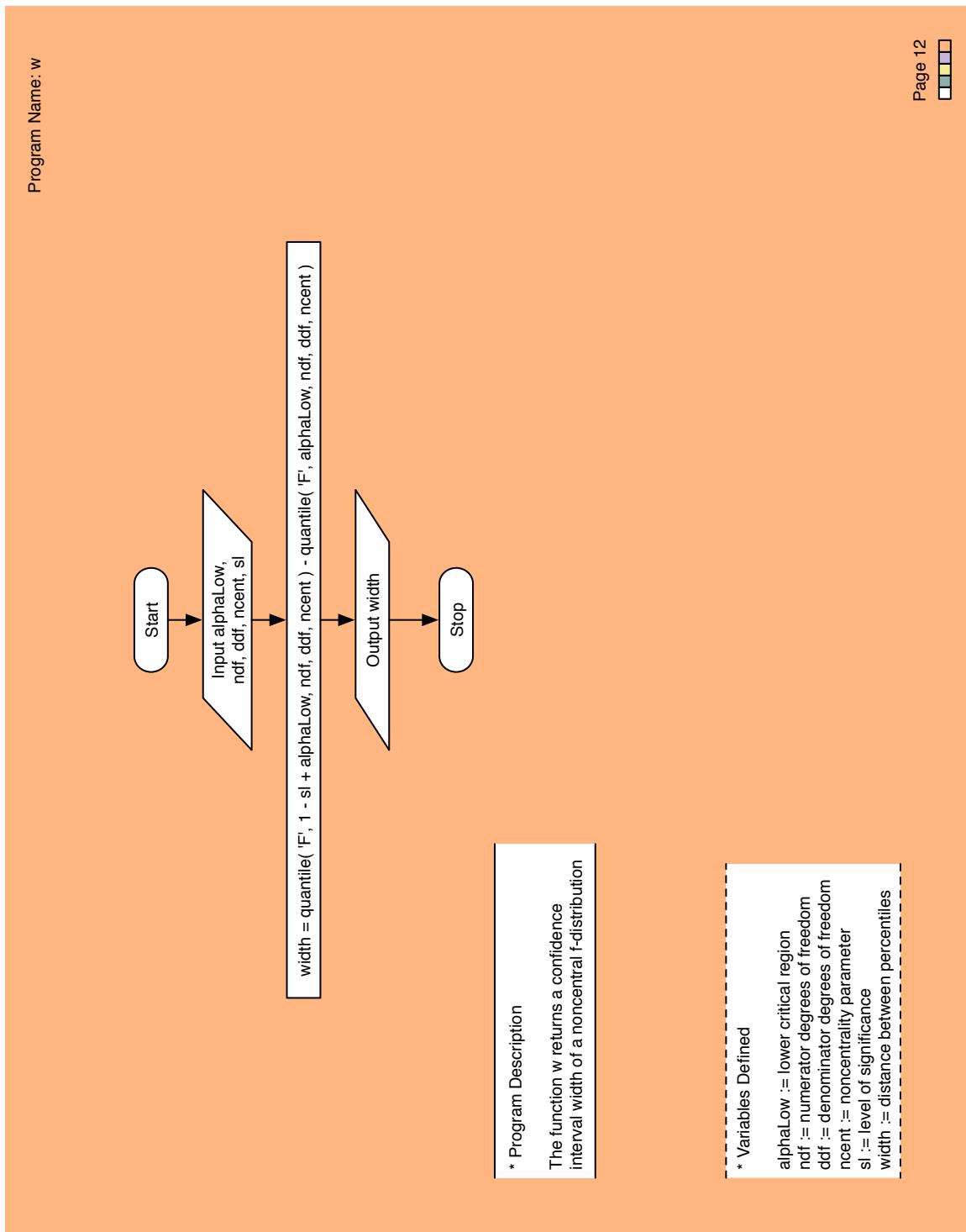


Appendix A: Four-Sample Flow Chart of Power Simulations

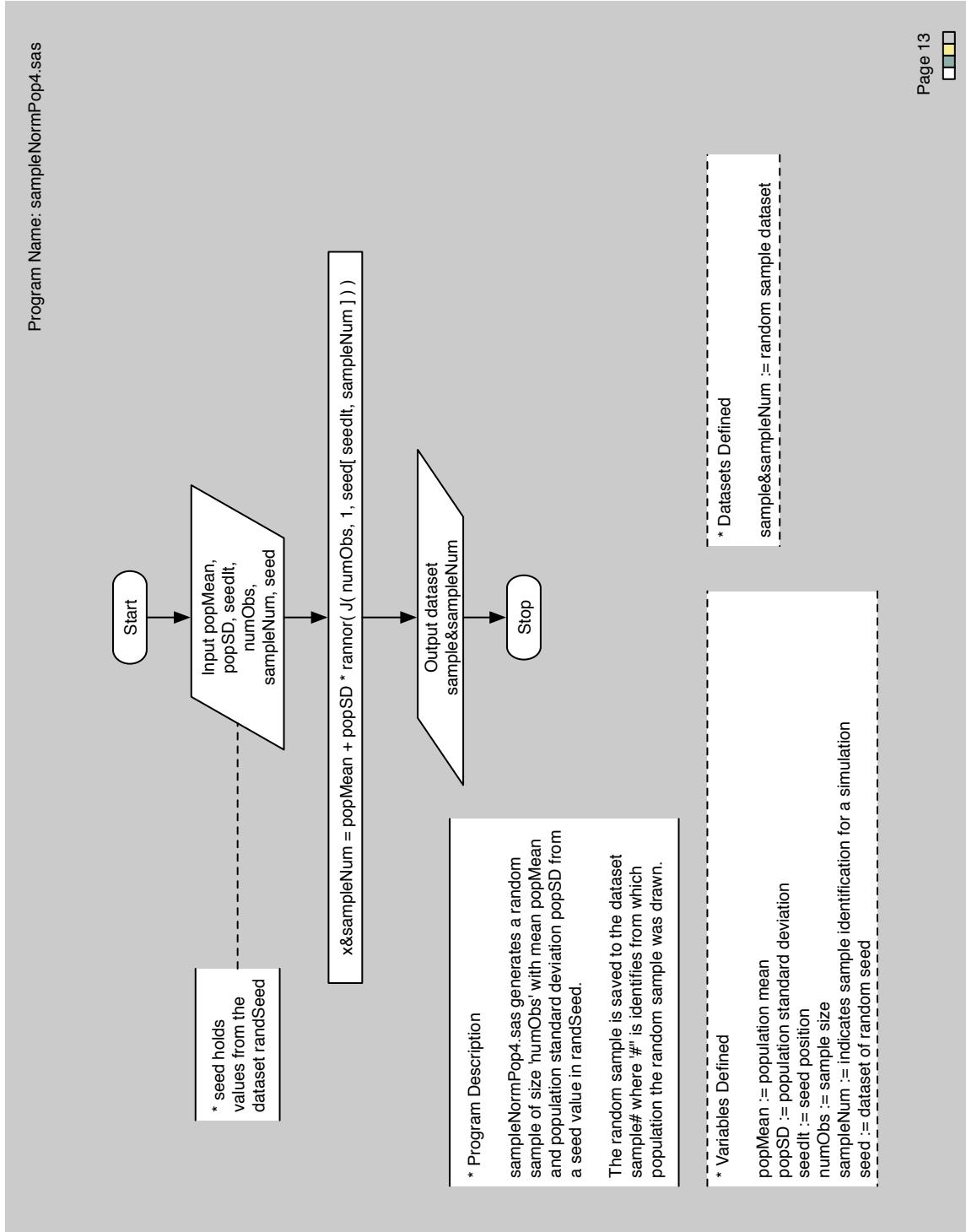
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wx1 := height associated with x1  
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```

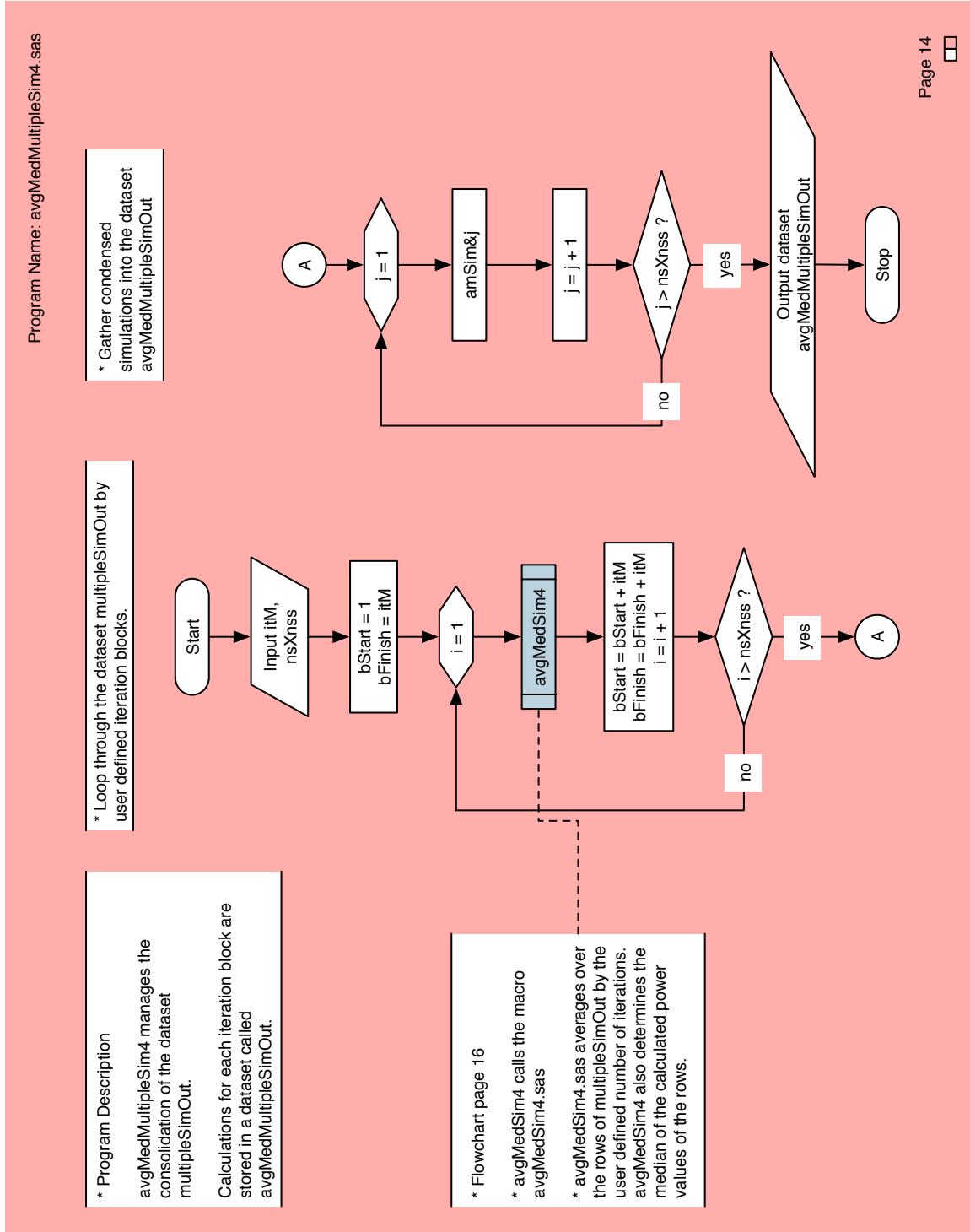
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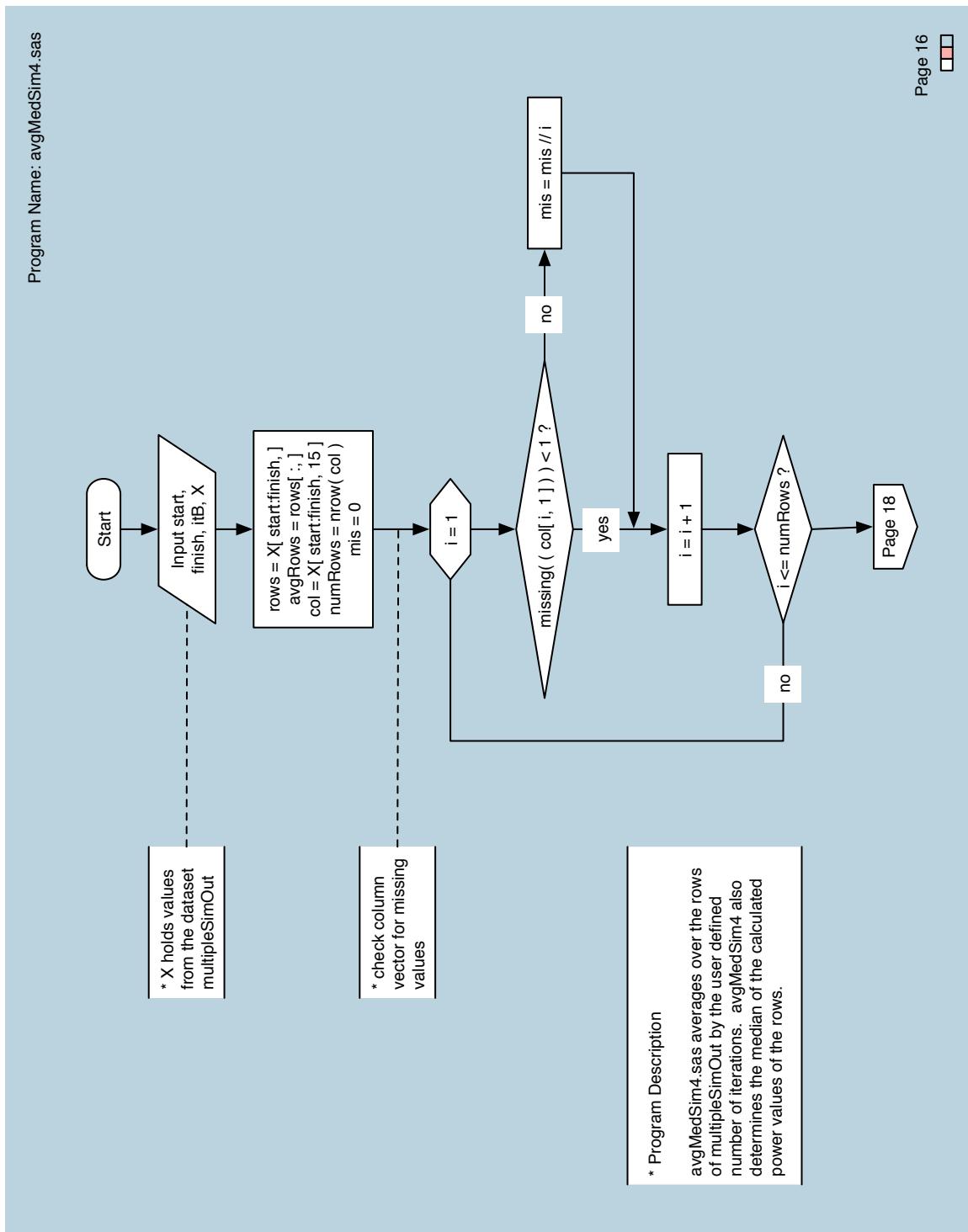
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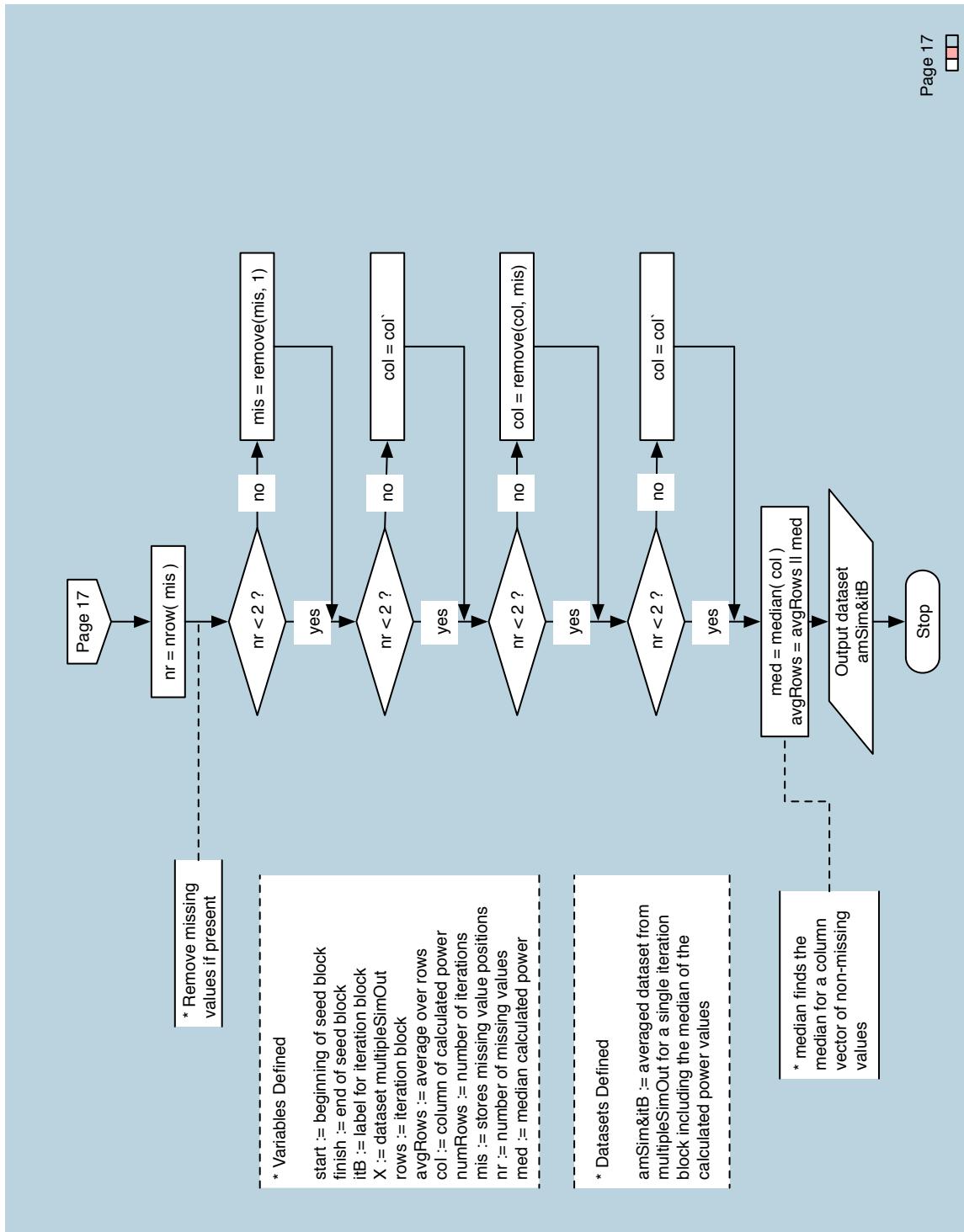
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* Variables Defined  
  
itM := maximum number of iterations  
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blocks including the median for the calculated power values in each iteration  
block
```

Appendix A: Four-Sample Flow Chart of Power Simulations



Appendix A: Four-Sample Flow Chart of Power Simulations



Appendix B: Two-Sample SAS Code for Power Simulations

```

1  /* twoSample *****/
2  /*
3   * Requirement(s):
4   * rand2.csv, multipleSim2.sas, avgMedMultipleSim2.sas
5   */
6  /* Program Description:
7   * twoSample.sas acts as a controller interface which accepts user
8   * defined values for two population means, upper and lower bounds
9   * for a range of population standard deviations, a maximum sample size,
10  /* a type I error rate, and the number of iterations per sample size
11  /* to population standard deviation combination.
12  */
13 *****/
14
15 proc printto print='\\Path\\to\\Output\\Folder\\twoSampleOut.lst' log='\\Path\\to\\Output\\Folder\\twoSampleOut.log';
16
17 run;
18
19 /* Input random seed file to the dataset randseed *****/
20 /*
21 */
22 /* rand2.csv is a two column file of random numbers generated from
23 /* http://www.random.org. For each iteration of multipleSim2 two random
24 /* values are used to seed two random samples.
25 */
26 *****/
27
28 data randSeed;
29   infile "\\Path\\to\\File\\rand2.csv" dlm=' ' ;
30   input seedCol1 seedCol2;
31
32 run;
33
34 /* Produce simulations *****/
35 /*
36 /* The multipleSim2 manages the production of simulations.
37 /* Calls are made to the macro simulation2.sas which performs
38 /* calculations based on Thomas 1997.
39 */
40 /* Calculations for each simulation are stored in a dataset called
41 /* multipleSimOut.
42 /*
43 *****/
44 %multipleSim2(mu1=30, mu2=40, sigmaMin=5, sigmaMax=40, itMax=100, sampleSizeMax=50, sigLevel=0.05);
45
46
47

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

48 /* Print multipleSimOut */
49
50 proc print data=multipleSimOut;
51 run;
52
53 /* Produce averages and medians for simulations *****/
54 /*
55 /* The avgMedMultipleSim2 manages the consolidation of simulations.
56 /* Calls are made to avgMedSim2 for each sigma by sample size
57 /* combination. avgMedSim2 averages over a user specified number
58 /* of rows from multipleSimOut. avgMedSim2 also determines the
59 /* median of the calculated power values of the rows.
60 /*
61 /* Calculations for each iteration block are stored in a dataset
62 /* called avgMedMultipleSimOut.
63 /*
64 /*
65 *****
66 %avgMedMultipleSim2(itM=100,nsXnss=300);
67
68 /* Print avgMedMultipleSimOut */
69
70 proc print data=avgMedMultipleSimOut;
71
72 run;
73
74 /*
75 /* Plot avgMedMultipleSimOut */
76
77
78 options ftext=swiss fttext=swiss;
79 symbol value=square interpol=join width=0.5 color=blue;
80 symbol value=star interpol=join width=0.5 color=red;
81 symbol value=triangle interpol=join width=0.5 color=yellow;
82 symbol value='x' interpol=join width=0.5 color=orange;
83 symbol value=circle interpol=join width=0.5 color=blue;
84 symbol value=dot interpol=join width=0.5 color=green;
85 title2 height=1.4 "sample size vs Power";
86 axis1 order=(0.0 to 1.0 by 0.1) label=(angle=90 'Power') minor=none;
87 axis2 order=(5 to 50 by 5) label=(angle=0 'Sample Size');
88 legend label=none value=(h=.8 'upper AVG Estimated Power' 'True Power' 'AVG Estimated Power' 'Median Power' 'Lower
89 "" AVG Estimated Power' 'AVG P-Value')
90 position=center;
91
92
93

```

Appendix B: Two-Sample SAS Code for Power Simulations

```
94 proc qplot data=avgMedMultipleSimOut;
95   plot powerU*sampleSize truePower*sampleSize estPower*sampleSize medPower*sampleSize powerL*sampleSize
96   ... pvalue*sampleSize/overlay vaxis=axis1 haxis=axis2 hminor=4 legend=legend1;
96   by sigma;
97 run;
```

Appendix B: Two-Sample SAS Code for Power Simulations

```

1  /* multiplesim2 *****/
2  /*
3   * Requirement(s):
4   * simulation2.sas
5   */
6  /*
7   * Program Description:
8   * multiplesim2 Iterates simulation2.sas and combines simulation output to a
9   * dataset.
10 */
11 /*
12  * Input:
13  * mu1          := population one mean
14  * mu2          := population two mean
15  * simgamin     := minimum population standard deviation controller
16  * sigmaMax     := maximum population standard deviation controller
17  * itMax        := number of iterations
18  * sampleSizeMax := largest sample size considered
19  * sigLevel     := level of significance
20 */
21 /*
22  * Note(s):
23  * In the subsloop iterations run in blocks. This is done to
24  * traverse the dataset randSeed, where by each call to the macro
25  * simulation2 gets a different random seed position.
26  */
27 /*
28  * This macro contains two main loops. The first main loop
29  * iterates from the minimum user define population standard deviation to 19
30  * by 1. The second main loop iterates from 20 to the maximum user defined
31  * population standard deviation by 20. Each main loop has two subloops.
32  * The first subplot iterates from a system defined minimum sample size of 5
33  * to 14 by 1. The second subplot iterates from 15 to the maximum user
34  * defined sample size by 5. Each subplot has a subsubplot.
35  * Each subplot iterates by a user defined number of iterations. After
36  * a subplot executes, the starting position of an iteration 'block' is
37  * increased by the user defined number of iterations.
38  */
39 %macro multiplesim2(mu1=,mu2=,simgamin=,sigmaMax=,itMax=,sampleSizeMax=,sigLevel=);
40 %local std sd nn it block s n i b;
41 %let block = 1;
42 %let b = 1;
43 %let count = 1;
44
45
46
47

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

48 /* Produce simulations */
49 %do std = &sigmaMin %to 19 %by 1;
50   %let sd = %sysvalf(&std/2);
51   %do nn = 5 %to 14 %by 1;
52     %do it = &block %to &block+&itMax-1 %by 1;
53       %simulation2(muA=&mul1,muB=&mul2,sigma=&sd,sampleSize=&it,seedNum=&nn,alpha=&sigLevel,simNum=&count);
54       %let count = %eval(&count+1);
55     %end;
56   %end;
57   %let block = &it;
58 %end;
59 %do nn = 15 %to &sampleSizeMax %by 5;
60   %do it = &block %to &block+&itMax-1 %by 1;
61     %simulation2(muA=&mul1,muB=&mul2,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
62     %let count = %eval(&count+1);
63   %end;
64   %let block = &it;
65 %end;
66 %end;
67 %do std = 20 %to &sigmaMax %by 20;
68   %let sd = %sysvalf(&std/2);
69   %do nn = 5 %to 14 %by 1;
70     %do it = &block %to &block+&itMax-1 %by 1;
71       %simulation2(muA=&mul1,muB=&mul2,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
72       %let count = %eval(&count+1);
73     %end;
74     %let block = &it;
75   %end;
76   %do nn = 15 %to &sampleSizeMax %by 5;
77     %do it = &block %to &block+&itMax-1 %by 1;
78       %simulation2(muA=&mul1,muB=&mul2,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
79       %let count = %eval(&count+1);
80     %end;
81     %let block = &it;
82   %end;
83 %end;
84
85
86
87
88
89
90
91
92
93
94

```

Appendix B: Two-Sample SAS Code for Power Simulations

```
95 /* Combine simulations into a dataset and name columns */
96
97 data multipleSimOut;
98   set
99     %do k = 1 %to &count-1 %by 1;
100    sim&k
101   %end;
102   ; /* Double do-loop to reference datasets.  ';' closes 'set' */
103
104 rename col1=mu1 col2=mu2 col3=sigma col4=sampleSize col5=sigLevel col6=pValue col7=trueLambda col8=truePower
105 ...
106 col9=estLambda col10=adjEstLambda col11=lambda col12=lambdal col13=estPower col14=powerL col15=powerU;
107
108
109 %mend multipleSim2;
```

Appendix B: Two-Sample SAS Code for Power Simulations

```

1  /* simulation2 *****/
2  /*
3   * Requirement(s):
4   * sampleNormPop2.sas
5   */
6  /* Program Description:
7   * simulation2.sas performs a hypothesis
8   * test of no difference for two population means. Random samples
9   * are normally distributed with respective means muA and muB with equal
10  * population standard deviation sigma. Random samples are generated
11  * using seed values from the dataset randSeed. Power calculations
12  * are based on methods outlined in Thomas 1997. Calculations are
13  * saved to a dataset named 'sim#', where # represents the
14  * simulation count.
15  */
16  /* Input:
17  * muA      := mean of population one
18  * muB      := mean of population two
19  * sigma     := standard deviation of populations one and two
20  * seedNum   := position for random seed
21  * sampleSize := sample size taken for populations one and two
22  * alpha     := level of significance
23  * simNum    := simulation ID
24  */
25  /* Note(s):
26  * Optimal lower significance level by Golden Section Search method
27  */
28  *****
29
30  %MACRO simulation2(muA=,muB=,sigma=,seedNum=,sampleSize=,alpha=,simNum=);
31
32  /* Generate random samples from two normal populations */
33
34  %sampleNormPop2(popMean=&muA,popSD=&sigma,seedIt=&seedNum,numObs=&sampleSize,sampleNum=1);
35  %sampleNormPop2(popMean=&muB,popSD=&sigma,seedIt=&seedNum,numObs=&sampleSize,sampleNum=2);
36
37
38  /* Merge datasets sample1 and sample2 to produce dataset sampleTwoPops */
39  data sampleTwoPops;
40  merge sample1 sample2;
41
42  run;
43
44
45
46
47

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

48 /* Main */
49
50 proc iml;
51   reset nolog;
52   use sampleTwoPops;
53   read all into X;
54
55   /* Function *****/
56   /*
57   /* Return the confidence interval width for a noncentral
58   /* f-dstribution
59   /* */
60   /*
61   ****
62
63 start w(alphaLow,ndf,ddf,ncent,s1);
64
65   width = quantile('F',1-s1+alphaLow,ndf,ddf,ncent)-quantile('F',alphaLow,ndf,ddf,ncent);
66
67   return(width);
68
69   finish w;
70
71   /* Function: optimal lower critical region *****/
72   /*
73   /* Finds the minimum confidence interval width using an
74   /* implementation of the Golden Section Search method
75   /* similar to Press et al. (2007).
76   /*
77   /* Note(s):
78   /* The Golden Section Search method finds a value which minimizes
79   /* a function. The function must be one dimensional and unimodal.
80   /*
81   /* Imagine a 'typical' pdf of an F distribution. Say we want to
82   /* setup an interval such that the area covered by the sum of the two
83   /* tails is alpha. For each alpha we can find the
84   /* associated quantile using the SAS function 'quantile'.
85   /* Since alpha is specified, knowing the lower area easily leads to
86   /* the upper area. Here we consider only the lower
87   /* area. As such we can relate interval width as a function of
88   /* lower alpha size. As we increase or decrease the size of the
89   /* lower area we increase or decrease the width of the interval.
90   /* Assuming a minimum interval width exists, we optimize over the
91   /* values [0.00, 0.05].
92   /*
93   /**
94

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

95      start alphaIU(ndf1,ddf2,ncp,s1Alpha,to1);
96      alphaA = 0;
97      alphaB = s1Alpha;
98
99      /* Initial lower bracket on optimal region */
100     /* Initial upper bracket on optimal region */
101
102    /* Note 1 *****/
103    /* For each iteration an interval */
104    /* [alphaA, alphaB] is retained */
105    /* *****/
106
107    /* Golden ratio constant reduction factor */
108    /* Set test bracket a1 */
109    /* Calculate Interval width for a1 */
110    /* Set test bracket a2 */
111    /* Calculate interval width for a2 */
112
113    /* Note 2 *****/
114    /* The initial setup invokes two function */
115    /* calls. The following loop uses one */
116    /* function call per iteration. */
117
118
119
120 do while (abs(alphaB-alphaA)>tol);
121
122
123
124
125
126
127
128
129
130
131
132
133
134
135
136
137
138
139
140
141

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

142      /* Note 5 *****/
143      /*
144      /* Within the interval [alphaA, a1] we have */
145      /* the width wa2. a2 is situated by the */
146      /* expression alphaA+phi*(a1-alphaA). */
147      /*
148      *****/
149
150      alphaB = a1;
151      a1 = a2;
152      wa1 = wa2;
153      a2 = alphaA+phi*phi*(alphaB-alphaA);
154      wa2 = w(a2,ndf1,ddf2,ncp,s1Alpha);
155      end;
156      else do;
157      end;
158
159      /* Note 6 *****/
160      /*
161      /* Within the interval [a2, alphaB] we have */
162      /* the width wa1. a1 is situated by the */
163      /* expression alphaA+phi*phi*(alphaB-a2). */
164      /*
165      *****/
166
167      alphaA = a2;
168      a2 = a1;
169      wa2 = wa1;
170      a1 = alphaA+phi*(alphaB-alphaA);
171      wa1 = w(a1,ndf1,ddf2,ncp,s1Alpha);
172      end;
173
174      out = (alphaA+alphaB)/2;
175      return(out);
176
177      finish alphaLU;
178
179
180      /* Calculations */
181
182      eps = 1e-4;
183      n = nrow(X);
184      p = ncol(X);
185      J = J(p,1);
186
187      xBar = X(|+,|`/n;
188
189      /* Sample means */

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

189      muG = (&muA+&muB)/p;
190      sst = n*xBar`*xBar-(n/p)*(xBar`*J)**2;
191      df1 = p-1;
192      mst = sst/df1;
193      sse = sst(X)-(xBar`*xBar)*n;
194      df2 = p*(n-1);
195      mse = sse/df2;
196      fSamp = mst/mse;
197
198      fCrit = quantile('F',1-&alpha,df1,df2);
199
200      pValue = 1-cdf('F',fSamp,df1,df2);
201
202      trueLambda = (n*(((&muA-&muG)**2)+((&muB-&muG)**2))/((&sigma)**2));
203
204      truePower = 1-cdf('F',fCrit,df1,df2,trueLambda);
205
206      estLambda = fSamp*(p-1);
207      adjEstLambda = (estLambda*(df2-2)/df2)-df1;
208
209      if adjEstLambda < 0 then adjEstLambda = 0;
210
211      /* Note 7 *****/
212      /* For some random seeds the */
213      /* noncentrality parameter is */
214      /* negative. */
215      /* A negative noncentrality */
216      /* parameter occurs when */
217      /* the expression */
218      /* estLambda*(df2-2)/df2 is */
219      /* less than df1. */
220
221      /* *****/
222
223      alphaL = alphalU(df1,df2,adjEstLambda,&alpha,eps);
224      alphaU = &alpha-&alphaL;
225
226
227
228
229
230
231
232
233
234
235

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

236 adjAlphaL = 1-alphaL;
237 probCenF = cdf( 'F' ,fsamp,df1,df2);
238
239 /* fnonct reports an error for a supplied */
240 /* probability greater than the */
241 /* probability from the cdf of a central */
242 /* F distribution.
243 */
244 /* As the probability supplied gets large */
245 /* fnonct tends toward zero. probcenF */
246 /* with fnonct is small around zero.
247 /* Using adjAlphaL would then be smaller */
248 /* about zero. Set to zero.
249 */
250
251
252 if adjAlphaL > probcenF then lambdaL = 0; /* See note 8 */
253 else lambdaL = fnonct(fsamp,df1,df2,adjAlphaL); /* See note 8 */
254 if alphaU > probcenF then lambdaU = 0;
255 else lambdaU = fnonct(fsamp,df1,df2,alphaU); /* See note 8 */
256
257 estPower = 1-cdf( 'F' ,fcrit,df1,df2,adjEstLambda);
258 powerL = 1-cdf( 'F' ,fcrit,df1,df2,lambdaL);
259 powerU = 1-cdf( 'F' ,fcrit,df1,df2,lambdaU);
260
261
262 /* Output simulation to a data set */
263
264 Y = {"&muA" "&muB" "&sigma" "&sampleSize" "&alpha" "."
265 "."
266 "."
267 "."
268 Y[6] = pvalue;
269 Y[7] = trueLambda;
270 Y[8] = truePower;
271 Y[9] = estLambda;
272 Y[10] = adjEstLambda;
273 Y[11] = lambdaL;
274 Y[12] = lambdaU;
275 Y[13] = estPower;
276 Y[14] = powerL;
277 Y[15] = powerU;
278
279 create simsimNum from Y;
280 append from Y;
281
282 quit;

```

Appendix B: Two-Sample SAS Code for Power Simulations

```
283 proc datasets lib=work nolist;
284   delete sample1 sample2 sampleTwoPops;
285   quit;
286   run;
287
288 %MEND simulation2;
289
```

Appendix B: Two-Sample SAS Code for Power Simulations

```

1 /* sampleNormPop2 *****/
2 /*
3 /* Program Description:
4 /* sampleNormPop2 generates a random sample from a normal population
5 /* and outputs the values to a dataset
6 /*
7 /* Input:
8 /* seed      := dataset of random seeds from randSeed
9 /* popMean   := population mean
10 /* popSD    := population standard deviation
11 /* seedIt    := seed value position for generating random observations
12 /* numObs   := sample size
13 /* sampleNum := identifies sample
14 /*
15 /* *****/
16 /*
17 %macro sampleNormPop2(popMean=,popSD=,seedIt=,numObs=,sampleNum=);
18 /*
19 proc iml;
20   reset nolog;
21   use randSeed;
22   read all into seed;
23   /*
24   x&sampleNum = &popMean+&popSD*rannor(J(&numObs,1,seed[&seedIt,&sampleNum]));
25   test = seed[&seedIt,&sampleNum];
26   /*
27   create sample&sampleNum from x&sampleNum[colname={obs&sampleNum}];
28   append from x&sampleNum;
29   /*
30   quit;
31   /*
32   %mend sampleNormPop2;
33 /*

```

Appendix B: Two-Sample SAS Code for Power Simulations

```

1  /* avgMedMultipleSim2 *****/
2  /*
3   * Requirement(s):
4   * avgMedSim2.sas
5   */
6  /* Program Description:
7   * avgMedMultipleSim2 calls avgMedSim2 for each sigma by sample size */
8  /* combination.
9   */
10 /* Input:
11  * itM          := number of iterations
12  * nsXnss       := number of iteration groups
13  *                  (sigmas times number of sample sizes)
14  */
15 /* *****/
16 /* *****/
17
18 %macro avgMedMultipleSim2(itM=,nsXnss=);
19
20 /* Produce average and median of simulations */
21
22 %let bStart = 1;
23 %let bFinish = &itM;
24
25 %do i = 1 %to &nsXnss %by 1;
26   %avgMedSim2(start=&bStart,finish=&bFinish,itB=&i);
27   %let bStart = &bStart+&itM;
28   %let bFinish = &bFinish+&itM;
29
30
31 /* Combine average simulations into a data set and name columns */
32
33 data avgMedMultipleSimOut;
34
35 set
36   %do j = 1 %to &nsXnss %by 1;
37   ansim&j
38   %end;
39   /* Double do-loop to reference data sets. ';' closes 'set' */
40   rename col1=mu1 col2=mu2 col3=sigma col4=sampleSize col5=sigLevel col6=pValue col7=trueLambda col8=truePower
41   col9=estLambda col10=adjEstLambda col11=lambda col12=lambdaD col13=estPower col14=powerL col15=powerU
42   ...
43
44 %mend avgMedMultipleSim2;
45

```

Appendix B: Two-Sample SAS Code for Power Simulations

Appendix B: Two-Sample SAS Code for Power Simulations

```

48      nr = nrow(mis);
49
50      /* Number of missing values */
51
52      /* Note 2 *****/
53      /* If 'nr' is greater than */
54      /* one then we have missing */
55      /* values. These values are */
56      /* removed from the column. */
57
58      /* Remove initialize position */
59      /* Transpose a column to row */
60      /* Remove missing values */
61      /* Transpose row to a column */
62
63      /* Find median */
64
65      med = median(col);
66
67      /* Concatinate median to average row */
68      avgRows=avgRows || med;
69
70
71      /* Output to a dataset */
72      create amSim&itB from avgRows;
73      append from avgRows;
74
75      quit;
76      %mend avgMedSim2;
77

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

1  /* threeSample *****/
2  /*
3   * Requirement(s):
4   * rand3.csv, multipleSim3.sas, avgMedMultipleSim3.sas
5   */
6  /* Program Description:
7   * threeSample.sas acts as a controller interface which accepts user
8   * defined values for three population means, upper and lower bounds
9   * for a range of population standard deviations, a maximum sample size,
10  /* a type I error rate, and the number of iterations per sample size
11  /* to population standard deviation combination.
12  */
13 /*******/
14
15 proc printto print='\\Path\\to\\Output\\Folder\\threeSampleOut.lst' log='\\Path\\to\\Output\\Folder\\threeSampleOut.log';
16
17 run;
18
19
20 /* Input random seed file to the dataset randseed *****/
21 /*
22 */
23 /* rand3.csv is a three column file of random numbers generated from
24 /* http://www.random.org. For each iteration of multipleSim3 three random
25 /* values are used to seed three random samples.
26 */
27 /*******/
28
29 data randSeed;
30   infile "\\Path\\to\\File\\rand3.csv" dlm=' , ';
31   input seedCol1 seedCol2 seedCol3;
32 run;
33
34 /* Produce simulations *****/
35 /*
36 */
37 /* The multipleSim3 manages the production of simulations.
38 /* Calls are made to the macro simulation3.sas which performs
39 /* calculations based on Thomas 1997.
40 */
41 /* Calculations for each simulation are stored in a dataset called
42 /* multipleSimOut.
43 */
44 /*******/
45 %multipleSim3(mu1=30,mu2=40,mu3=35,sigmaMin=5,sigmaMax=50,itMax=5,sampleSizeMax=50,sigLevel=0.05);
46
47

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

48 /* Print multipleSimOut */
49
50 proc print data=multipleSimOut;
51 run;
52
53
54 /* Produce averages and medians for simulations *****/
55 /*
56 /*
57 /* The avgMedMultipleSim3 manages the consolidation of simulations.
58 /* Calls are made to avgMedSim3 for each sigma by sample size
59 /* combination. avgMedSim3 averages over a user specified number
60 /* of rows from multipleSimOut. avgMedSim3 also determines the
61 /* median of the calculated power values of the rows.
62 /*
63 /* Calculations for each iteration block are stored in a dataset
64 /* called avgMedMultipleSimOut.
65 /*
66 *****/
67 %avgMedMultipleSim3(itM=5,nsXnss=306);
68
69
70 /* Print avgMedMultipleSimOut */
71
72 proc print data=avgMedMultipleSimOut;
73 run;
74
75
76 /* Plot avgMedMultipleSimOut */
77
78 options ftitle=swiss ftext=swiss;
79 symbol1 value=square interpol=join width=0.5 color=blue;
80 symbol2 value=star interpol=join width=0.5 color=red;
81 symbol3 value=triangle interpol=join width=0.5 color=yellow;
82 symbol4 value='x' interpol=join width=0.5 color=orange;
83 symbol5 value=circle interpol=join width=0.5 color=blue;
84 symbol6 value=dot interpol=join width=0.5 color=green;
85 title2 height=1.4 "Sample Size vs Power";
86 axis1 order=(0.0 to 1.0 by 0.1) label=(angle=90 'Power') minor=none;
87 axis2 order=(5 to 50 by 5) label=(angle=0 'Sample Size');
88 legend label=none value=(h=.8 'Upper Avg Estimated Power' 'True Power' 'AVG Estimated Power' 'Median Power' 'Lower
... AVG Estimated Power' 'AVG P-Value')
position=center;
89
90
91
92
93

```

Appendix B: Three-Sample SAS Code for Power Simulations

```
94 proc qplot data=avgMedMultipleSimOut;
95   plot powerU*sampleSize truePower*sampleSize estPower*sampleSize medPower*sampleSize powerL*sampleSize
96   ... pvalue*sampleSize/overlay vaxis=axis1 haxis=axis2 hminor=4 legend=legend1;
96   by sigma;
97 run;
98
```

Appendix B: Three-Sample SAS Code for Power Simulations

```

1 /* multiplesim3 *****/
2 /*
3 /* Requirement(s):
4 /* simulation3.sas
5 /*
6 /* Program Description:
7 /* multiplesim3 Iterates simulation3.sas and combines simulation output to a */
8 /* dataset.
9 /*
10 /* Input:
11 /* mu1      := population one mean
12 /* mu2      := population two mean
13 /* mu3      := population three mean
14 /* sigmaMin := minimum population standard deviation controller
15 /* sigmaMax := maximum population standard deviation controller
16 /* itMax    := number of iterations
17 /* sampleSizeMax := largest sample size considered
18 /* sigLevel1 := level of significance
19 /*
20 /* Note(s):
21 /* In the subsubloop iterations run in blocks. This is done to
22 /* traverse the dataset randSeed, where by each call to the macro
23 /* simulation2 gets a different random seed position.
24 /*
25 /* This macro contains two main loops. The first main loop
26 /* iterates from the minimum user define population standard deviation to 19
27 /* by 1. The second main loop iterates from 20 to the maximum user defined
28 /* population standard deviation by 20. Each main loop has two subloops.
29 /* The first subloop iterates from 5 to the maximum sample size of 5
30 /* to 14 by 1. The second subloop iterates from 15 to the maximum user
31 /* defined sample size by 5. Each subloop has a subsubloop.
32 /* Each subsubloop iterates by a user defined number of iterations. After
33 /* a subsubloop executes, the starting position of an iteration 'block' is
34 /* increased by the user defined number of iterations.
35 /*
36 /******
37 /*
38 %macro multiplesim3(mu1=,mu2=,mu3=,sigmaMin=,sigmaMax=,itMax=,sampleSizeMax=,sigLevel1=);
39 /*
40 %local std sd nn it block s n i b;
41 %let block = 1;
42 %let b = 1;
43 %let count = 1;
44 /*
45 /*
46 /*
47 
```

Appendix B: Three-Sample SAS Code for Power Simulations

```

48 /* Produce simulations */
49
50 %do std = &sigmaMin %to 19 %by 1;
51   %let sd = %sysvalf(&std/2);
52   %do nn = 5 %to 14 %by 1;
53     %do it = &block %to &block+&itMax-1 %by 1;
54
55   %simulation3(muA=&mu1,muB=&mu2,muC=&mu3,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
56   ...
57   %let count = %eval(&count+1);
58
59   %let block = &it;
60
61   %do nn = 15 %to &sampleSizeMax %by 5;
62     %do it = &block %to &block+&itMax-1 %by 1;
63
64   %simulation3(muA=&mu1,muB=&mu2,muC=&mu3,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
65   ...
66   %let count = %eval(&count+1);
67
68   %end;
69   %do std = 20 %to &sigmaMax %by 20;
70     %let sd = %sysvalf(&std/2);
71     %do nn = 5 %to 14 %by 1;
72       %do it = &block %to &block+&itMax-1 %by 1;
73
74   %simulation3(muA=&mu1,muB=&mu2,muC=&mu3,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
75   ...
76   %let count = %eval(&count+1);
77
78   %let block = &it;
79
80   %do nn = 15 %to &sampleSizeMax %by 5;
81     %do it = &block %to &block+&itMax-1 %by 1;
82
83   %simulation3(muA=&mu1,muB=&mu2,muC=&mu3,sigma=&sd,seedNum=&it,sampleSize=&nn,alpha=&sigLevel,simNum=&count);
84   ...
85
86
87
88
89
90

```

Appendix B: Three-Sample SAS Code for Power Simulations

```
91 /* Combine simulations into a dataset and name columns */
92
93 data multipleSimOut;
94   set
95     %do k = 1 %to &count-1 %by 1;
96     sim&k
97   %end;
98   /* Double do-loop to reference datasets.  ';' closes 'set' */
99
100 rename col1=mu1 col2=mu2 col3=mu3 col4=sigma col5=sampleSize col6=sigLevel col7=pValue col8=trueLambda
101 col9=truePower col10=estLambda col11=adjEstLambda col12=lambda col13=lambda col14=estPower col15=powerL
102 col16=powerU;
103
104 run;
105
106 %mend multipleSim;
```

Appendix B: Three-Sample SAS Code for Power Simulations

```

1  /* simulation3 *****/
2  /*
3  /* Requirement(s):
4  /* sampleNormPop3.sas
5  /*
6  /* Program Description:
7  /* simulation3.sas performs a hypothesis
8  /* test of no difference for three population means. Random samples
9  /* are normally distributed with respective means muA, muB and muC with equal
10 /* population standard deviation sigma. Random samples are generated
11 /* using seed values from the dataset randSeed.
12 /* are based on methods outlined in Thomas 1997. Calculations are
13 /* saved to a dataset named 'sim#' where # represents the
14 /* simulation count.
15 /*
16 /* Input:
17 /* muA          := mean of population one
18 /* muB          := mean of population two
19 /* muC          := mean of population three
20 /* sigma         := standard deviation of populations one, two, and three
21 /* seedNum       := position for random seed
22 /* samplesize    := sample size taken for populations one, two, and three
23 /* alpha         := level of significance
24 /* simnum        := simulation ID
25 /*
26 /* Note(s):
27 /* Optimal lower significance level by Golden Section Search method
28 /*
29 /***** *****
30 /*
31 %MACRO simulation3(muA=,muB=,muC=,sigma=,seedNum=,samplesize=,alpha=,simNum=);
32 /*
33 /* Generate random samples from three normal populations */
34 /*
35 %sampleNormPop3(popMean=&muA,popSD=&sigma,seedIt=&seedNum,numObs=&sampleSize,sampleNum=1);
36 %sampleNormPop3(popMean=&muB,popSD=&sigma,seedIt=&seedNum,numObs=&sampleSize,sampleNum=2);
37 %sampleNormPop3(popMean=&muC,popSD=&sigma,seedIt=&seedNum,numObs=&sampleSize,sampleNum=3);
38 /*
39 /*
40 /* Merge datasets sample1, sample2 and sample3 to produce dataset sampleThreePops */
41 /*
42 data sampleThreePops;
43 merge sample1 sample2 sample3;
44 run;
45 /*
46 /*
47 /*
48 /*

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

49 /* Main */
50
51 proc iml;
52   reset nolog;
53   use sampleThreePops;
54   read all into X;
55
56 /* Function *****/
57 /*
58  * Return the confidence interval width for a noncentral
59  * f-distribution
60  */
61 /*
62  */
63 start w(alphalow,ndf,ddf,ncent,s1);
64
65   width = quantile('F',1-s1+alphalow,ndf,ddf,ncent)-quantile('F',alphalow,ndf,ddf,ncent);
66
67   return(width);
68
69 finish w;
70
71 /* Function: optimal lower critical region *****/
72 /*
73  */
74 /* Finds the minimum confidence interval width using an
75  * implementation of the Golden Section Search method
76  * similar to Press et al. (2007). */
77 /*
78  */
79 /* Note(s):
80  * The Golden Section Search method finds a value which minimizes
81  * a function. The function must be one dimensional and unimodal.
82  */
83 /* Imagine a 'typical' pdf of an F distribution. Say we want to
84  * setup an interval such that the area covered by the sum of the two
85  * tails is alpha. For each alpha we can find the
86  * associated quantile using the SAS function 'quantile'.
87  */
88 /* Since alpha is specified, knowing the lower area easily leads to
89  * the upper area. Here we consider only the lower
90  */
91 /* area. As such we can relate interval width as a function of
92  * lower alpha size. As we increase or decrease the size of the
93  * lower area we increase or decrease the width of the interval.
94  */
95 /* Assuming a minimum interval width exists, we optimize over the
96  */
97 /*
98  */
99 /*
100 */
101 /*
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292 /*
293 */
294 /*
295 */
296 /*
297 */
298 /*
299 */
299 */

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

97      start alphaLU(ndf1,ddf2,ncp,sAlpha,to1);
98
99      alphaA = 0;
100     alphaB = sAlpha;
101
102     /* Initial lower bracket on optimal region */
103     /* Initial upper bracket on optimal region */
104
105     /* Note 1 *****/
106     /* For each iteration an interval */
107     /* [alphaA, alphaB] is retained */
108
109     /* *****/
110     /* Golden ratio constant reduction factor */
111     /* Set test bracket a1 */
112     /* Calculate Interval width for a1 */
113     /* Set test bracket a2 */
114     /* Calculate interval width for a2 */
115
116     /* Note 2 *****/
117     /* */
118     /* The initial setup invokes two function */
119     /* calls. The following loop uses one */
120     /* function call per iteration. */
121
122     /* *****/
123
124     do while (abs(alphaB-alphaA)>tol);
125
126     /* We continue until the absolute difference */
127     /* between an upper and a lower bracket is */
128     /* negligible. Tolerance is set tol=4. */
129
130     /* *****/
131
132     /* Note 4 */
133
134     /* Four brackets are maintained at any */
135     /* given time. */
136
137
138     if wa1>wa2 then do;
139
140     /* If wa1 > wa2 then minimum width is between */
141     /* alphaA and a1 */
142
143
144

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

145      /* Note 5 *****/
146      /*
147      /* Within the interval [alphaA, a1] we have */
148      /* the width wa1. a2 is situated by the */
149      /* expression alphaA+phi*(a1-alphaA). */
150      /*
151      /*
152      *****/
153
154      alphaB = a1;
155      a1 = a2;
156      wa1 = wa2;
157      a2 = alphaA+phi*phi*(alphaB-alphaA);
158      wa2 = w(a2,ndf1,ddf2,ncp,s1Alpha);
159
160      else do;
161
162
163      /* Note 6 *****/
164      /*
165      /* Within the interval [a2, alphaB] we have */
166      /* the width wal. a1 is situated by the */
167      /* expression alphaA+phi*phi*(alphaB-a2). */
168      /*
169      *****/
170
171      alphaA = a2;
172      a2 = a1;
173      wa2 = wal;
174      a1 = alphaA+phi*(alphaB-alphaA);
175      wal = w(a1,ndf1,ddf2,ncp,s1Alpha);
176
177      end;
178
179      out = (alphaA+alphaB)/2;
180
181      return(out);
182
183      finish alphalU;
184
185      /* Calculations */
186
187      eps = 1e-4;
188      n = nrow(X);
189      p = ncol(X);
190      J = J(p,1);
191      muG = (fmuA+fmuB+fmuC)/p;
192      xBar = x(|+,|`/n;

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

193   sst = n*xBar`*xBar - (n/p)*(xBar`*J)**2;
194   df1 = p-1;
195   mst = sst/df1;
196   sse = ssq(X)-(xBar`*xBar)*n;
197   df2 = p*(n-1);
198   mse = sse/df2;
199   fSamp = mst/mse;
200
201   fcrit = quantile('F',1-&alpha,df1,df2);
202
203   pValue = 1-cdf('F',fSamp,df1,df2);
204
205   trueLambda = (n*(((&muA-&muG)**2)+((&muB-&muG)**2)+((&muC-&muG)**2))/((&sigma)**2)); /* trueLambda */
206
207   truePower = 1-cdf('F',fcrit,df1,df2,trueLambda); /* truePower */
208
209   estLambda = fSamp*(p-1);
210   adjEstLambda = (estLambda*(df2-2)/df2)-df1;
211
212   if adjEstLambda < 0 then adjEstLambda = 0;
213
214
215
216
217
218
219
220
221
222
223
224
225
226
227
228
229
230
231
232
233
234
235
236
237
238
239
240

```

```

/* Sum of squares for treatments */
/* Degrees of freedom for treatment */
/* Mean square for treatment */
/* Sum of squares for error */
/* Degrees of freedom for error */
/* Mean square for error */
/* Observed F */

/* F critical value */

/* P-Value */

/* truePower */

/* Estimated lambda */
/* Adjusted estimated lambda */

/* Note 7 **** */
/* For some random seeds the */
/* noncentrality parameter is */
/* negative. */
/* A negative noncentrality */
/* parameter occurs when */
/* the expression */
/* estLambda*(df2-2)/df2 is */
/* less than df1. */

/* Lower significance */
/* Upper significance */

alphaL = alphaLU(df1,df2,adjEstLambda,&alpha,eps);
alphaU = &alpha-alphaL;

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

241 adjAlphaL = 1-alphaL;
242 probCenF = cdf('F',fsamp,df1,df2);
243
244 /* fnonct reports an error for a supplied */
245 /* probability greater than the */
246 /* probability from the cdf of a central */
247 /* F distribution. */
248 /* As the probability supplied gets large */
249 /* fnonct tends toward zero. probCenF */
250 /* with fnonct is small around zero. */
251 /* Using adjAlphaL would then be smaller */
252 /* about zero. Set to zero. */
253 /* */
254 /* **** */
255
256 if adjAlphaL > probCenF then lambdaL = 0;
257 else lambdaL = fnonct(fsamp,df1,df2,adjAlphaL);
258 if alphaU > probCenF then lambdaU = 0;
259 else lambdaU = fnonct(fsamp,df1,df2,alphaU);
260
261 estPower = 1-cdf('F',fcrit,df1,df2,adjBstLambda);
262 powerL = 1-cdf('F',fcrit,df1,df2,lambdaL);
263 powerU = 1-cdf('F',fcrit,df1,df2,lambdaU);
264
265 /* Output simulation to a data set */
266
267 /* Output simulation to a data set */
268 Y = {"&muA" "&muB" "&muC" "&sigma" "&sampleSize" "&alpha"};
269 Y = num(Y);
270
271 Y[7] = pValue;
272 Y[8] = trueLambda;
273 Y[9] = estPower;
274 Y[10] = estLambda;
275 Y[11] = adjBstLambda;
276 Y[12] = lambdaL;
277 Y[13] = lambdaU;
278 Y[14] = estPower;
279 Y[15] = powerL;
280 Y[16] = powerU;
281
282 create sim&simNum from Y;
283 append from Y;
284
285 quit;
286
287
288

```

Appendix B: Three-Sample SAS Code for Power Simulations

```
289 proc datasets lib=work nolist;
290   delete sample1 sample2 sample3 sampleThreePops;
291   quit;
292   run;
293
294 %MEND simulation3;
295
```

Appendix B: Three-Sample SAS Code for Power Simulations

```

1 /* sampleNormPop3 **** */
2 /*
3 /* Program Description:
4 /* sampleNormPop3 generates a random sample from a normal population
5 /* and outputs the values to a dataset
6 /*
7 /* Input:
8 /* seed      := dataset of random seeds from randSeed
9 /* popMean   := population mean
10 /* popSD    := population standard deviation
11 /* seedIt    := seed value position for generating random observations
12 /* numObs   := sample size
13 /* sampleNum := identifies sample
14 /*
15 /* **** */
16 /*
17 %macro sampleNormPop3 (popMean=,popSD=,seedIt=,numObs=,sampleNum=);
18 /*
19 proc iml;
20   reset nolog;
21   use randSeed;
22   read all into seed;
23   x&sampleNum = &popMean+&popSD*ranmor(J(&numObs,1,seed[&seedIt,&sampleNum]));
24   test = seed[&seedIt,&sampleNum];
25   create sample&sampleNum from x&sampleNum[colname={obs&sampleNum}];
26   append from x&sampleNum;
27   quit;
28   %mend sampleNormPop3;

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

1  /* avgMedMultipleSim3 *****/
2  /*
3   * Requirement(s):
4   * avgMedSim3.sas
5   */
6  /* Program Description:
7   * avgMedMultipleSim3 calls avgMedSim3 for each sigma by sample size */
8  /* combination.
9   */
10 /* Input:
11  /* itM          := number of iterations
12  /* nsXnss       := number of iteration groups
13  /*             (sigmas times number of sample sizes)
14  */
15  */
16  */
17
18 %macro avgMedMultipleSim3(itM=,nsXnss=);
19
20 /* Produce average and median of simulations */
21
22 %let bStart = 1;
23 %let bFinish = &itM;
24
25 %do i = 1 %to &nsXnss %by 1;
26   %avgMedSim3(start=&bStart,finish=&bFinish,itB=&i);
27   %let bStart = &bStart+&itM;
28   %let bFinish = &bFinish+&itM;
29
30
31 /* Combine average simulations into a data set and name columns */
32
33 data avgMedMultipleSimOut;
34
35 set
36   %do j = 1 %to &nsXnss %by 1;
37     ansim&j
38   %end;
39   ; /* Double do-loop to reference data sets. ';' closes 'set' */
40   rename col1=mu1 col2=mu2 col3=mu3 col4=sigma col5=sampleSize col6=sigLevel col7=pValue col8=trueLambda
41   col9=truePower col10=estLambda col11=adjEstLambda col12=lambda col13=1ambda col14=estPower col15=powerL
42   ...
43
44 %mend avgMedMultipleSim3;
45

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

1  /* avgMedSim3 **** */
2  /*
3   * Program Description:
4   * avgMedSim3 averages over a user specified number of rows from
5   * multiplesimOut. avgMedSim3 also determines the median
6   * of the calculated power values of the rows.
7   */
8  /* Input:
9   * X          := gets dataset multipleSimOut
10  /* start     := beginning of iteration block
11  /* finish    := end of iteration block
12  /* itB       := number of iterations
13  */
14  **** */
15
16 %macro avgMedSim3(start=&finish=,itB=);
17
18 proc iml;
19
20 reset nolog;
21 use multipleSimOut;
22 read all into X;
23
24 /* Average rows for an iteration block */
25
26 rows = X[&start:&finish,];
27 avgRows = rows[:,];
28
29
30 /* Median for column calculated power */
31 col = X[&start:&finish,14];
32
33 numRows = nrow(col);
34
35
36 /* Storage for positions of missing values */
37 mis = 0;
38 i = 1;
39
40 do while (i <=numRows);
41   if missing(col[i,1]) > 0 then mis=mis/i;
42   i=i+1;
43 end;
44
45
46
47 **** */

```

Appendix B: Three-Sample SAS Code for Power Simulations

```

48      nr = nrow(mis);
49
50      /* Number of missing values */
51      /* Note 2 *****/
52      /*
53      /* If 'nr' is greater than */
54      /* one then we have missing */
55      /* values. These values are */
56      /* removed from the column. */
57      /*
58      ****
59
60      if nr>1 then mis = remove(mis,1);
61      if nr>1 then col = col`;
62      if nr>1 then col = remove(col,mis);
63      if nr>1 then col = col`;
64
65      med = median(col);
66
67      /* Concatinate median to average row */
68      avgRows=avgRows || med;
69
70      avgRows=avgRows || med;
71
72      create amSim&itB from avgRows;
73      append from avgRows;
74
75      quit;
76      %mend avgMedSim3;
77

```

Appendix B: Four-Sample SAS Code for Power Simulations

```

1  /* fourSample *****/
2  /*
3   * Requirement(s):
4   * rand4.csv, multipleSim4.sas, avgMedMultipleSim4.sas
5   */
6  /* Program Description:
7   * fourSample.sas acts as a controller interface which accepts user
8   * defined values for four population means, upper and lower bounds
9   * for a range of population standard deviations, a maximum sample size,
10  /* a type I error rate, and the number of iterations per sample size
11  /* to population standard deviation combination.
12  */
13 *****/
14
15 proc printto print='\\Path\\to\\Output\\Folder\\fourSampleOut.lst'
16 log='\\Path\\to\\Output\\Folder\\fourSampleOut.log';
18 run;
19
20
21 /* Input random seed file to the dataset randseed *****/
22 /*
23 /*
24 /* rand4.csv is a four column file of random numbers generated from
25 /* http://www.random.org. For each iteration of multipleSim4 four random
26 /* values are used to seed four random samples.
27 */
28 *****/
29
30 data randSeed;
31 infile "\\Path\\to\\File\\rand4.csv" dlm=',';
32 input seedCol1 seedCol2 seedCol3 seedCol4;
33 run;
34
35 /* Produce simulations *****/
36 /*
37 /*
38 /* The multipleSim4 manages the production of simulations.
39 /* Calls are made to the macro simulation4.sas which performs
40 /* calculations based on Thomas 1997.
41 /*
42 /* Calculations for each simulation are stored in a dataset called
43 /* multipleSimOut.
44 /*
45 *****/
46 %multipleSim4(mu1=30,mu2=40,mu3=35,mu4=35,sigmaMin=5,sigmaMax=40,itMax=100,sampleSizeMax=50,sigLevel=0.05);
47

```

Appendix B: Four-Sample SAS Code for Power Simulations

```

48
49 /* Print multipleSimOut */
50
51 proc print data=multipleSimOut;
52 run;
53
54
55 /* Produce averages and medians for simulations *****/
56 /*
57 /*
58 /* The avgMedMultipleSim4 manages the consolidation of simulations.
59 /* Calls are made to avgMedSim4 for each sigma by sample size
60 /* combination. avgMedSim4 averages over a user specified number
61 /* of rows from multipleSimOut. avgMedSim4 also determines the
62 /* median of the calculated power values of the rows.
63 /*
64 /* Calculations for each iteration block are stored in a dataset
65 /* called avgMedMultipleSimOut.
66 /*
67 *****/
68
69 %avgMedMultipleSim4(itM=100,nsXnss=306);
70
71 /* Print avgMedMultipleSimOut */
72
73 proc print data=avgMedMultipleSimOut;
74 run;
75
76
77 /* Plot avgMedMultipleSimOut */
78
79 options ftitle=swiss ftext=swiss;
80 symbol1 value=square interpol=join width=0.5 color=blue;
81 symbol2 value=star interpol=join width=0.5 color=red;
82 symbol3 value=triangle interpol=join width=0.5 color=yellow;
83 symbol4 value='x' interpol=join width=0.5 color=orange;
84 symbol5 value=circle interpol=join width=0.5 color=blue;
85 symbol6 value=dot interpol=join width=0.5 color=green;
86 title2 height=1.4 "Sample Size vs Power";
87 axis1 order=(0.0 to 1.0 by 0.1) label=(angle=90 'Power') minor=none;
88 axis2 order=(5 to 50 by 5) label=(angle=0 'Sample Size');
89 legend label=none value=(h=.8 'Upper AVG Estimated Power' 'True Power' 'AVG Estimated Power' 'Median Power'
90 'Lower AVG Estimated Power' 'AVG P-Value')
91 position=center;
92
93
94

```

Appendix B: Four-Sample SAS Code for Power Simulations

```
95 proc qplot data=avgMedMultipleSimOut;
96   plot powerU*sampleSize truePower*sampleSize estPower*sampleSize powerL*sampleSize
97   pValue*sampleSize/overlay vaxis=axis1 haxis=axis2 hminor=4 legend=legend1;
98   by sigma;
99   run;
100
```

Appendix B: Four-Sample SAS Code for Power Simulations

```

1 /* multipleSim4 **** */
2 /*
3 /* Requirement(s):
4 /* simulation4.sas
5 /*
6 /* Program Description:
7 /* multipleSim4 Iterates simulation4.sas and combines simulation output to a
8 /* dataset.
9 /*
10 /* Input:
11 /* mu1      := population one mean
12 /* mu2      := population two mean
13 /* mu3      := population three mean
14 /* mu4      := population four mean
15 /* sigmaMin := minimum population standard deviation controller
16 /* sigmaMax := maximum population standard deviation controller
17 /* itMax    := number of iterations
18 /* sampleSizeMax := largest sample size considered
19 /* sigLevel := level of significance
20 /*
21 /* Note(s):
22 /* In the subsubloop iterations run in blocks. This is done to
23 /* traverse the dataset randseed, where by each call to the macro
24 /* simulation4 gets a different random seed position.
25 /*
26 /* This macro contains two main loops. The first main loop
27 /* iterates from the minimum user define population standard deviation to 19
28 /* by 1. The second main loop iterates from 20 to the maximum user defined
29 /* population standard deviaiton by 20. Each main loop has two subloops.
30 /* The first subloop iterates from a system defined minimum sample size of 5
31 /* to 14 by 1. The second subloop iterates from 15 to the maximum user
32 /* defined sample size by 5. Each subloop has a subsubloop.
33 /* Each subsubloop iterates by a use defined number of iterations. After
34 /* a subsubloop executes, the starting position of an iteration 'block' is
35 /* increased by the the user defined number of iterations.
36 /*
37 ****
38 /*
39 %macro multipleSim4(mu1=,mu2=,mu3=,mu4=,sigmaMin=,sigmaMax=,itMax=,sampleSizeMax=,sigLevel=);
40 /*
41 %local std sd nn it block s n i b;
42 %let block = 1;
43 %let b = 1;
44 %let count = 1;
45 /*
46

```

Appendix B: Four-Sample SAS Code for Power Simulations

```

48 /* Produce simulations */
49 %do std = &sigmaMin %to 19 %by 1;
50   %let sd = %sysvalf(&std/2);
51   %do nn = 5 %to 14 %by 1;
52     %do it = &block %to &block+&itMax-1 %by 1;
53
54   %simulation4(muA=&mu1,muB=&mu2,muC=&mu3,muD=&mu4,sigma=&sd,seedNum=&it,sampleSize=&it,seedNum=&sigLevel,simNum=&count)
55   ...
56   %let count = %eval(&count+1);
57   %end;
58   %let block = &it;
59   %end;
60   %do nn = 15 %to &sampleSizeMax %by 5;
61     %do it = &block %to &block+&itMax-1 %by 1;
62
63   %simulation4(muA=&mu1,muB=&mu2,muC=&mu3,muD=&mu4,sigma=&sd,seedNum=&it,sampleSize=&it,seedNum=&sigLevel,simNum=&count)
64   ...
65   %end;
66   %let count = %eval(&count+1);
67   %end;
68   %let block = &it;
69   %end;
70   %do it = &block %to &block+&itMax-1 %by 1;
71
72   %simulation4(muA=&mu1,muB=&mu2,muC=&mu3,muD=&mu4,sigma=&sd,seedNum=&it,sampleSize=&it,seedNum=&sigLevel,simNum=&count)
73   ...
74   %let count = %eval(&count+1);
75   %end;
76   %do nn = 15 %to &sampleSizeMax %by 5;
77     %do it = &block %to &block+&itMax-1 %by 1;
78
79   %simulation4(muA=&mu1,muB=&mu2,muC=&mu3,muD=&mu4,sigma=&sd,seedNum=&it,sampleSize=&it,seedNum=&sigLevel,simNum=&count)
80   ...
81   %let count = %eval(&count+1);
82   %end;
83   %end;
84
85
86

```

Appendix B: Four-Sample SAS Code for Power Simulations

```
87 /* Combine simulations into a dataset and name columns */
88
89 data multipleSimOut;
90   set
91     %do k = 1 %to &count-1 %by 1;
92     sim&k
93   %end;
94   ; /* Double do-loop to reference datasets.  ';' closes 'set' */
95
96 rename col1=mu1 col2=mu2 col3=mu3 col4=mu4 col5=sigma col6=sampleSize col7=sigLevel col8=pValue
97 col9=trueLambda col10=truePower col11=estLambda col12=adjEstLambda col13=lambdaU col14=lambdaD col15=estPower
98 col16=powerL col17=powerU;
99
100 run;
101 %mend multipleSim4;
102
```

Appendix B: Four-Sample SAS Code for Power Simulations

```

1  /* simulation4 *****/
2  /*
3  /* Requirement(s):
4  /* sampleNormPop4.sas
5  /*
6  /* Program Description:
7  /* simulation4.sas performs a hypothesis
8  /* test of no difference for four population means. Random samples
9  /* are normally distributed with respective means muA, muB, muC and muD
10 /* with equal population standard deviation sigma. Random samples are
11 /* generated using seed values from the dataset randSeed. Power
12 /* calculations are based on methods outlined in Thomas 1997.
13 /* Calculations are saved to a dataset named 'sim#' where # represents
14 /* the simulation count.
15 /*
16 /* Input:
17 /* muA      := mean of population one
18 /* muB      := mean of population two
19 /* muC      := mean of population three
20 /* muD      := mean of population four
21 /* sigma     := standard deviation of populations one and two
22 /* seedNum   := position for random seed
23 /* sampleSize := sample size taken for populations one and two
24 /* alpha     := level of significance
25 /* simNum    := simulation ID
26 /*
27 /* Note(s):
28 /* Optimal lower significance level by Golden Section Search method
29 /*
30 *****
31
32 %MACRO simulation4(muA=,muB=,muC=,muD=,sigma=,seedNum=,sampleSize=,alpha=,simNum=);
33
34 /* Generate random samples from four normal populations */
35
36 %sampleNormPop4(popMean=&muA, popSD=&sigma, seedIt=&seedNum, numObs=&sampleSize, sampleNum=1);
37 %sampleNormPop4(popMean=&muB, popSD=&sigma, seedIt=&seedNum, numObs=&sampleSize, sampleNum=2);
38 %sampleNormPop4(popMean=&muC, popSD=&sigma, seedIt=&seedNum, numObs=&sampleSize, sampleNum=3);
39 %sampleNormPop4(popMean=&muD, popSD=&sigma, seedIt=&seedNum, numObs=&sampleSize, sampleNum=4);
40
41
42 /* Merge datasets sample1, sample2, sample3 and sample4 to produce dataset sampleFourPops */
43
44 data sampleFourPops;
45 merge sample1 sample2 sample3 sample4;
46 run;
47

```

Appendix B: Four-Sample SAS Code for Power Simulations

```

48
49      /* Main */
50
51      proc iml;
52          reset nolog;
53          use sampleTourPops;
54          read all into X;
55
56          /* Function *****/
57          /*
58          /* Return the confidence interval width for a noncentral
59          /* f-dsitrubution
60          /* */
61          /*
62          /* *****/
63
64          start w(alphaLow,ndf,ddf,ncent,s1);
65
66          width = quantile('F',1-s1+alphaLow,ndf,ddf,ncent)-quantile('F',alphaLow,ndf,ddf,ncent);
67
68          return(width);
69
70          finish w;
71          /* Function: optimal lower critical region *****/
72          /*
73          /* Finds the minimum confidence interval width using an
74          /* implementation of the Golden Section Search method
75          /* similar to Press et al. (2007).
76          /*
77          /* Note(s):
78          /* The Golden Section Search method finds a value which minimizes
79          /* a function. The function must be one dimensional and unimodal.
80          /*
81          /* Imagine a 'typical' pdf of an F distribution. Say we want to
82          /* setup an interval such that the area covered by the sum of the two
83          /* tails is alpha. For each alpha we can find the
84          /* associated quantile using the SAS function 'quantile'.
85          /*
86          /* Since alpha is specified, knowing the lower area easily leads to
87          /* the upper area. Here we consider only the lower
88          /* area. As such we can relate interval width as a function of
89          /* lower alpha size. As we increase or decrease the size of the
90          /* lower area we increase or decrease the width of the interval.
91          /* Assuming a minimum interval width exists, we optimize over the
92          /* values [0.00, 0.05].
93          /*
94          /* *****/

```

Appendix B: Four-Sample SAS Code for Power Simulations

```

95      start alphaIU(ndf1,ddf2,ncp,s1Alpha,to1);
96      alphaA = 0;
97      alphaB = s1Alpha;
98
99      /* Initial lower bracket on optimal region */
100     /* Initial upper bracket on optimal region */
101
102    /* Note 1 *****/
103    /* For each iteration an interval */
104    /* [alphaA, alphaB] is retained */
105    /* *****/
106
107    /* Golden ratio constant reduction factor */
108    /* Set test bracket a1 */
109    /* Calculate Interval width for a1 */
110    /* Set test bracket a2 */
111    /* Calculate interval width for a2 */
112
113    /* Note 2 *****/
114    /* The initial setup invokes two function */
115    /* calls. The following loop uses one */
116    /* function call per iteration. */
117
118
119
120 do while (abs(alphaB-alphaA)>tol);
121
122
123
124
125
126
127
128
129
130
131
132
133
134
135
136
137
138
139
140
141

```

Appendix B: Four-Sample SAS Code for Power Simulations

Appendix B: Four-Sample SAS Code for Power Simulations

```

189     sst = n*xBar`*xBar-(n/p)*(xBar`*J)**2;          /* Sum of squares for treatments */
190     df1 = p-1;                                       /* Degrees of freedom for treatment */
191     mst = sst/df1;                                     /* Mean square for treatment */
192     sse = ssq(X)-(xBar`*xBar)*n;                     /* Sum of squares for error */
193     df2 = p*(n-1);                                    /* Degrees of freedom for error */
194     mse = sse/df2;                                     /* Mean square for error */
195     fSamp = mst/mse;                                   /* Observed F */
196
197     fcrit = quantile('F',1-salpha,df1,df2);           /* F critical value */
198
199     pValue = 1-cdf('F',fSamp,df1,df2);               /* P-Value */
200
201     trueLambda = (n*(((&muA-&muG)**2)+((&muB-&muG)**2)+((&muC-&muG)**2)+((&muD-&muG)**2))/((&sigma)**2); /* trueLambda */
202
203     truePower = 1-cdf('F',fcrit,df1,df2,trueLambda); /* truePower */
204
205     estLambda = fsamp*(P-1);                          /* Estimated lambda */
206
207     adjEstLambda = (estLambda*(df2-2)/df2)-df1;      /* Adjusted estimated lambda */
208
209     if adjEstLambda < 0 then adjEstLambda = 0;        /* Note 7 **** */
210
211     /* For some random seeds the */
212     /* noncentrality parameter is */
213     /* negative. */
214
215     /* A negative noncentrality */
216     /* parameter occurs when */
217     /* the expression */
218     /* estLambda*(df2-2)/df2 is */
219     /* less than df1. */
220
221     /* **** */
222
223     alphaL = alphalU(df1,df2,adjEstLambda,&alpha,eps); /* Lower significance */
224     alphaU = &alpha-alphaL;                            /* Upper significance */
225
226
227
228
229
230
231
232
233
234
235

```

Appendix B: Four-Sample SAS Code for Power Simulations

Appendix B: Four-Sample SAS Code for Power Simulations

```
283 proc datasets lib=work nolist;
284   delete sample1 sample2 sample3 sample4 sampleFourPops;
285   quit;
286   run;
287
288 %MEND simulation4;
289
```

Appendix B: Four-Sample SAS Code for Power Simulations

```
1 /* sampleNormPop4 **** */
2 /*
3 /* Program Description:
4 /* sampleNormPop4 generates a random sample from a normal population
5 /* and outputs the values to a dataset
6 /*
7 /* Input:
8 /* seed      := dataset of random seeds from randSeed
9 /* popMean   := population mean
10 /* popSD    := population standard deviation
11 /* seedIt    := seed value position for generating random observations
12 /* numObs   := sample size
13 /* sampleNum := identifies sample
14 /*
15 /* ****
16 /*
17 %macro sampleNormPop4 (popMean=,popSD=,seedIt=,numObs=,sampleNum=);
18 /*
19 proc iml;
20   reset nolog;
21   use randSeed;
22   read all into seed;
23
24 x&sampleNum = &popMean+&popSD*rannor(J(&numObs,1,seed[&seedIt,&sampleNum]));
25 test = seed[&seedIt,&sampleNum];
26
27 create sample&sampleNum from x&sampleNum[colname={obs&sampleNum}];
28 append from x&sampleNum;
29
30 quit;
31
32 %mend sampleNormPop4;
```

Appendix B: Four-Sample SAS Code for Power Simulations

```

1  /* avgMedMultipleSim4 *****/
2  /*
3   * Requirement(s):
4   * avgMedSim4.sas
5   */
6  /* Program Description:
7   * avgMedMultipleSim4 calls avgMedSim4 for each sigma by sample size */
8  /* combination.
9  */
10 /* Input:
11  * itM          := number of iterations
12  * nsXnss       := number of iteration groups
13  *                  (sigmas times number of sample sizes)
14  */
15 /* *****/
16 /* *****/
17
18 %macro avgMedMultipleSim4(itM=,nsXnss=);
19
20 /* Produce average and median of simulations */
21
22 %let bStart = 1;
23 %let bFinish = &itM;
24
25 %do i = 1 %to &nsXnss %by 1;
26   %avgMedSim4(start=&bStart,finish=&bFinish,itB=&i);
27   %let bStart = &bStart+&itM;
28   %let bFinish = &bFinish+&itM;
29
30
31 /* Combine average simulations into a data set and name columns */
32
33 data avgMedMultipleSimOut;
34
35 set
36   %do j = 1 %to &nsXnss %by 1;
37     ansim&j
38   %end;
39   /* Double do-loop to reference data sets. ';' closes 'set' */
40   rename col1=mu1 col2=mu2 col3=mu3 col4=mu4 col5=sigma col6=sampleSize col7=sigLevel col8=pValue
41   col9=trueLambda col10=truePower col11=estLambda col12=adjEstLambda col13=lambdaU col14=lambdaD col15=estPower
42   col16=powerL col17=powerU col18=medPower;
43
44 %mend avgMedMultipleSim4;
45

```

Appendix B: Four-Sample SAS Code for Power Simulations

```

1  /* avgMedSim4 **** */
2  /*
3  /* Program Description:
4  /* avgMedSim4 averages over a user specified number of rows from
5  /* multiplesimOut. avgMedSim4 also determines the median
6  /* of the calculated power values of the rows.
7  /*
8  /* Input:
9  /* X          := gets dataset multipleSimOut
10 /* start     := beginning of iteration block
11 /* finish    := end of iteration block
12 /* itB       := number of iterations
13 /*
14 /* **** */
15 /*
16 %macro avgMedSim4(start=&finish=,itB=);
17
18 proc iml;
19
20  reset nolog;
21  use multipleSimOut;
22  read all into X;
23
24  /* Average rows for an iteration block */
25
26  rows = X[&start:&finish,];
27  avgRows = rows[:,];
28
29
30  /* Median for column calculated power */
31  col = X[&start:&finish,15];
32
33  /* Column of calculated power */
34
35  numRows = nrow(col);
36
37  mis = 0;
38  i = 1;
39
40  do while (i <=numRows);
41    if missing(col[i,1]) > 0 then mis=mis/i;
42    i=i+1;
43  end;
44
45  /* Storage for positions of missing values */
46
47  /* Note 1 **** */
48  /* Loop through column of
49  /* of calculated power values.
50  /* If a missing value is found
51  /* record array position.
52  */
53  /* **** */

```

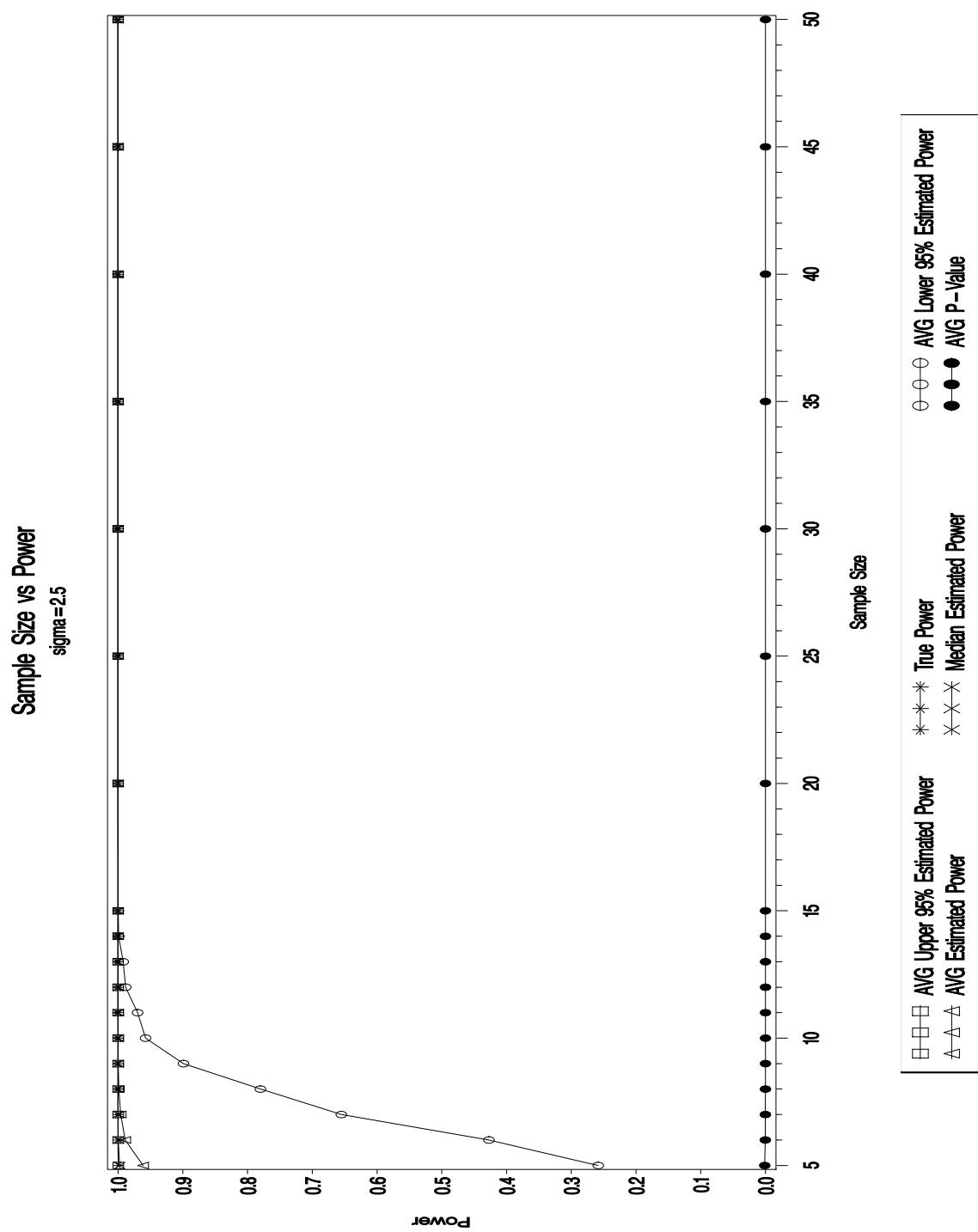
Appendix B: Four-Sample SAS Code for Power Simulations

```

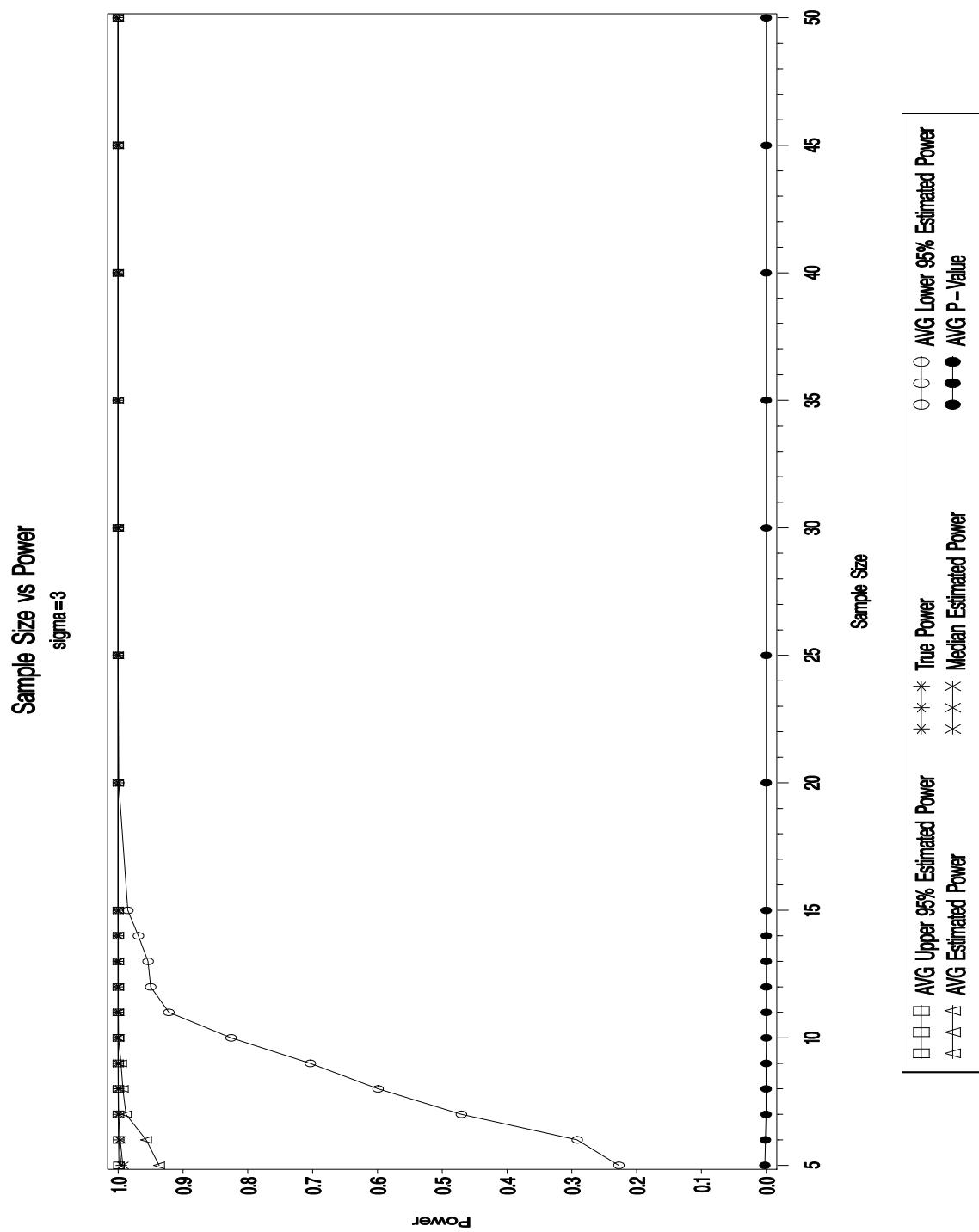
48      nr = nrow(mis);
49
50      /* Number of missing values */
51
52      /* Note 2 *****/
53      /*
54      * If 'nr' is greater than
55      * one then we have missing
56      * values. These values are
57      * removed from the column.
58      */
59
60      if nr>1 then mis = remove(mis,1);
61      if nr>1 then col = col`;
62      if nr>1 then col = remove(col,mis);
63      if nr>1 then col = col`;
64
65      med = median(col);
66
67      /* Concatinate median to average row */
68      avgRows=avgRows || med;
69
70      avgRows=avgRows || med;
71
72      create amSim&itB from avgRows;
73      append from avgRows;
74
75      quit;
76      %mend avgMedSim4;
77

```

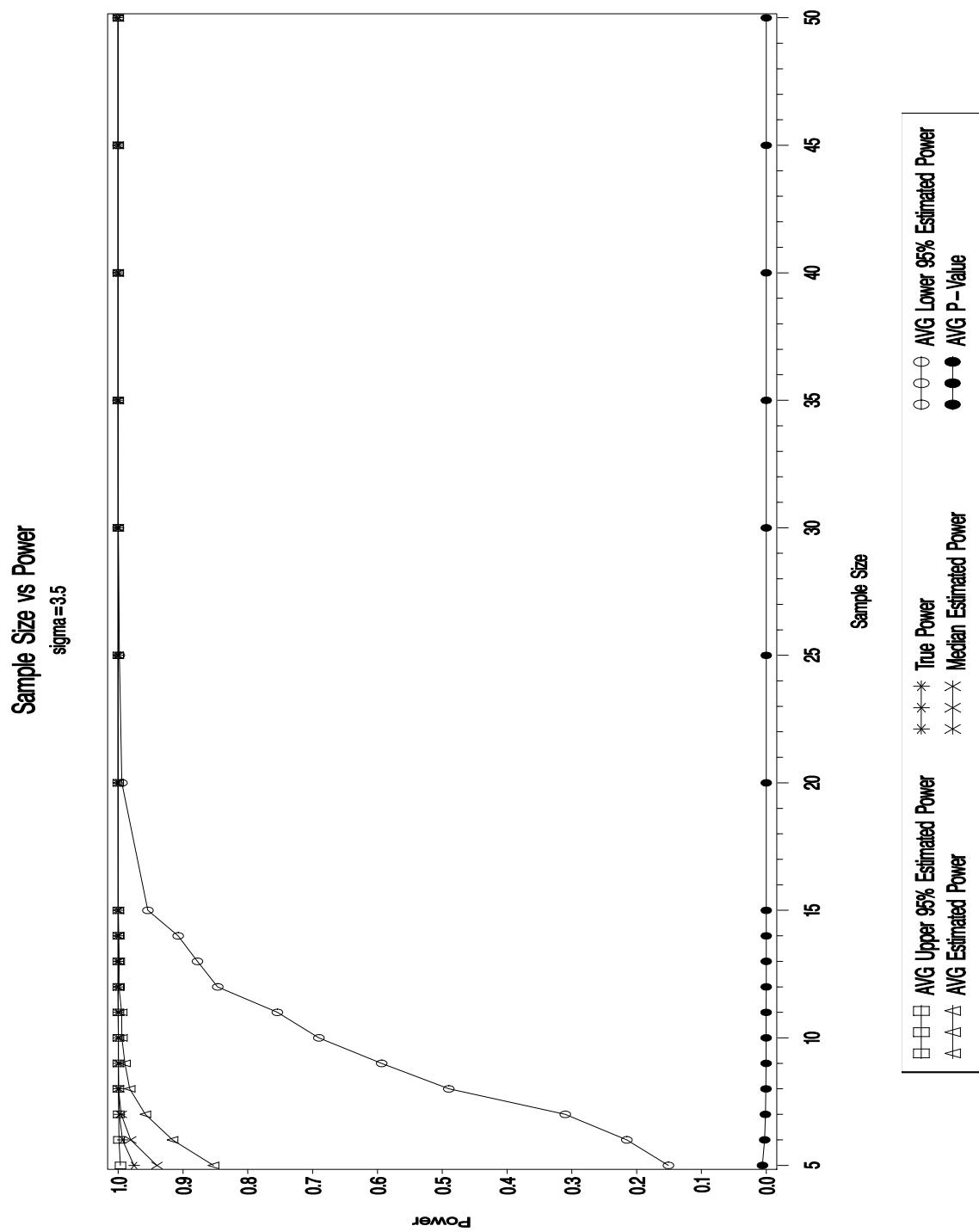
Appendix C: Graphs of Treatment Arrangement 30, 40



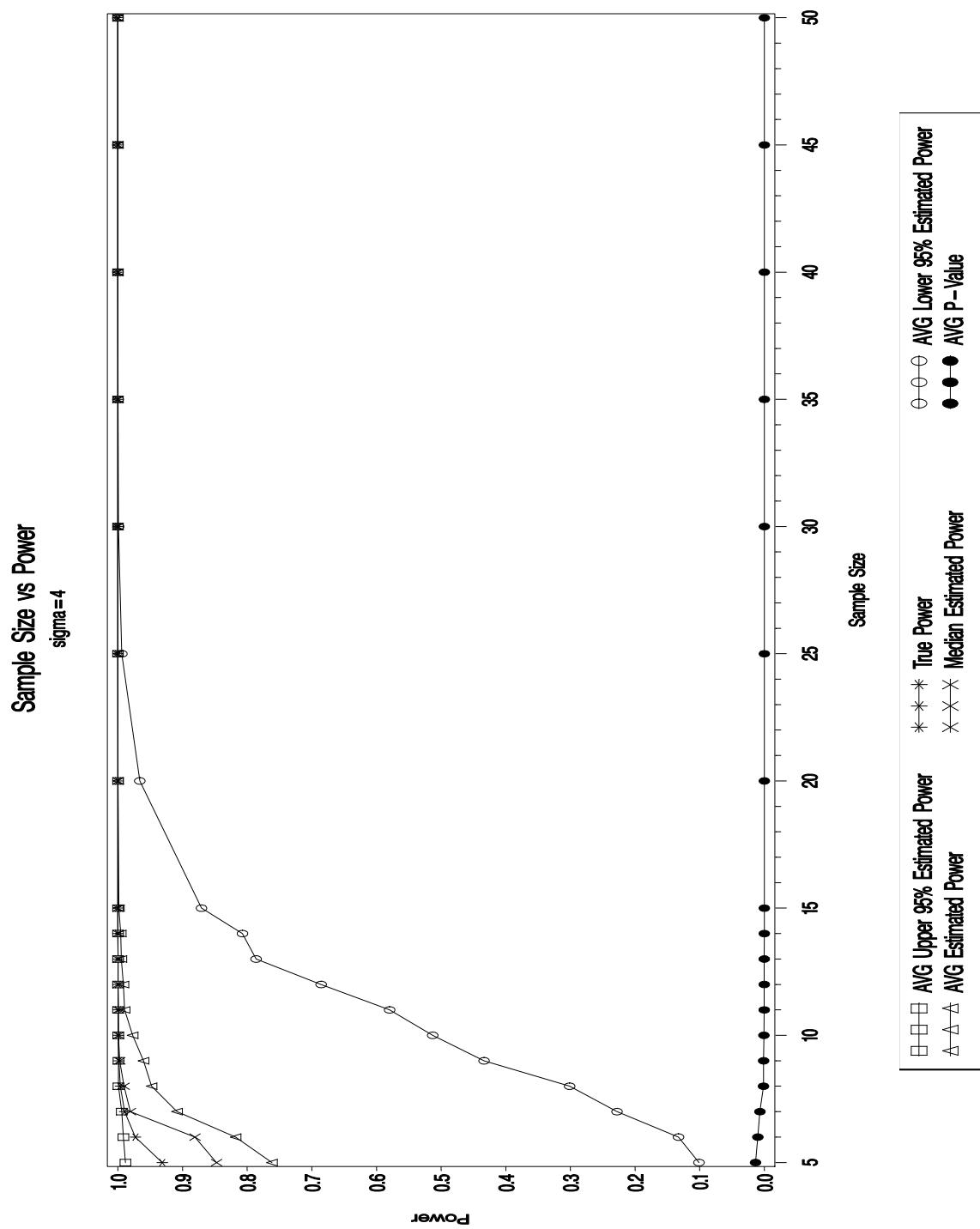
Appendix C: Graphs of Treatment Arrangement 30, 40



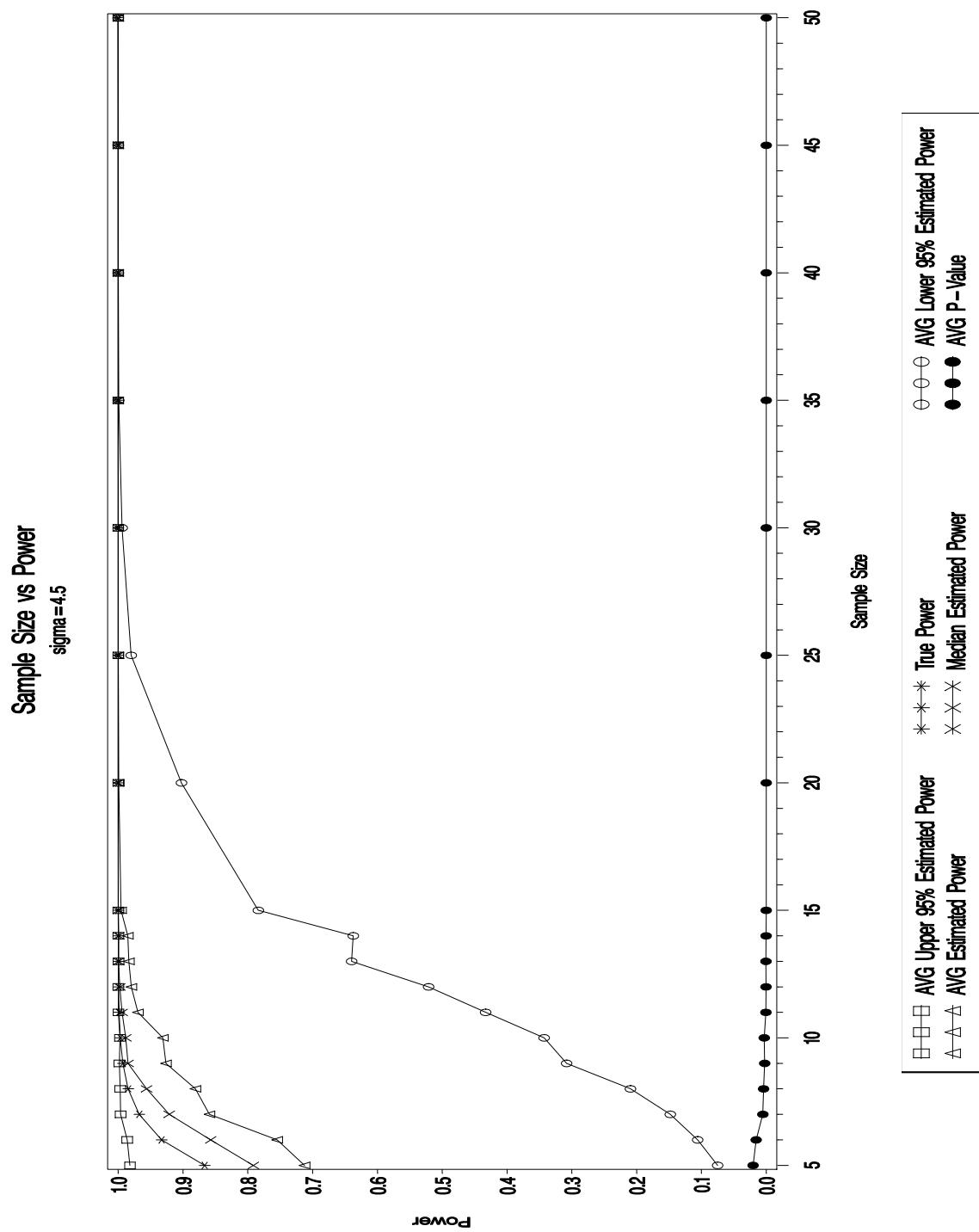
Appendix C: Graphs of Treatment Arrangement 30, 40



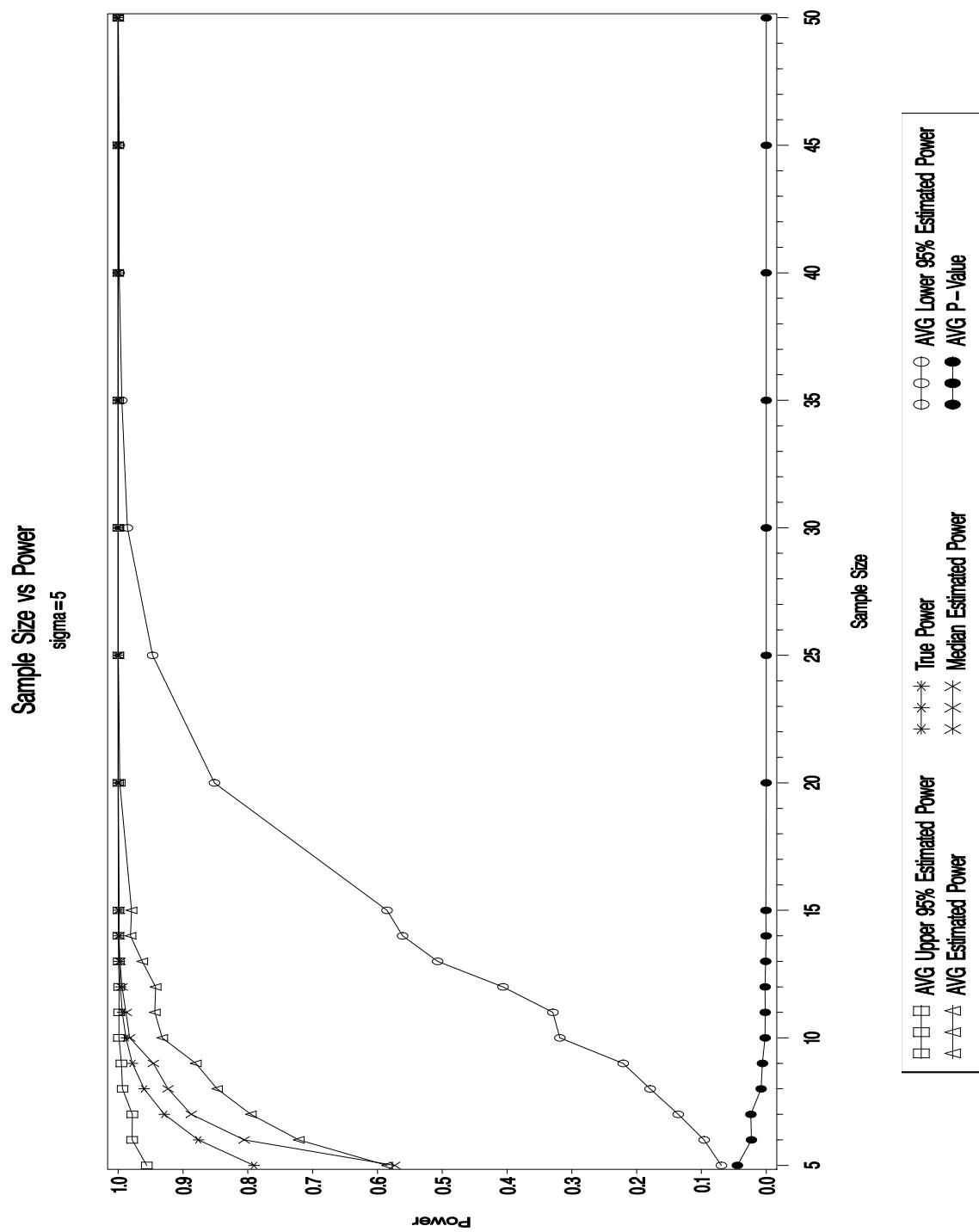
Appendix C: Graphs of Treatment Arrangement 30, 40



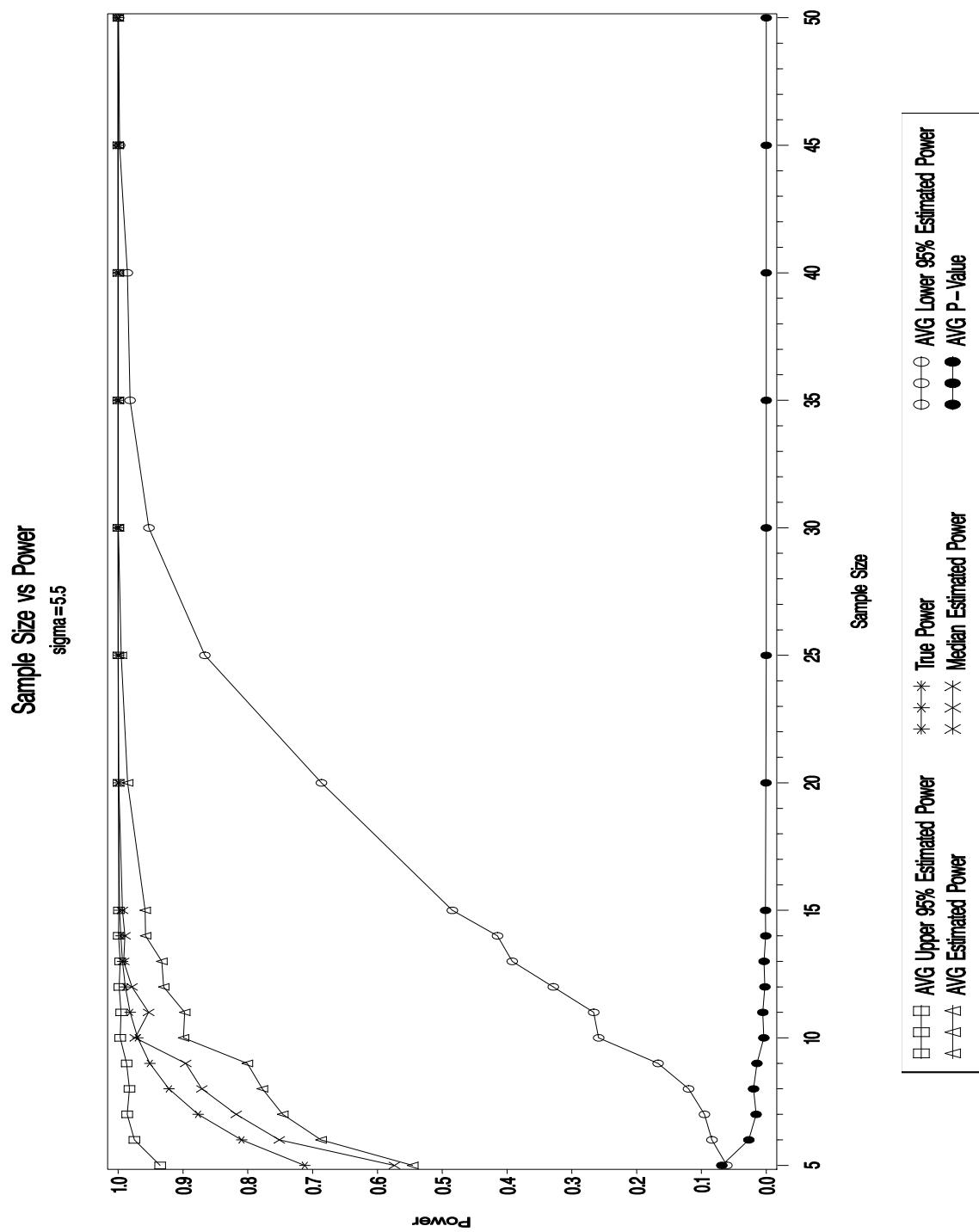
Appendix C: Graphs of Treatment Arrangement 30, 40



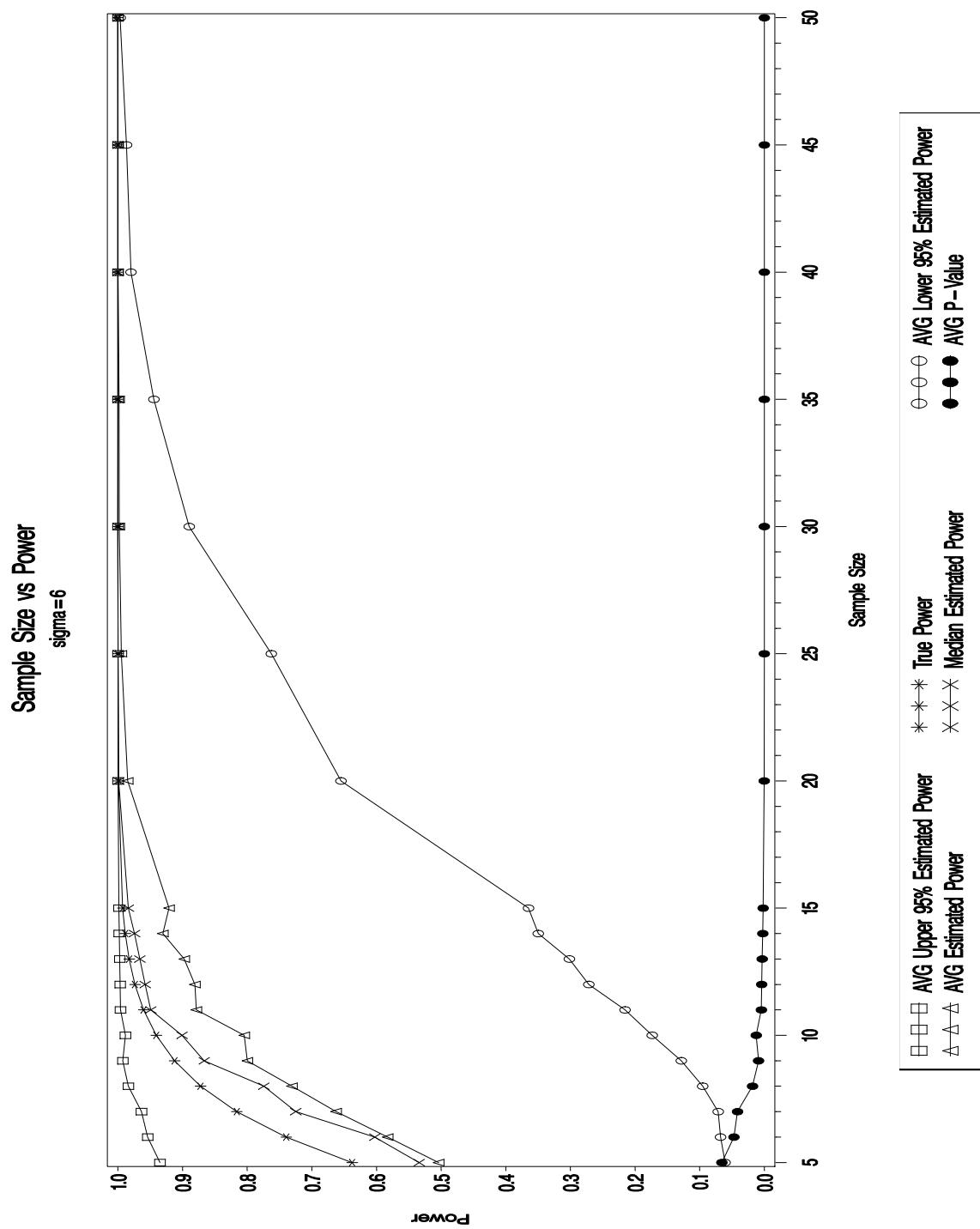
Appendix C: Graphs of Treatment Arrangement 30, 40



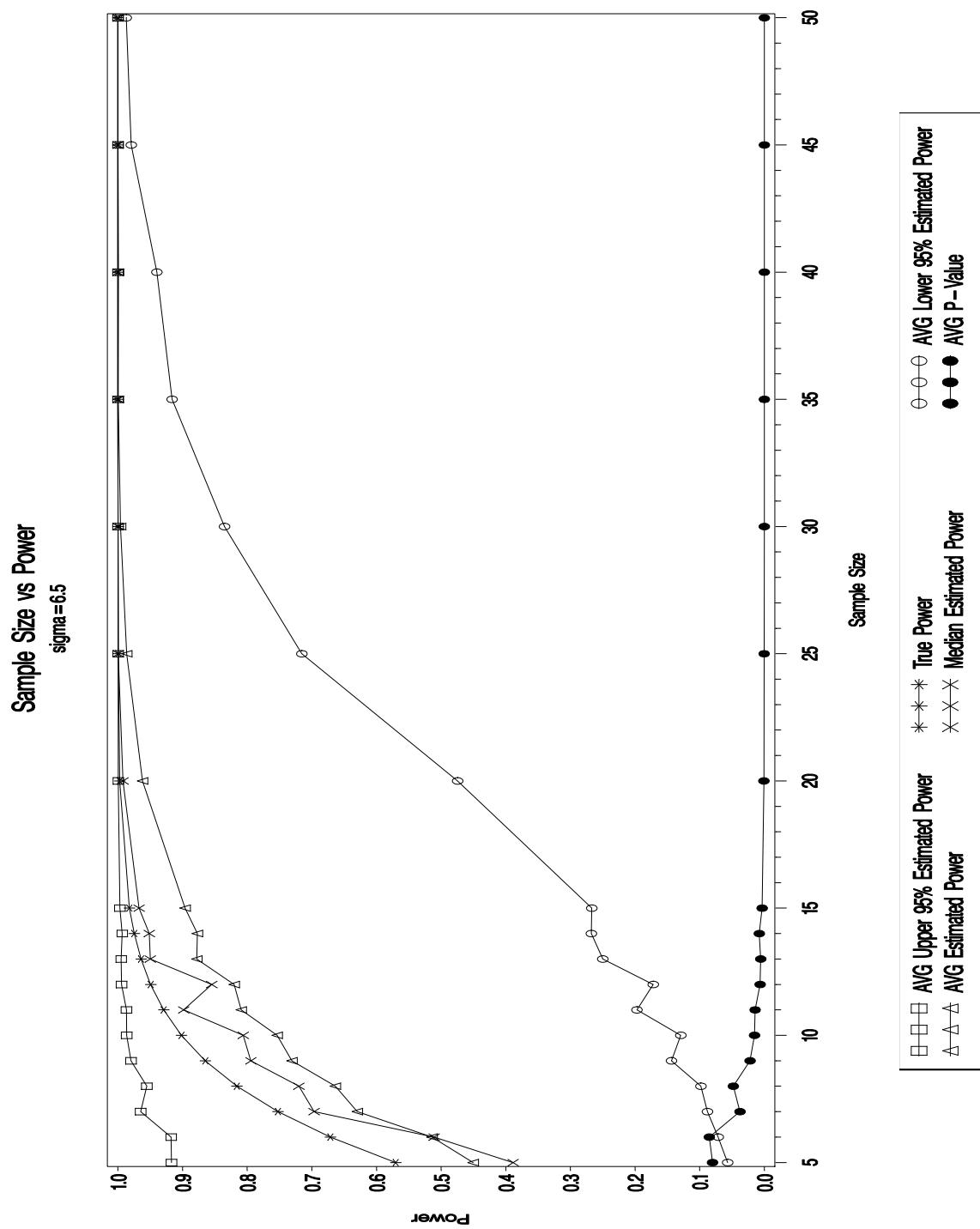
Appendix C: Graphs of Treatment Arrangement 30, 40



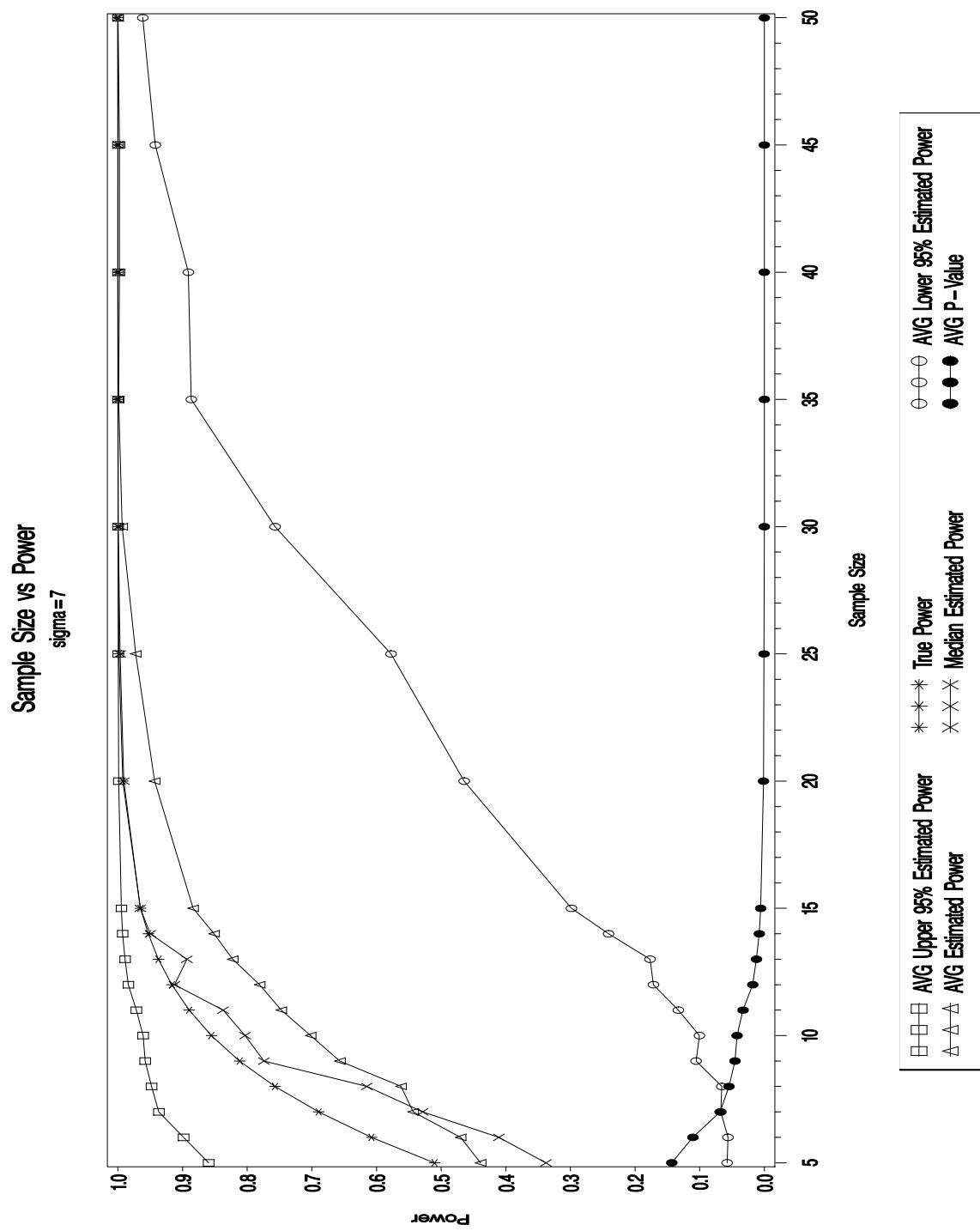
Appendix C: Graphs of Treatment Arrangement 30, 40



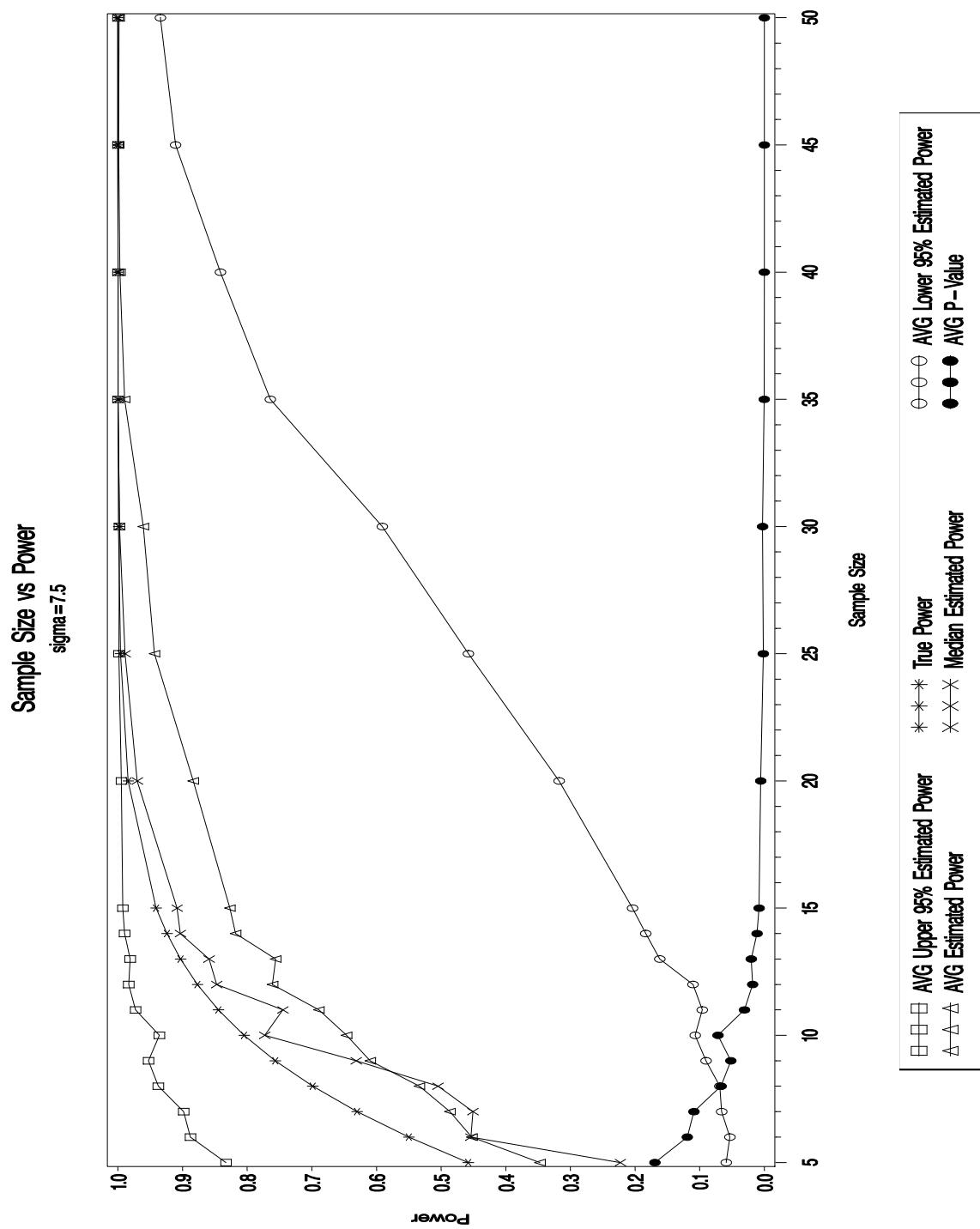
Appendix C: Graphs of Treatment Arrangement 30, 40



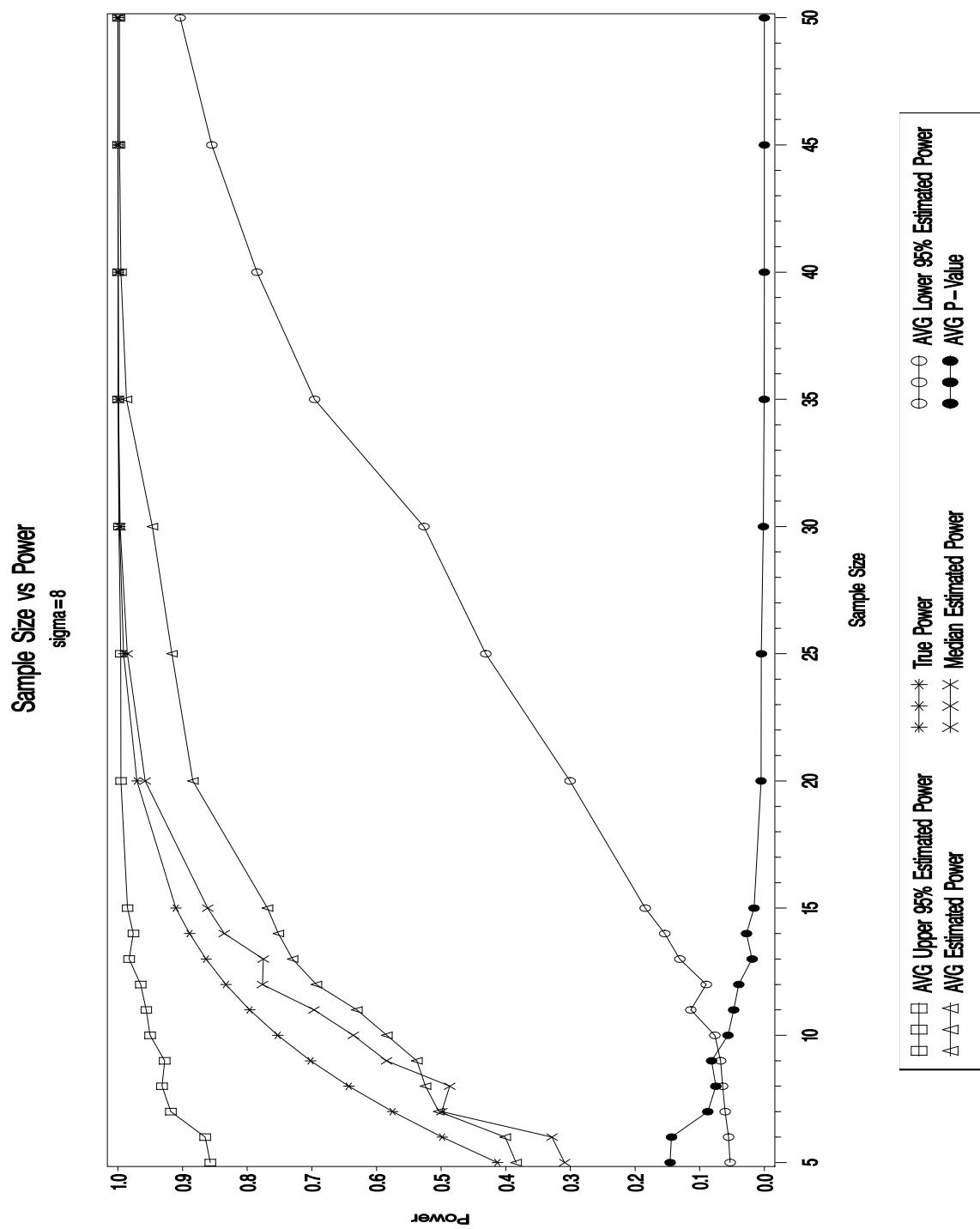
Appendix C: Graphs of Treatment Arrangement 30, 40



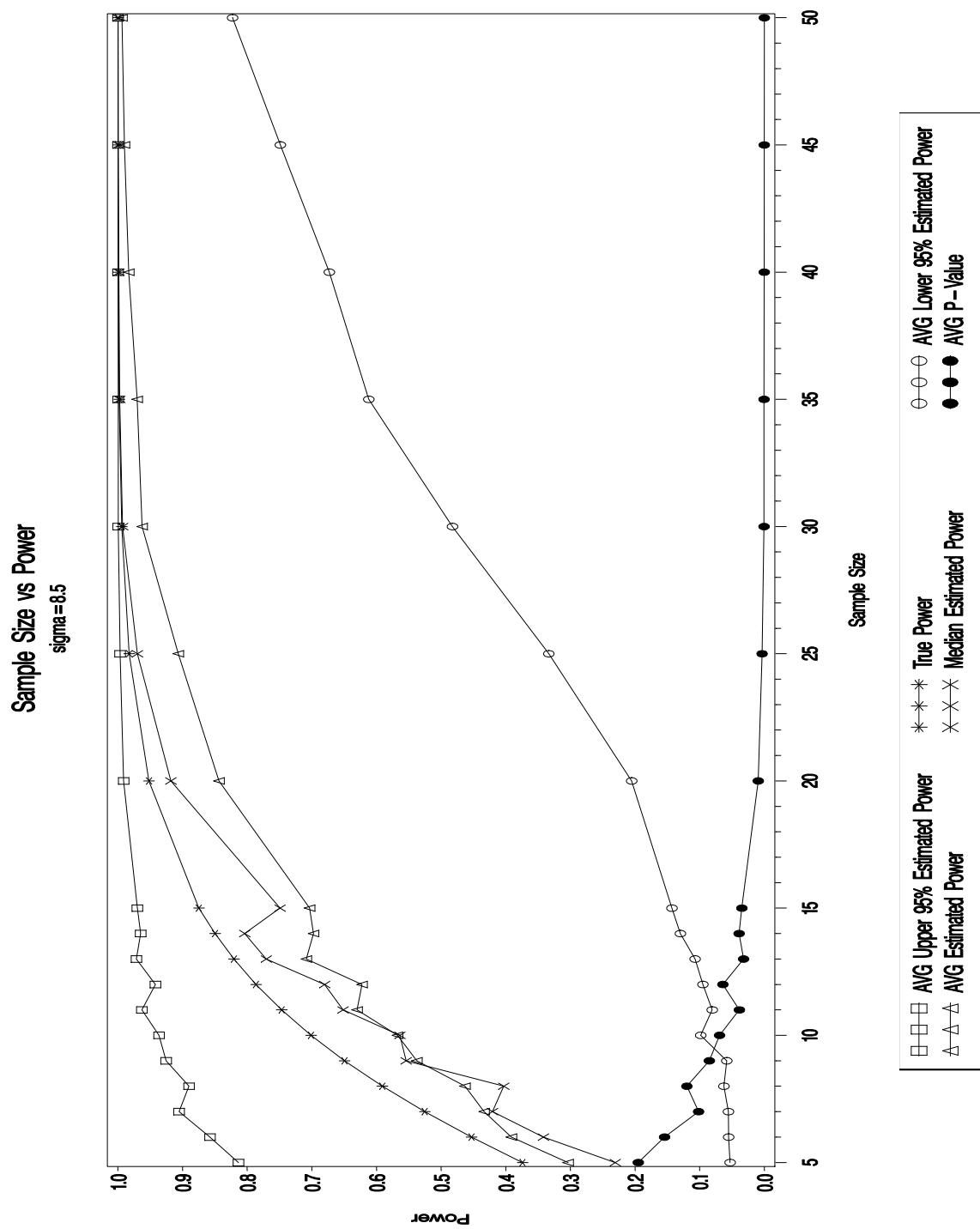
Appendix C: Graphs of Treatment Arrangement 30, 40



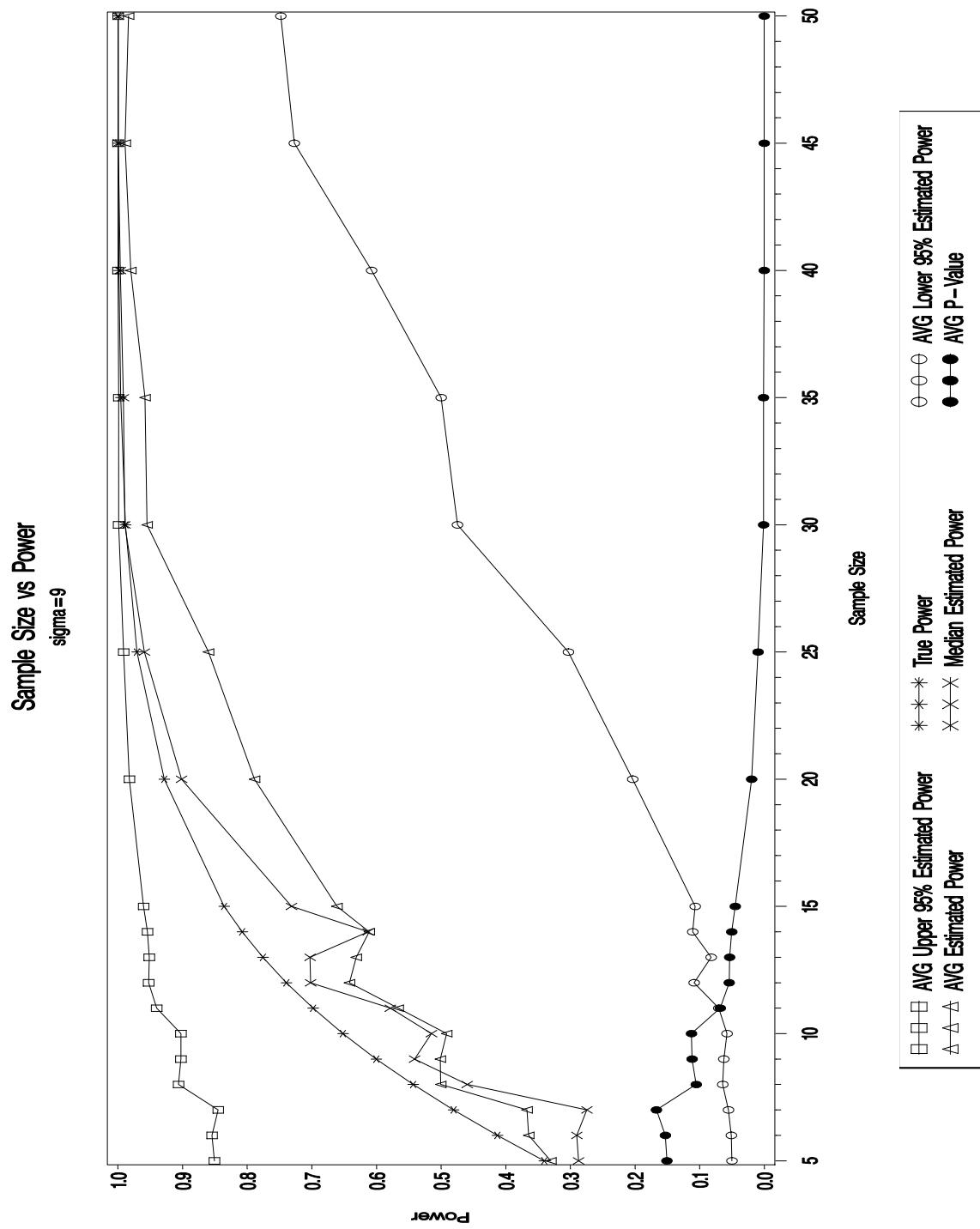
Appendix C: Graphs of Treatment Arrangement 30, 40



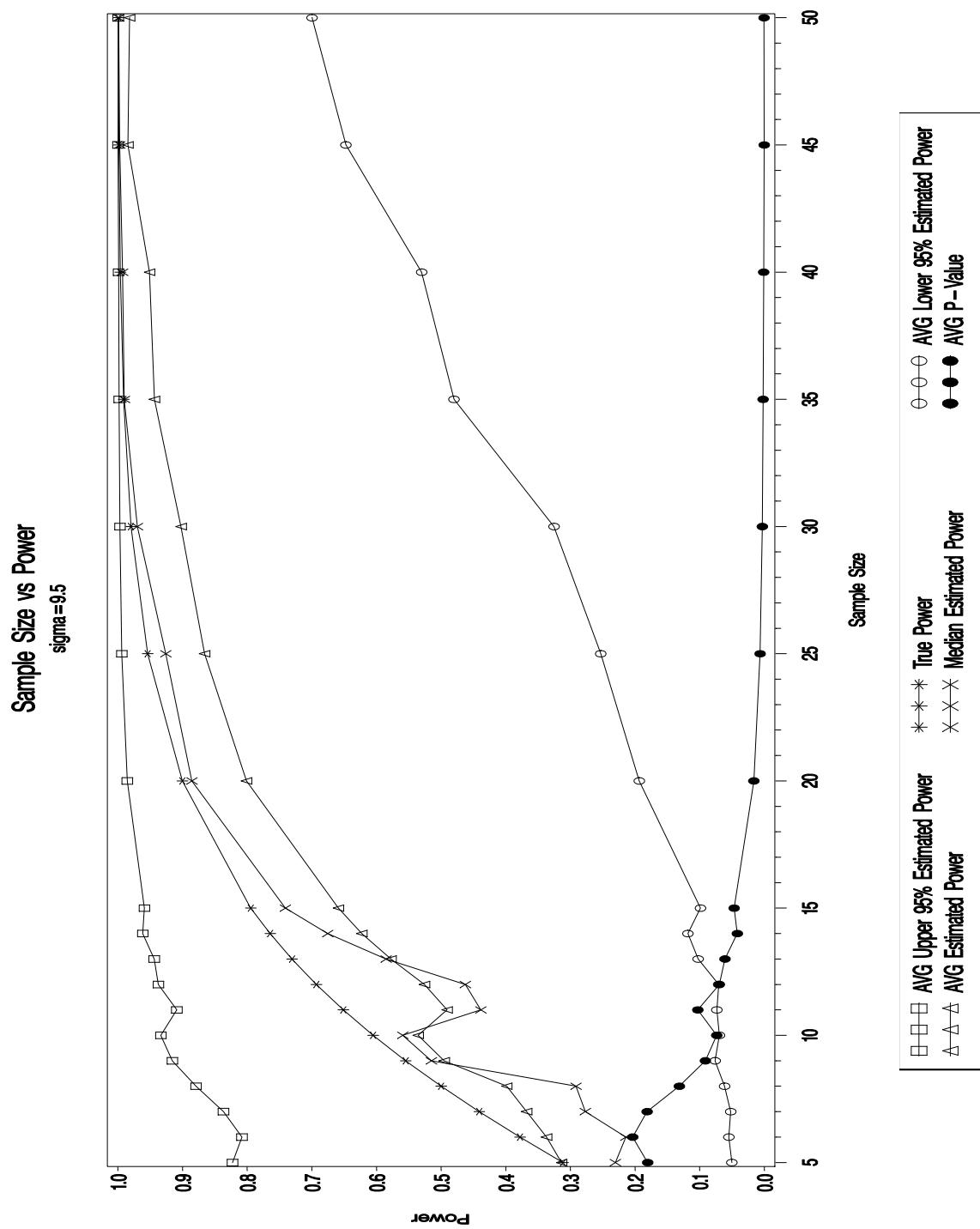
Appendix C: Graphs of Treatment Arrangement 30, 40



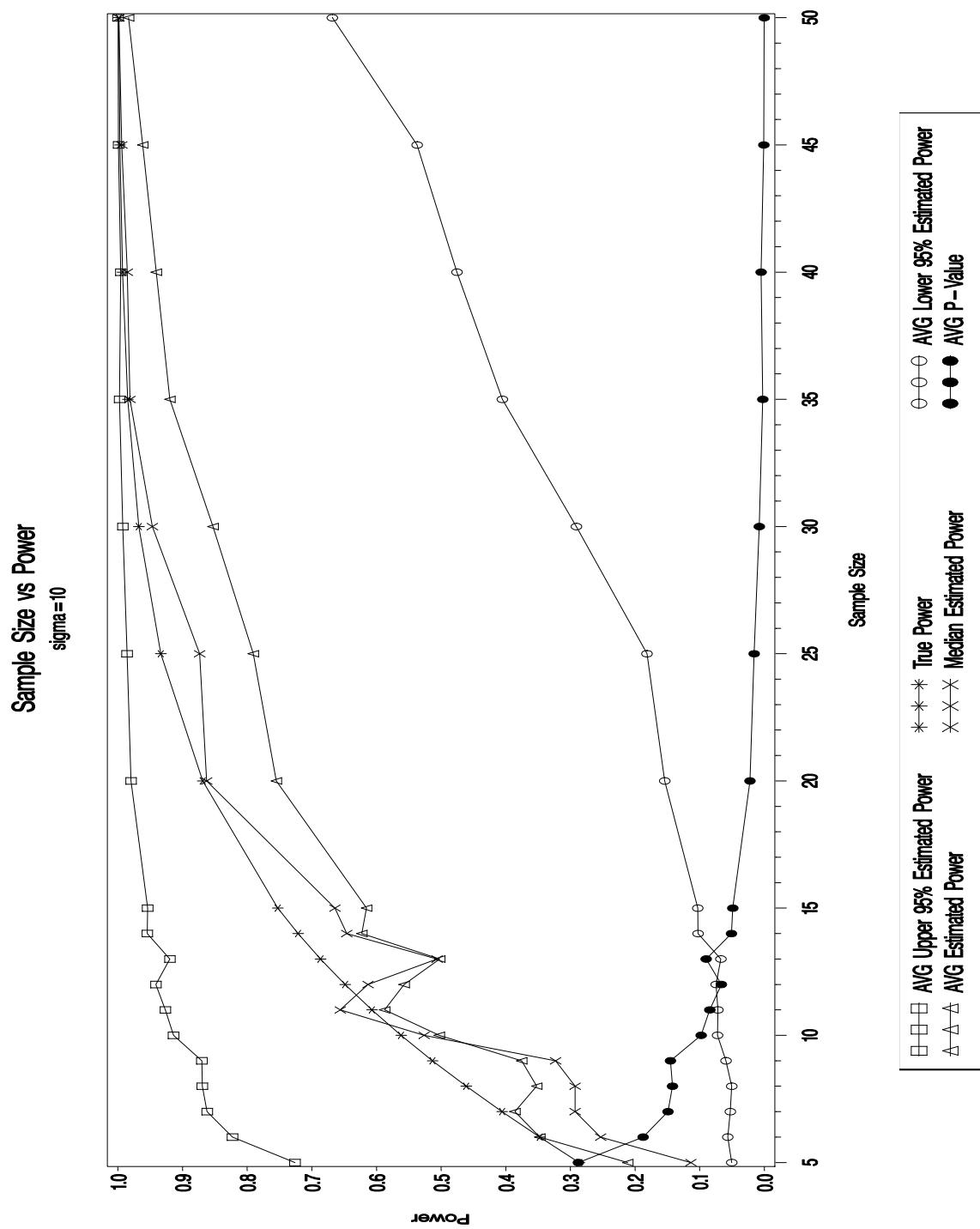
Appendix C: Graphs of Treatment Arrangement 30, 40



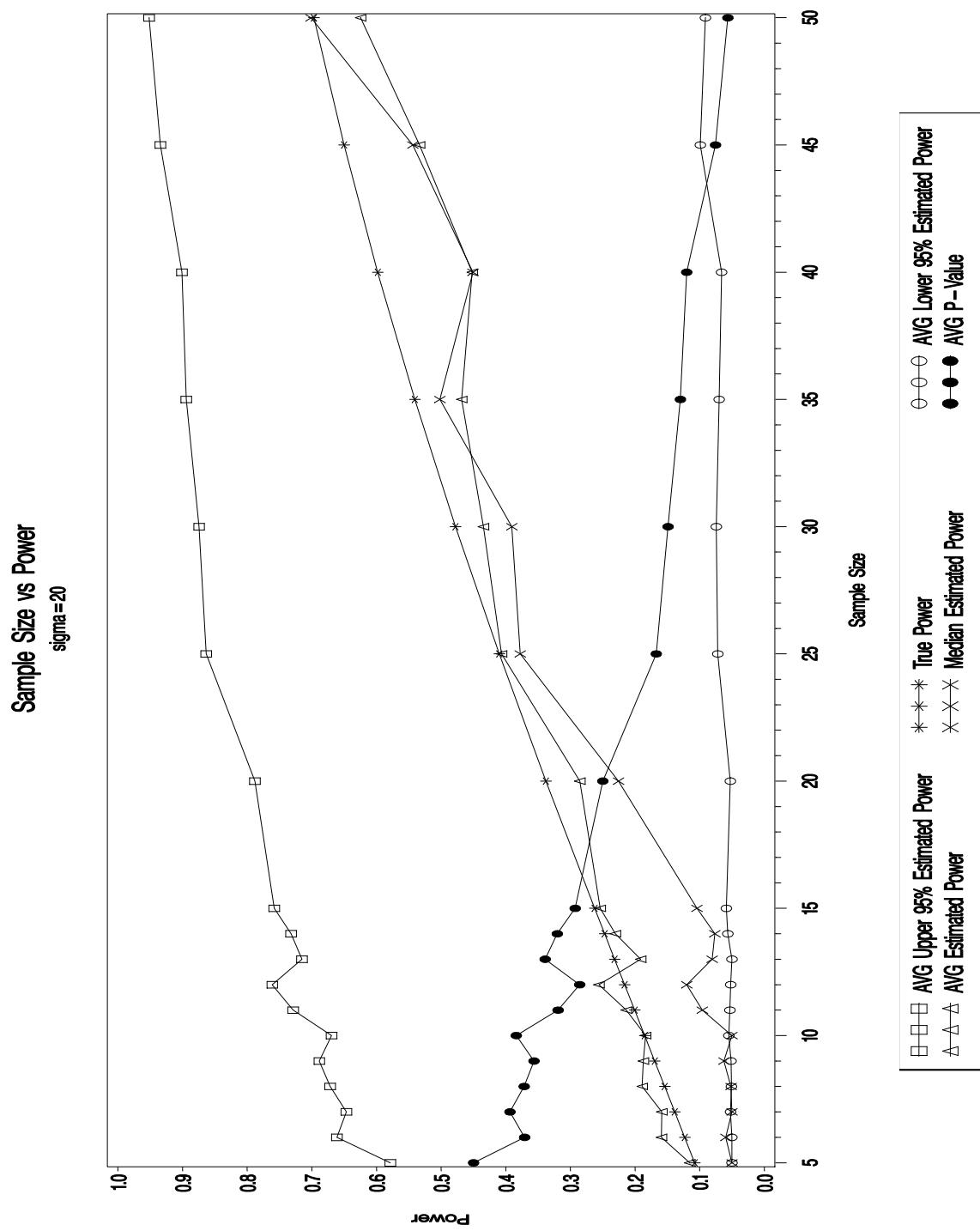
Appendix C: Graphs of Treatment Arrangement 30, 40



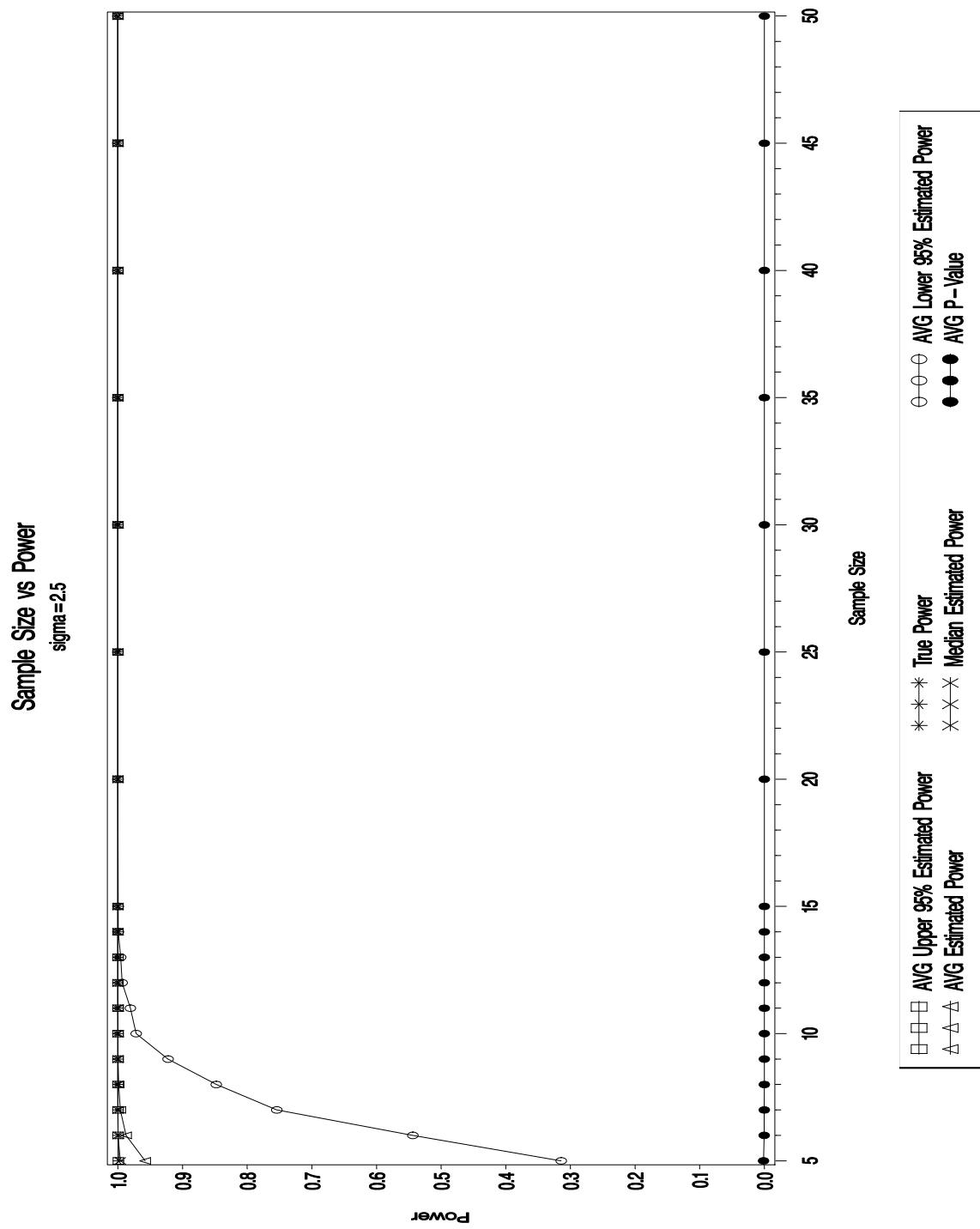
Appendix C: Graphs of Treatment Arrangement 30, 40



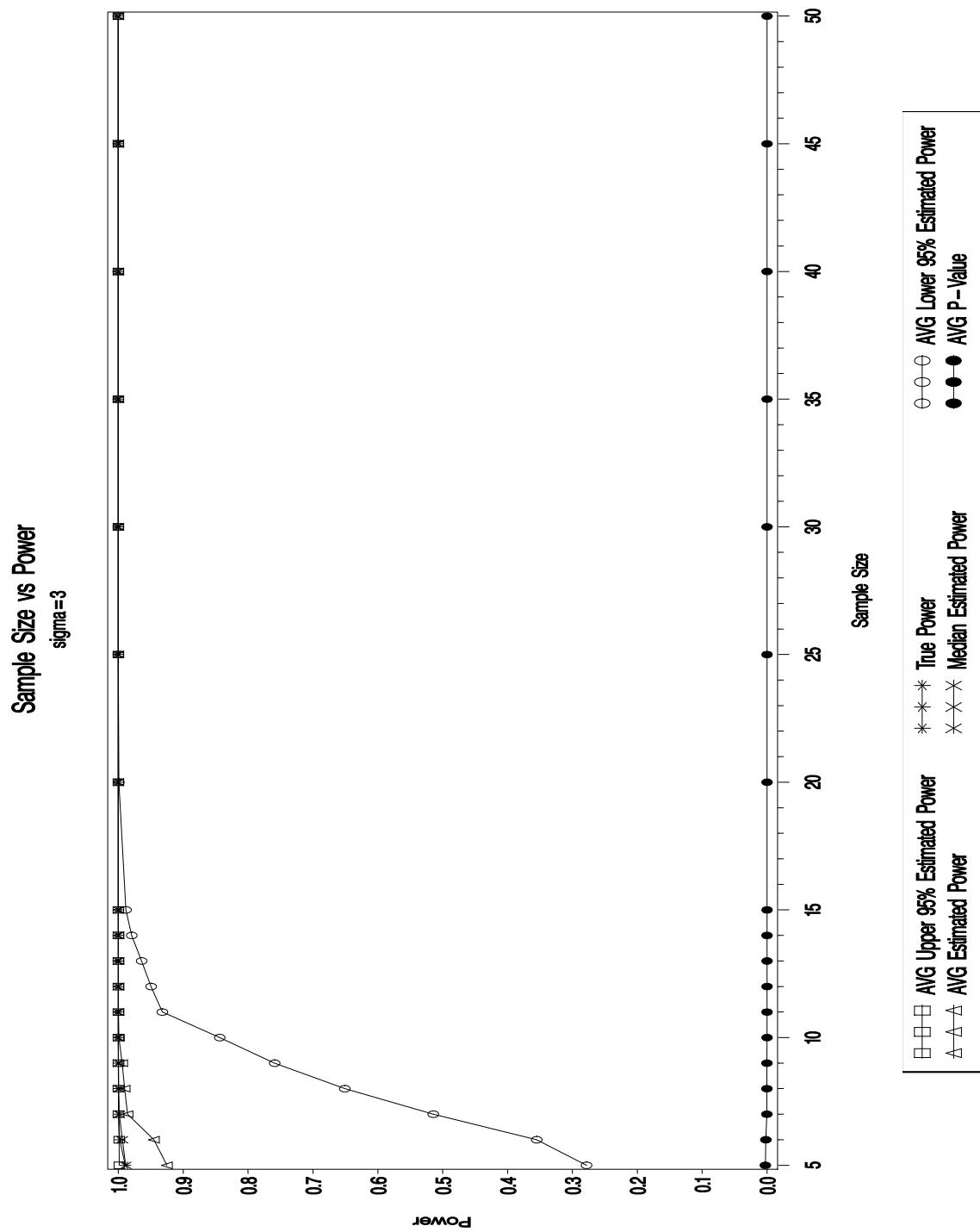
Appendix C: Graphs of Treatment Arrangement 30, 40



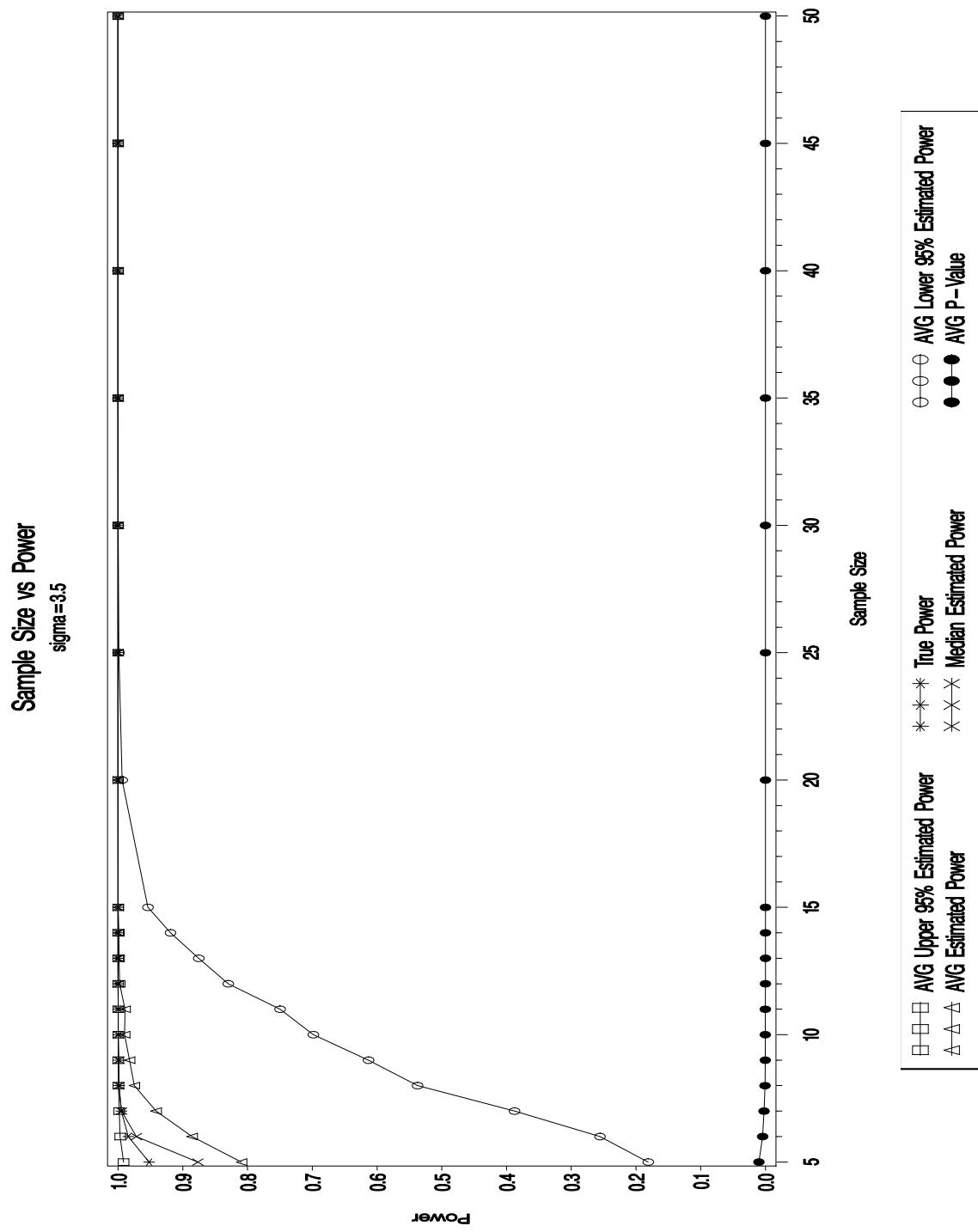
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



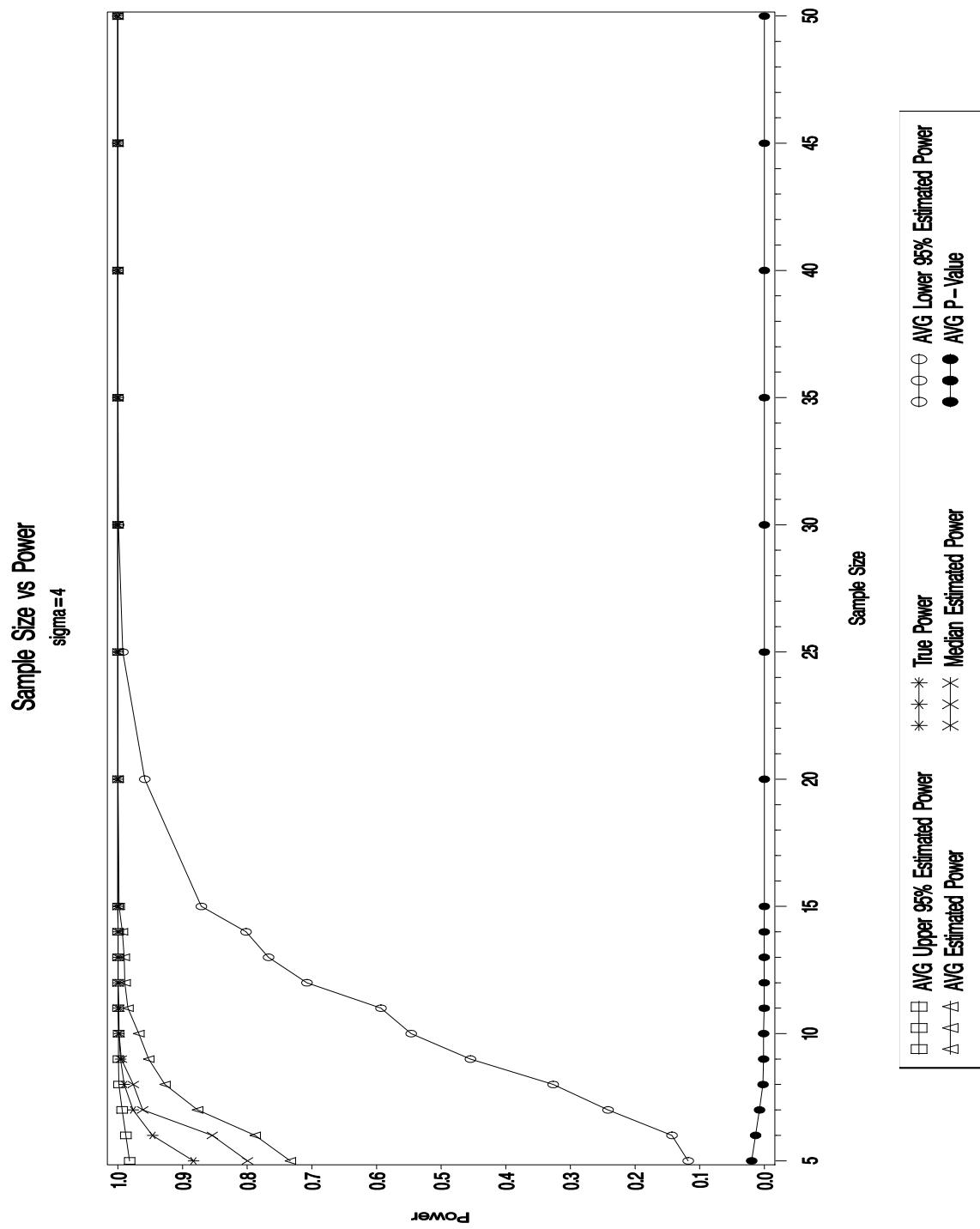
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



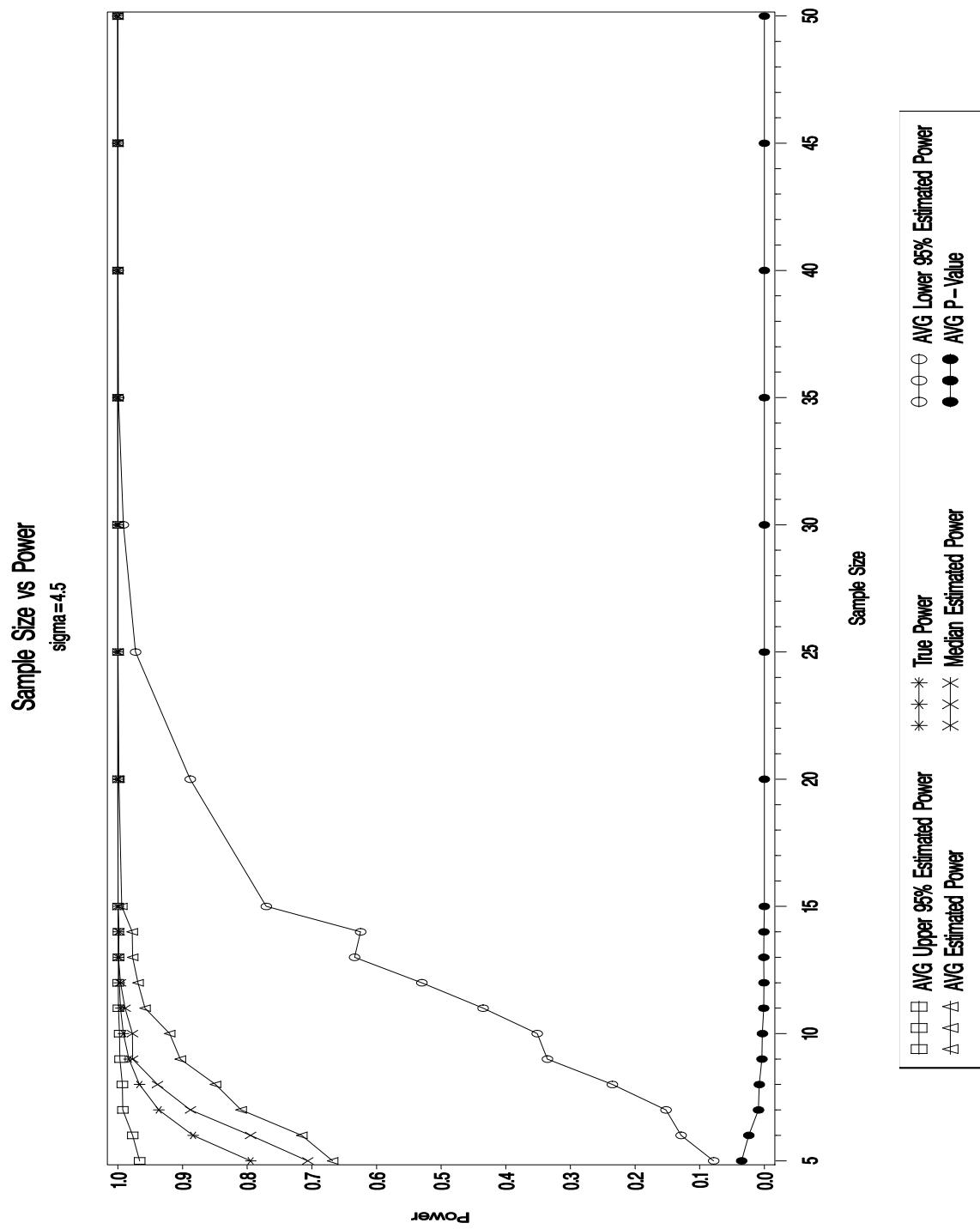
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



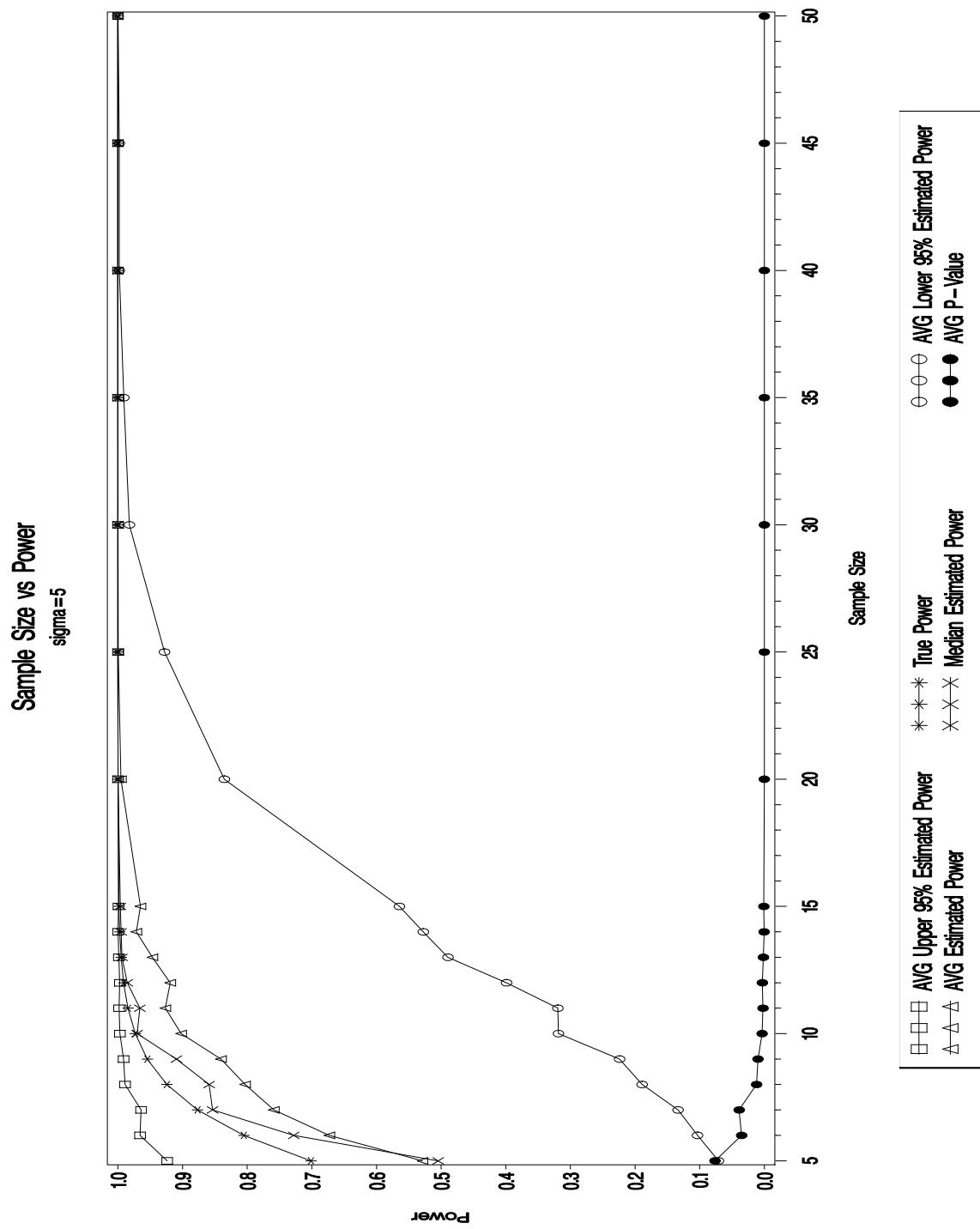
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



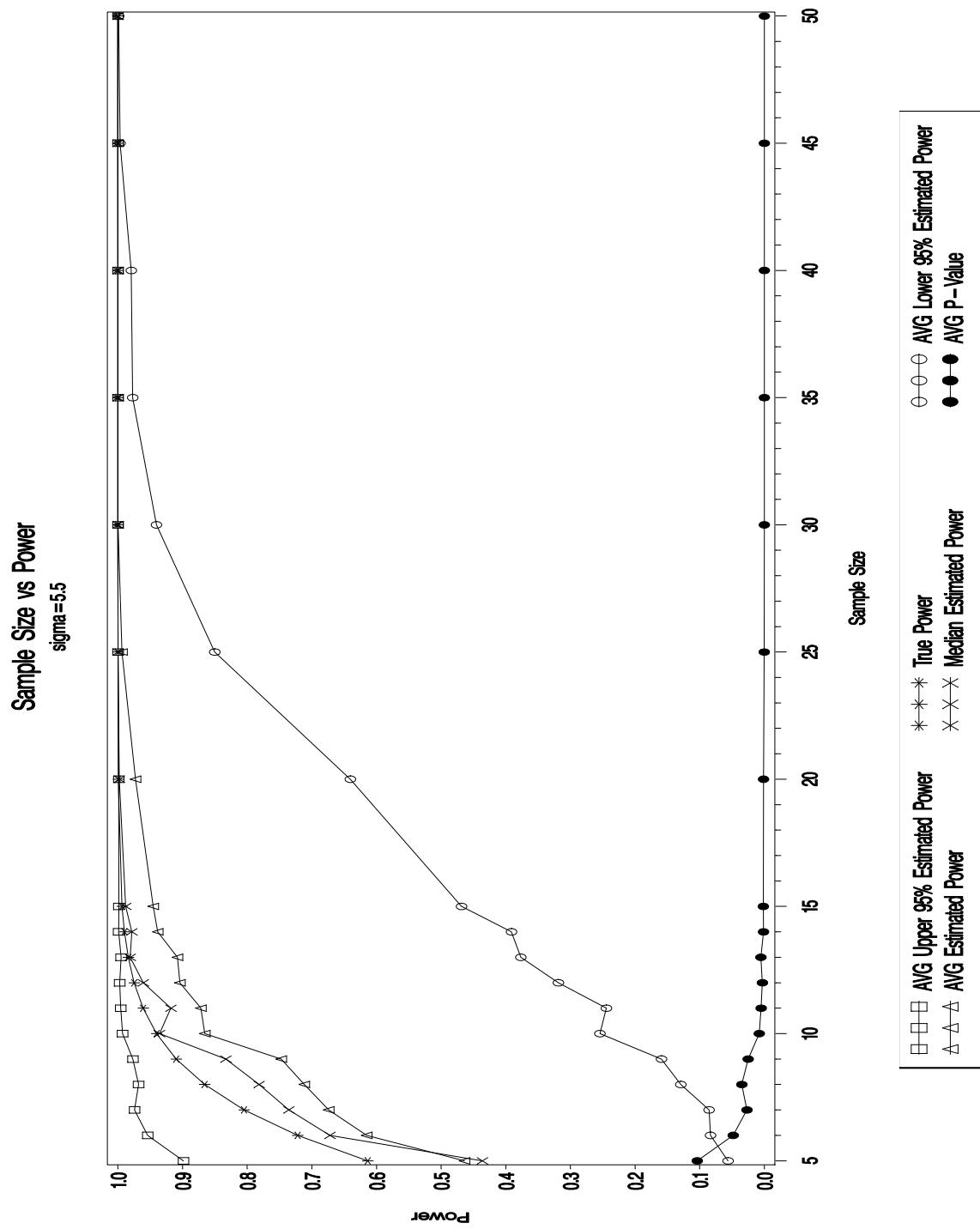
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



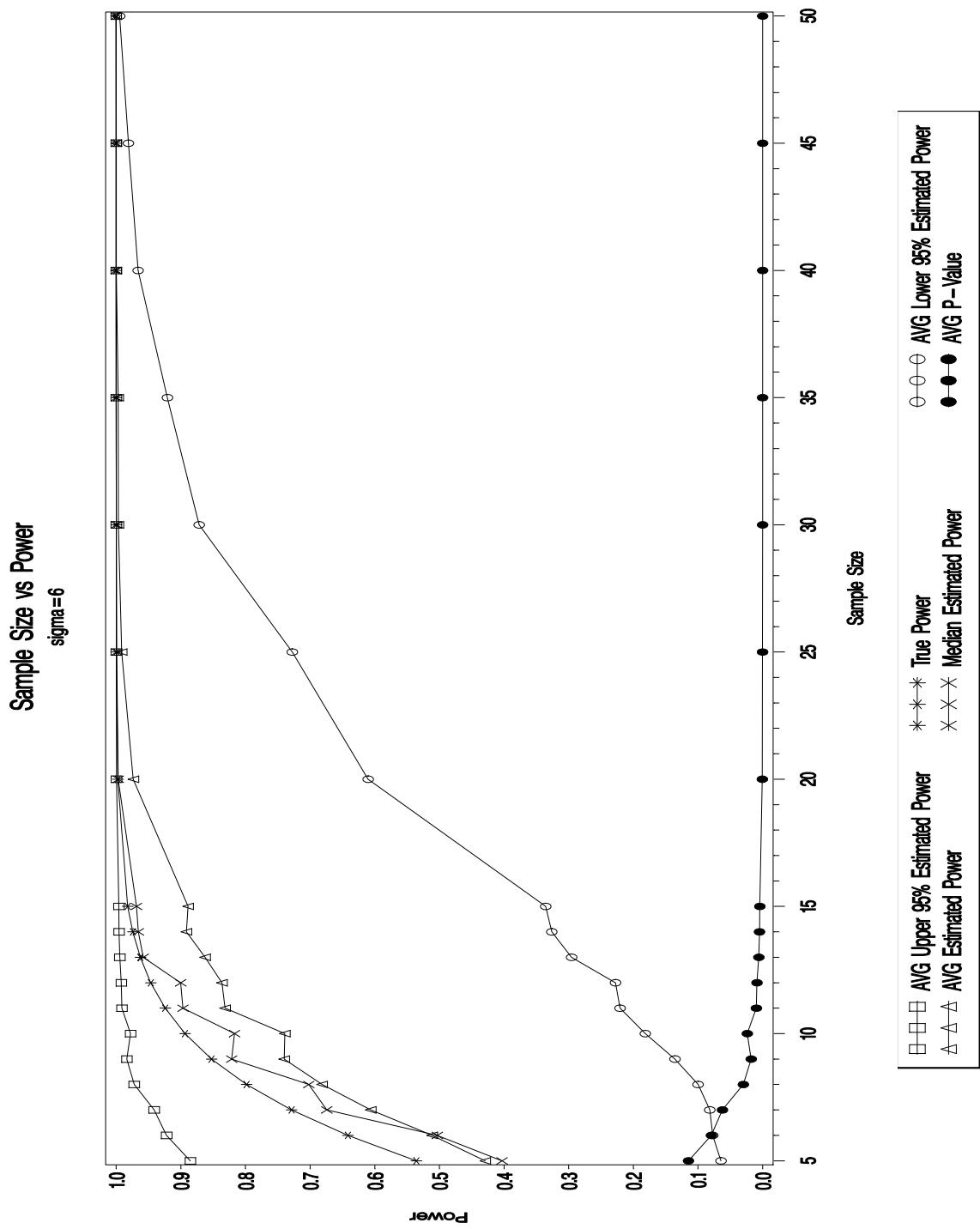
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



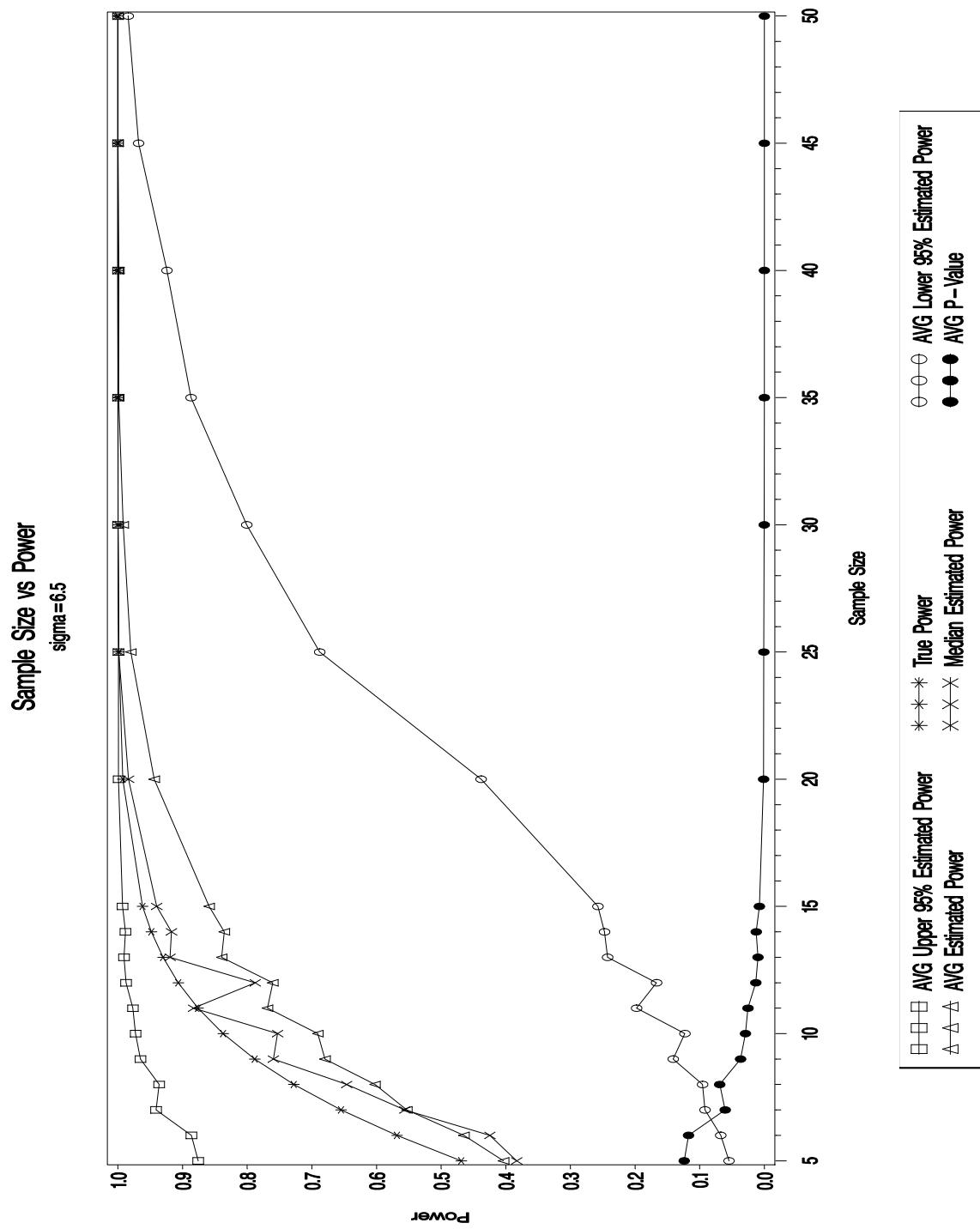
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



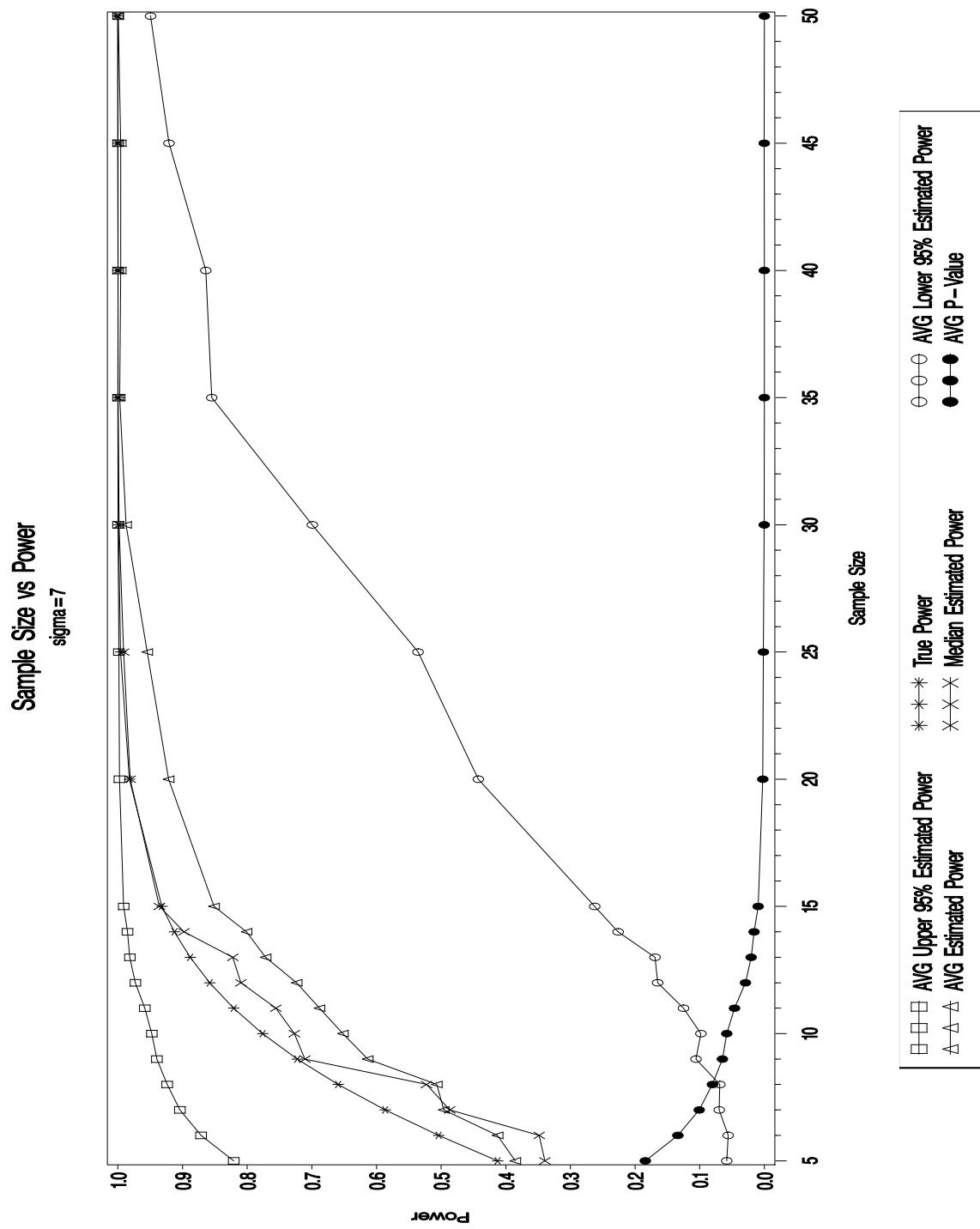
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



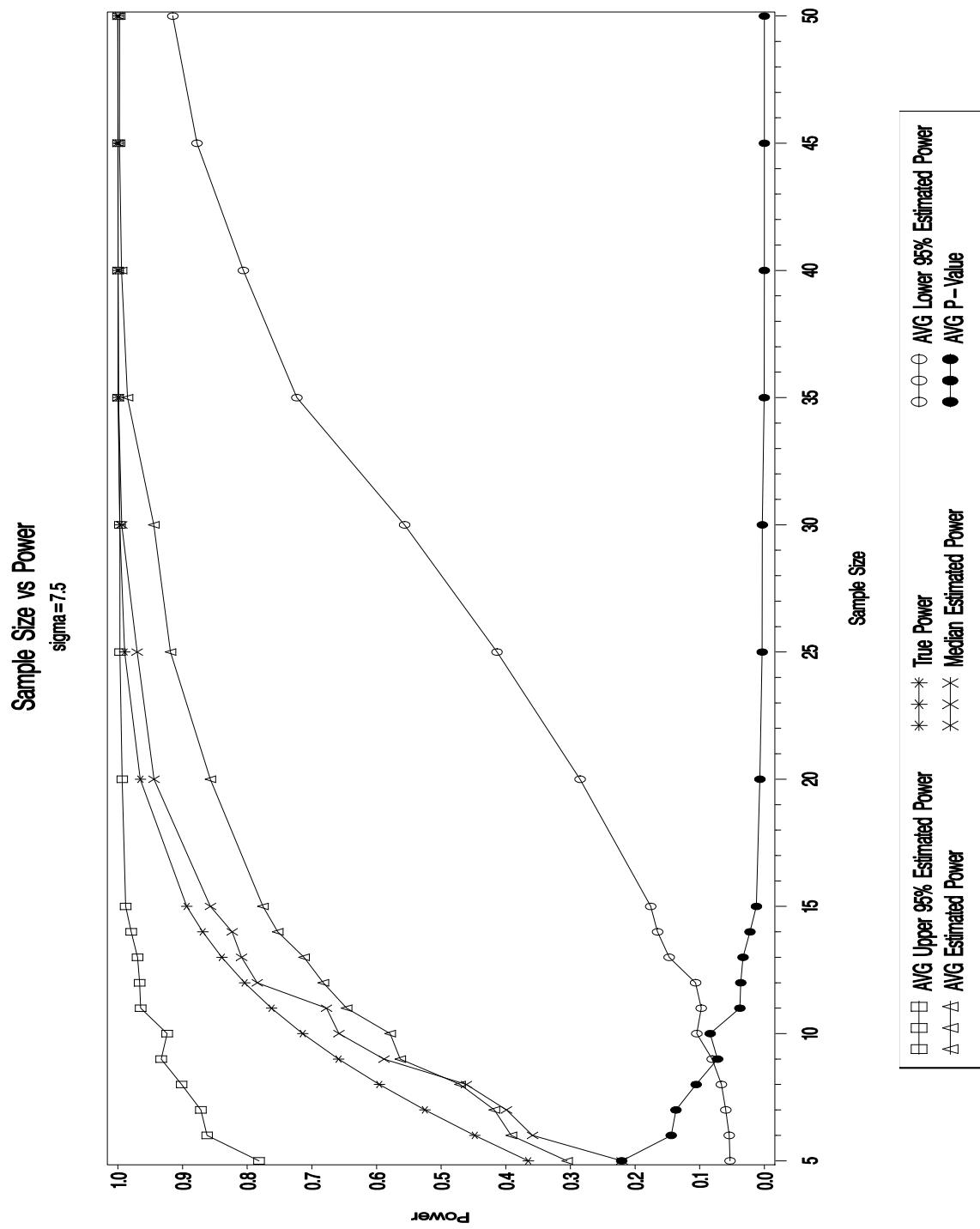
Appendix C: Graphs of Treatment Arrangement 30, 35, 40



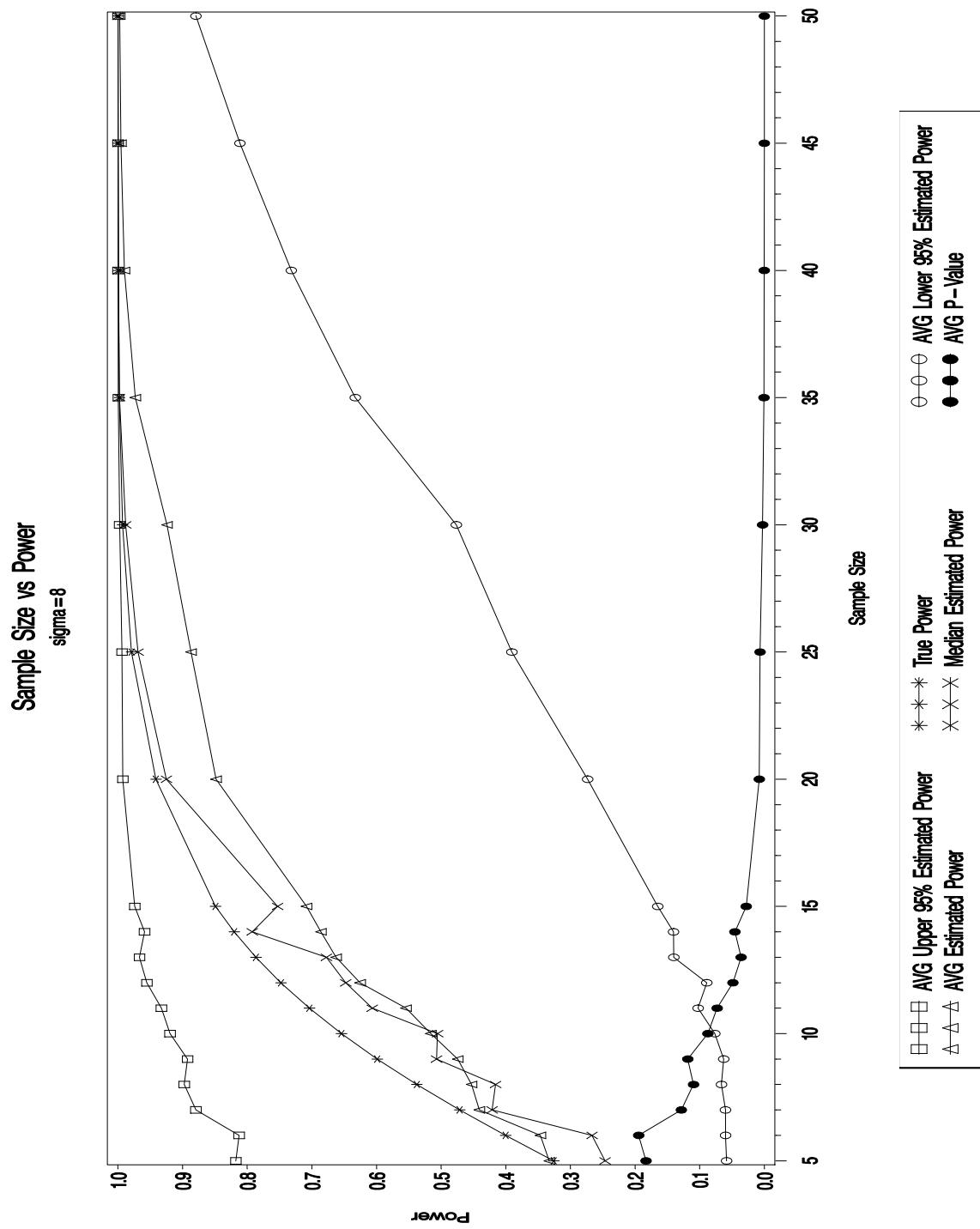
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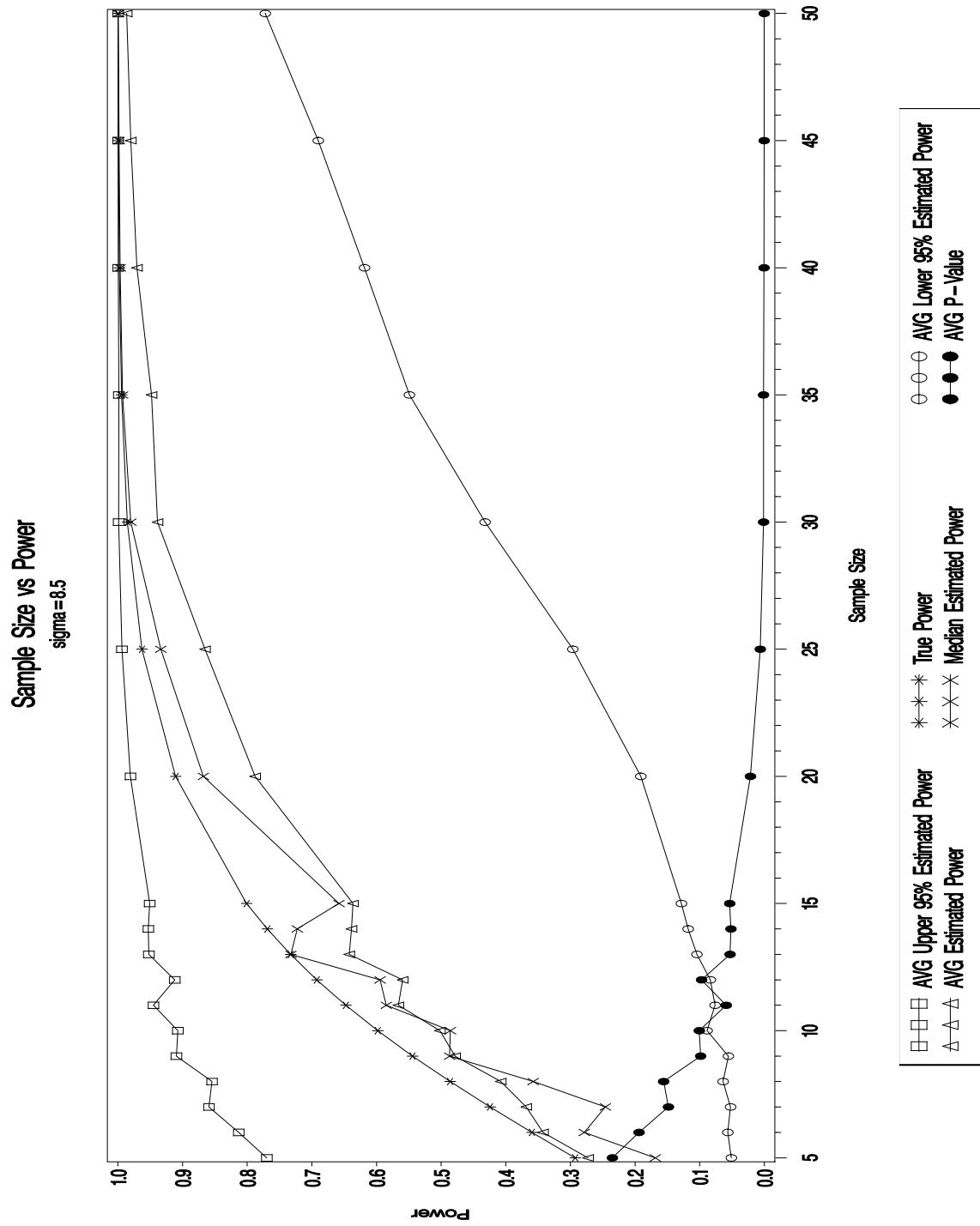
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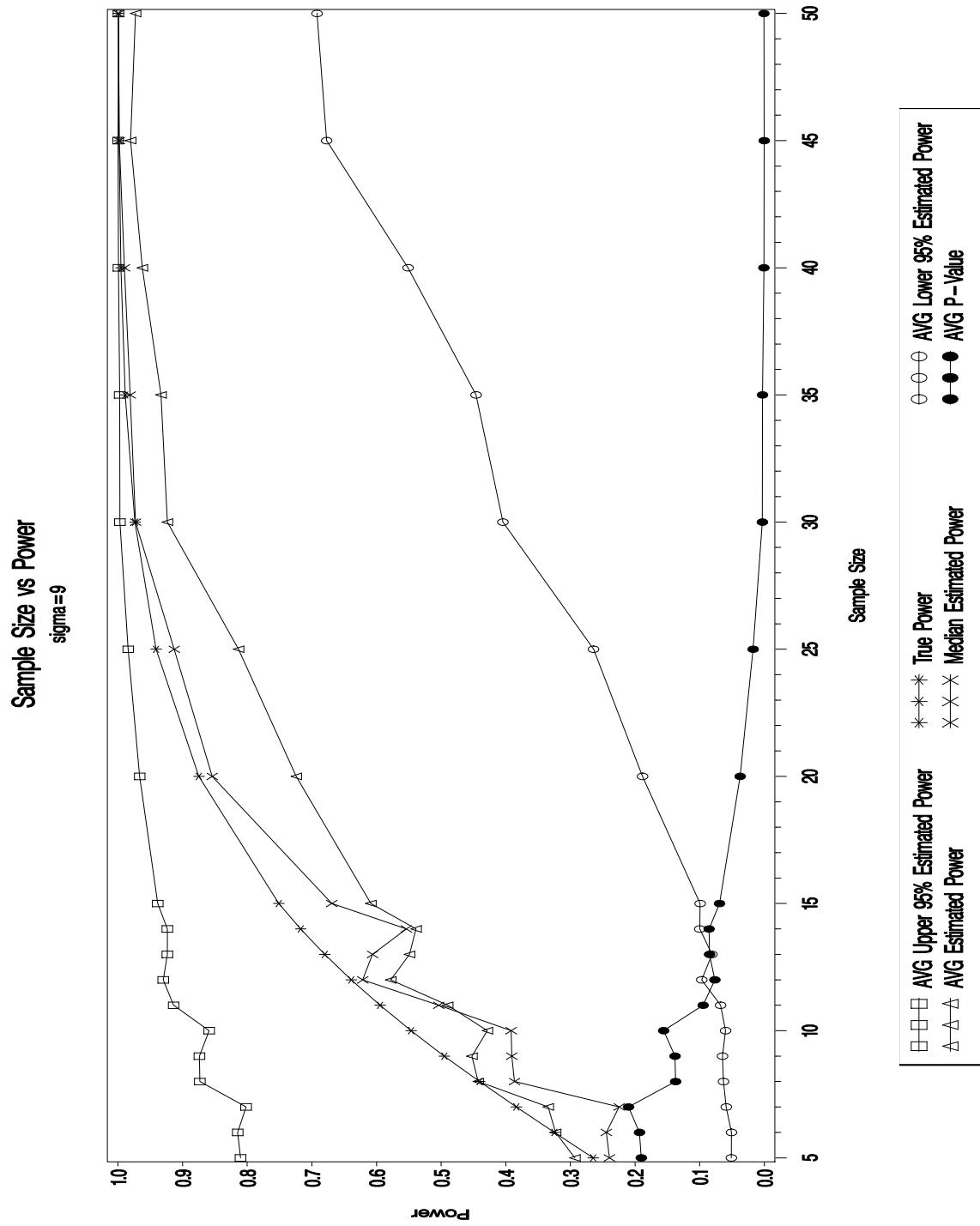
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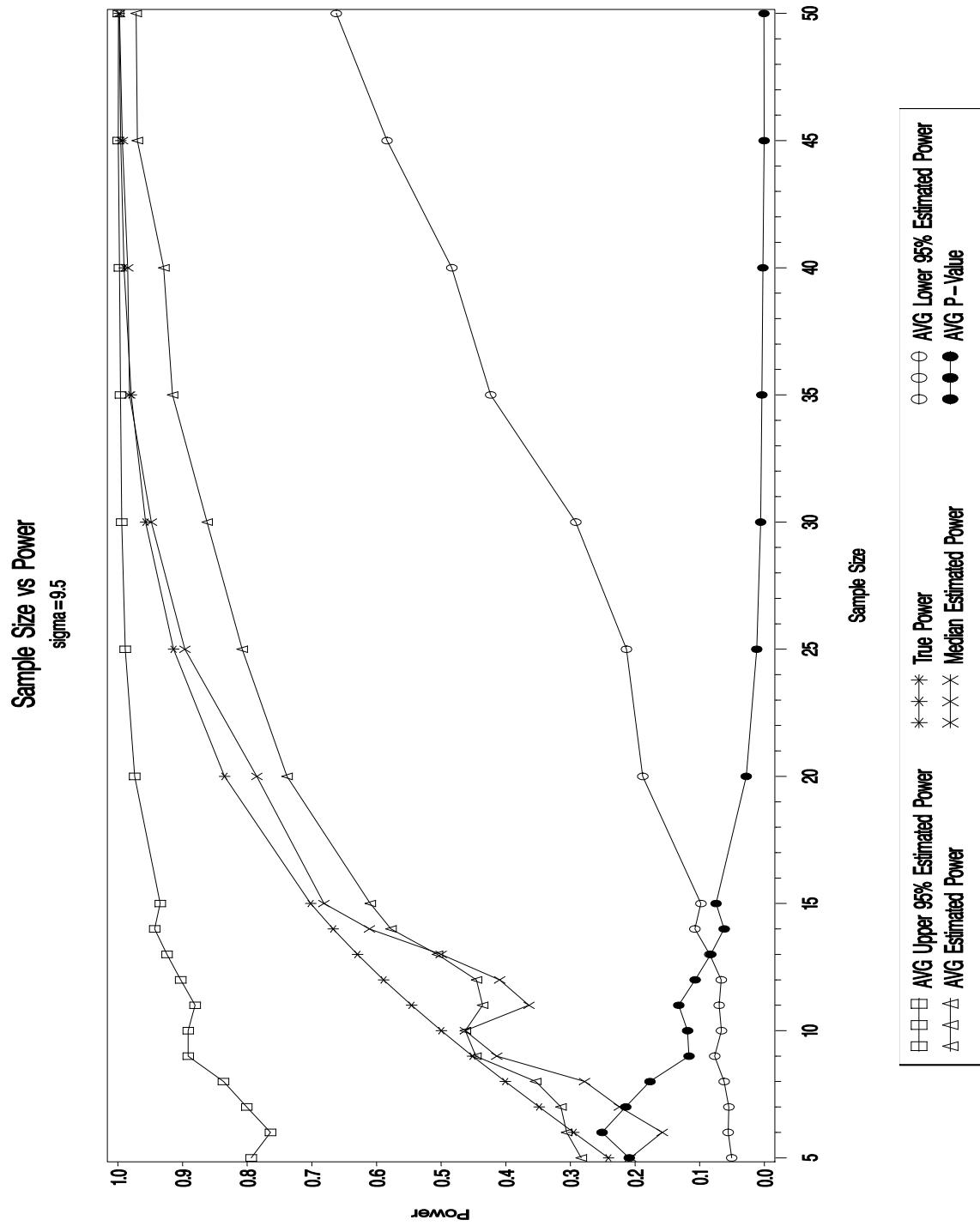
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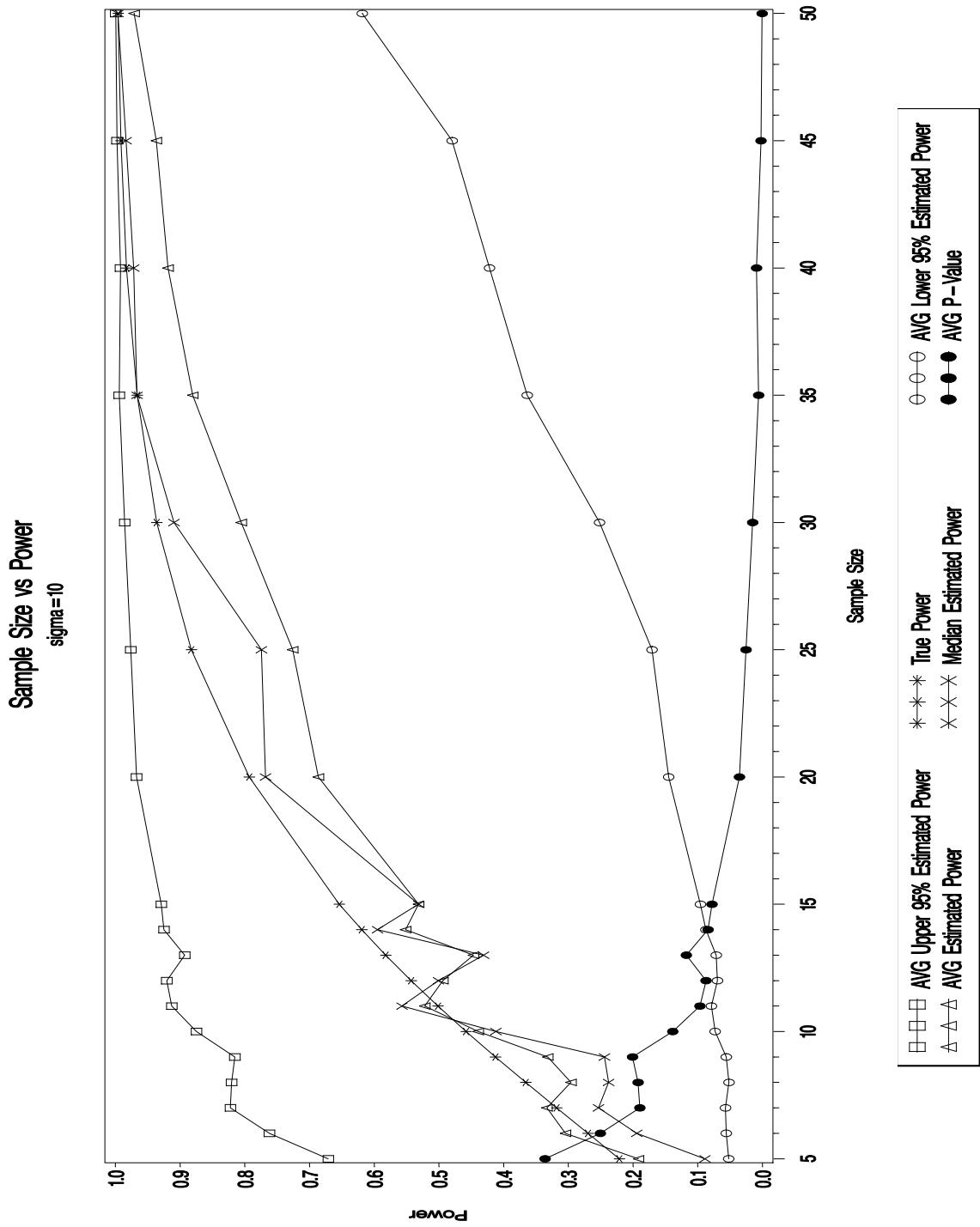
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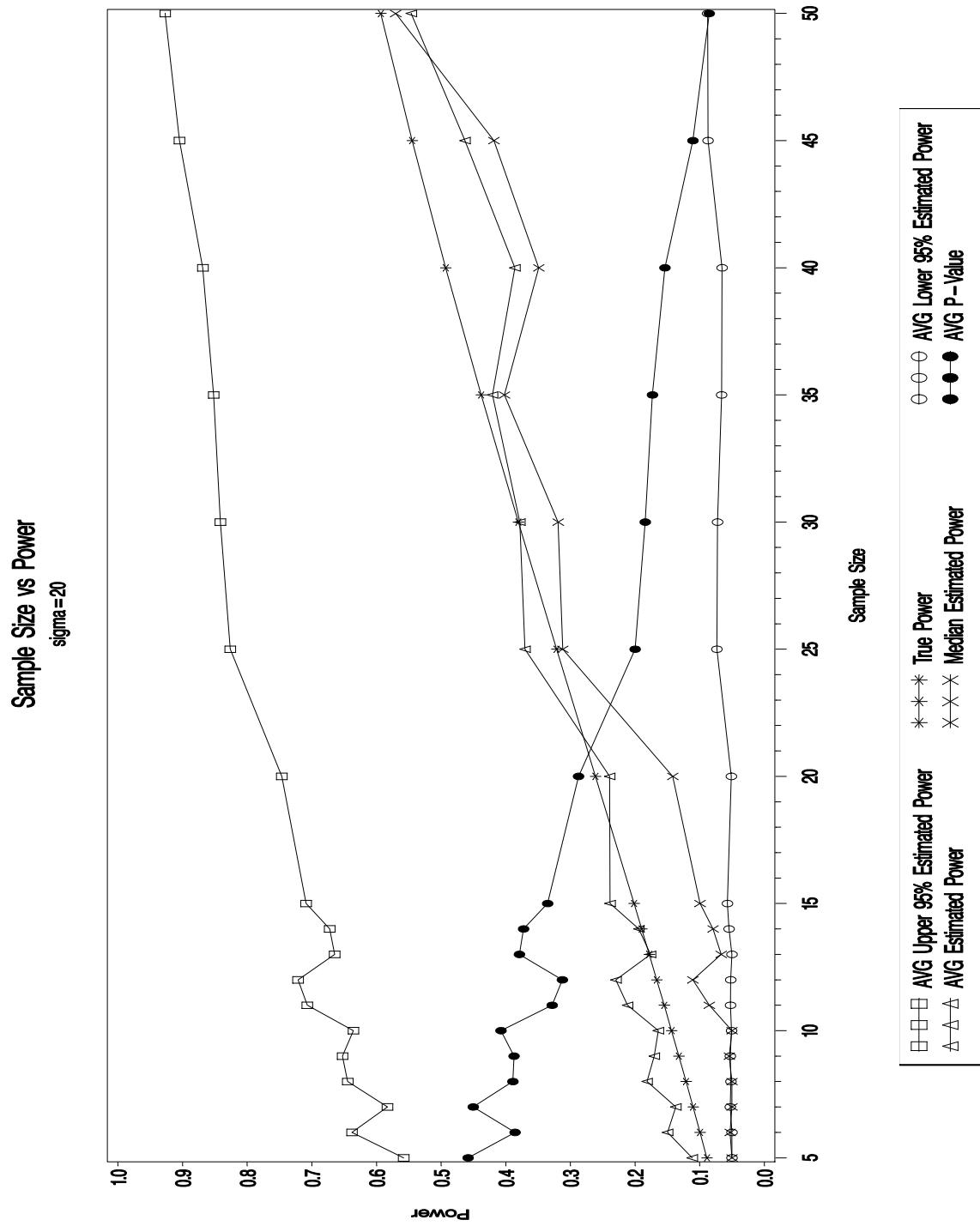
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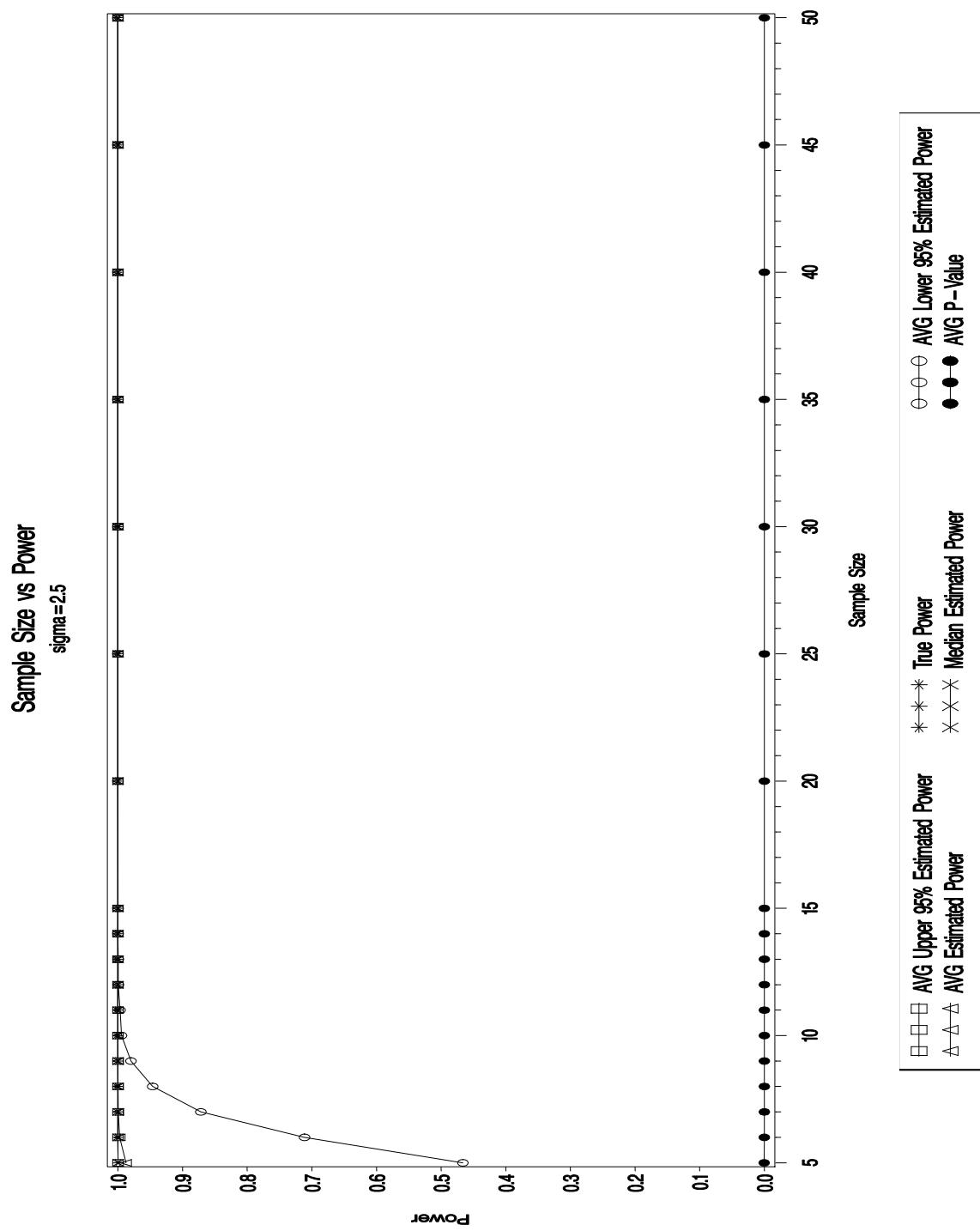
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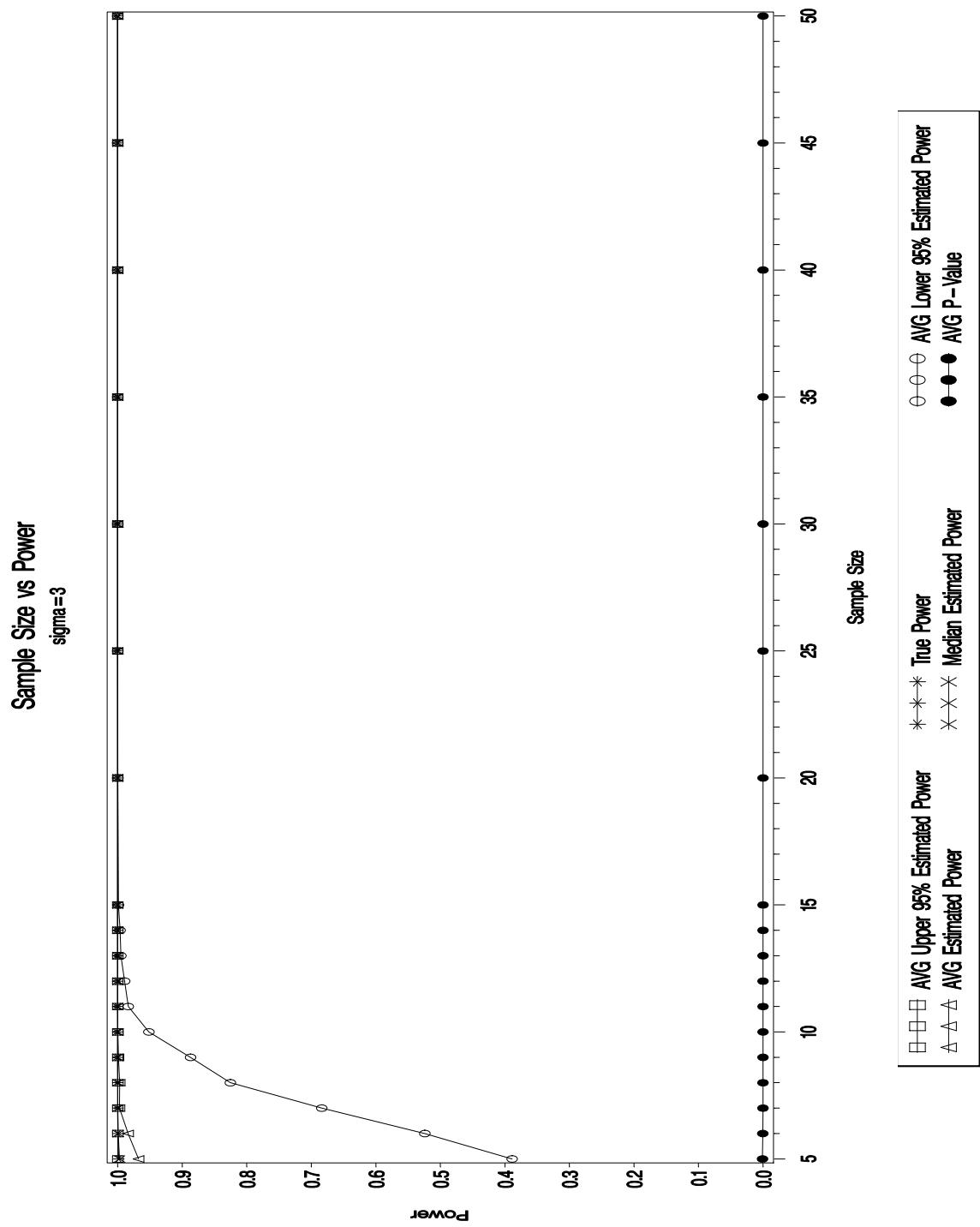
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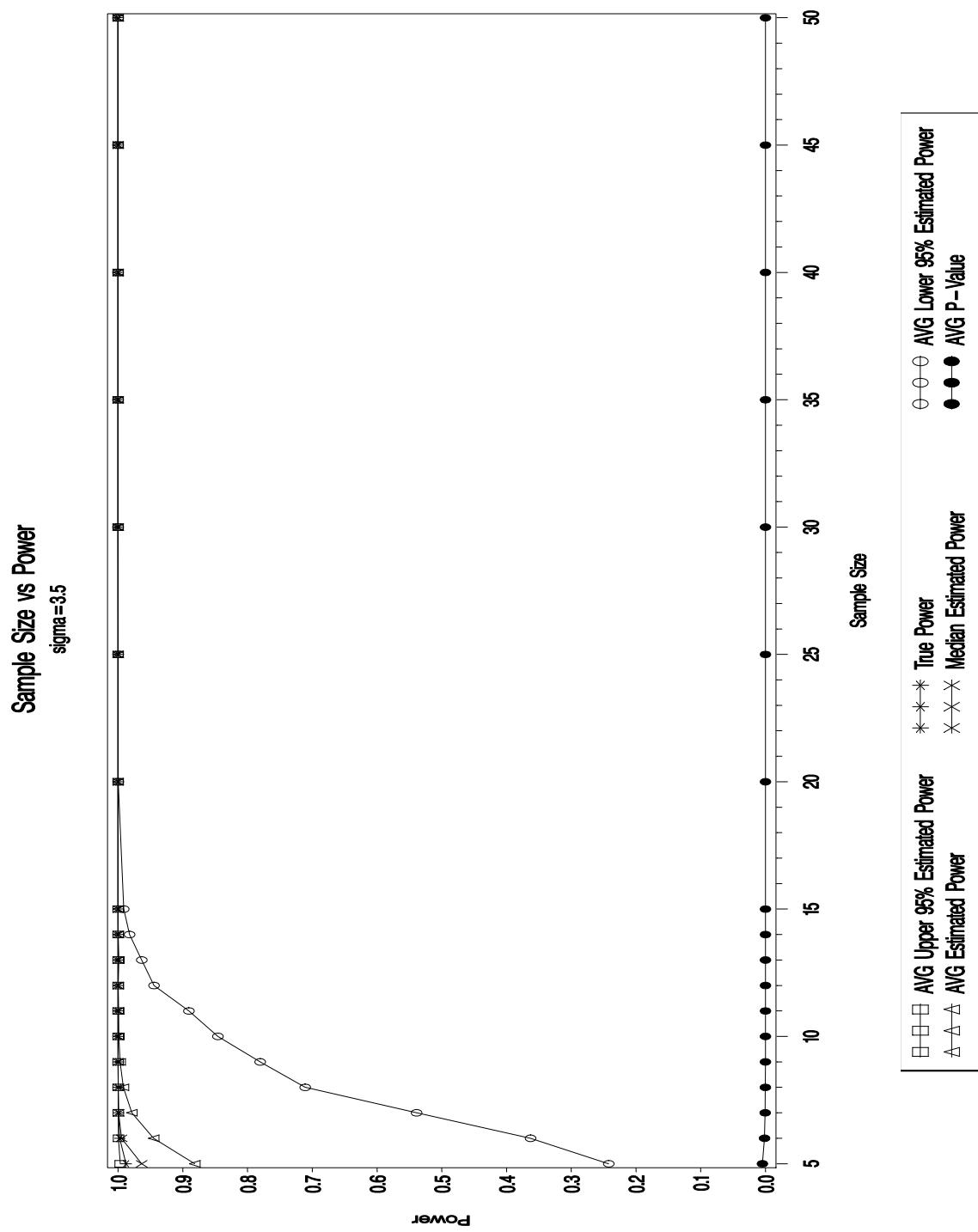
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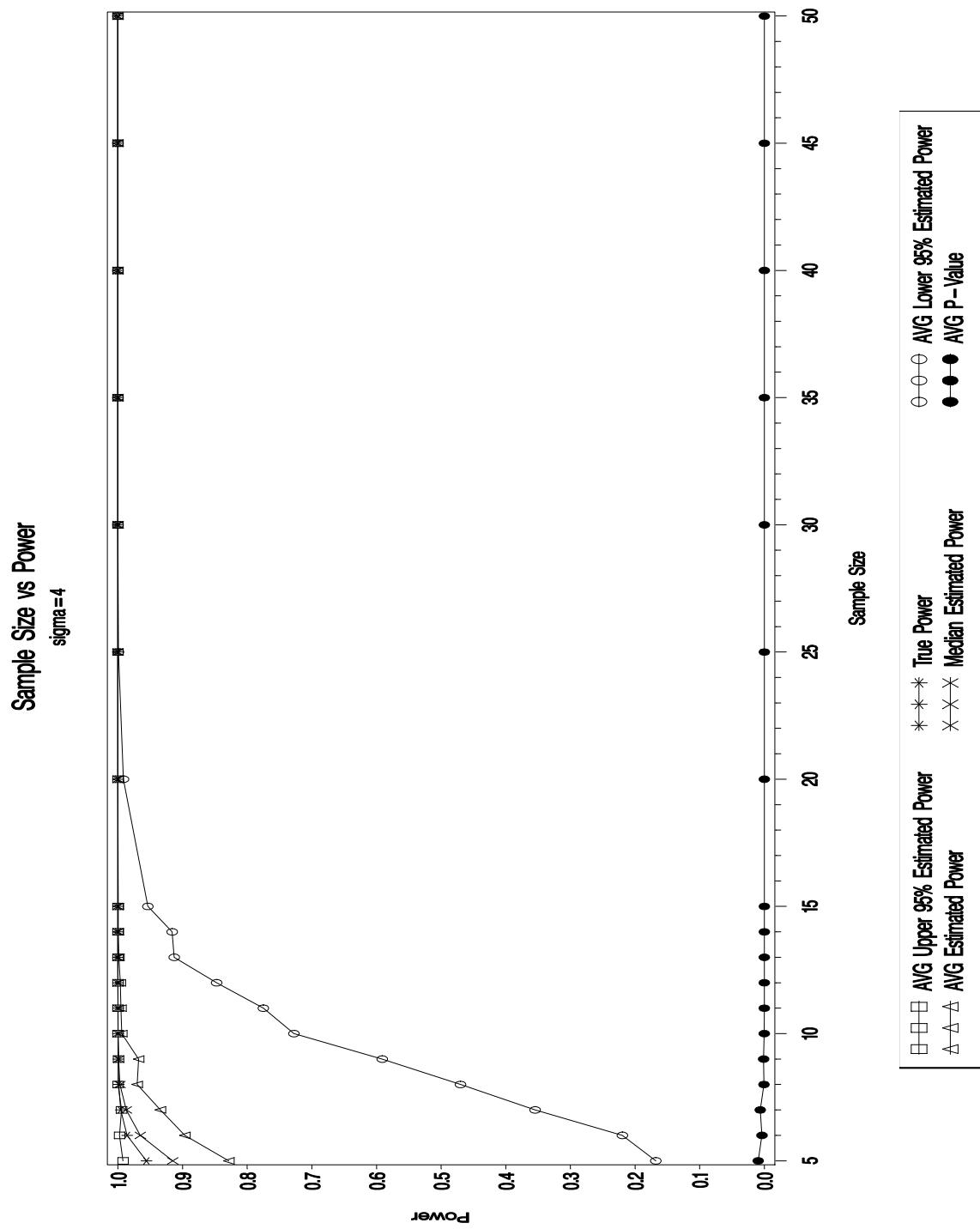
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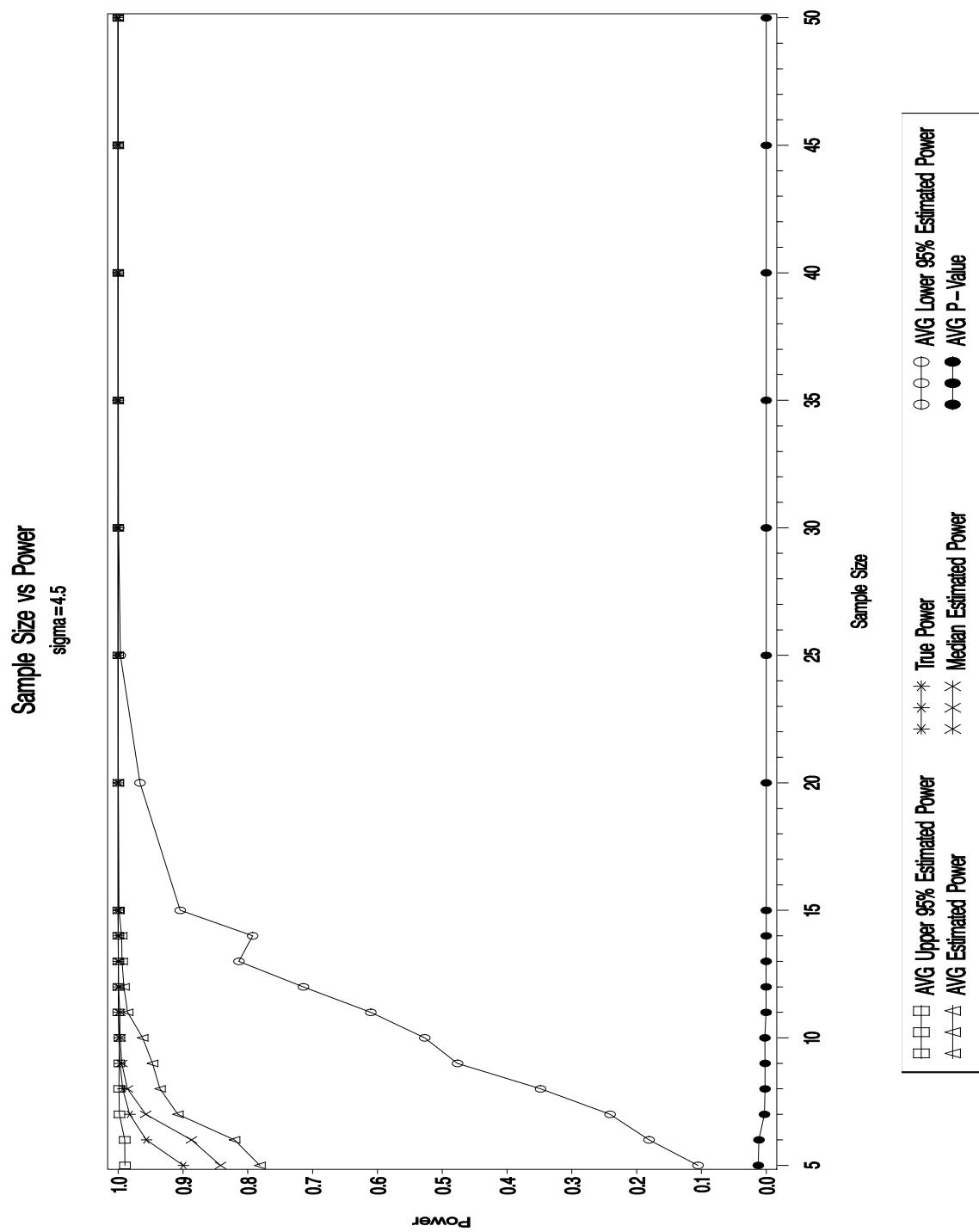
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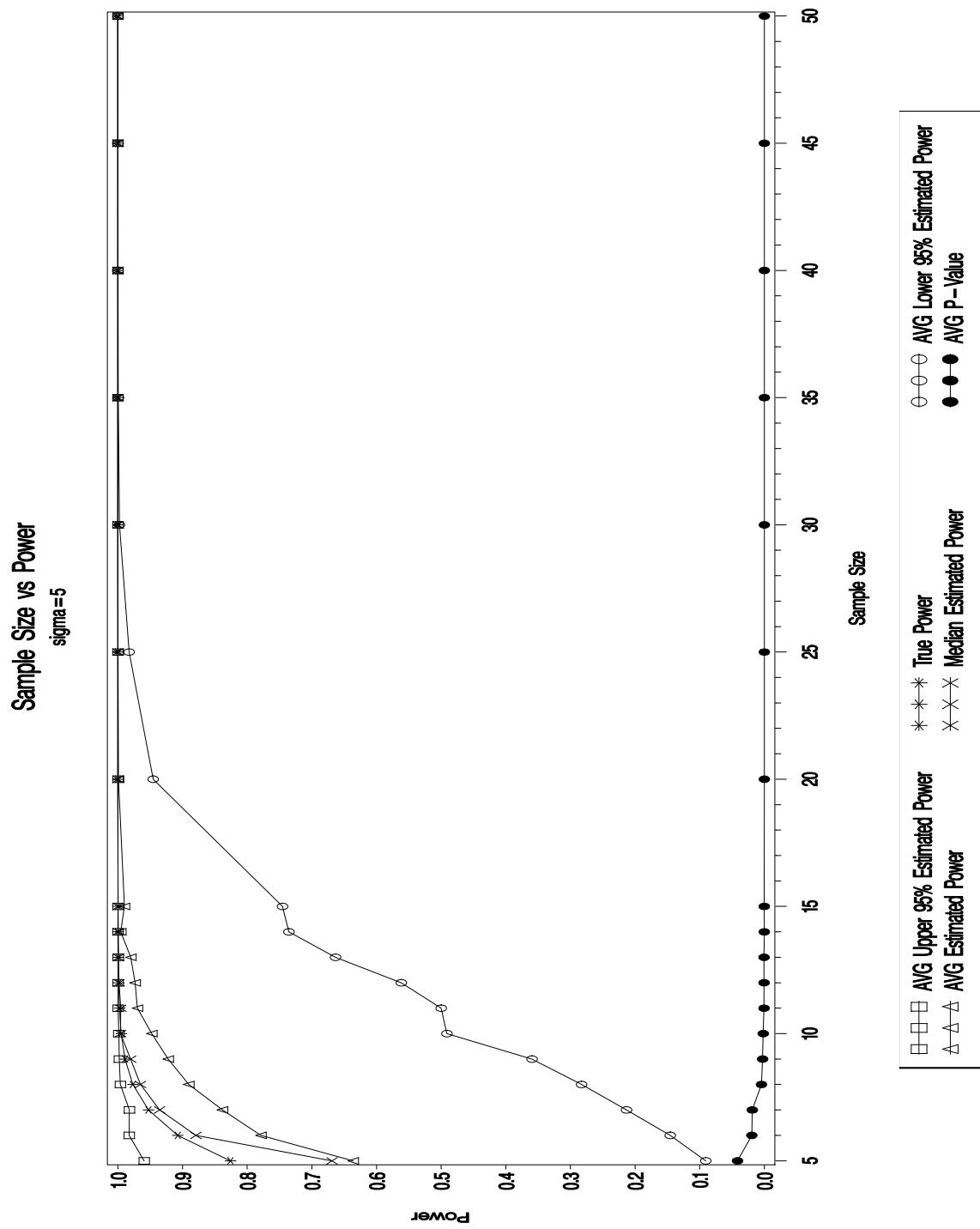
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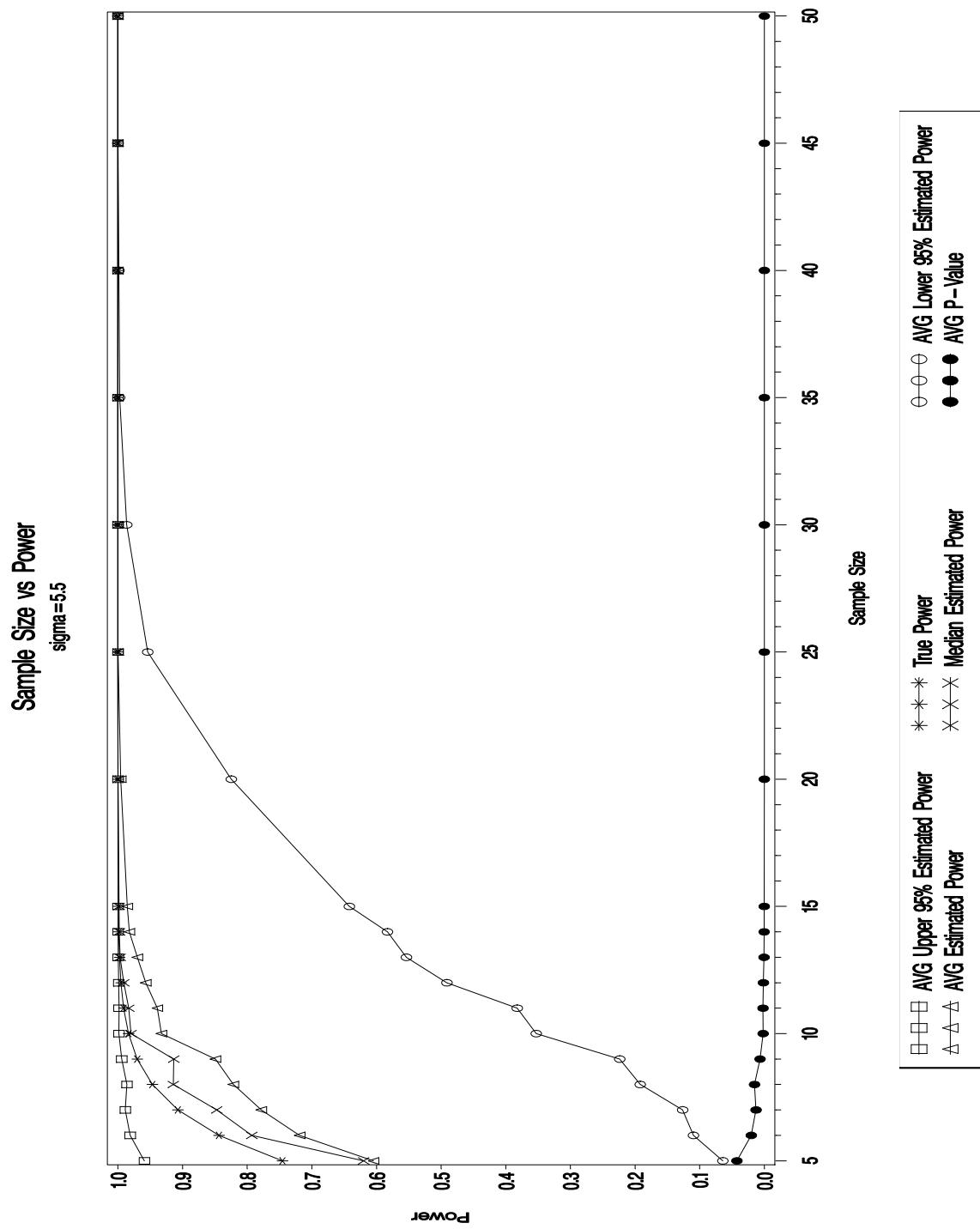
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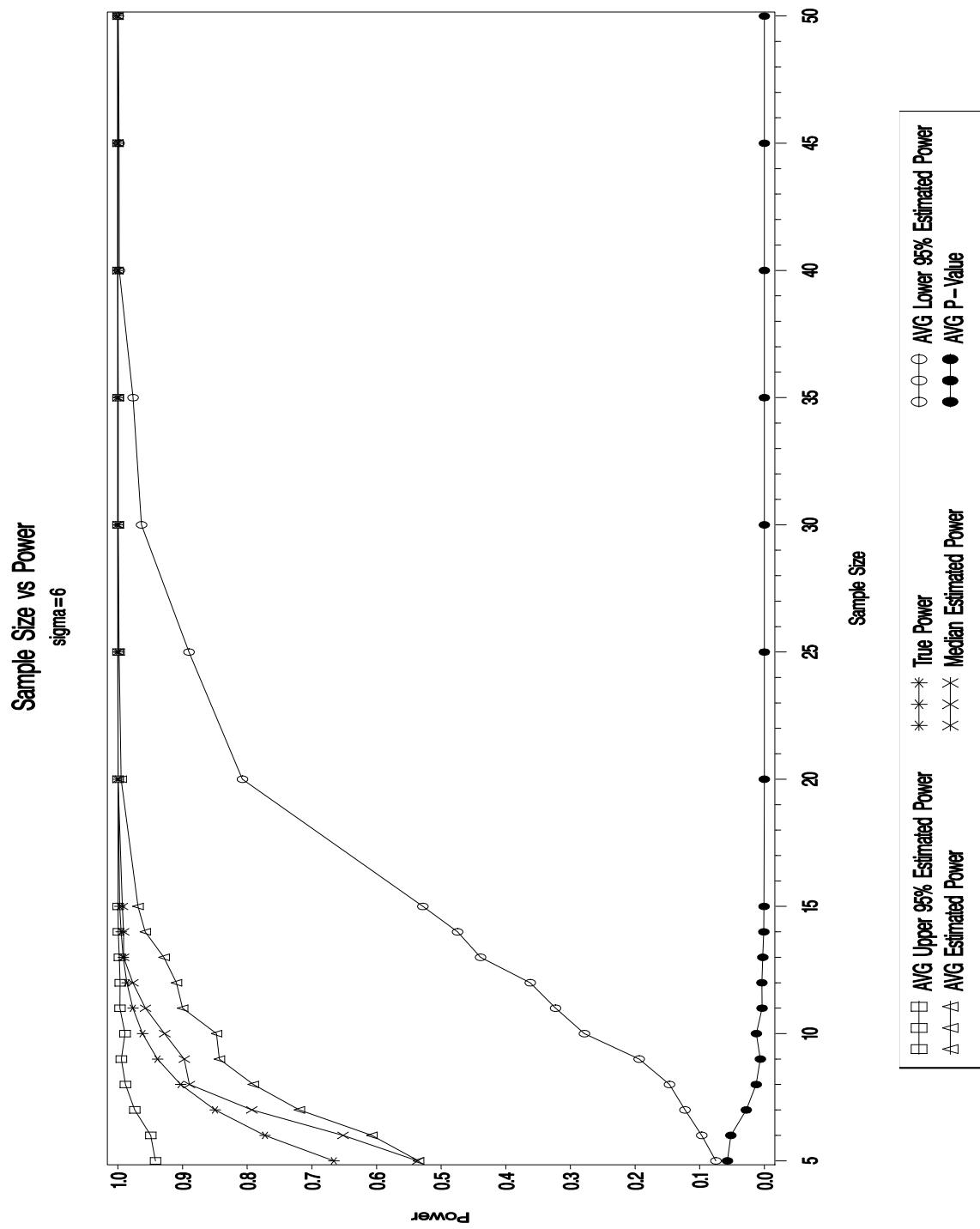
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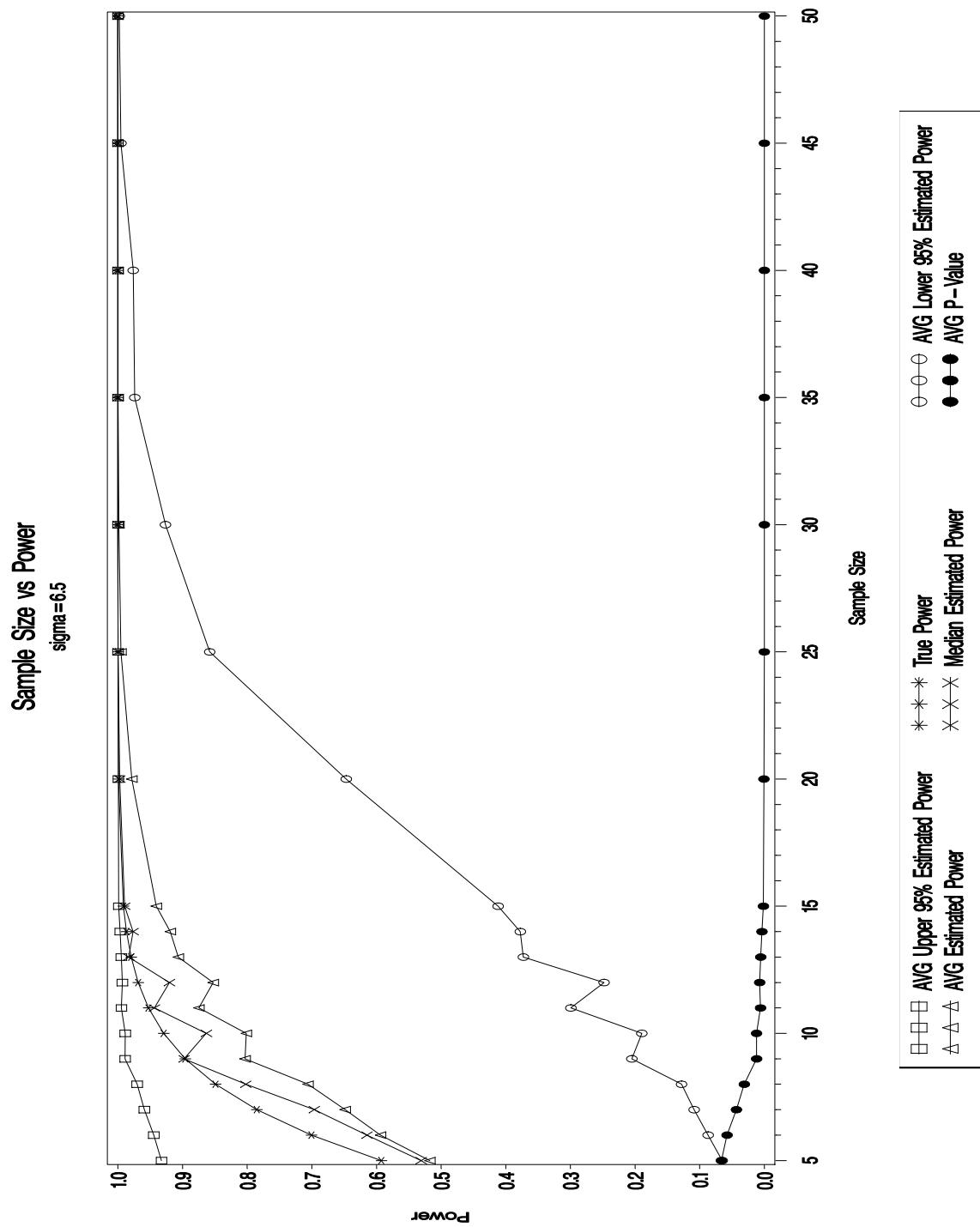
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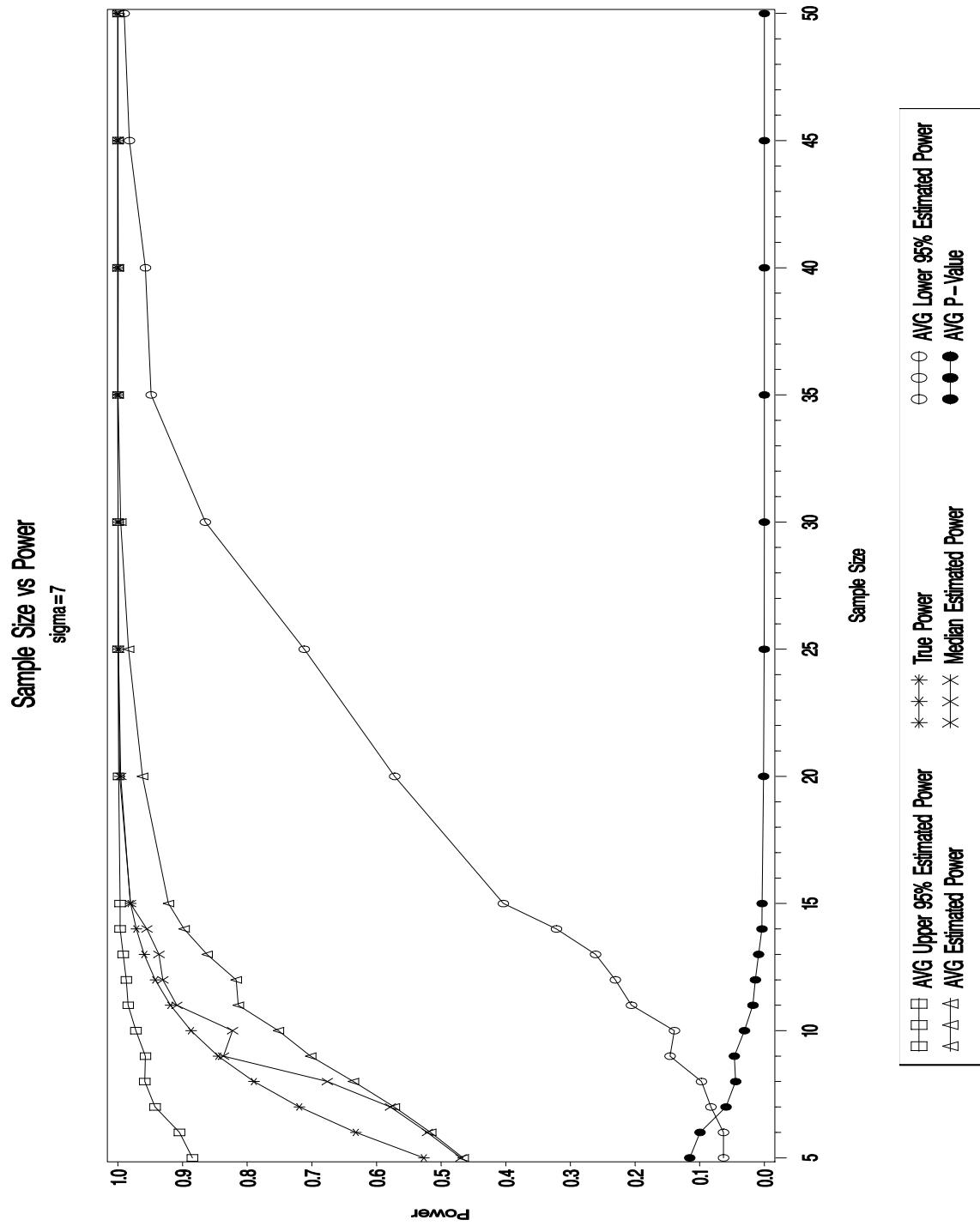
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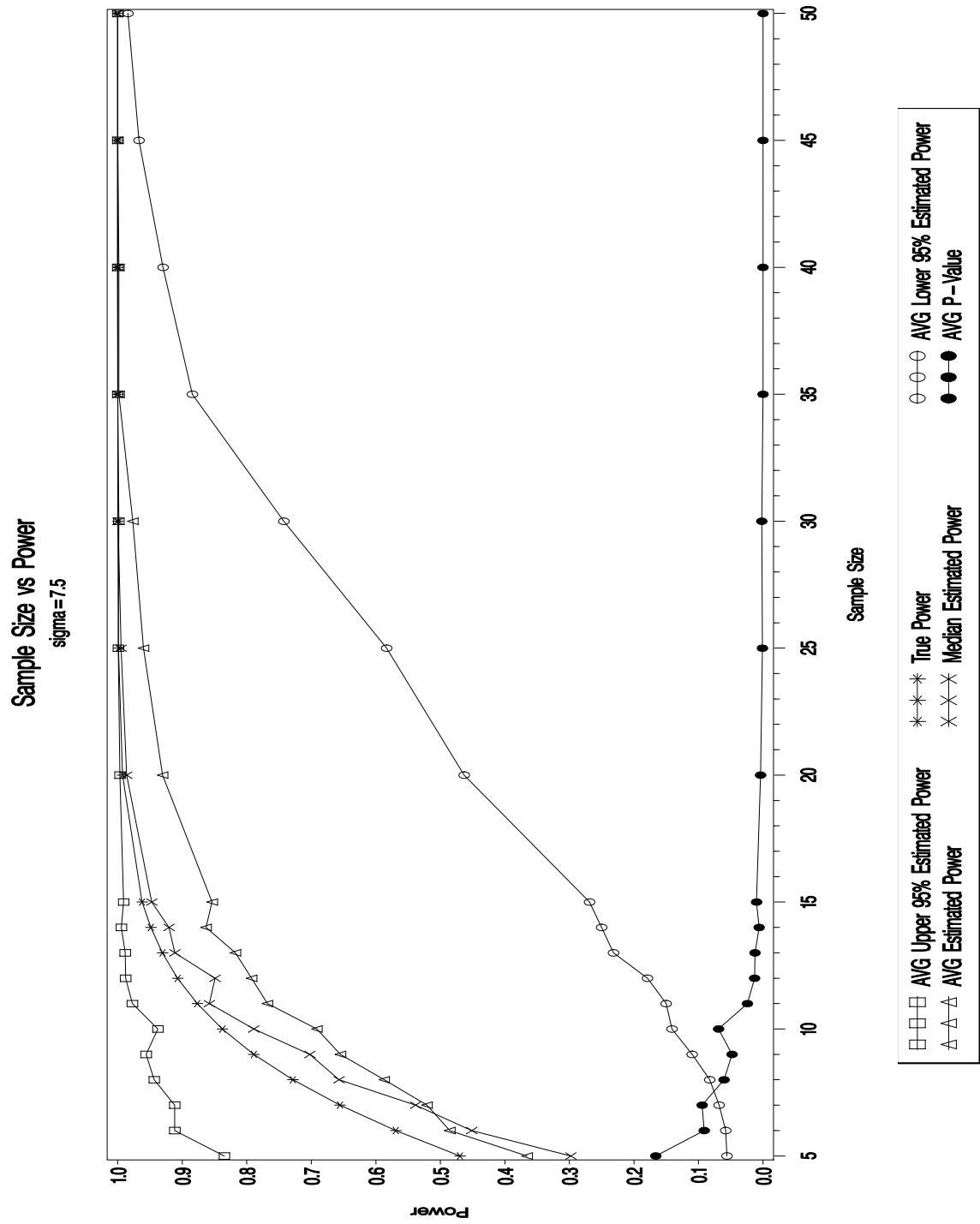
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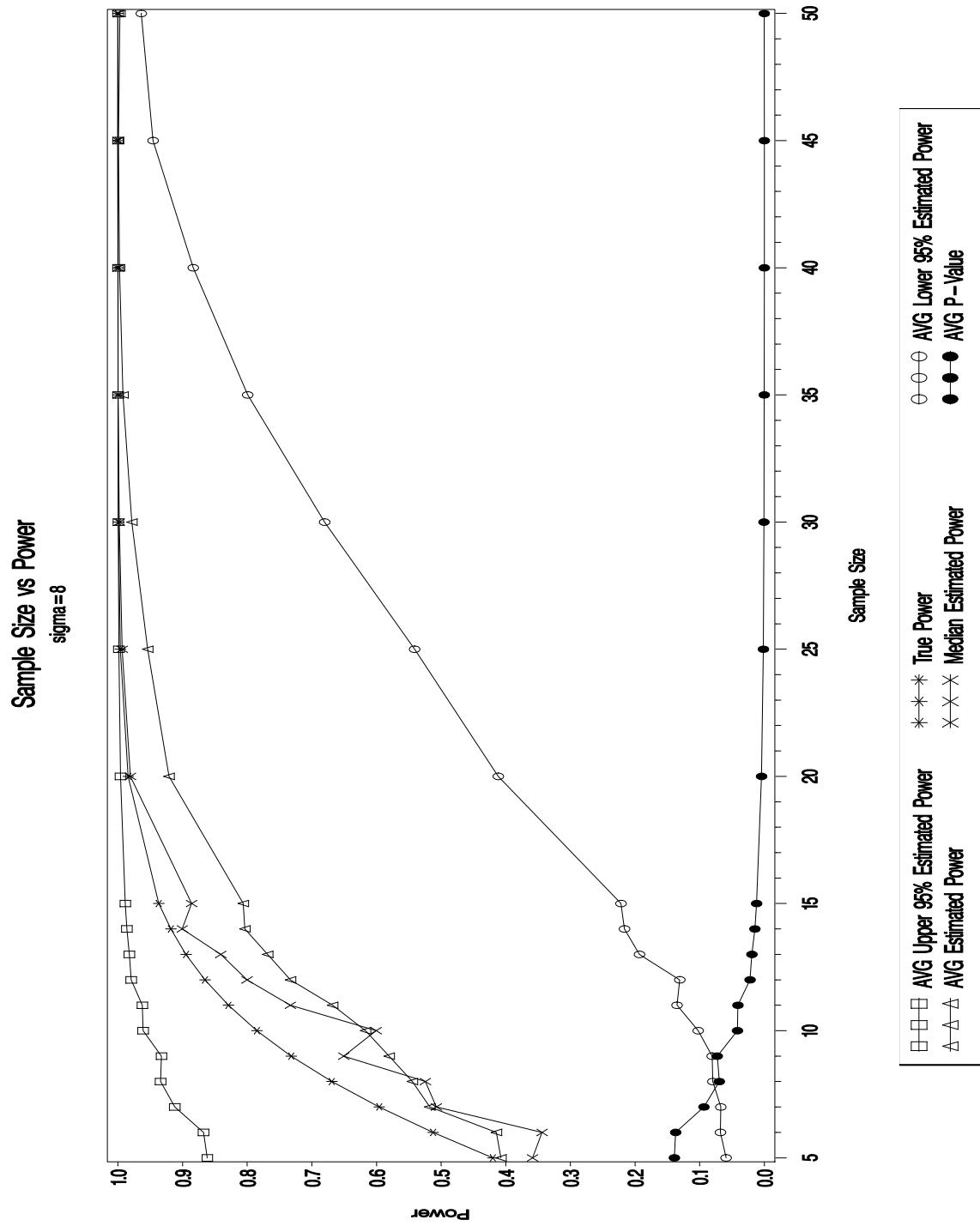
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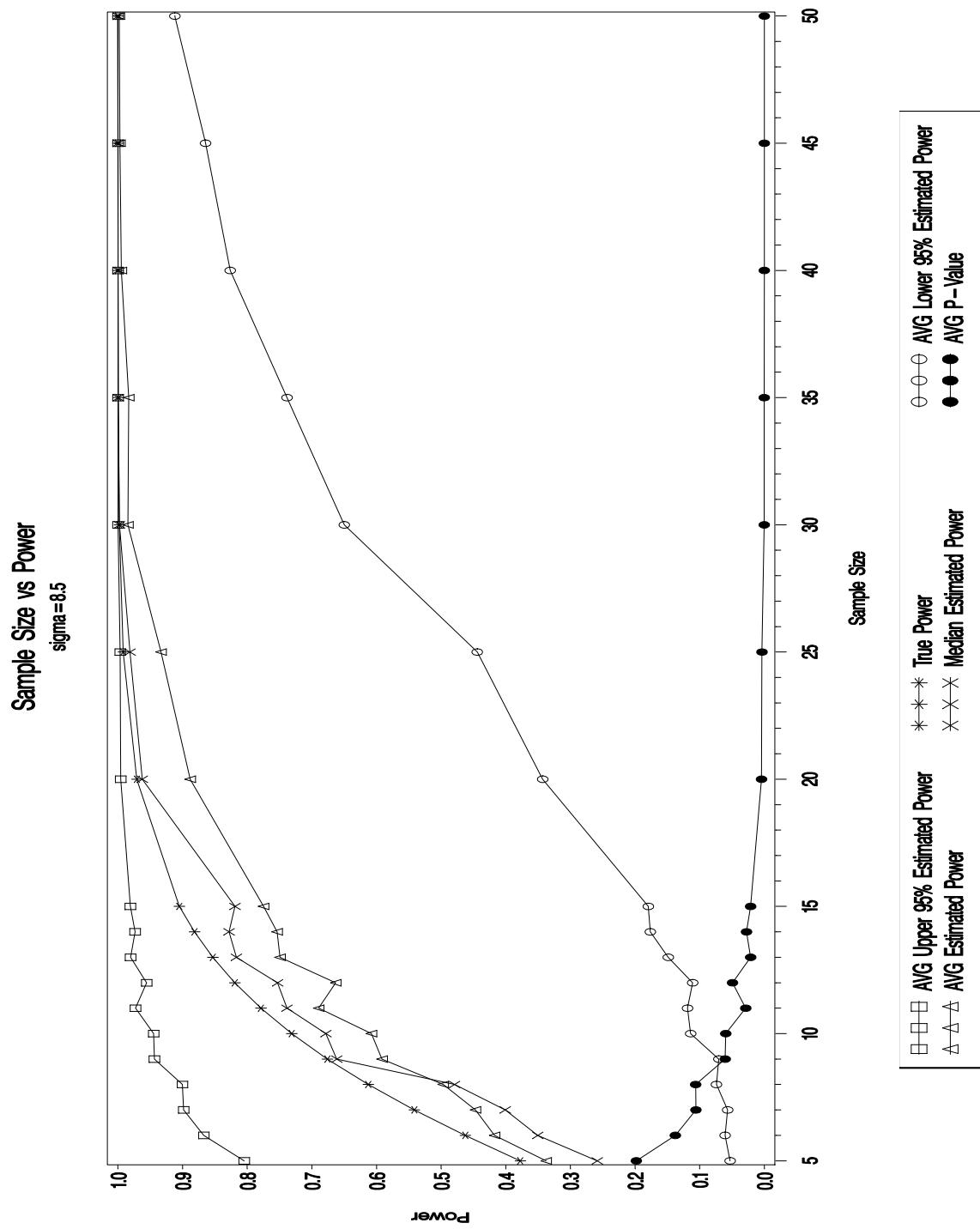
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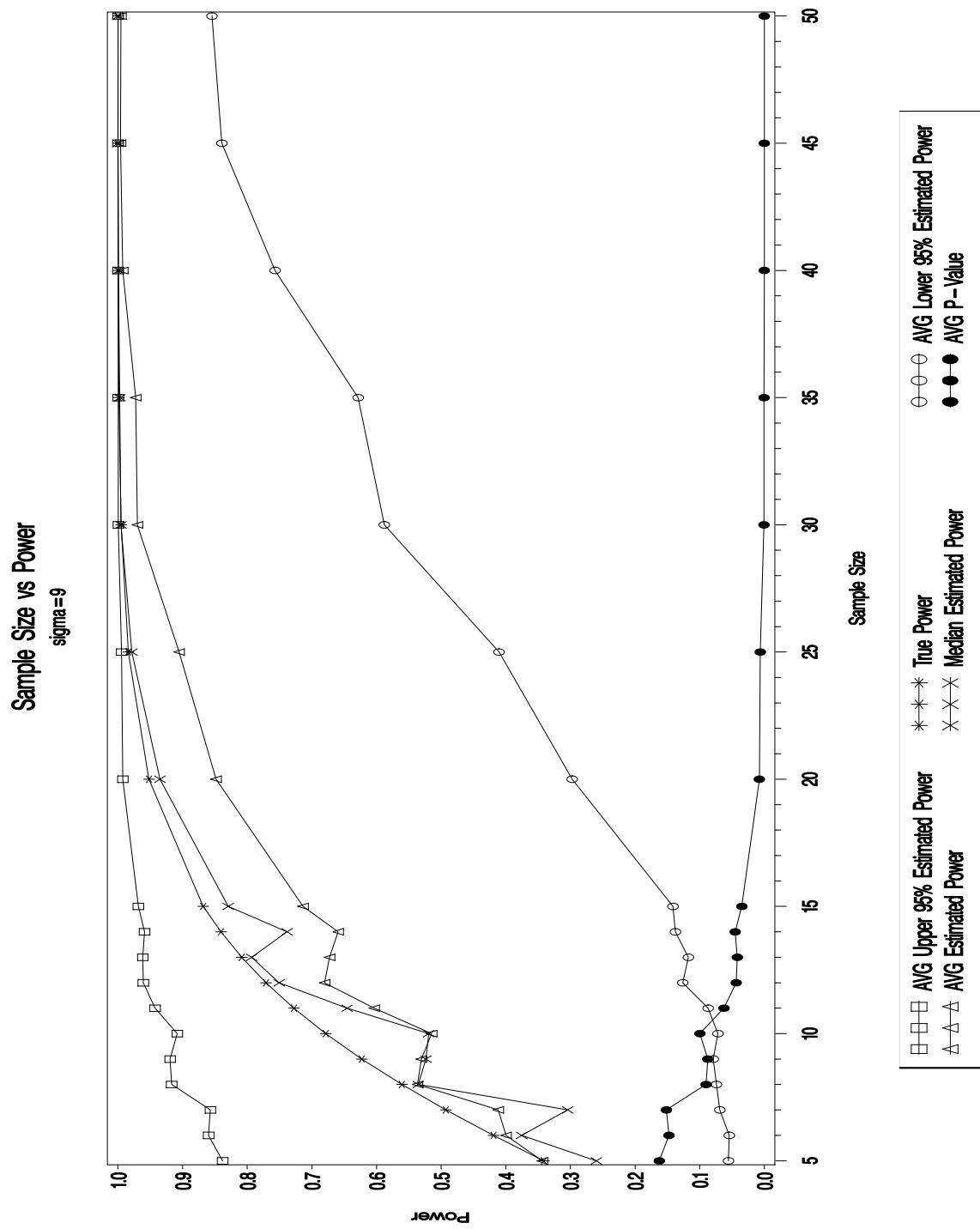
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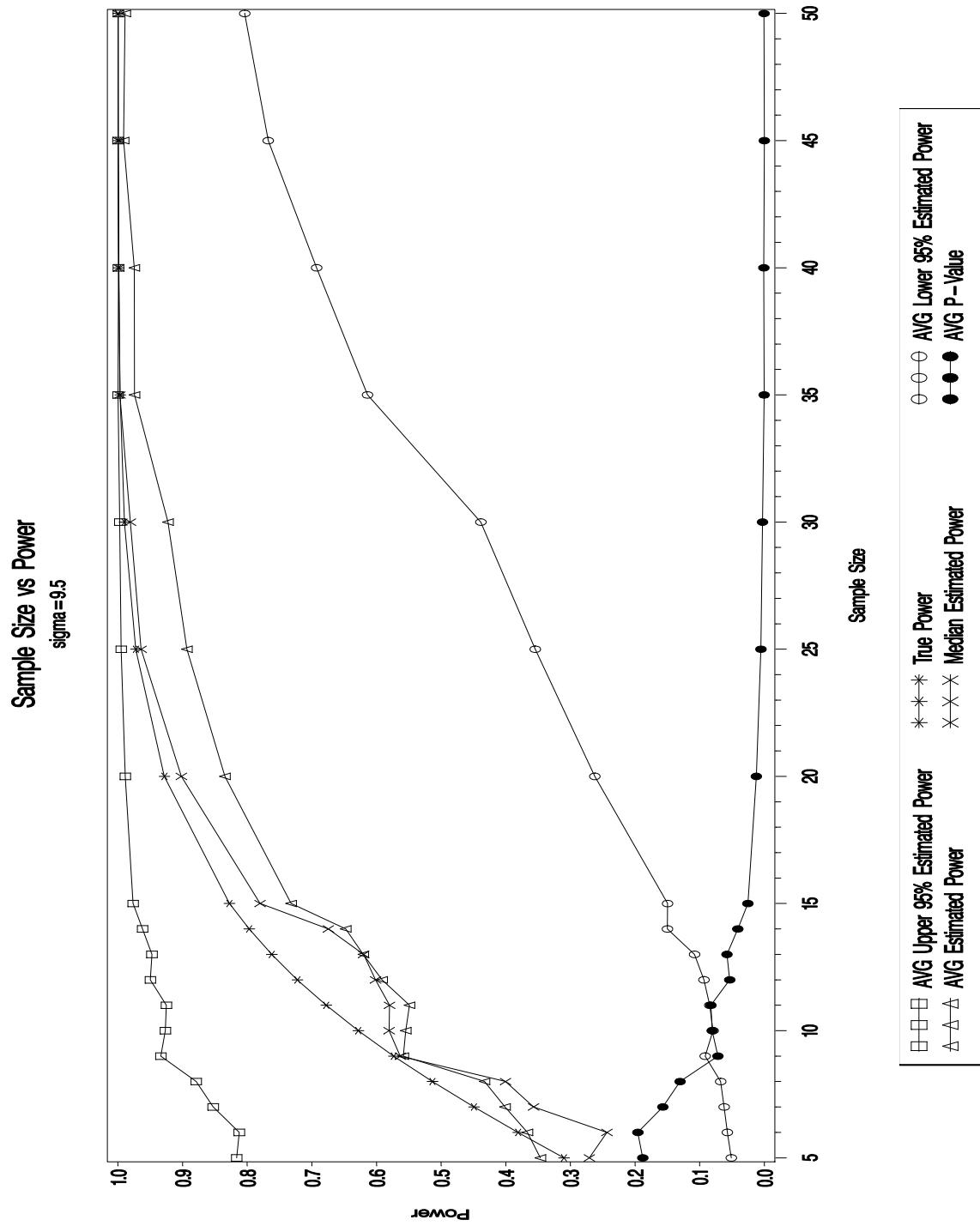
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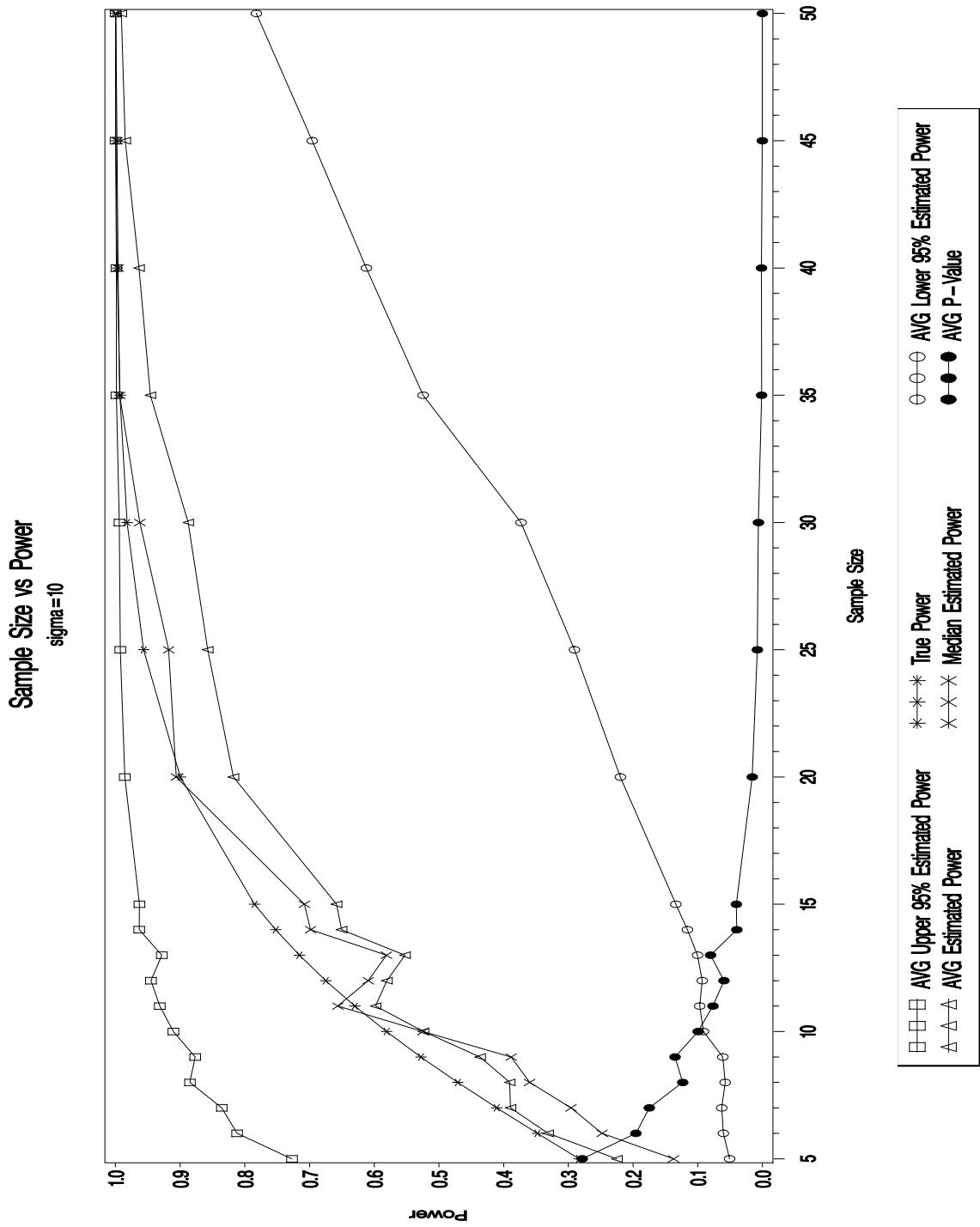
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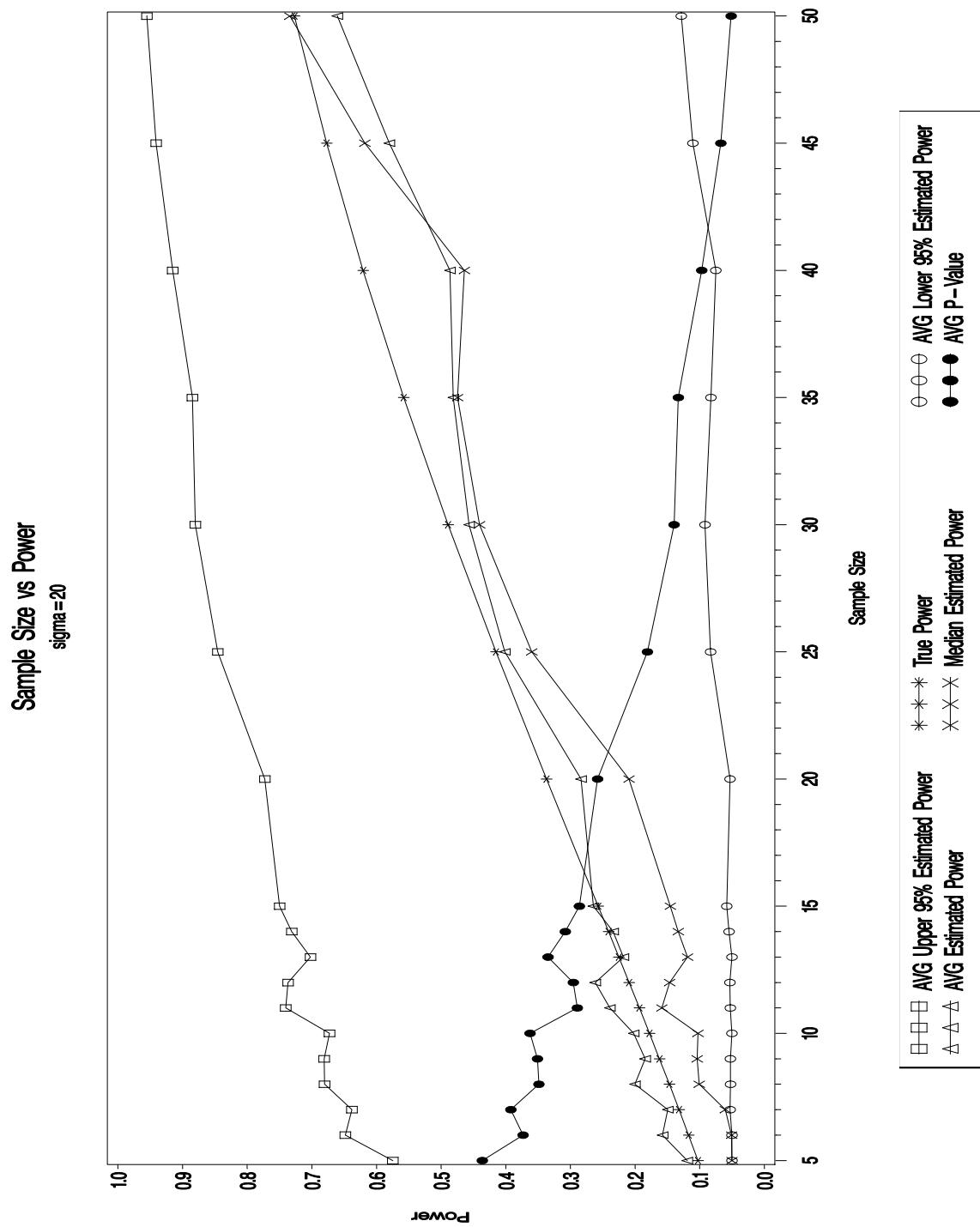
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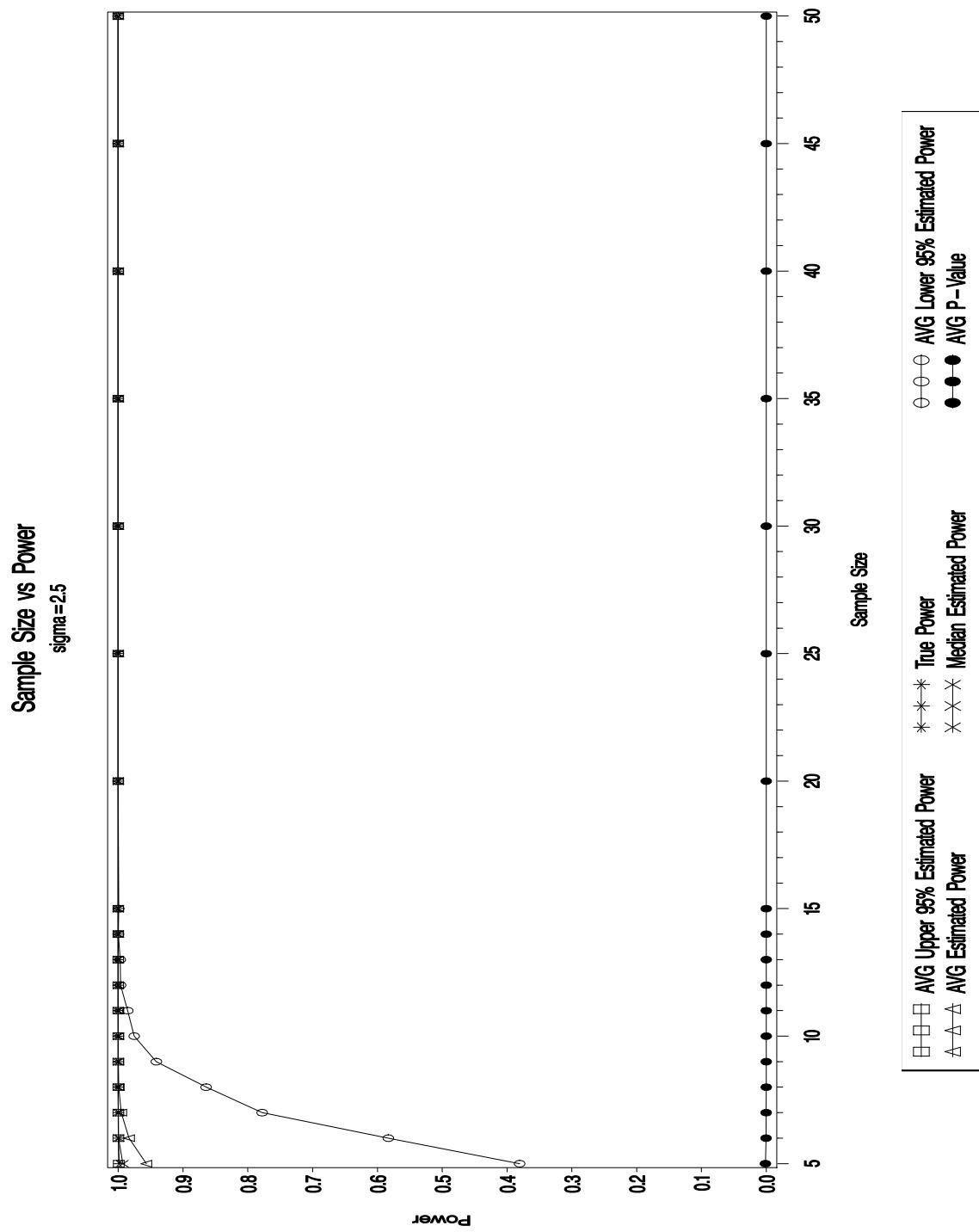
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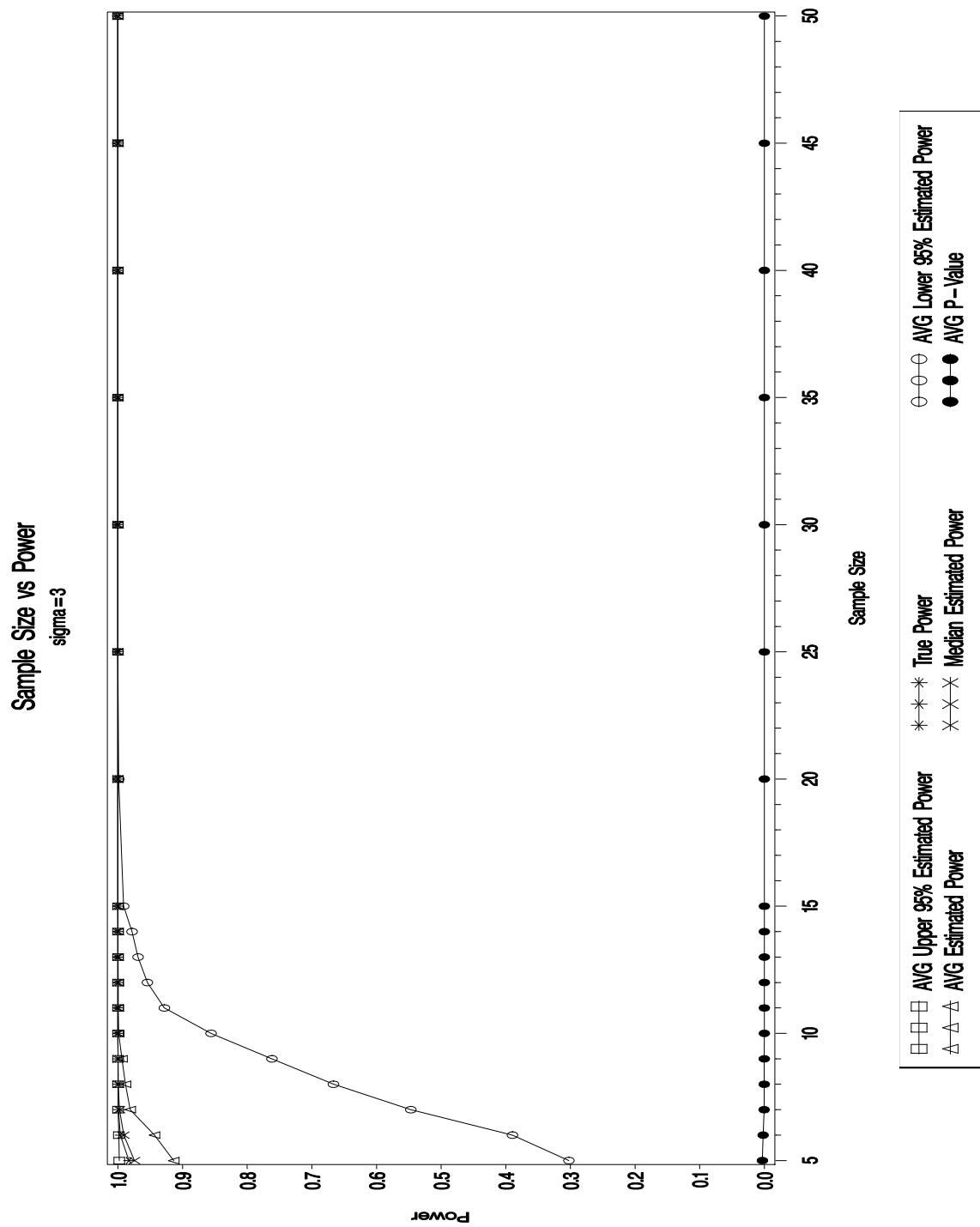
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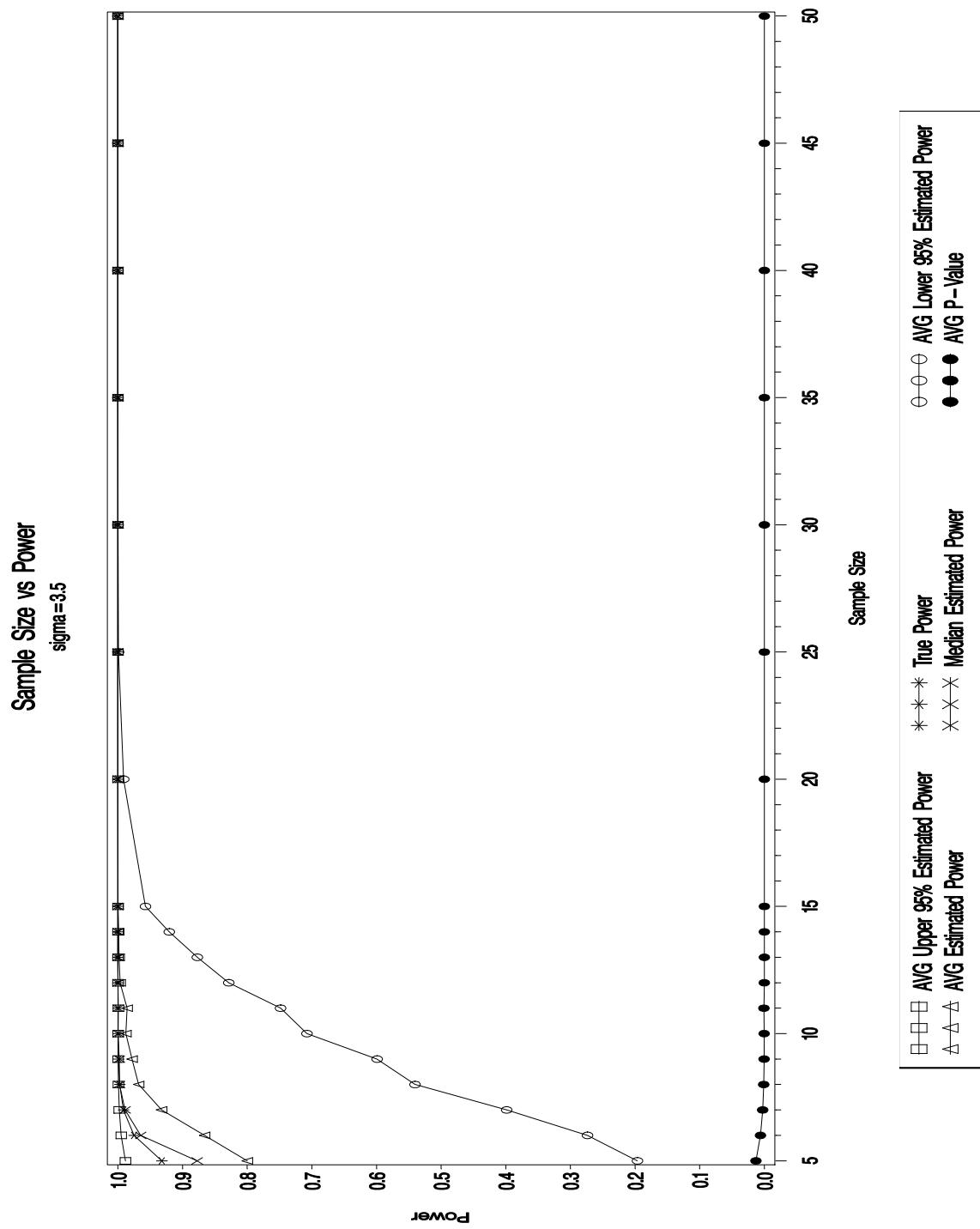
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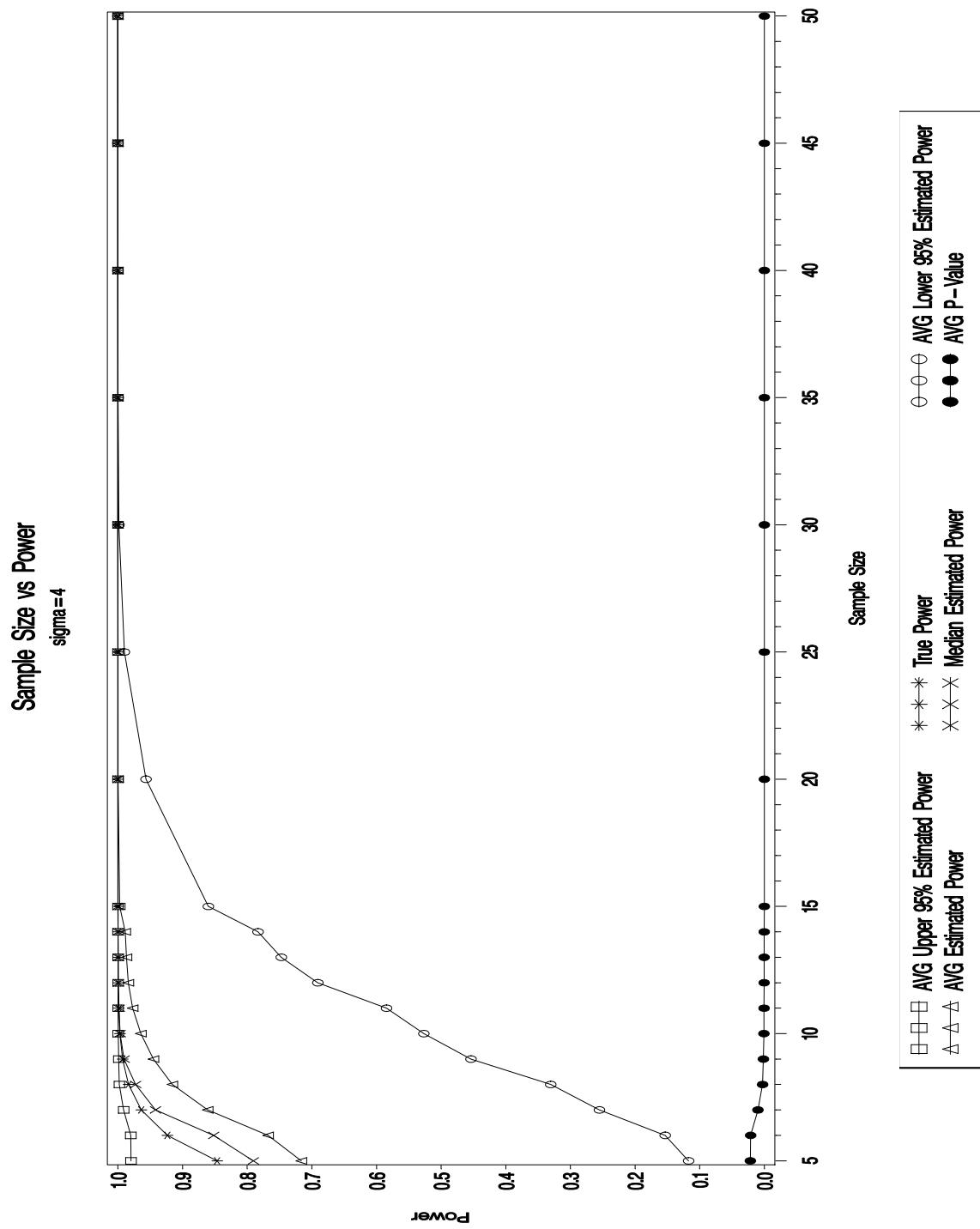
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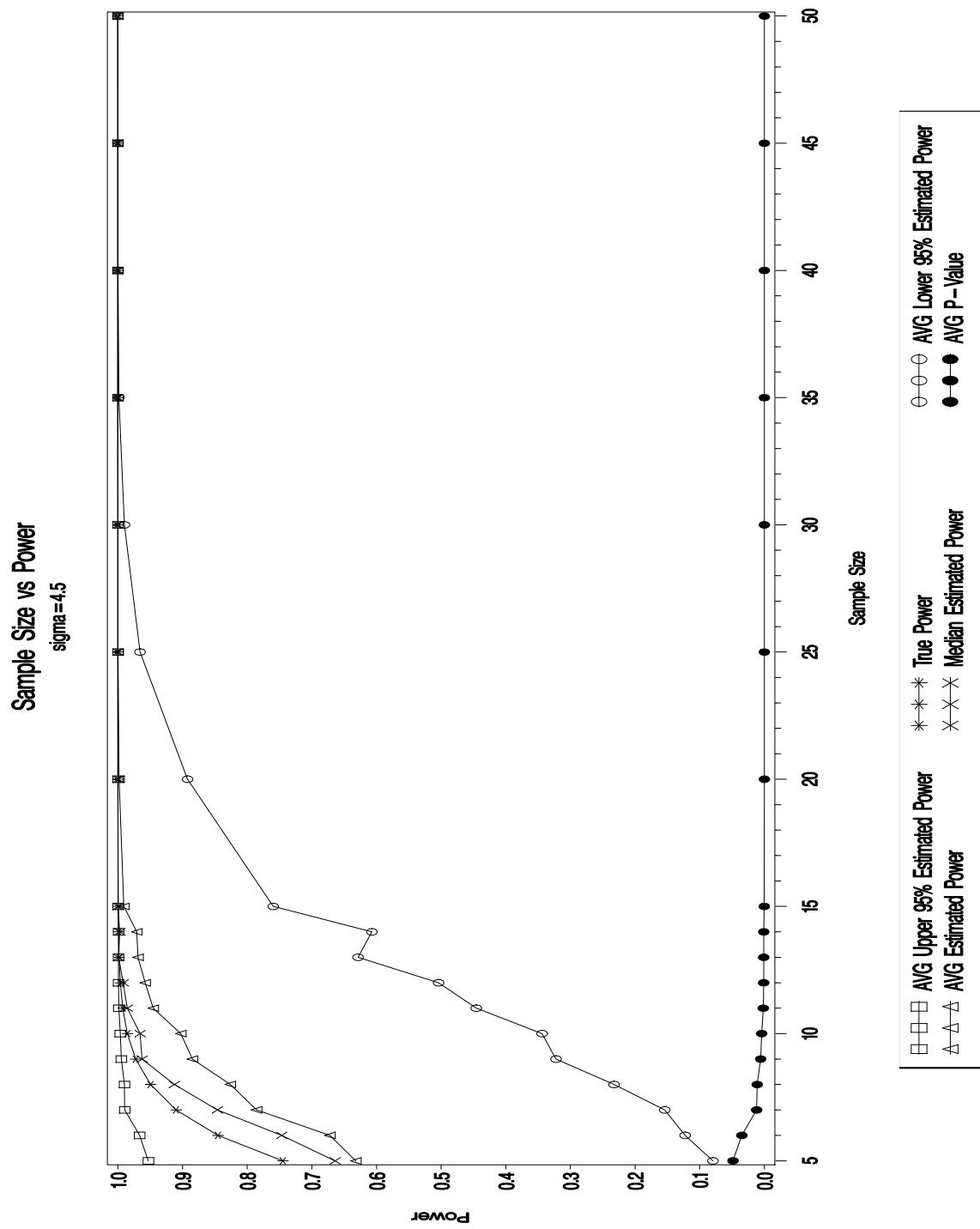
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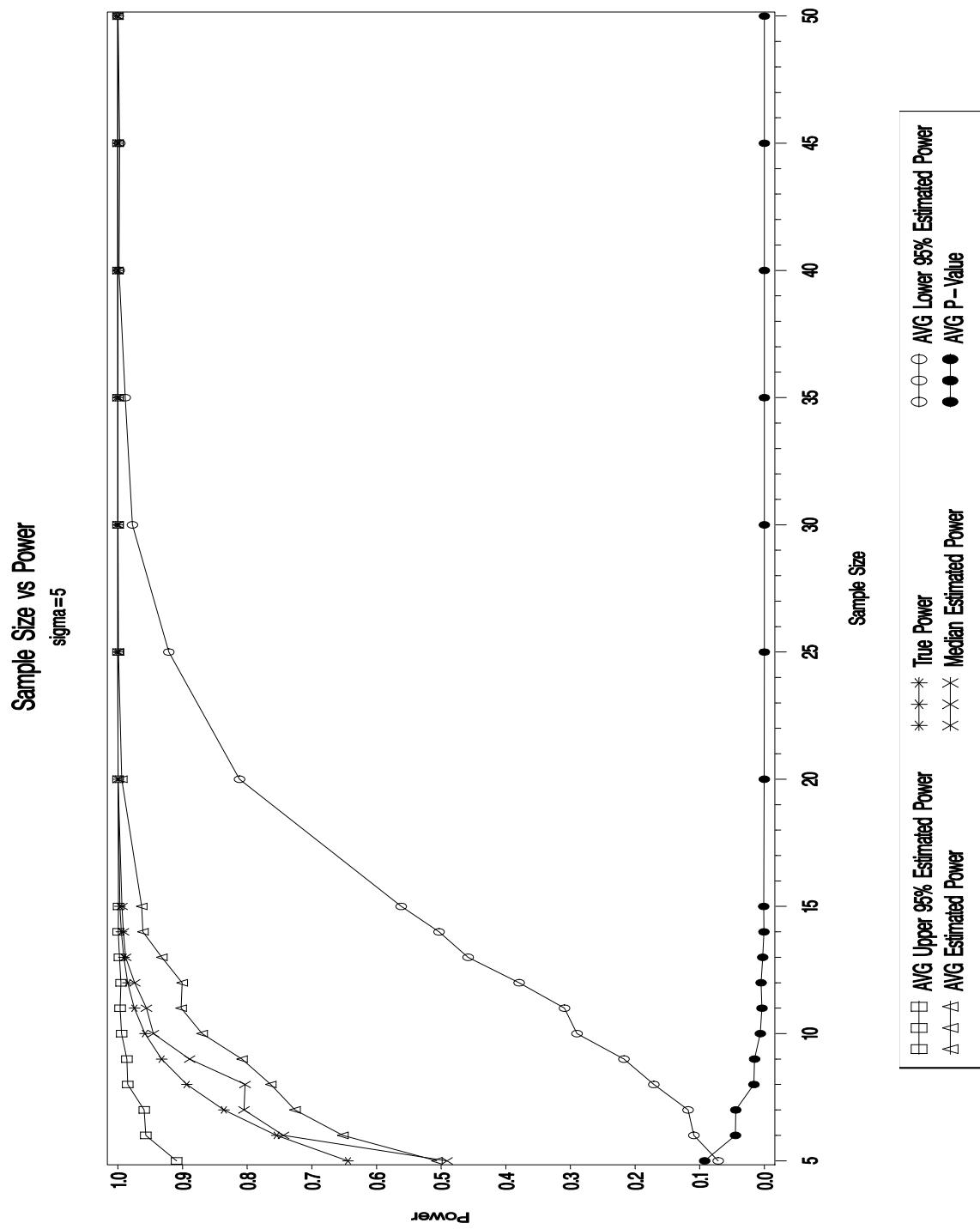
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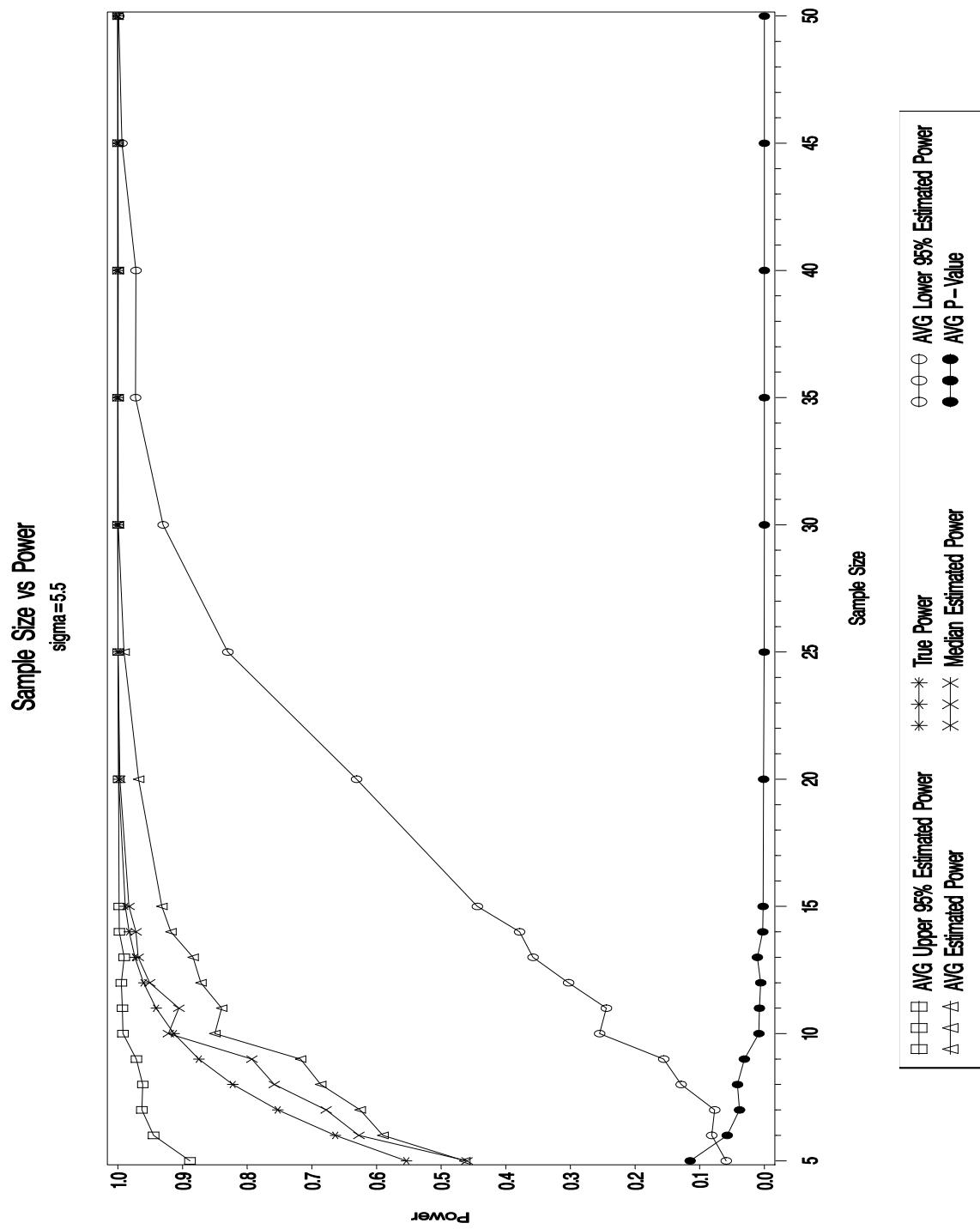
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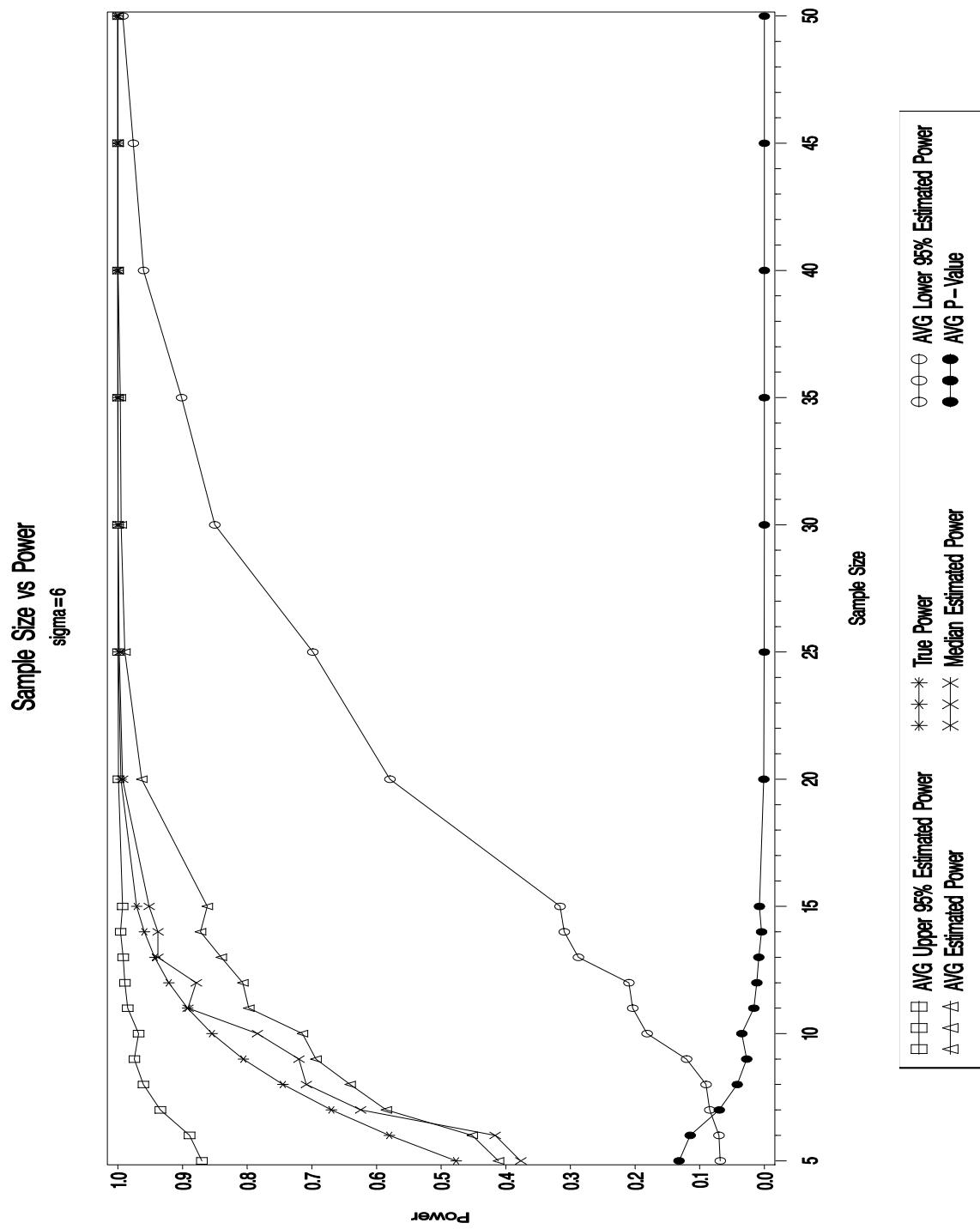
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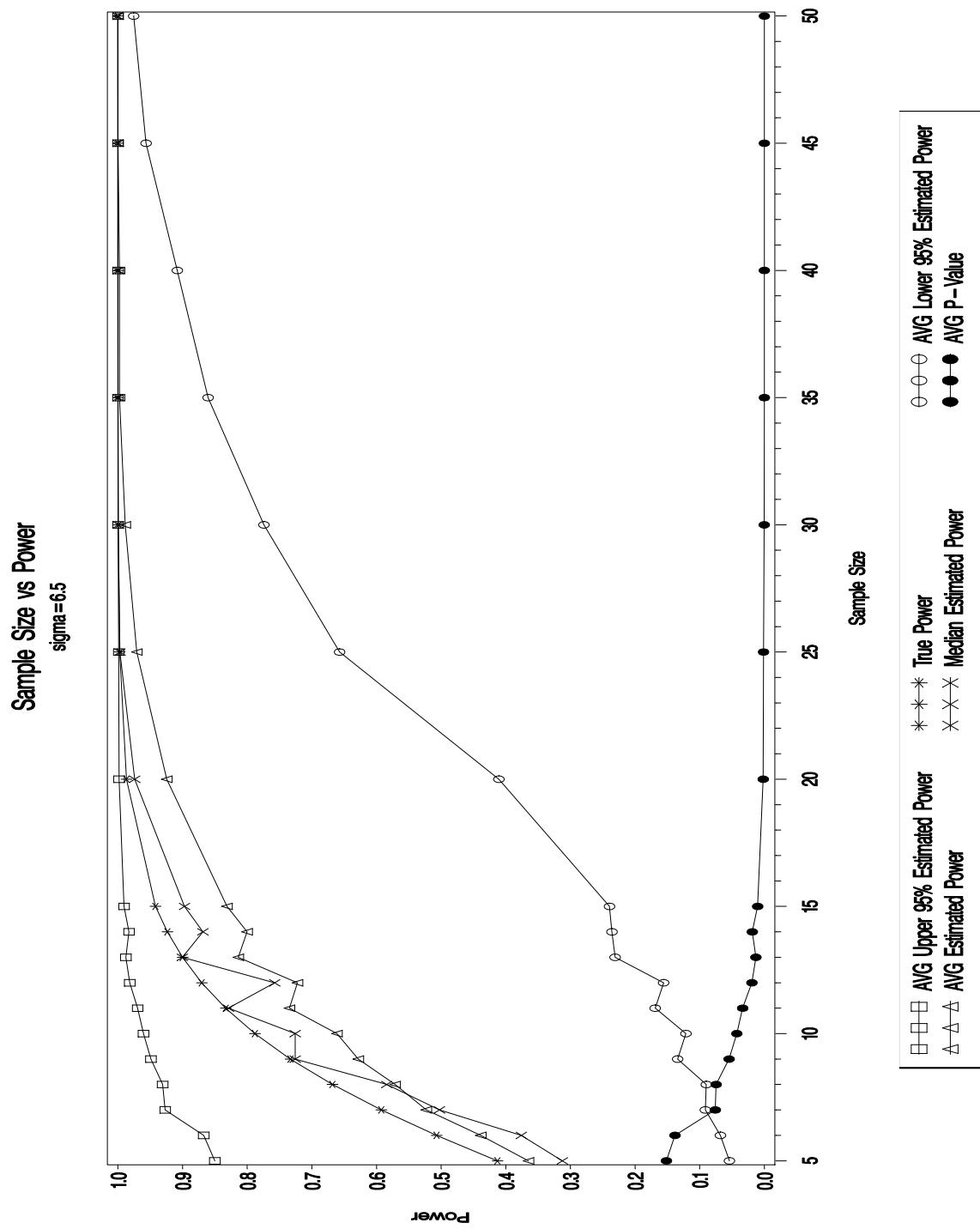
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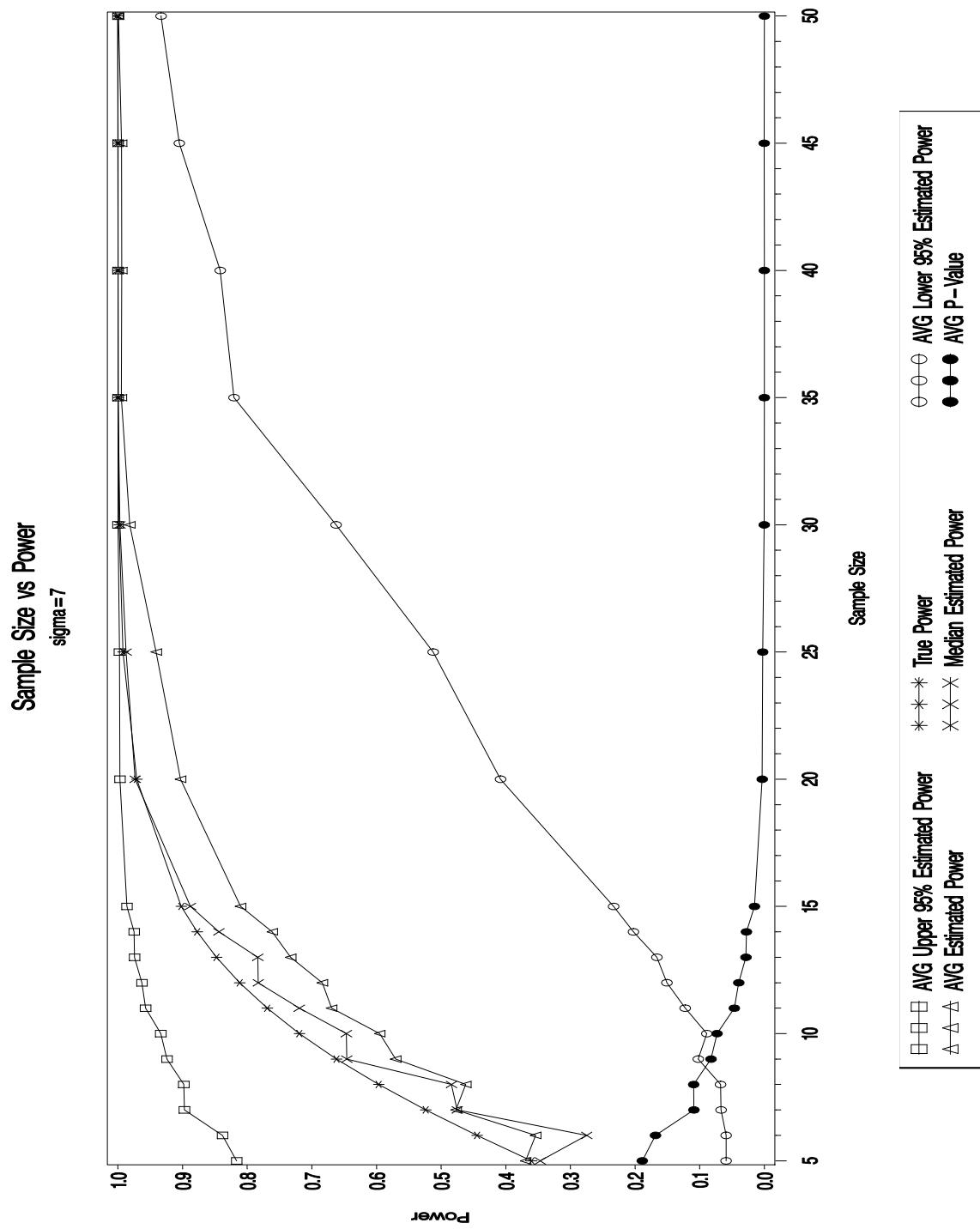
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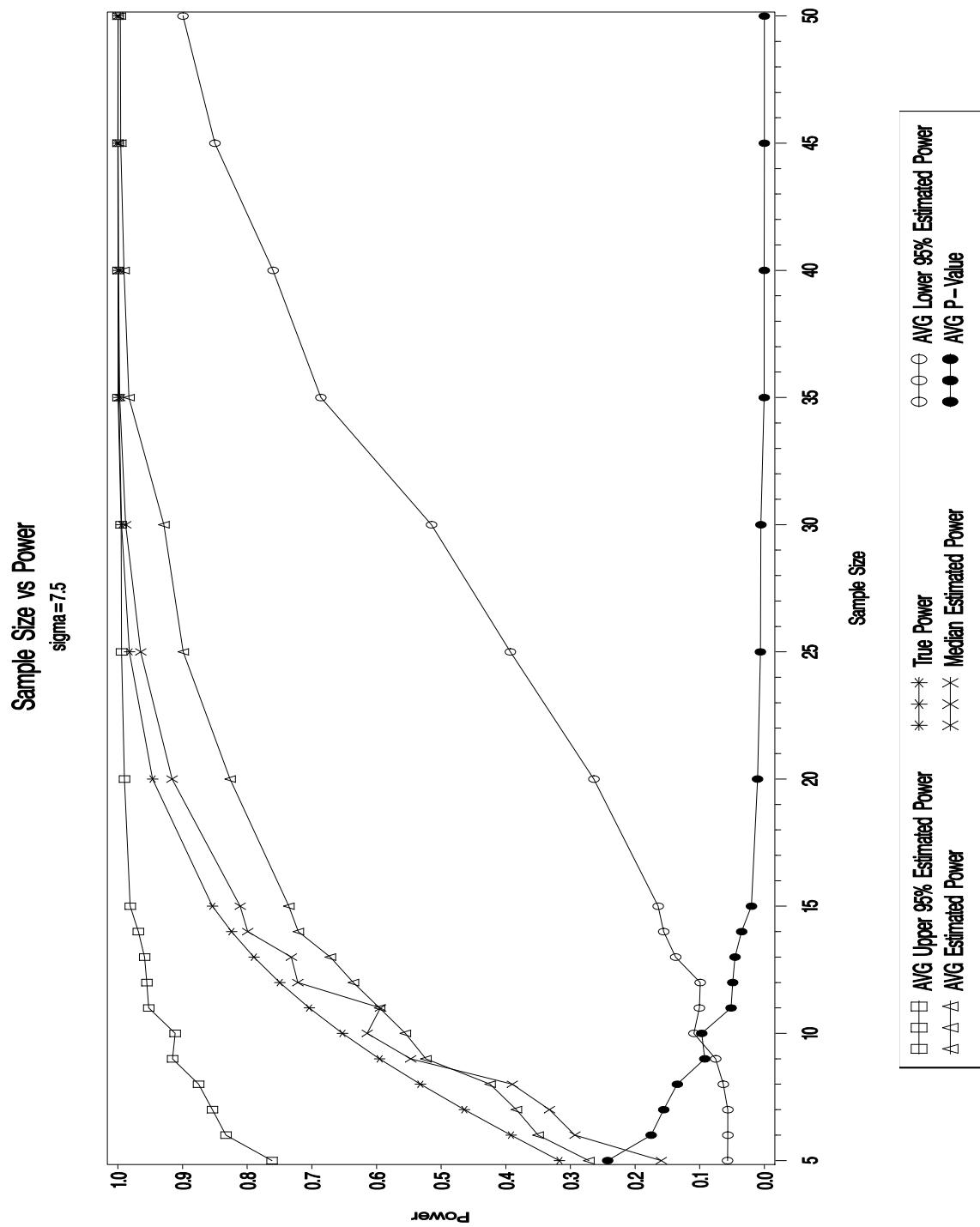
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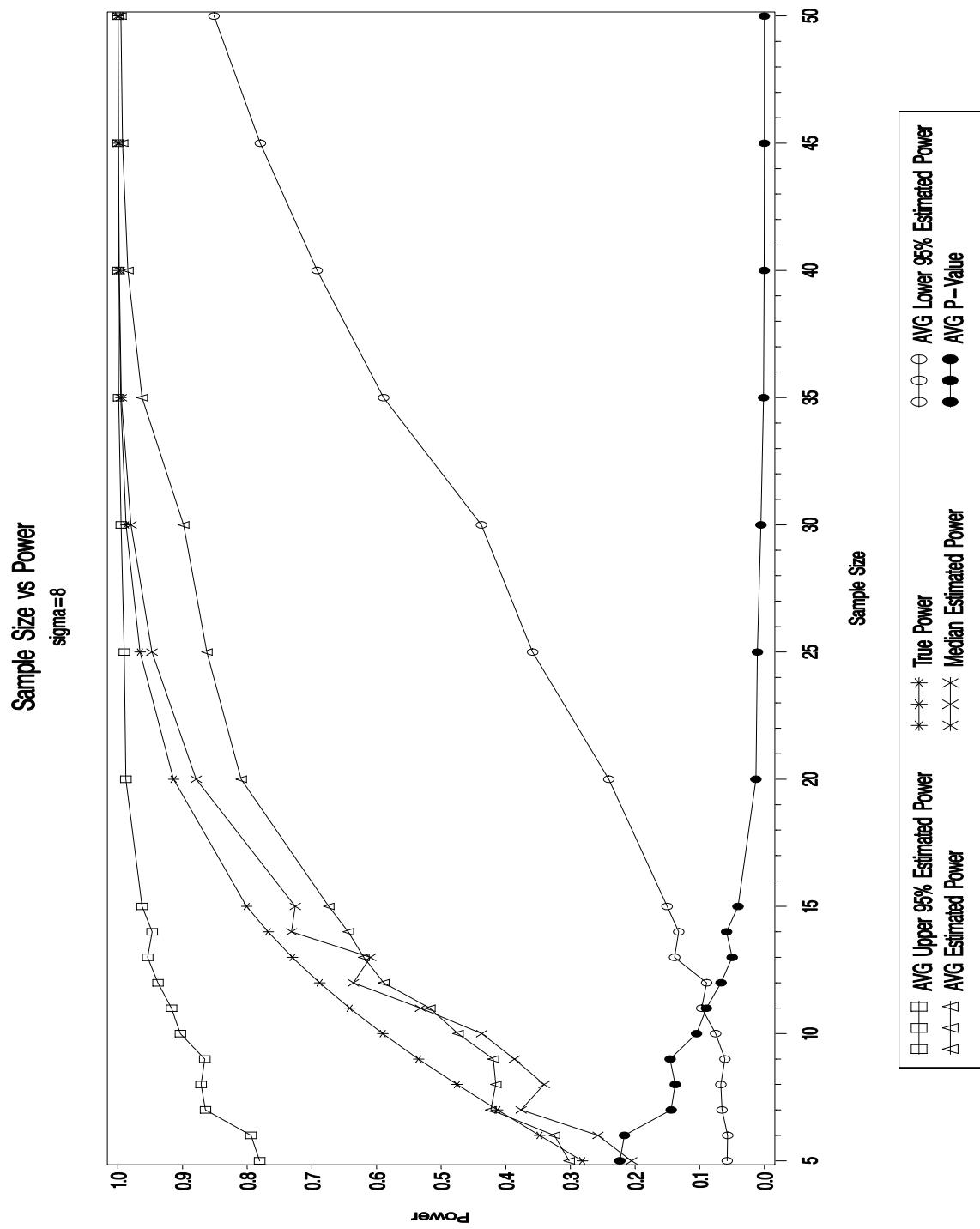
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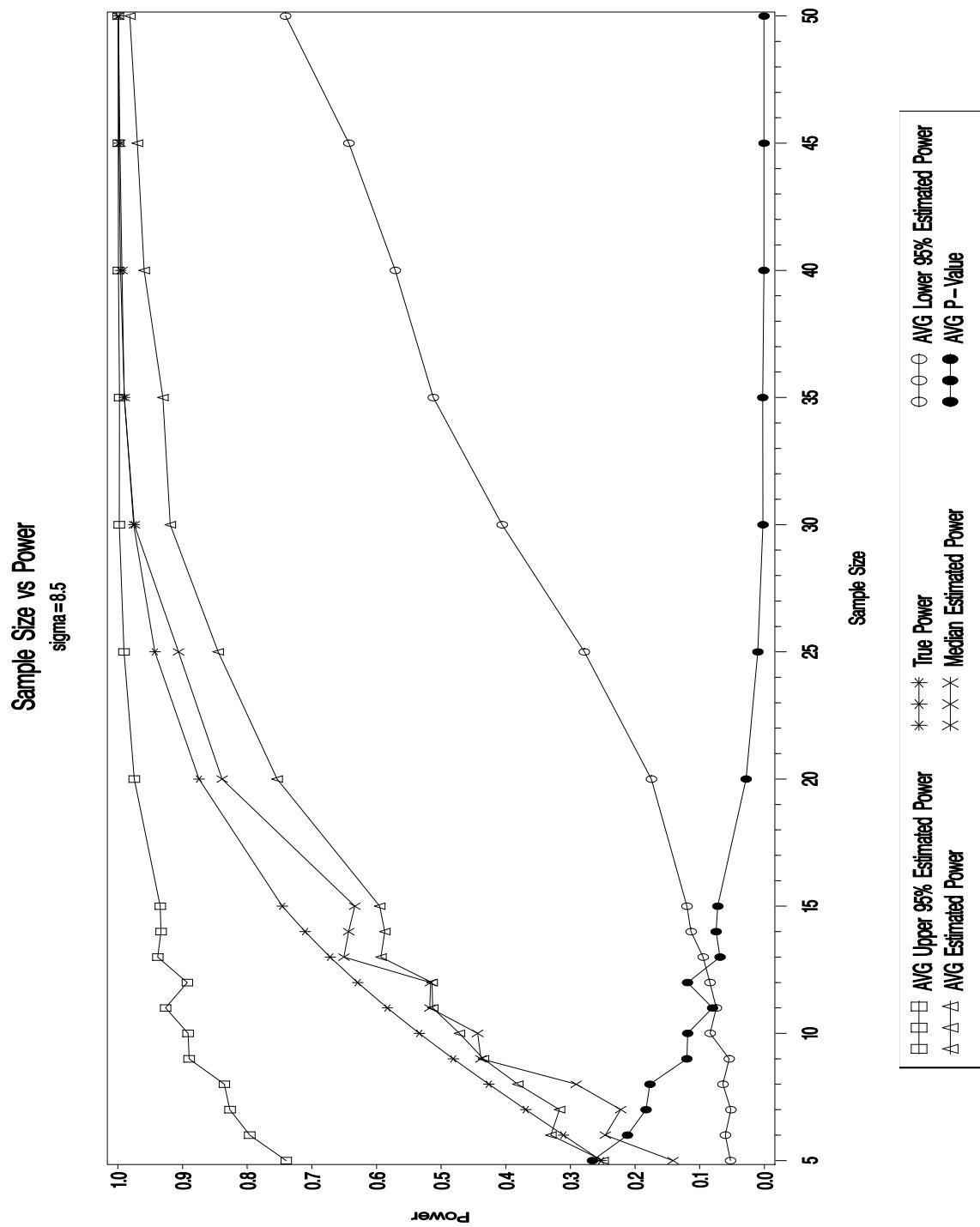
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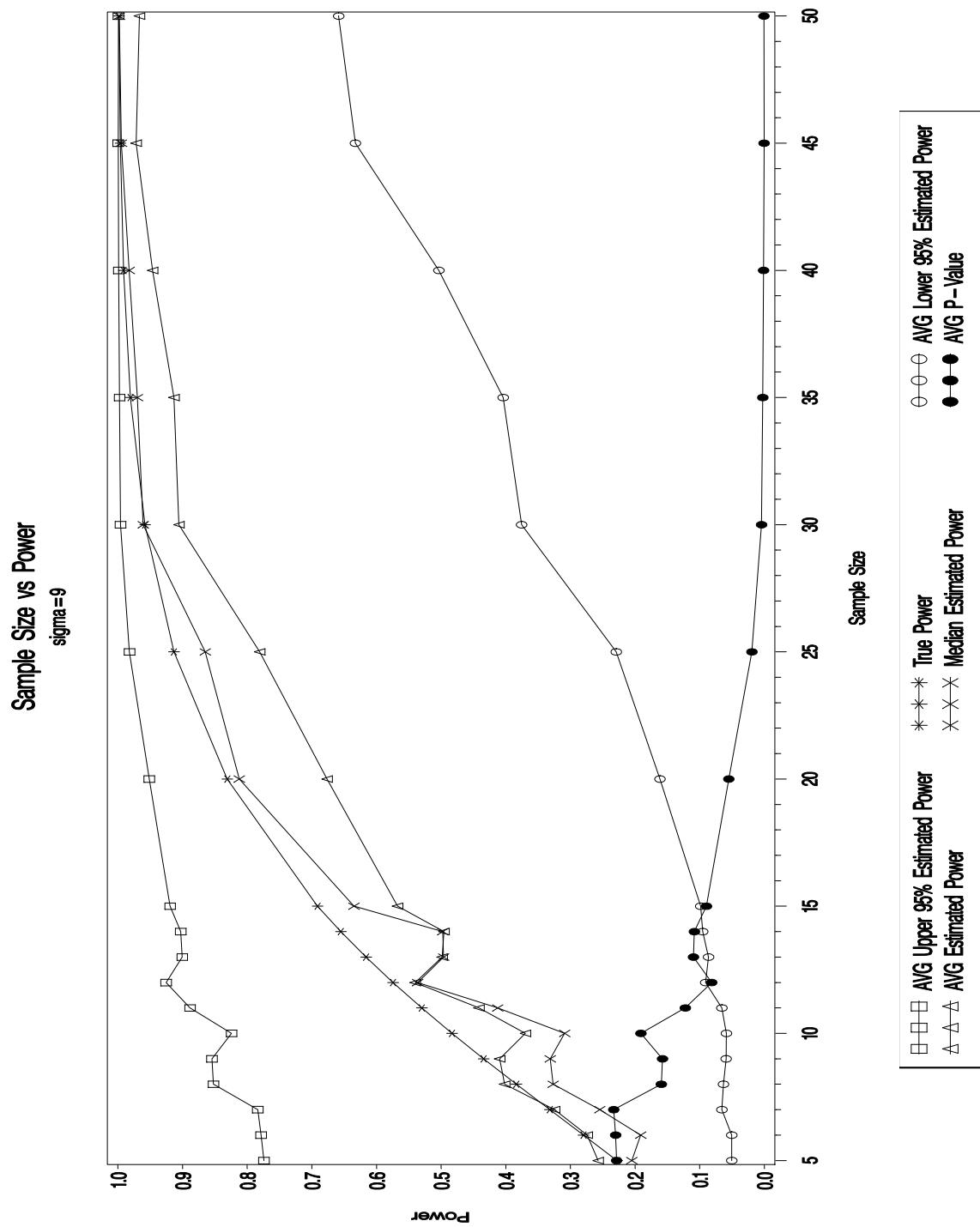
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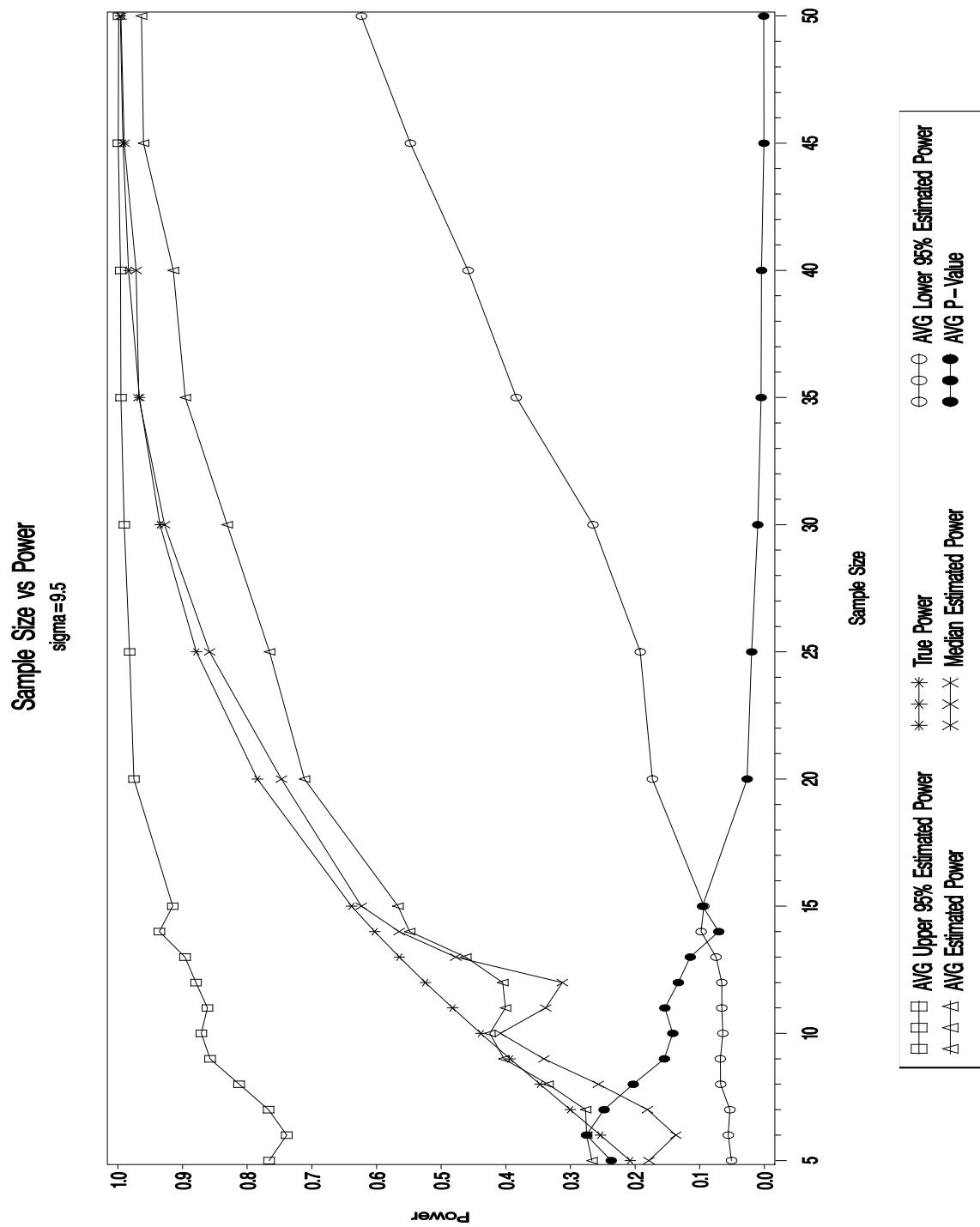
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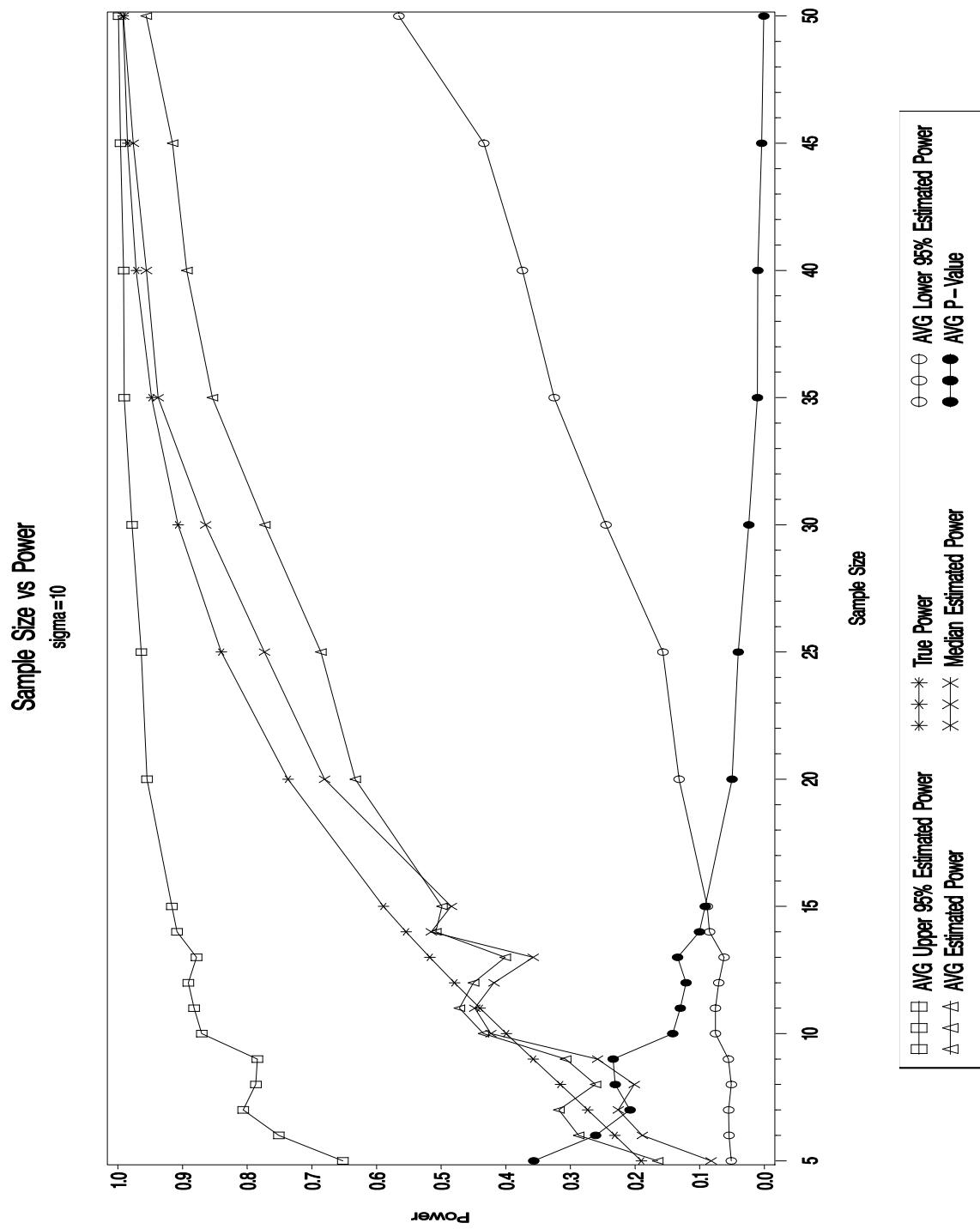
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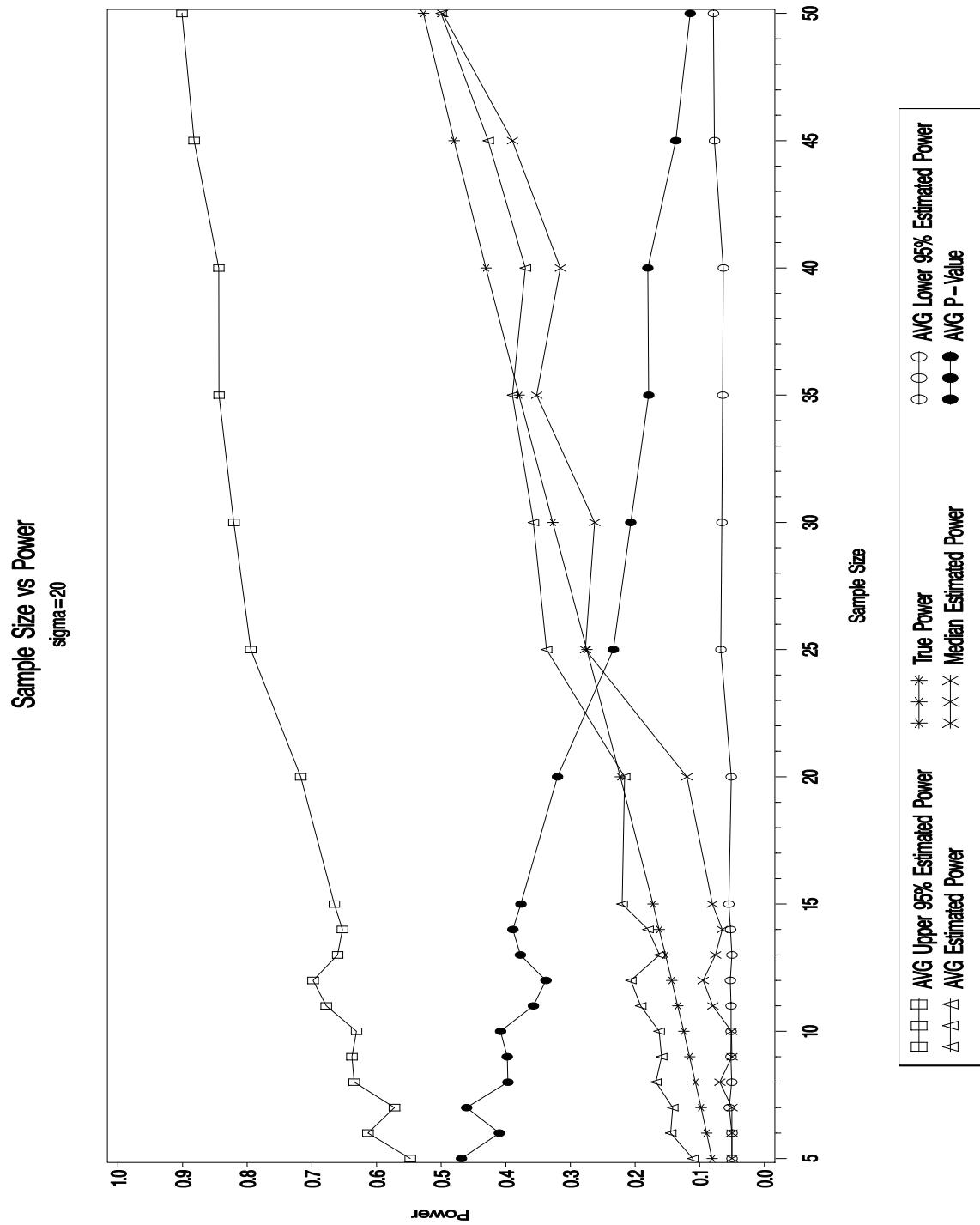
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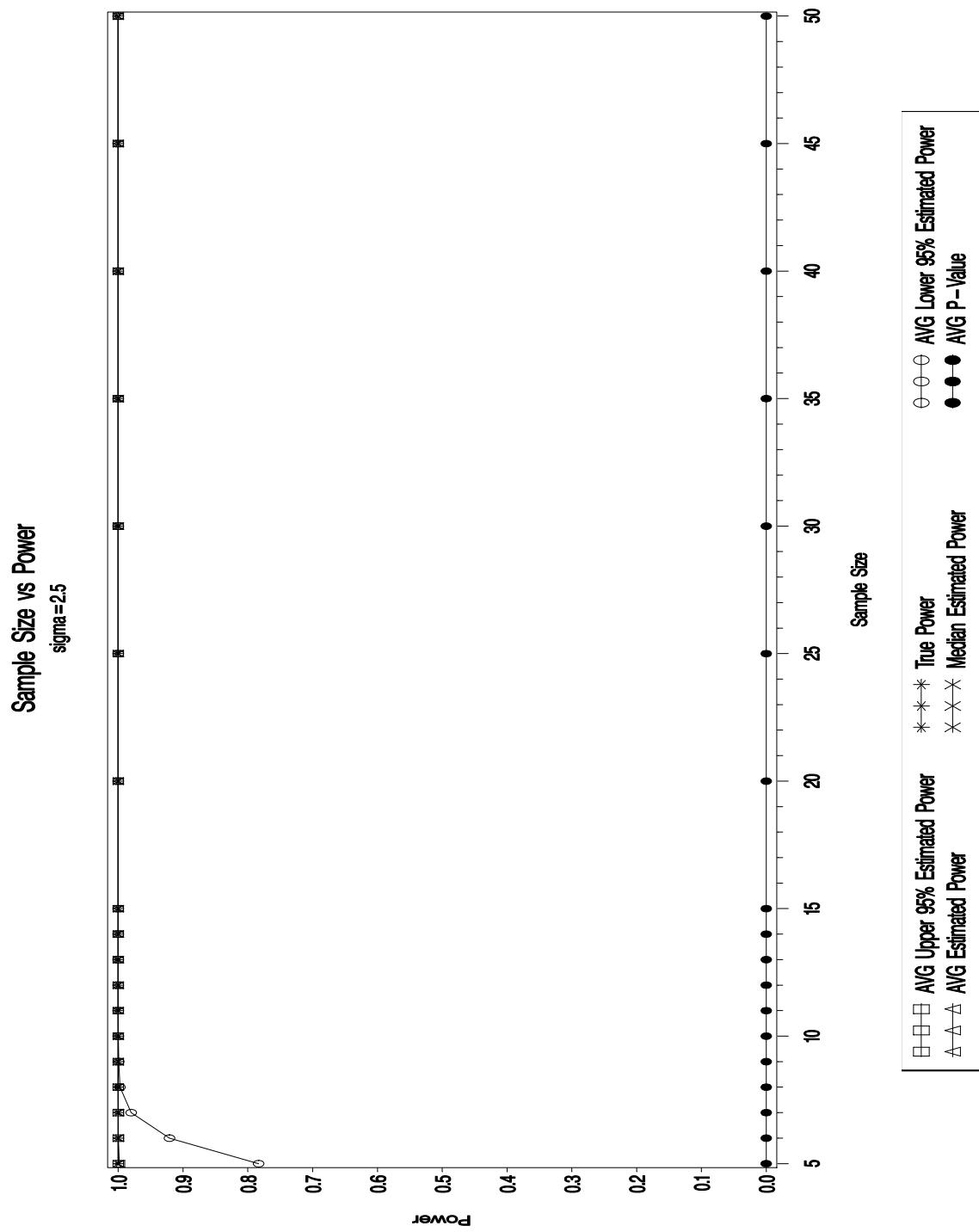
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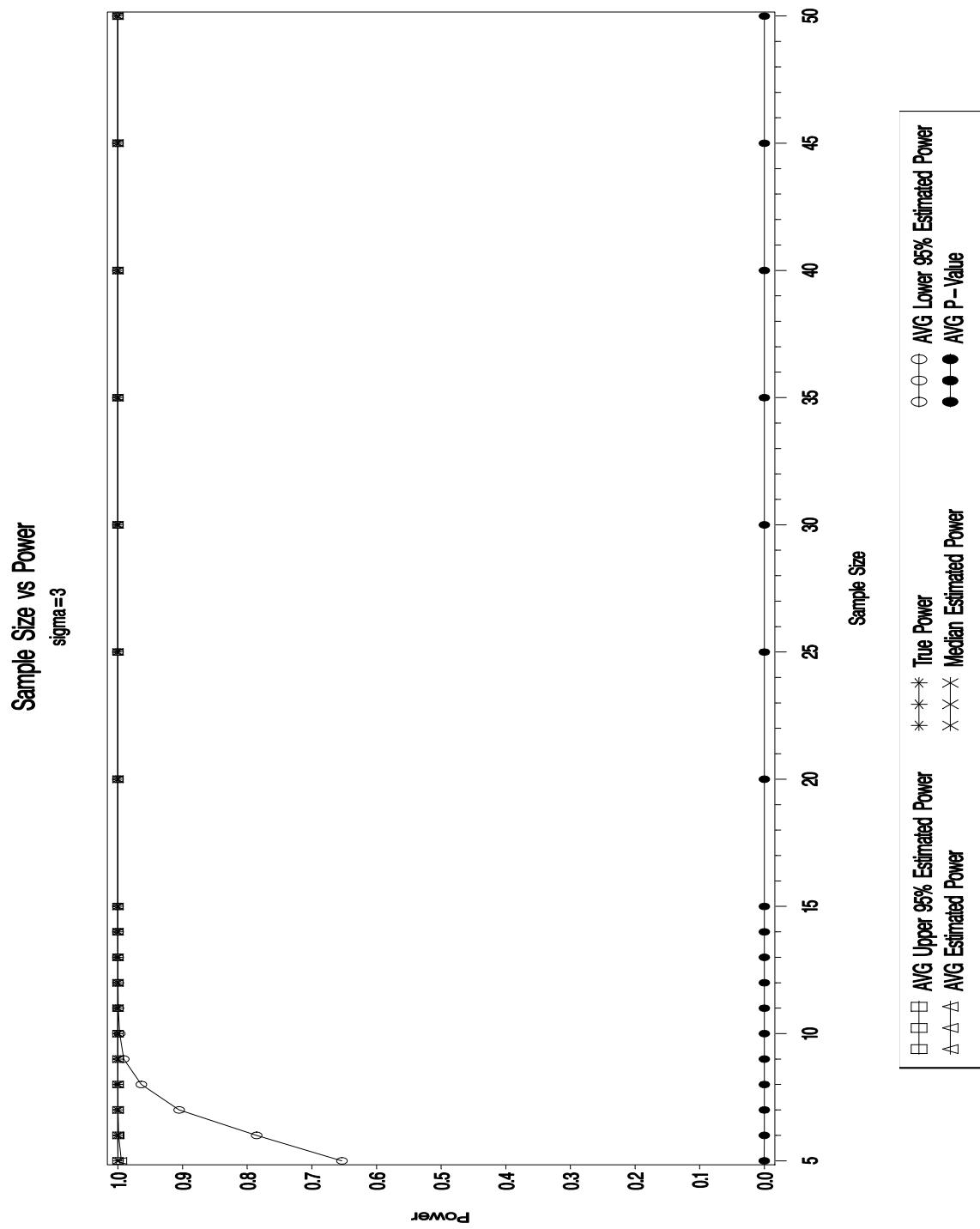
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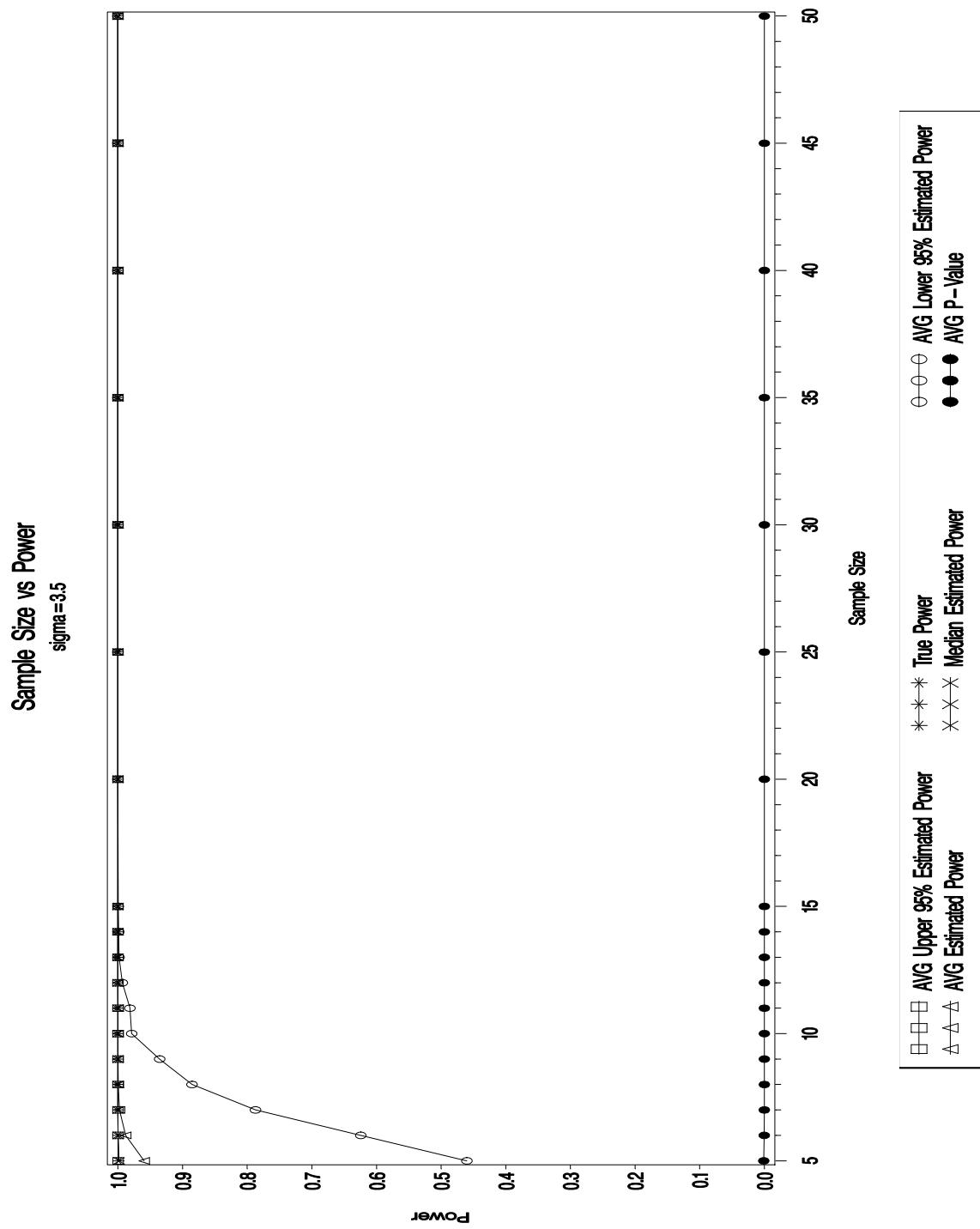
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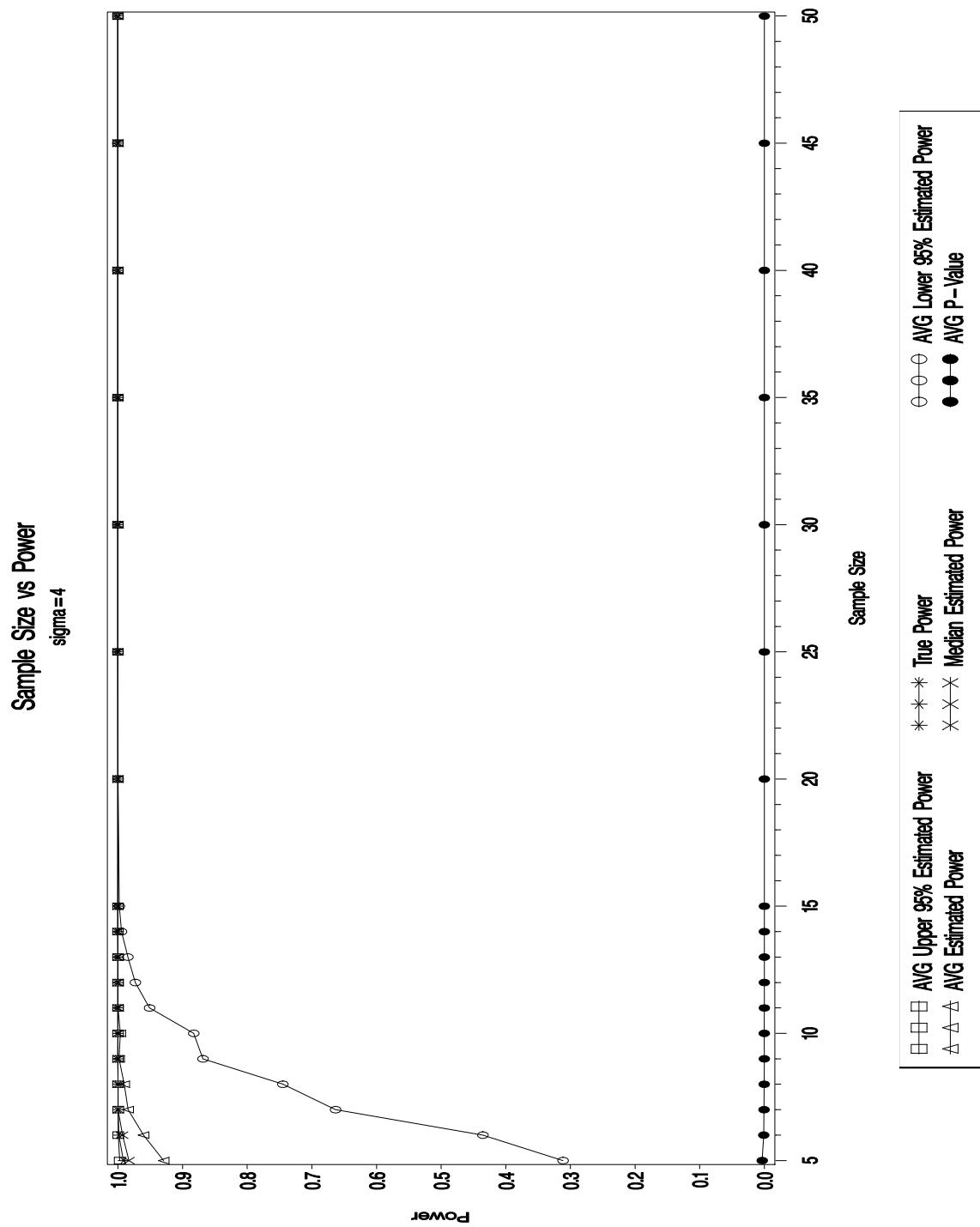
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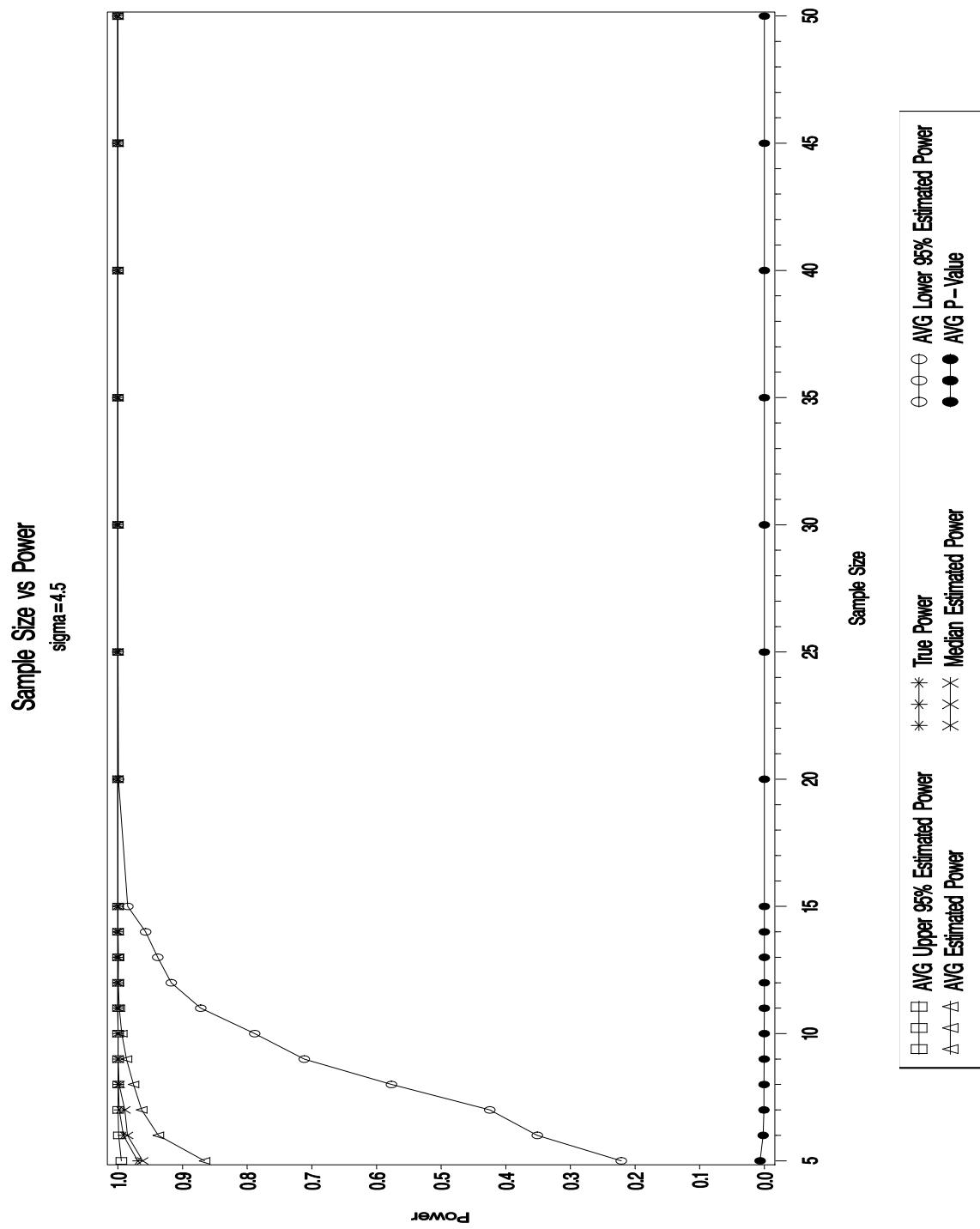
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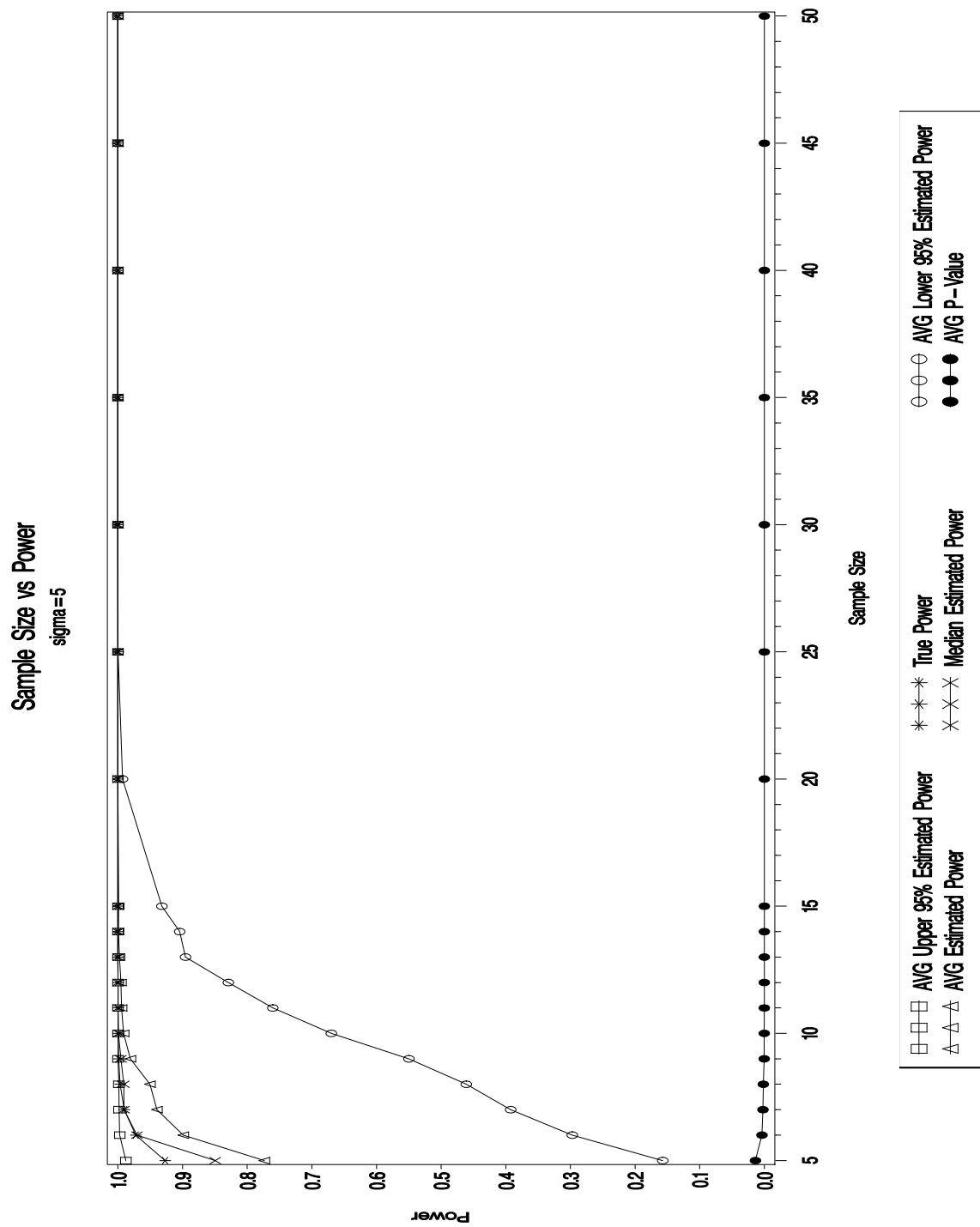
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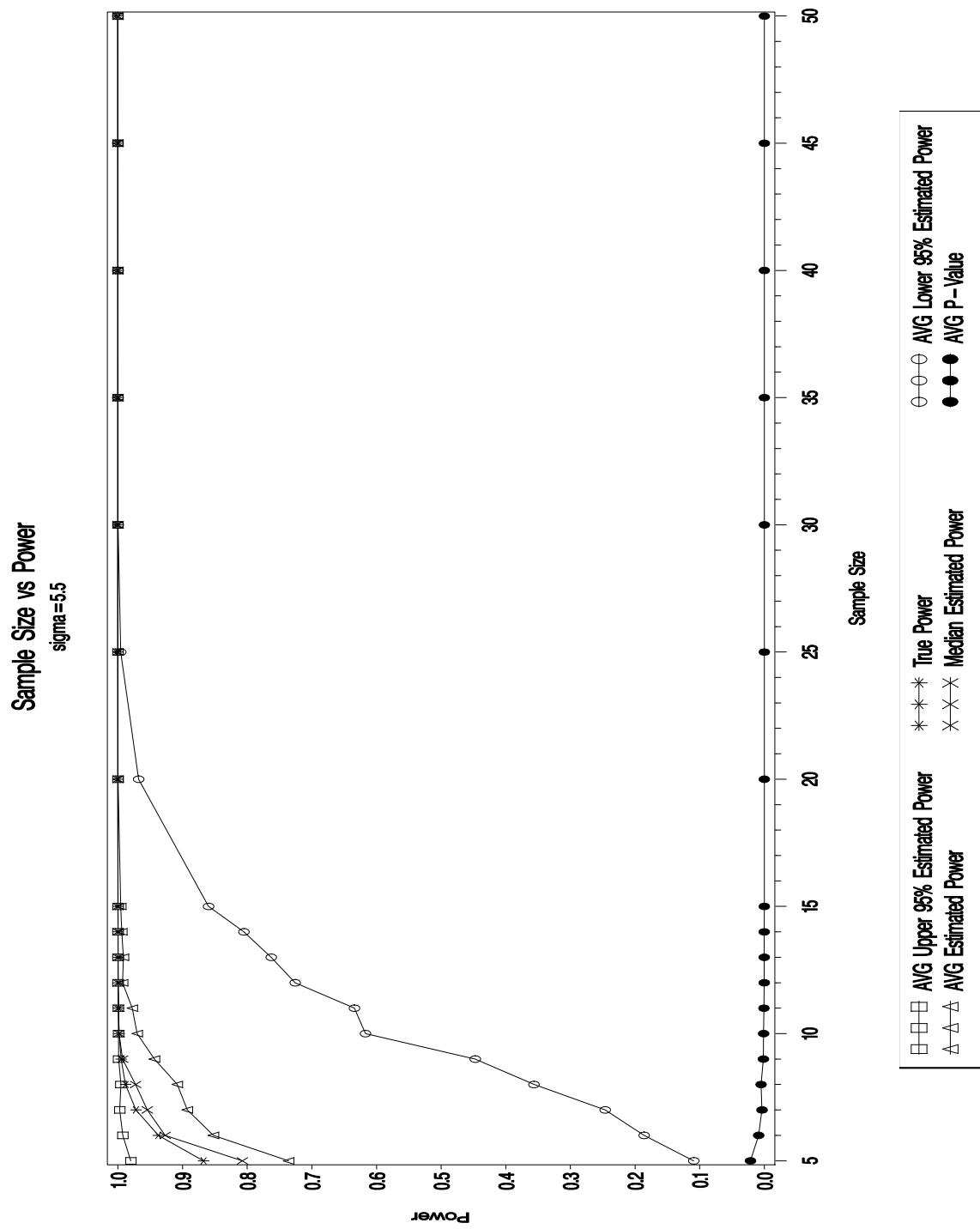
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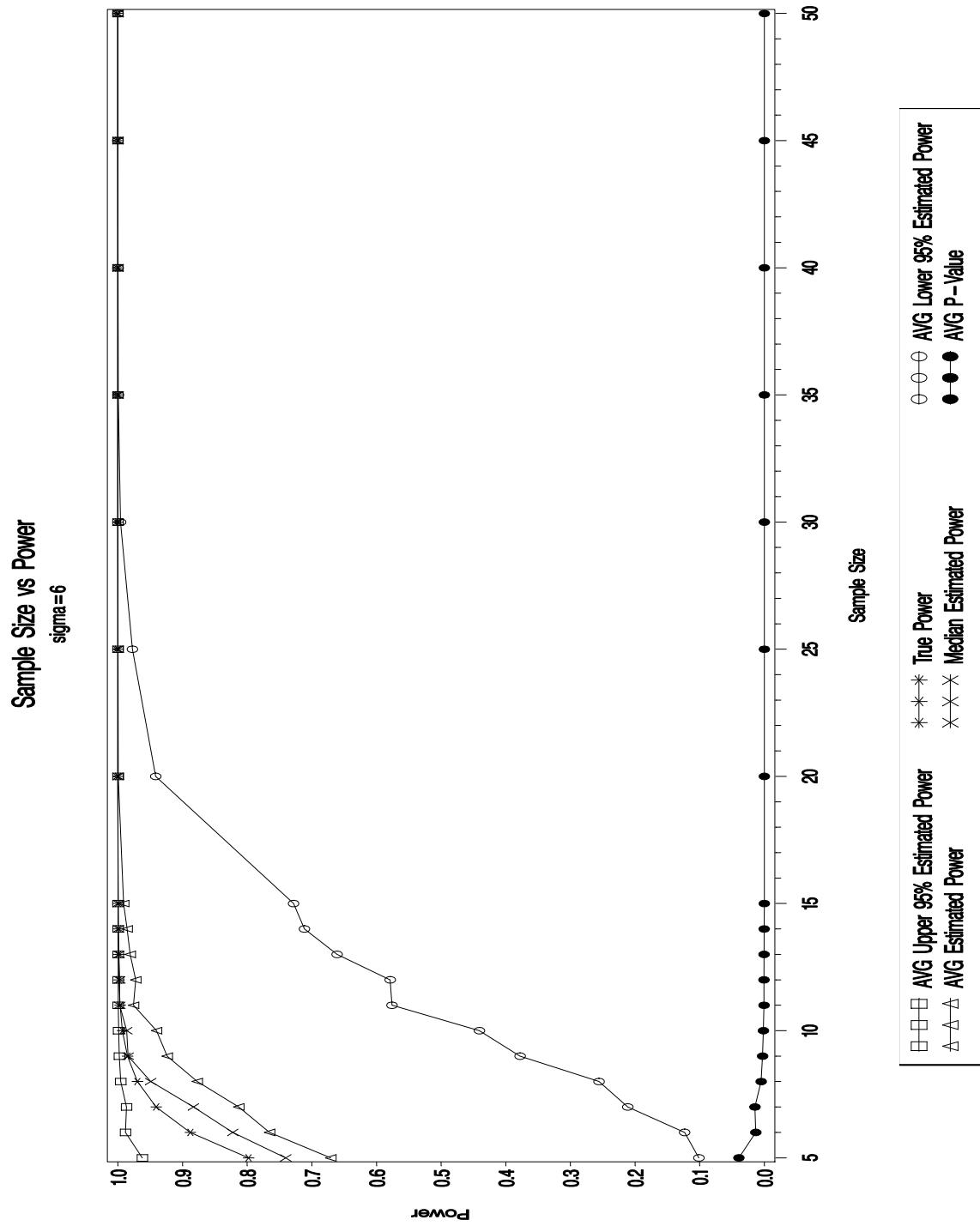
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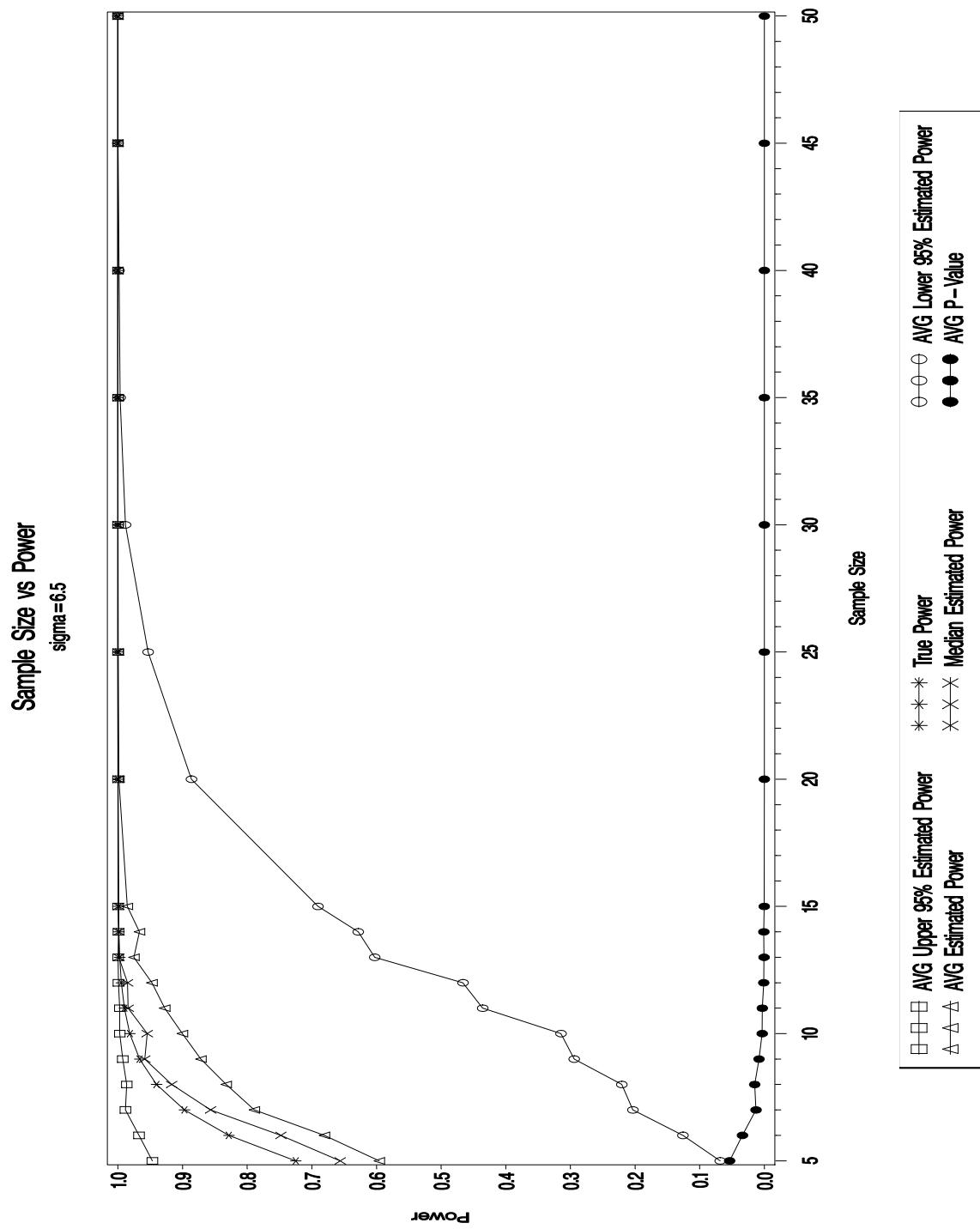
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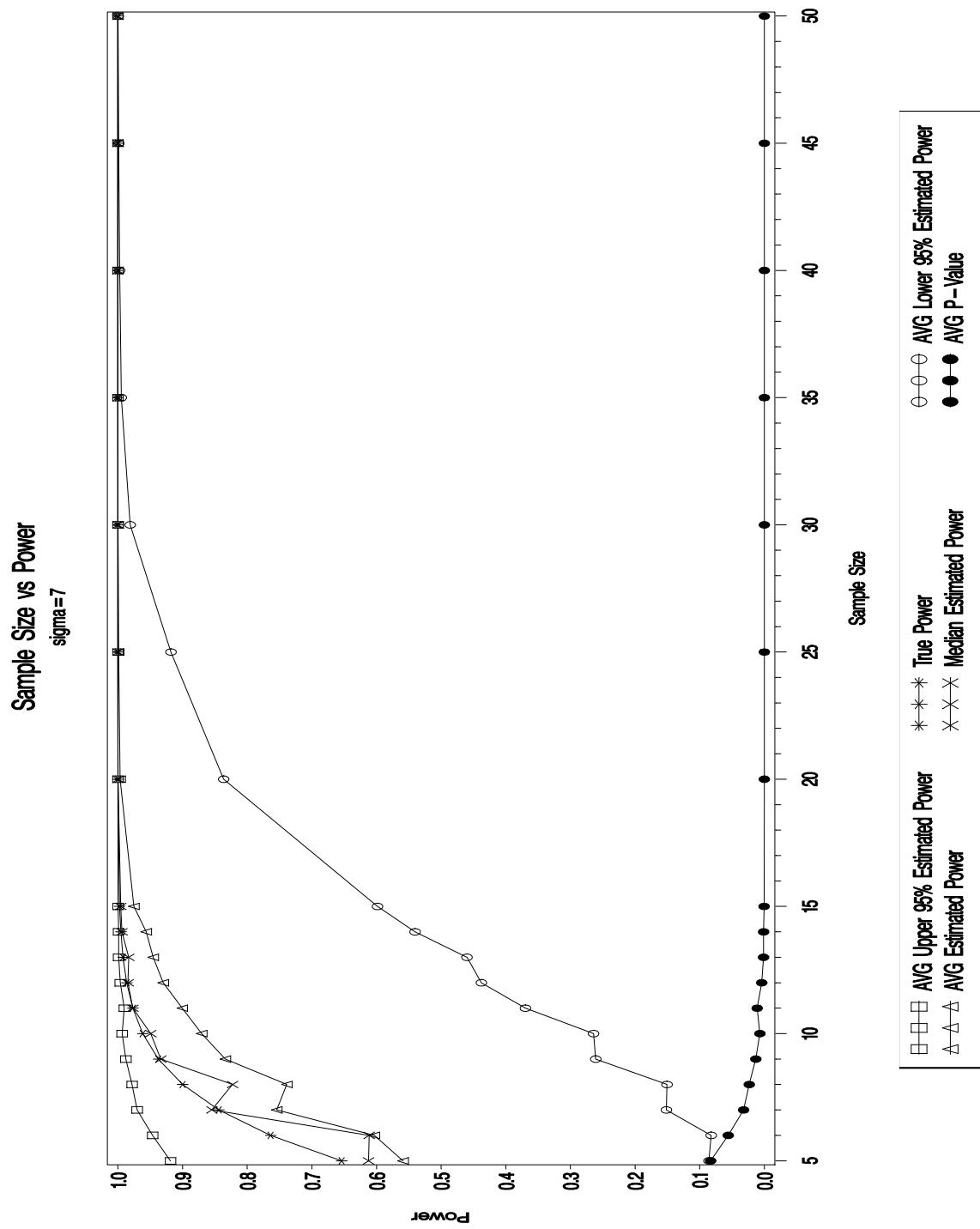
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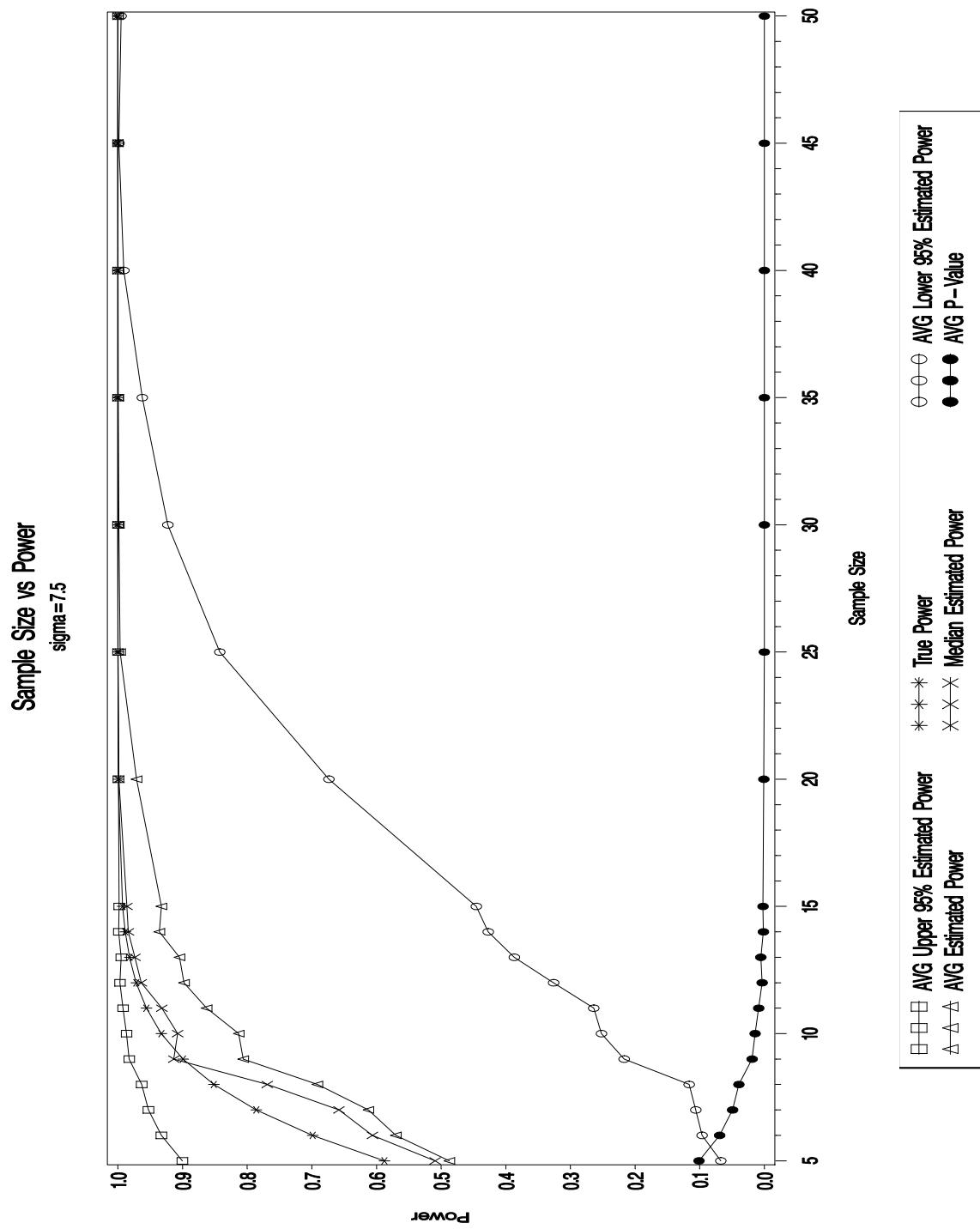
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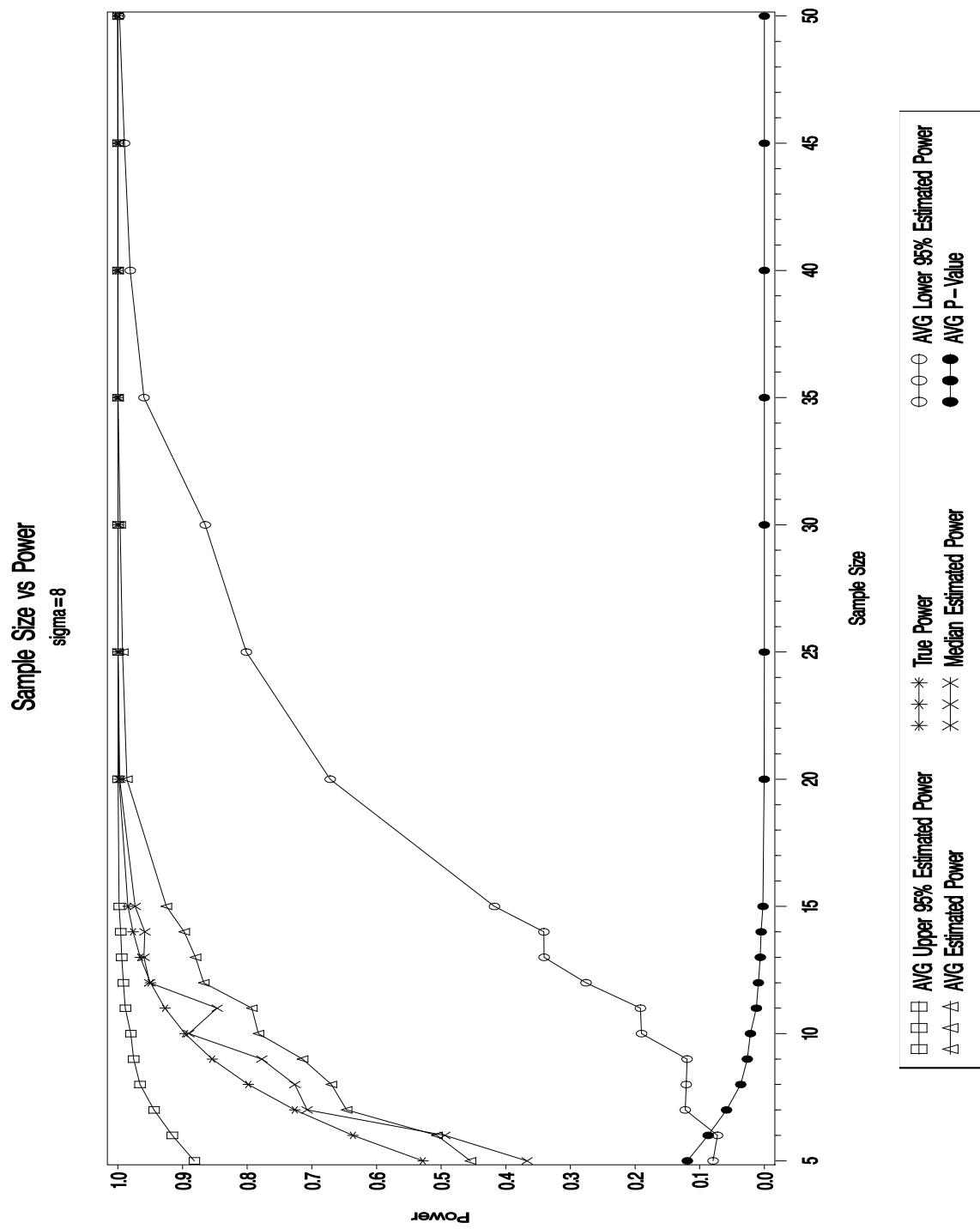
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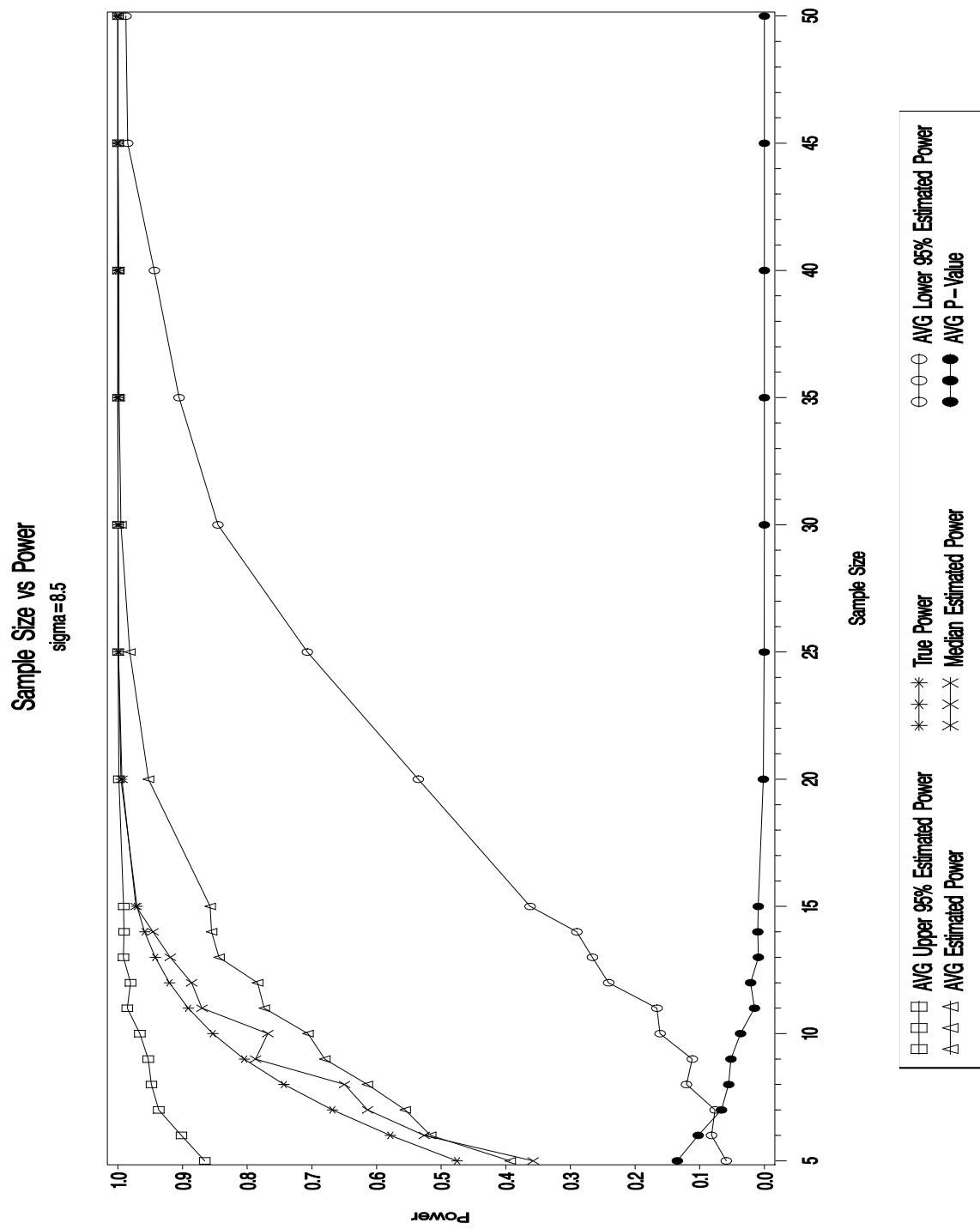
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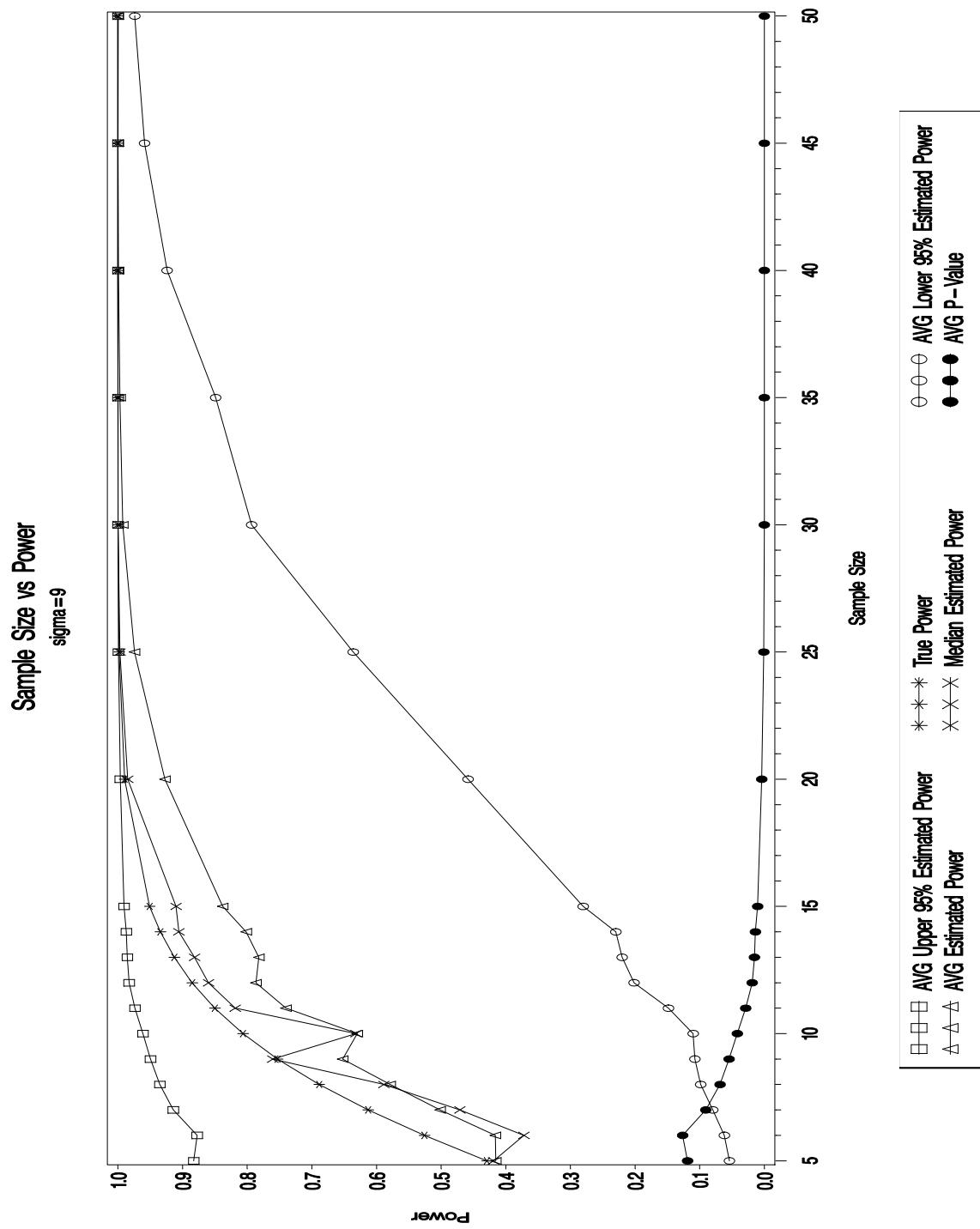
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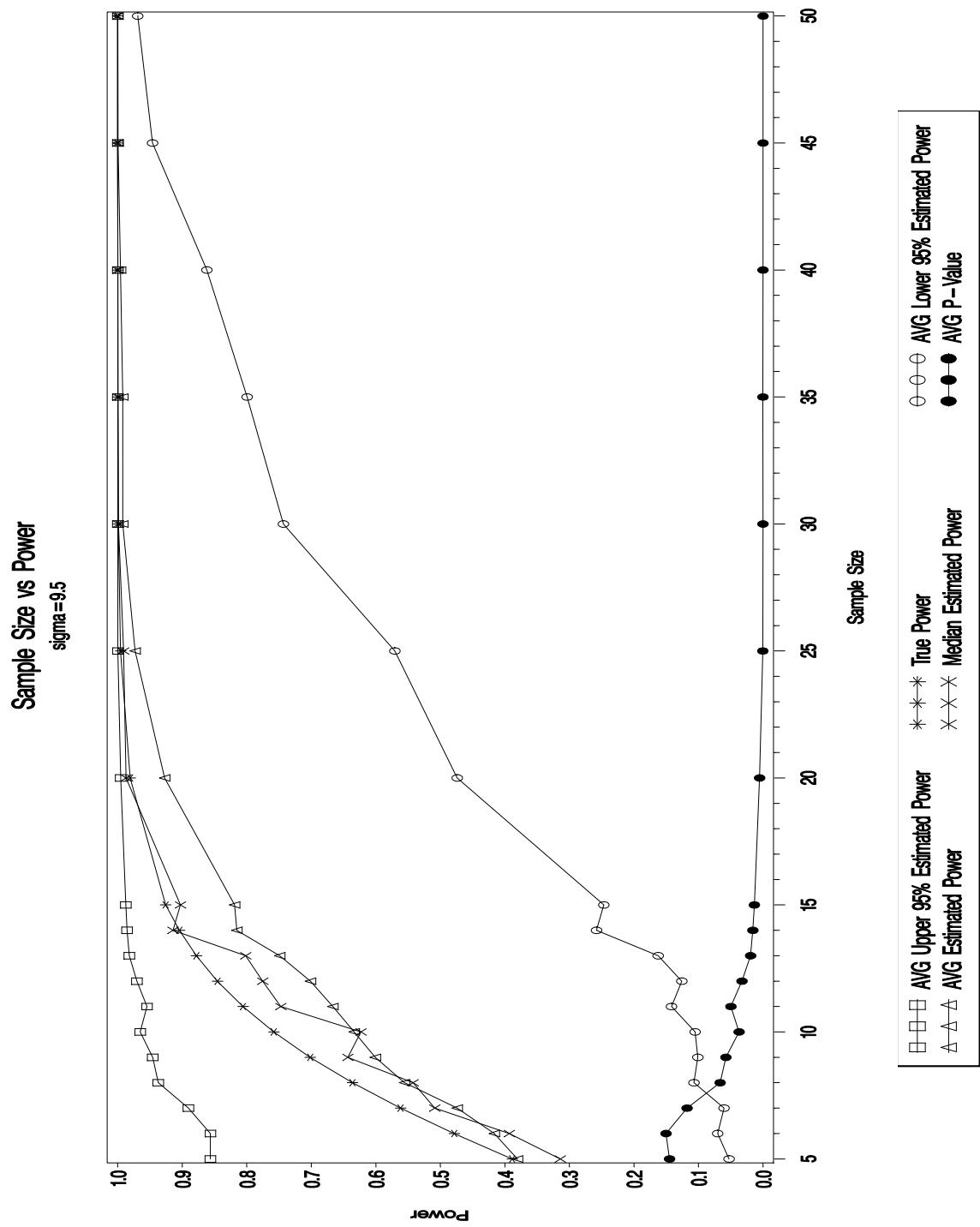
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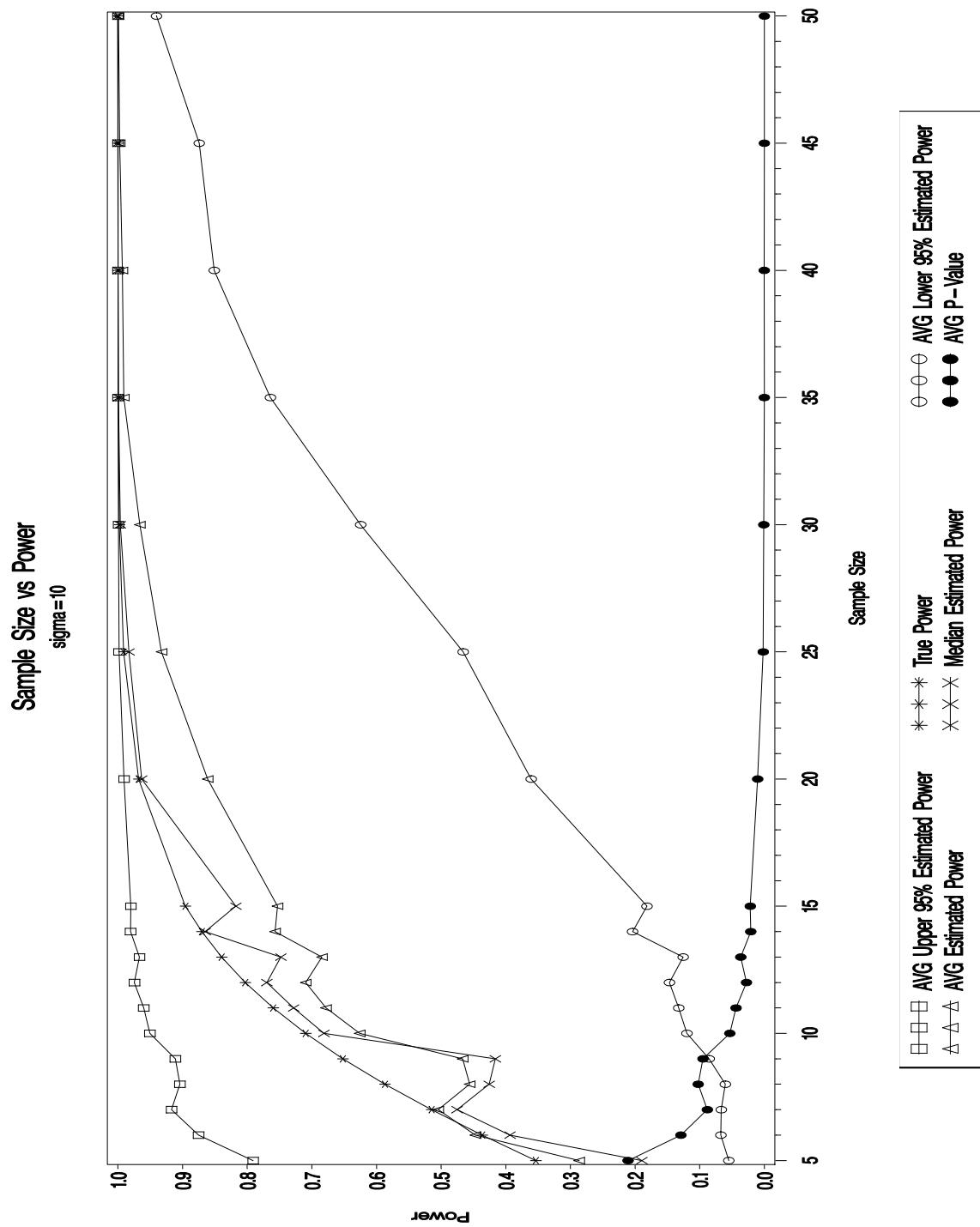
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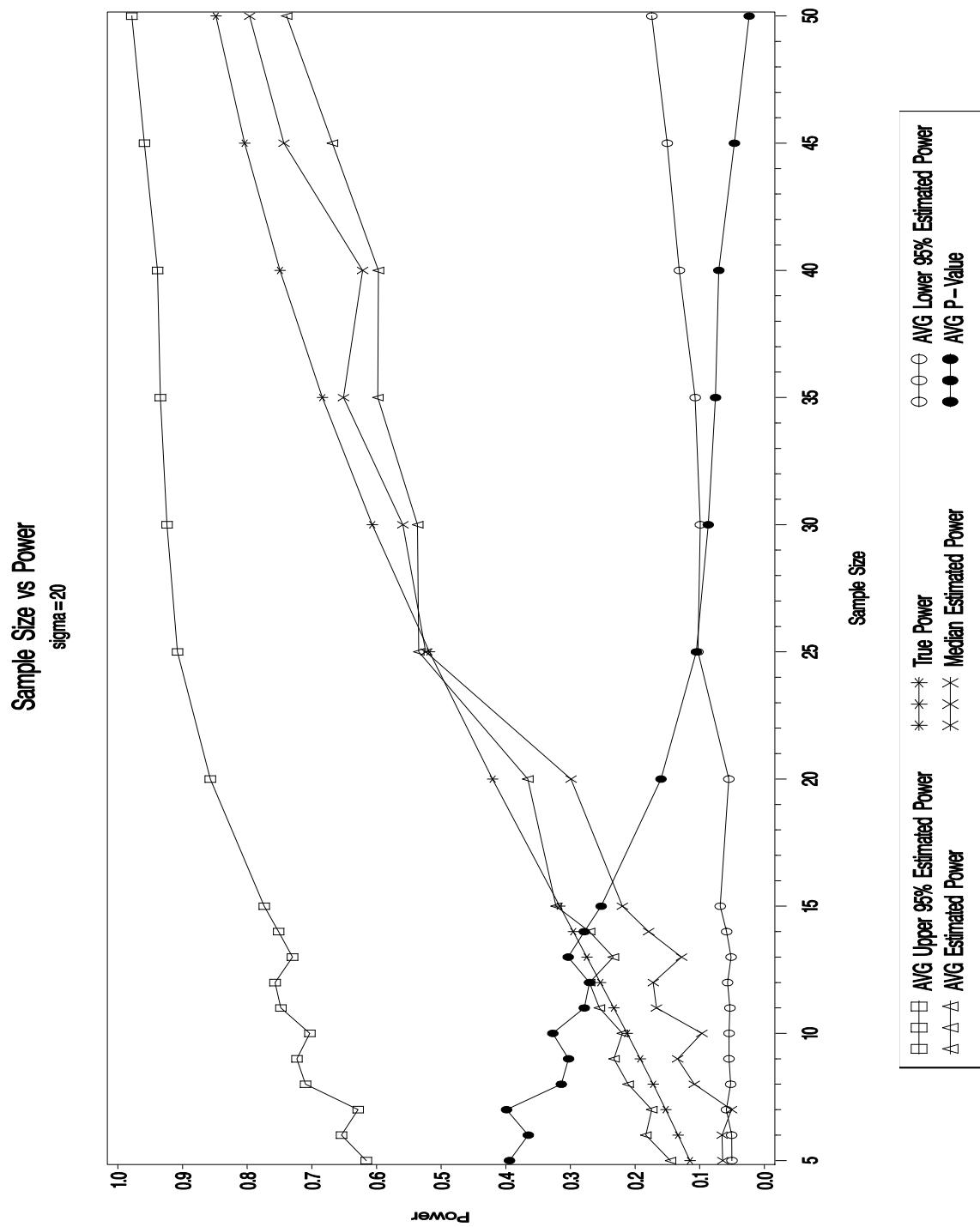
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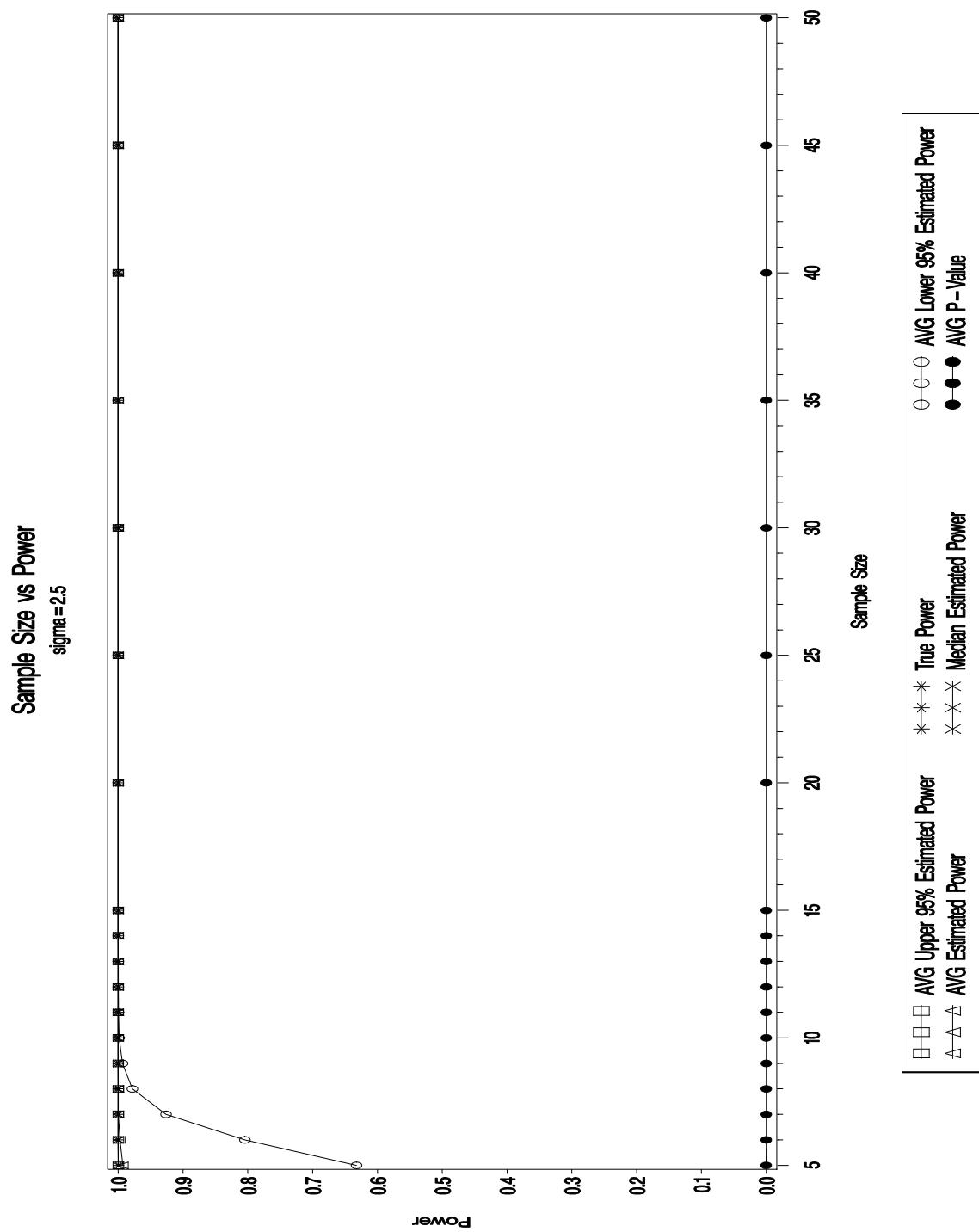
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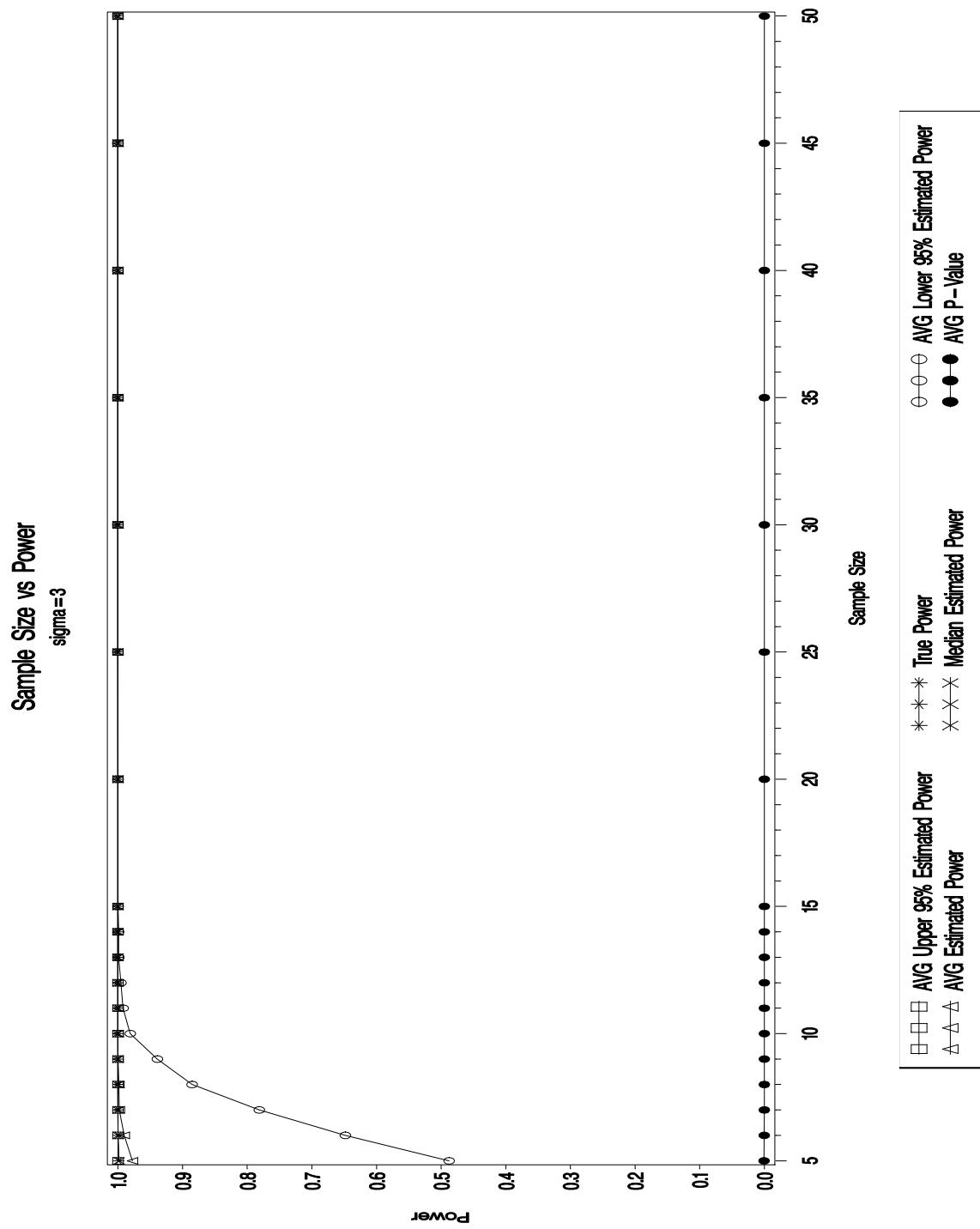
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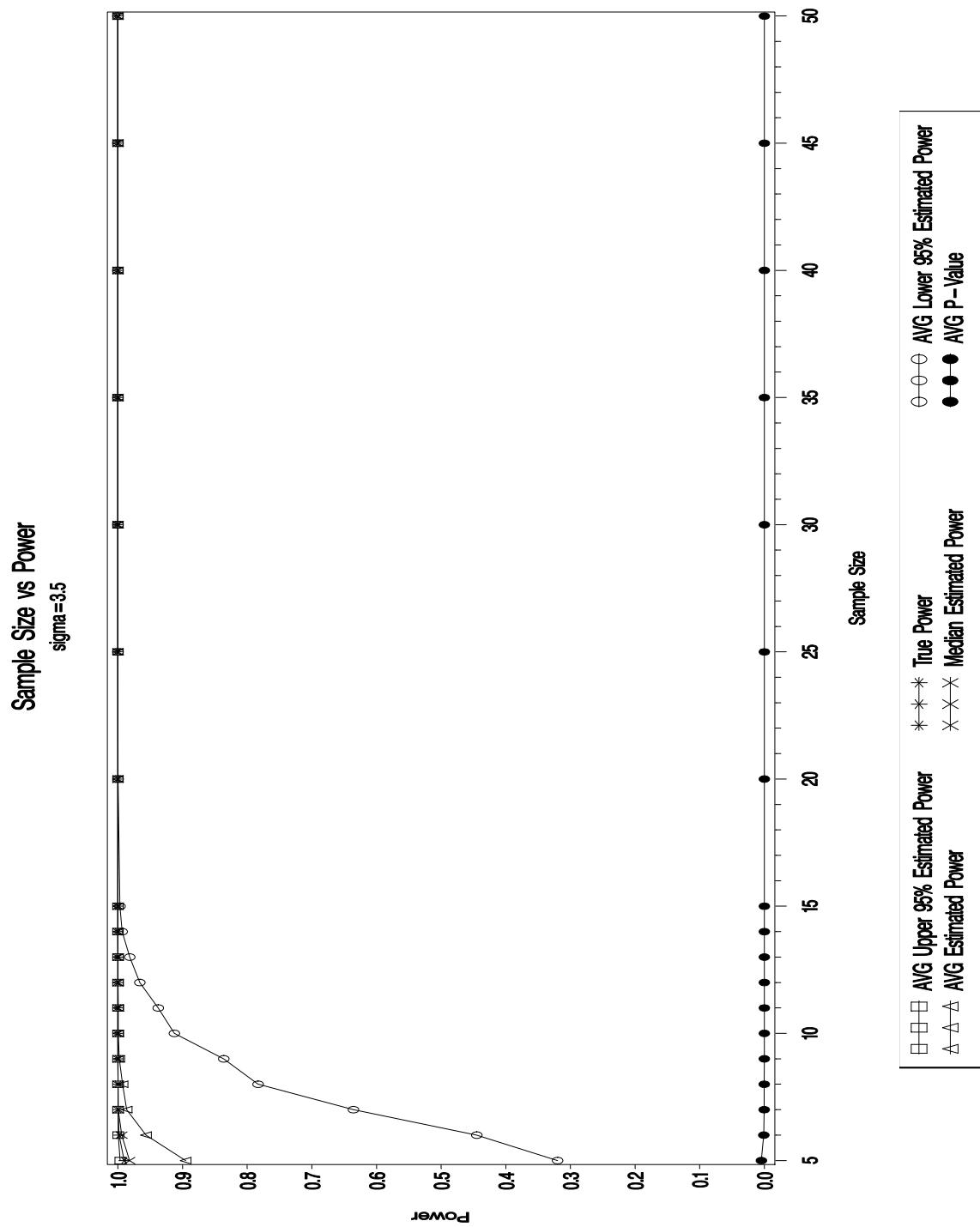
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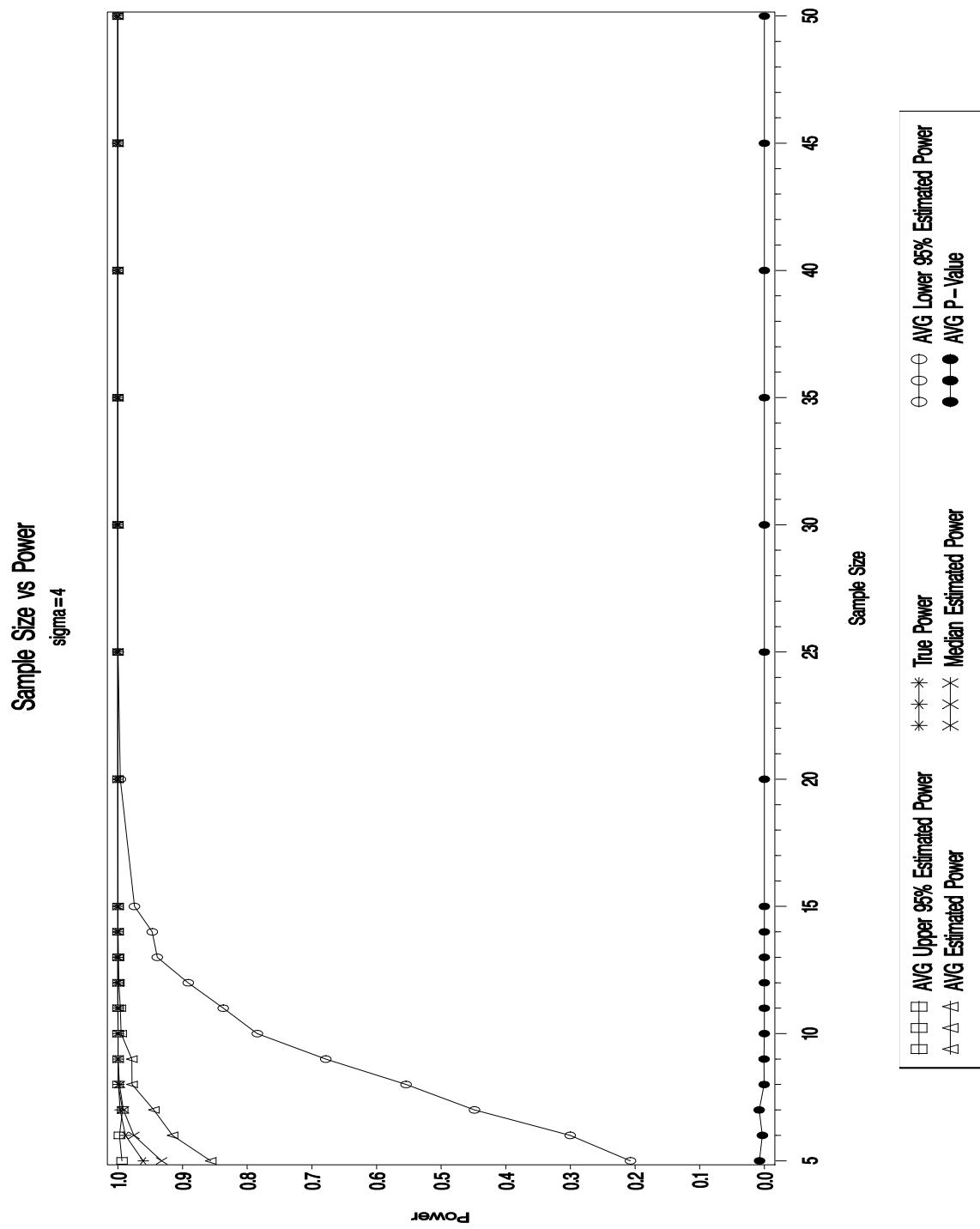
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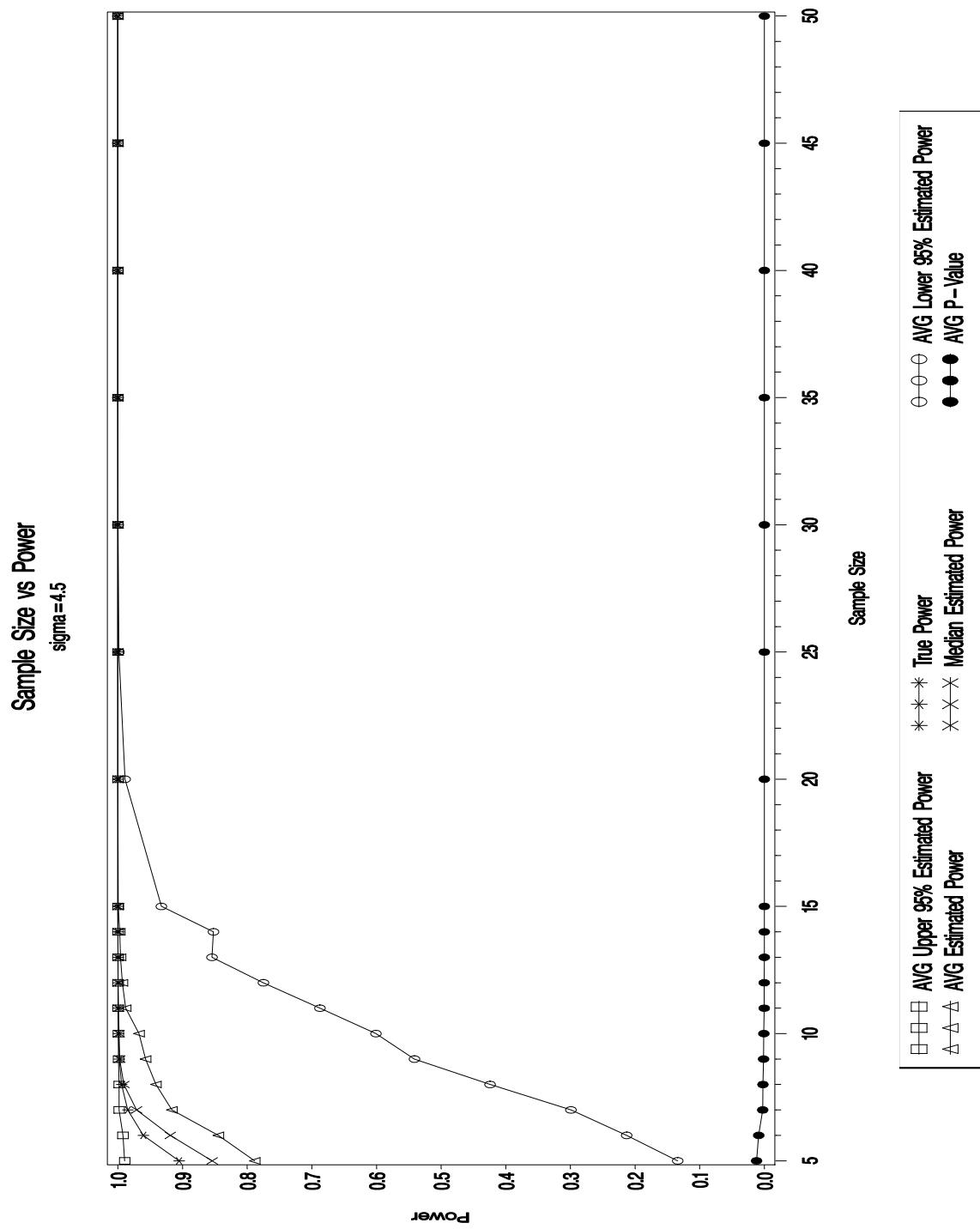
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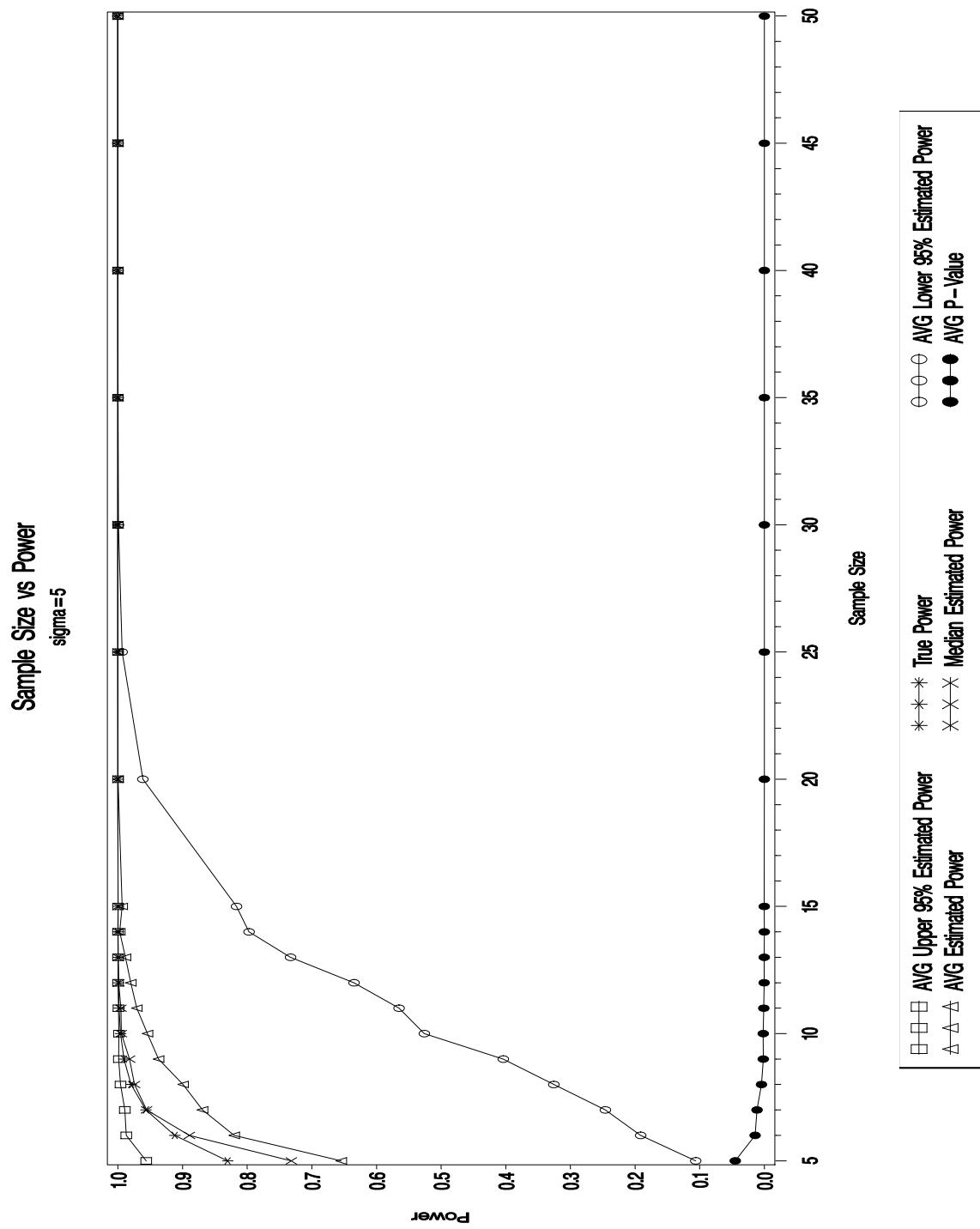
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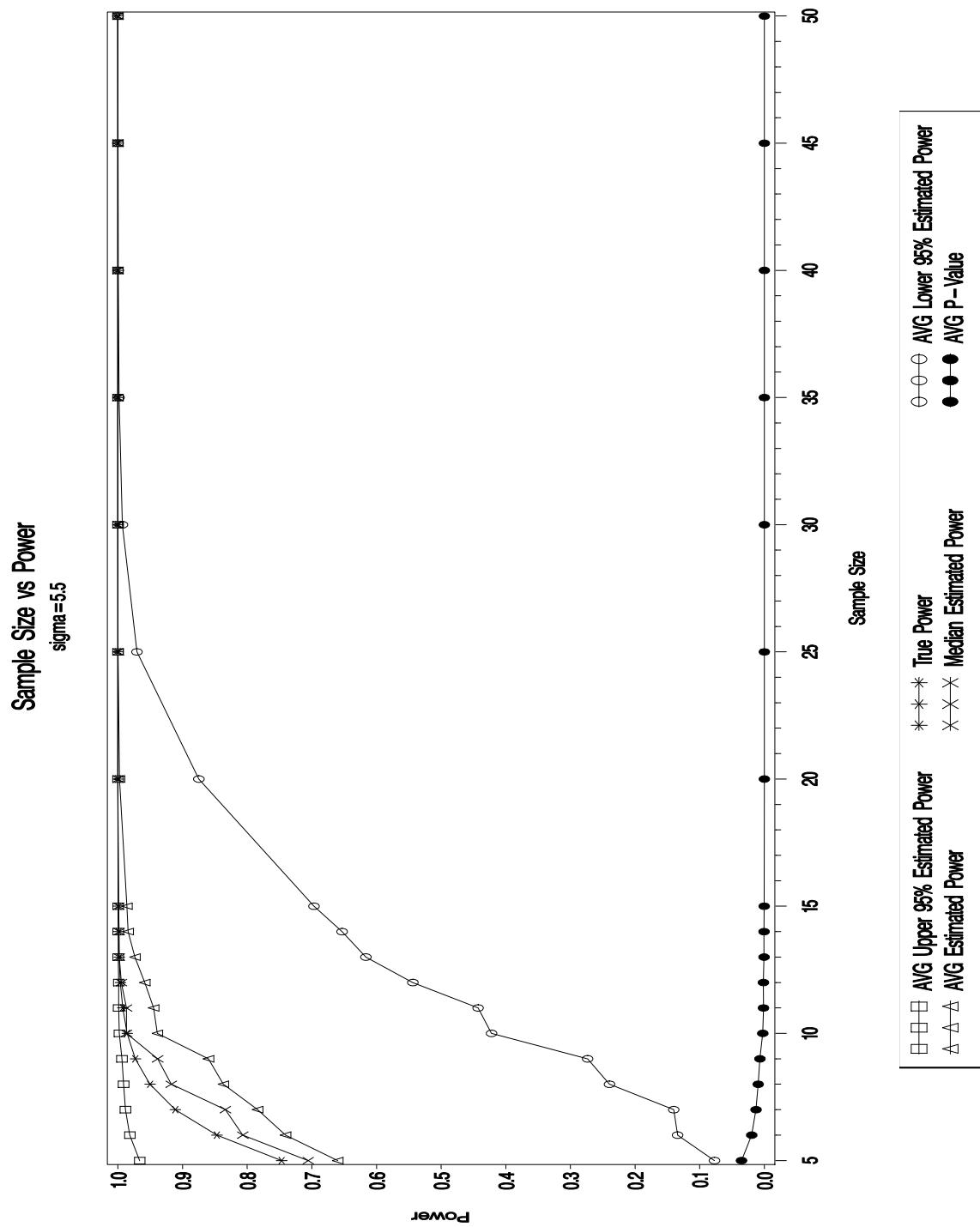
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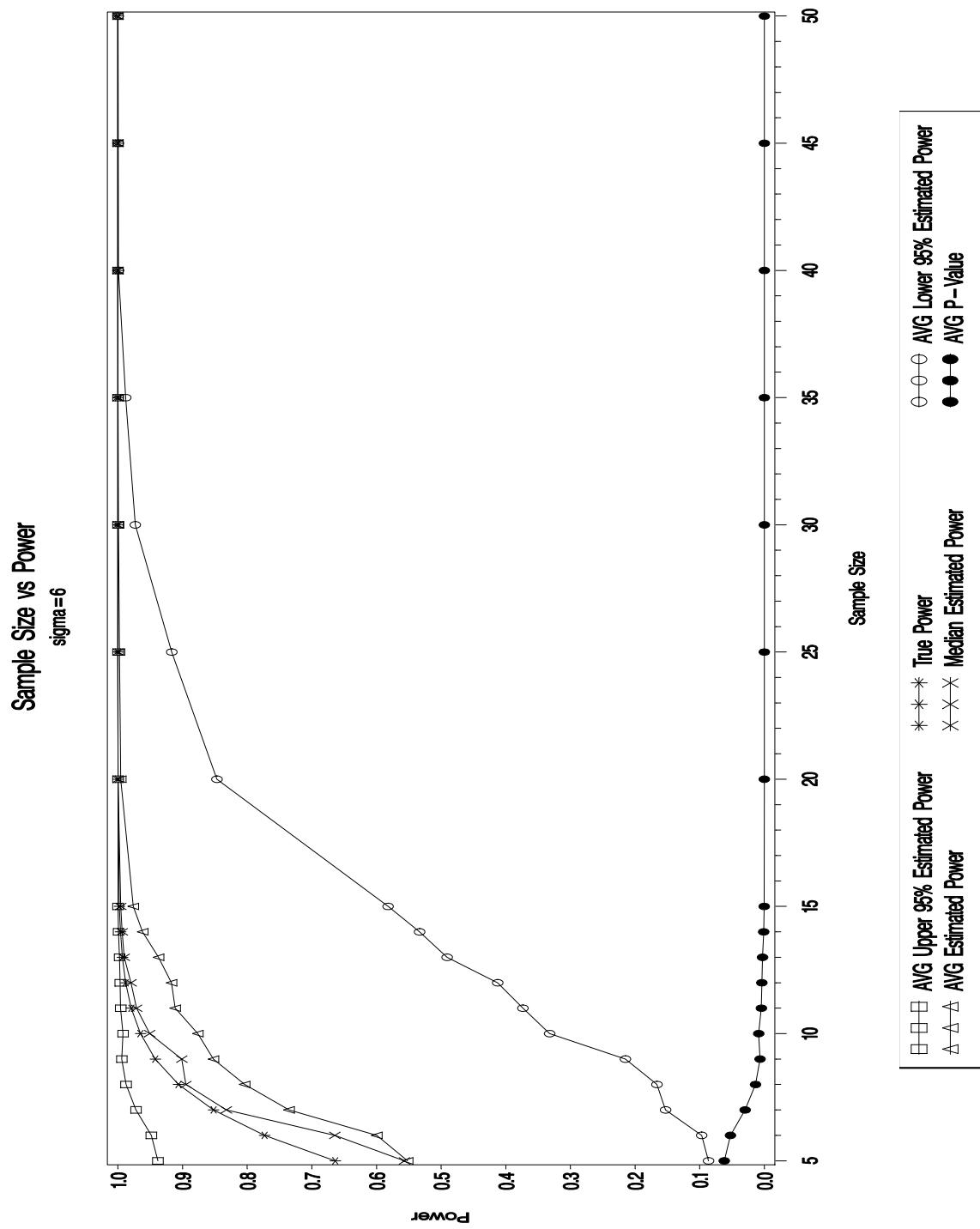
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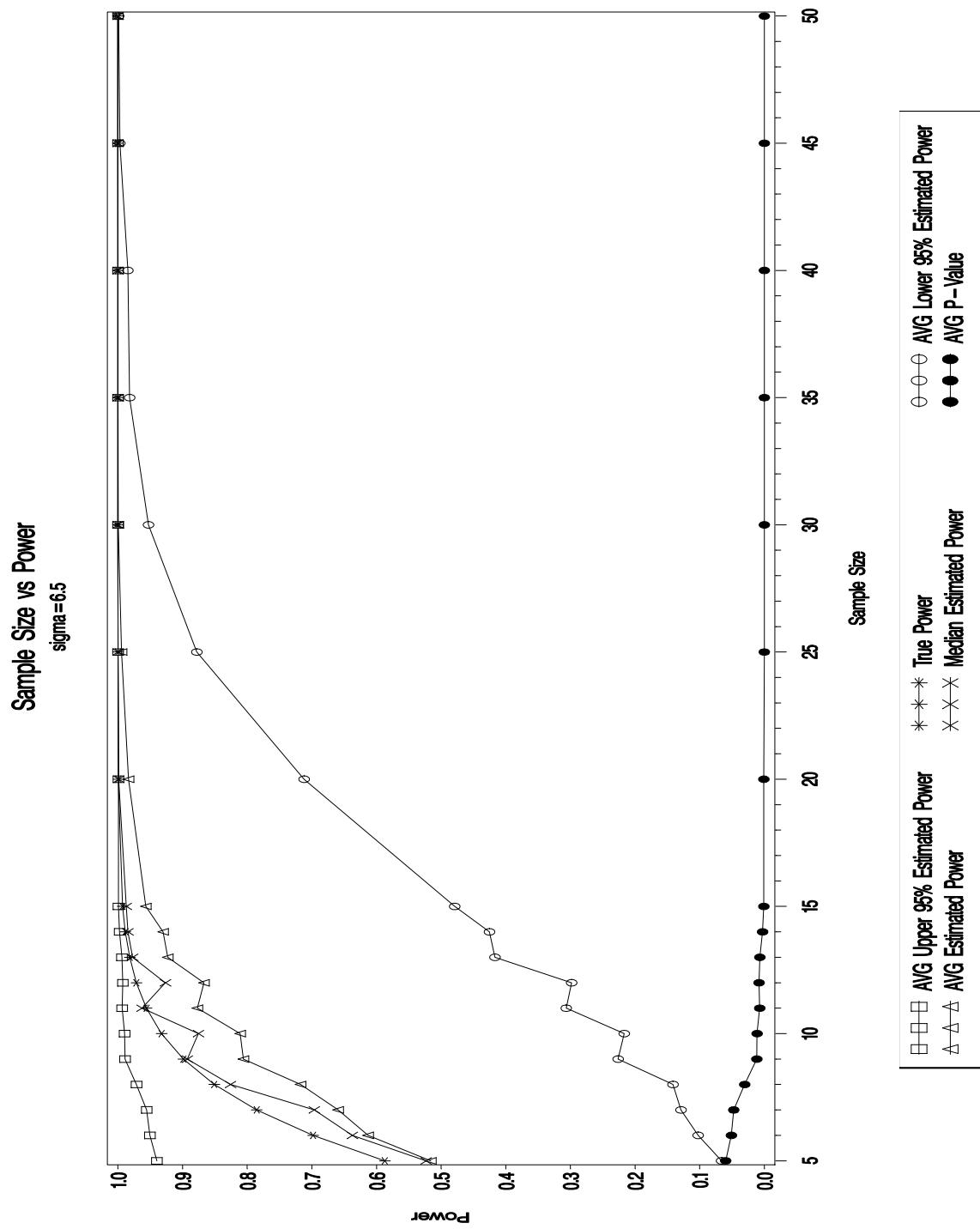
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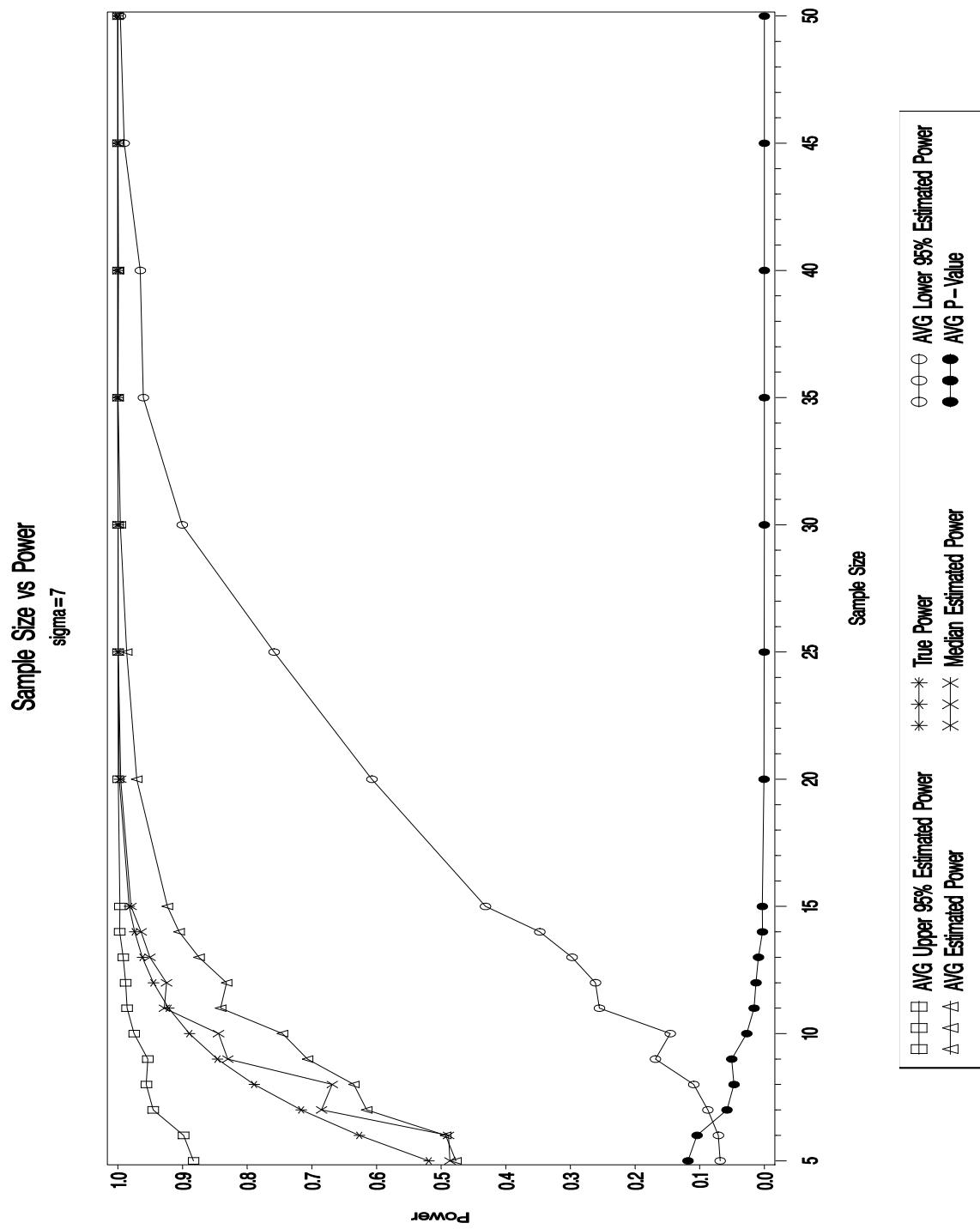
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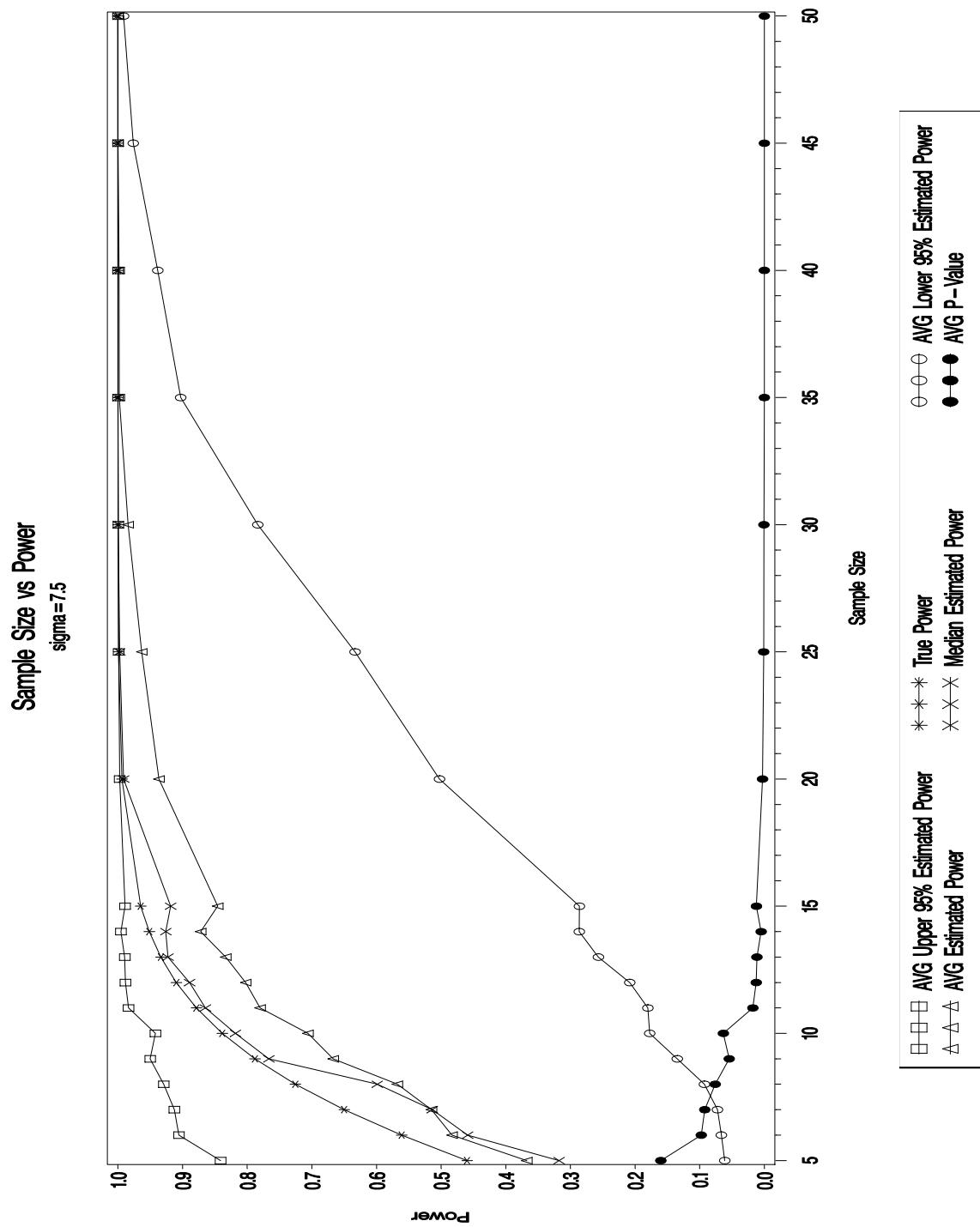
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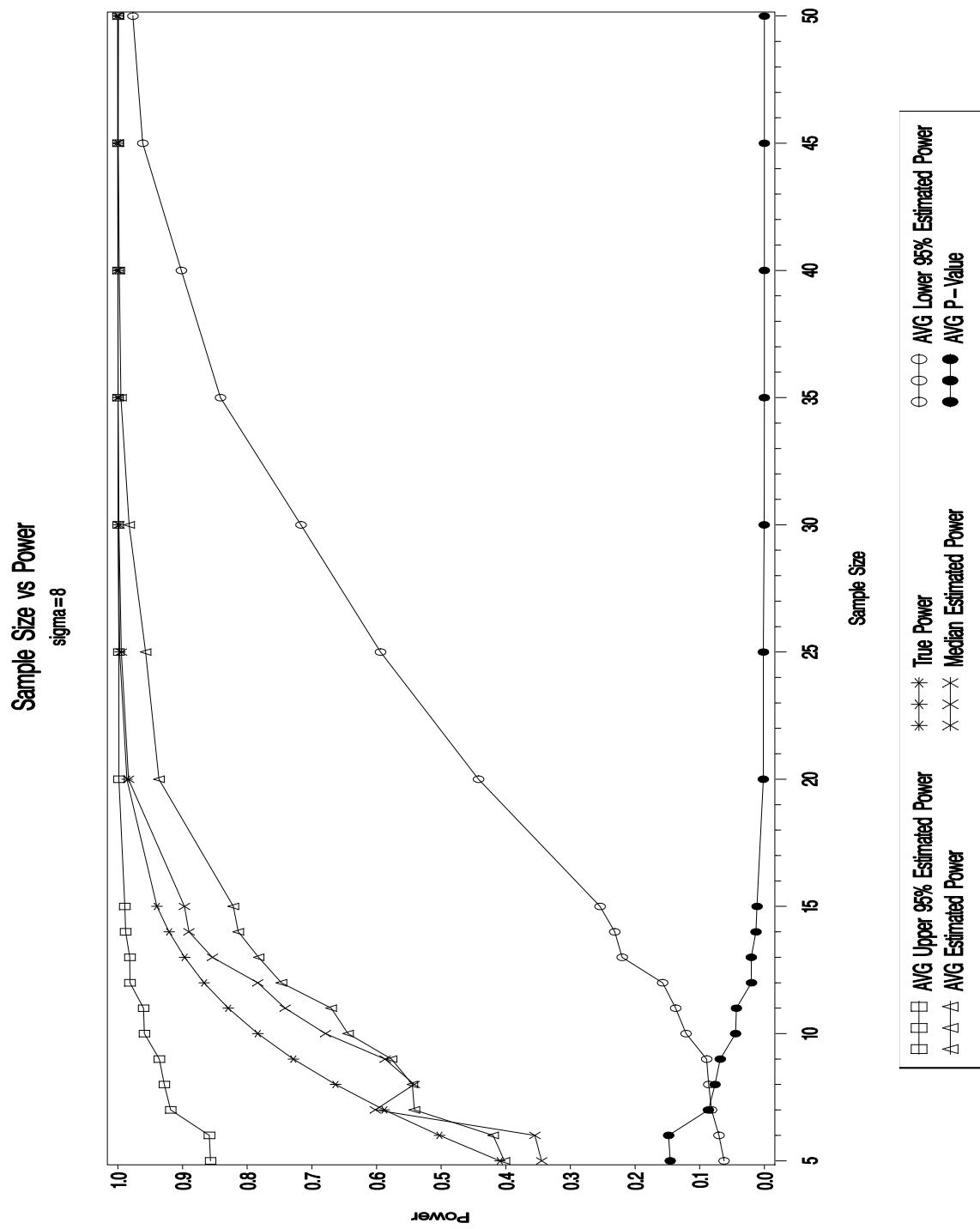
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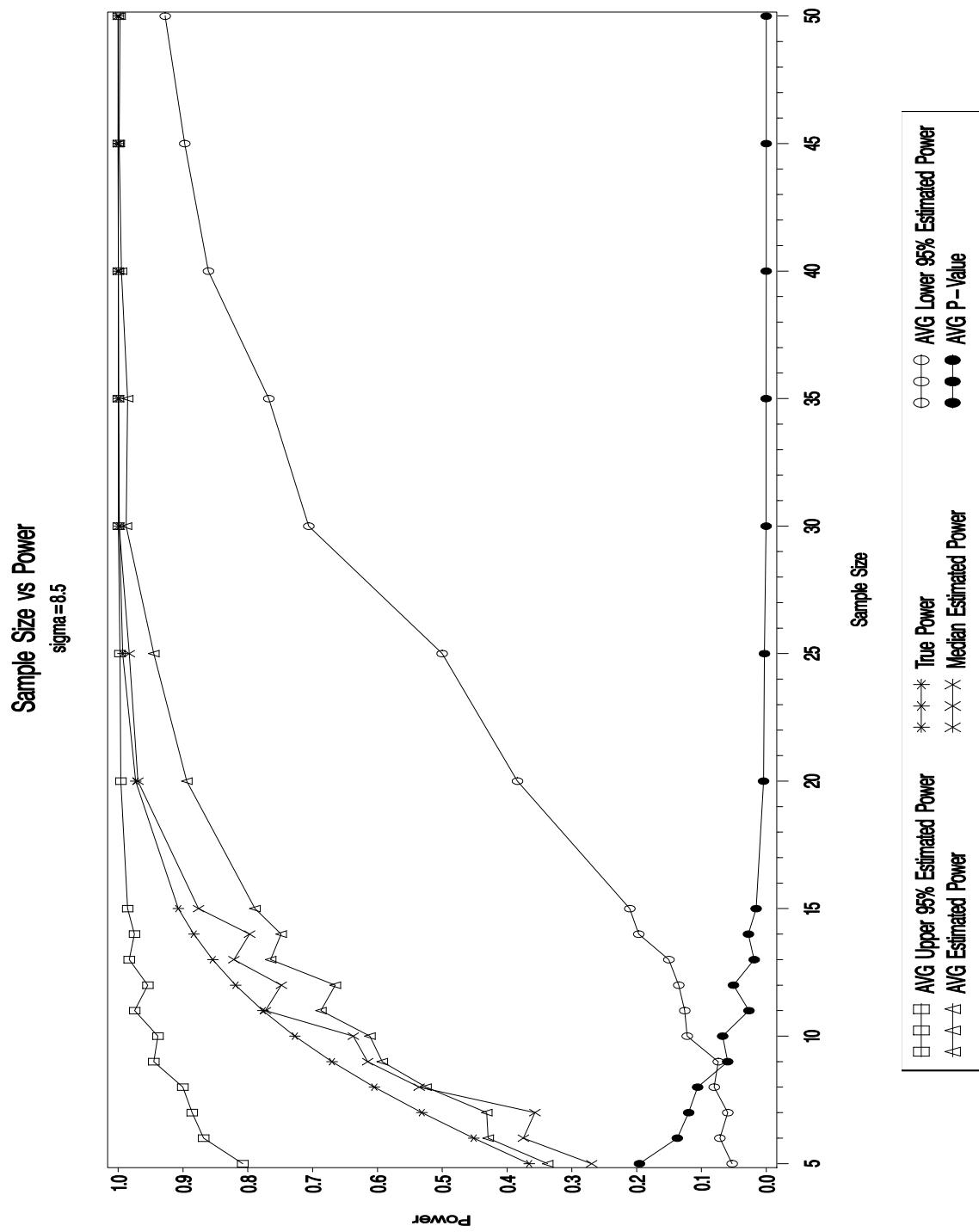
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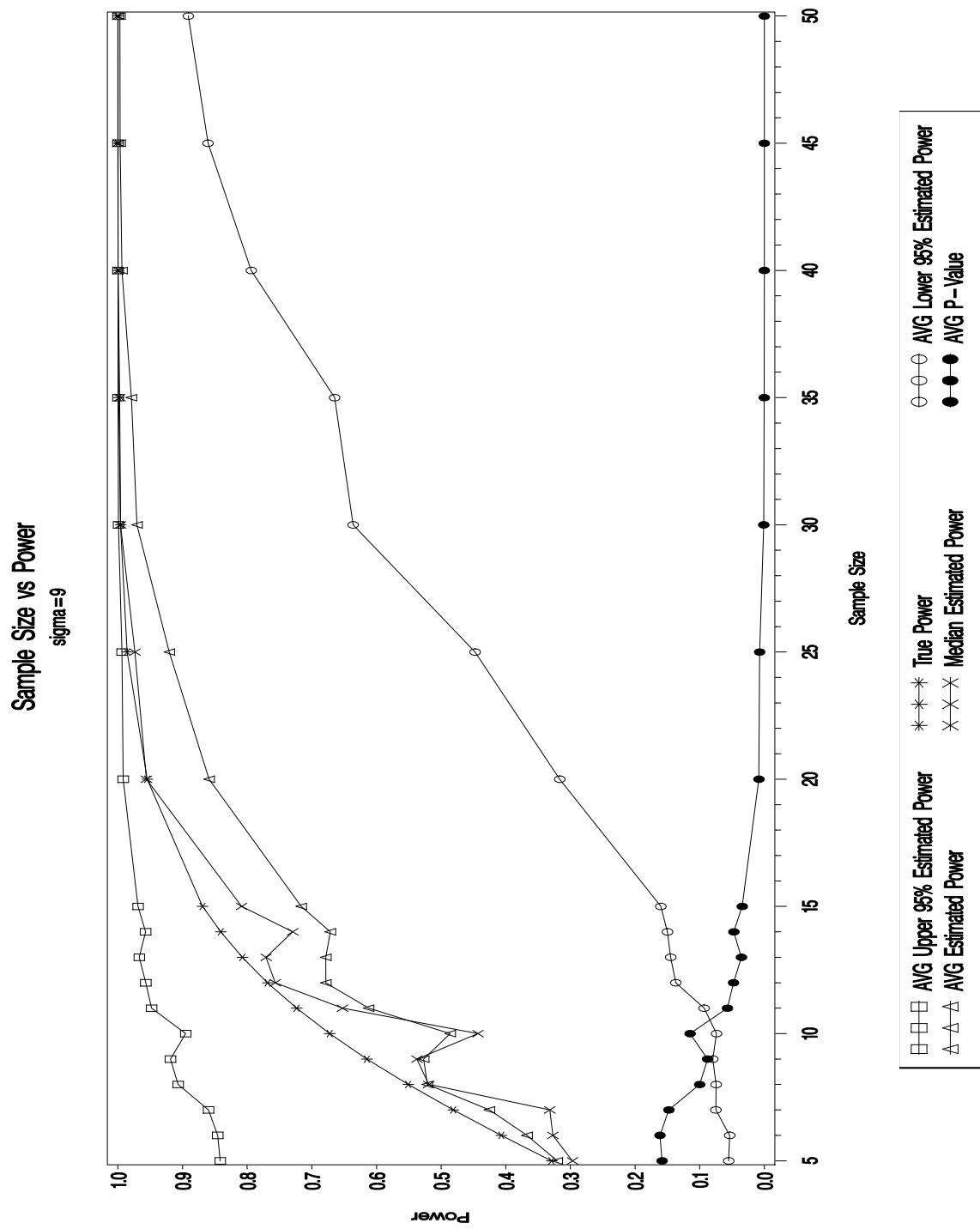
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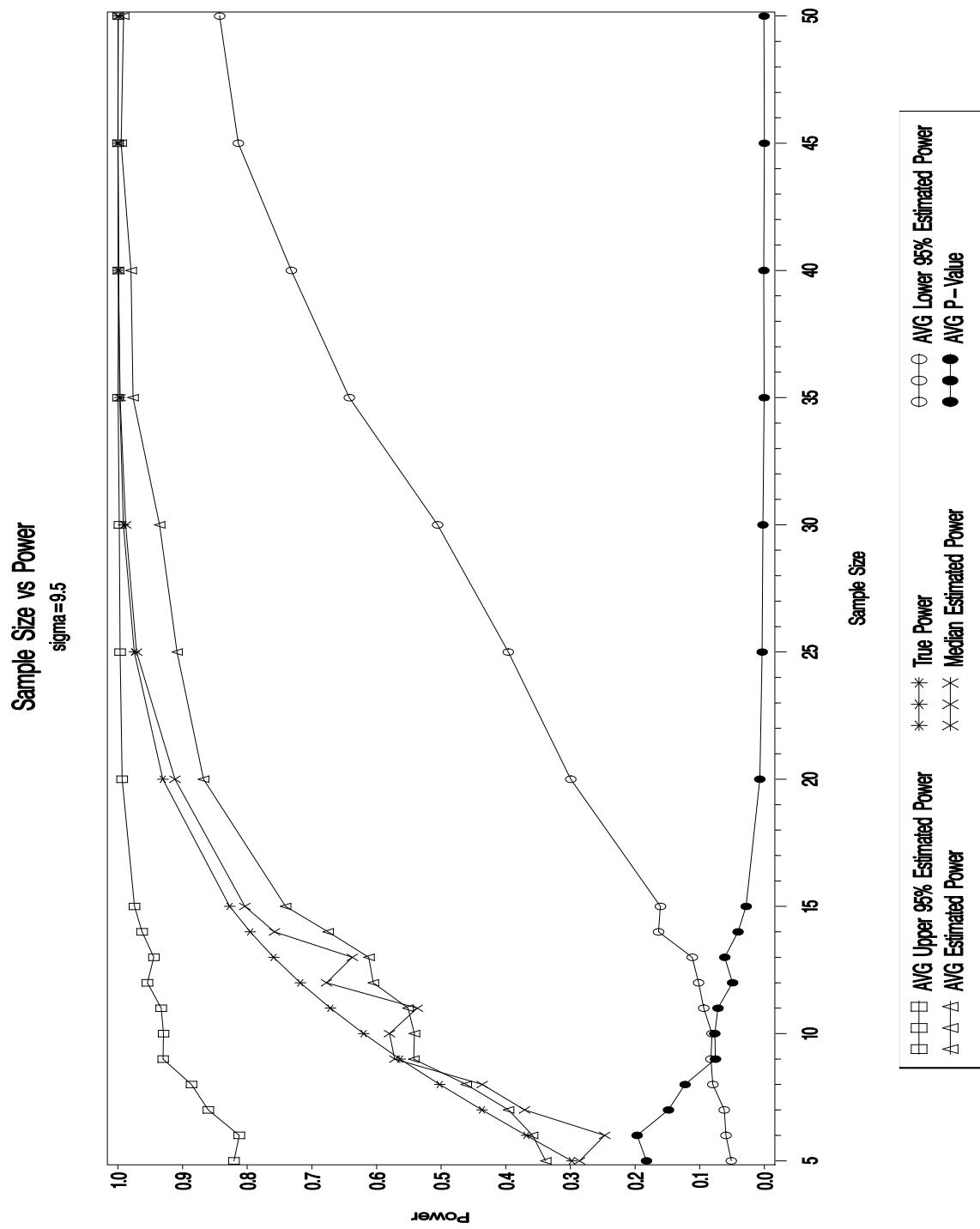
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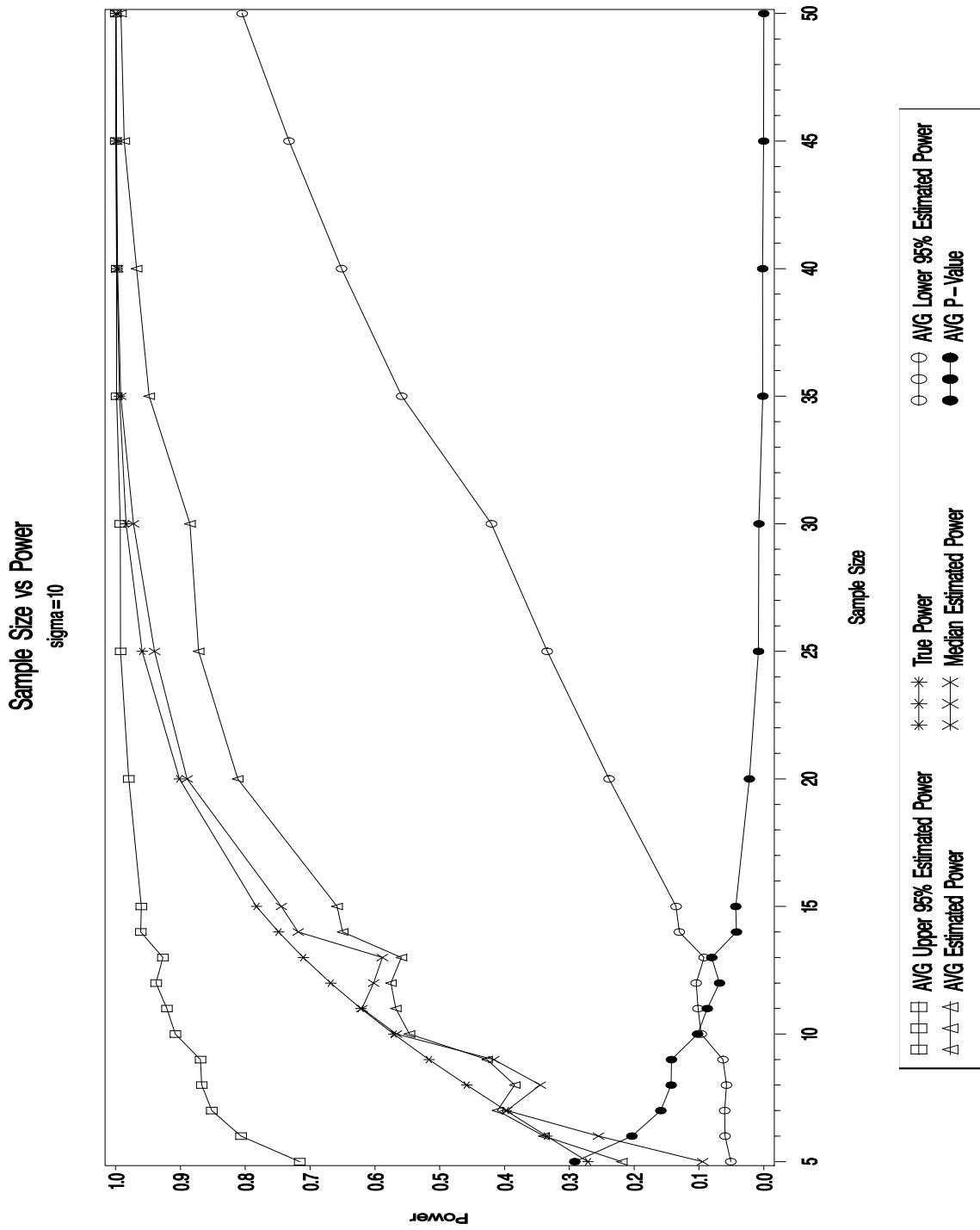
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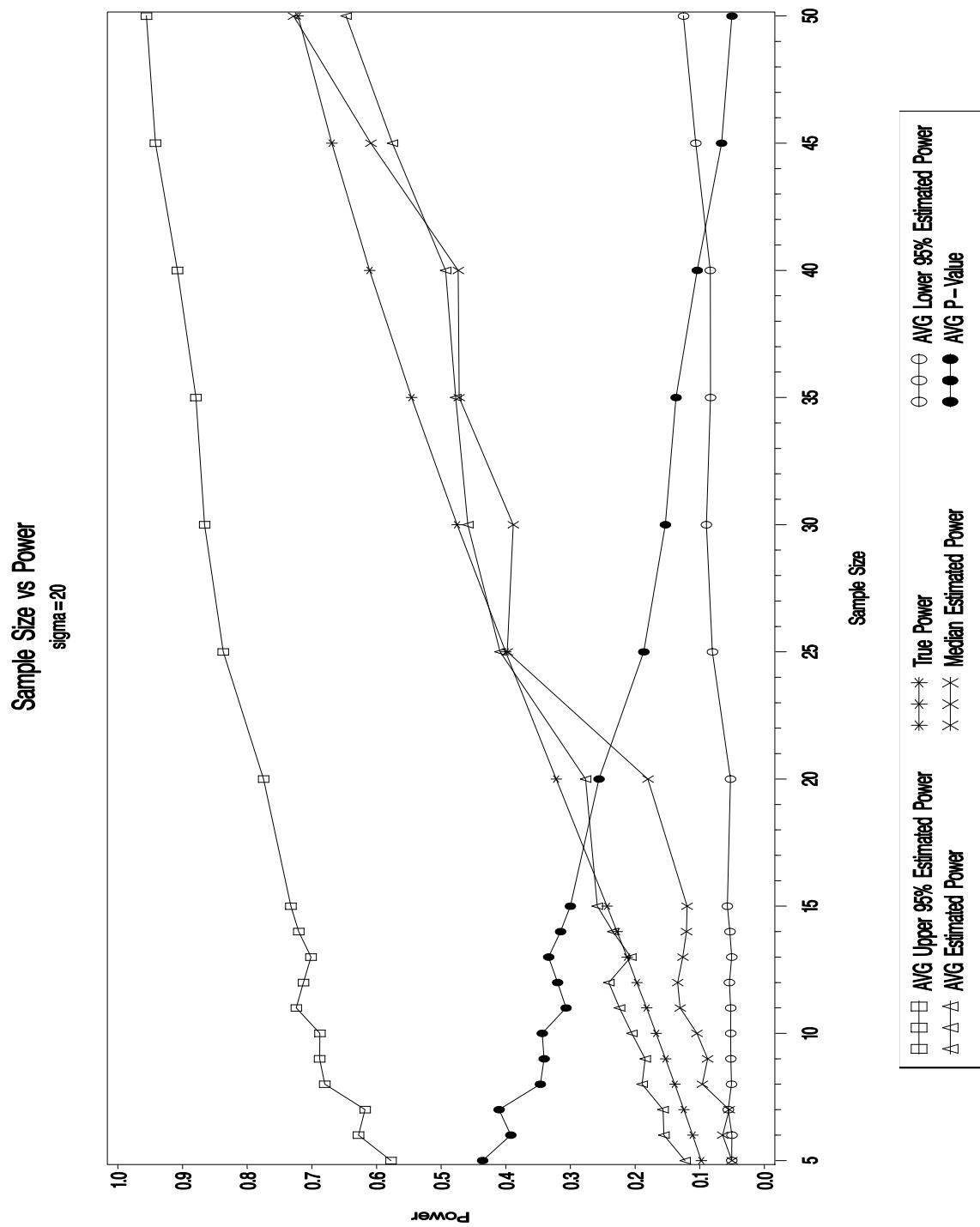
Appendix C: Graphs of Treatment Arrangement 30, 40, 40, 40



Appendix C: Graphs of Treatment Arrangement 30, 40, 40, 40



Appendix C: Graphs of Treatment Arrangement 30, 40, 40, 40



Appendix D: SAS Code for Estimating Power for an Observed F-Statistic

```

1  /* Application *****/
2  /*
3   * Input: significance level, observed F-statistic, and
4   * numerator and denominator degrees of freedom.
5   */
6  /* Output: estimate for power (1-alpha)% confidence interval for power, and */
7  /* p-value of the test
8  */
9  /*********************************************************************/
10
11 %macro application1(F=,df1=,df2=,alpha=);
12
13 proc iml;
14   reset nolog;
15
16   /* Function *****/
17   /* Return the confidence interval width for a noncentral
18   * f-dstribution
19   */
20
21   start w(alphaLow,ndf,ddf,ncent,s1);
22
23   width = quantile('F',1-s1+alphaLow,ndf,ddf,ncent)-quantile('F',alphaLow,ndf,ddf,ncent);
24
25   return(width);
26
27   finish w;
28
29
30
31
32
33
34
35
36
37
38
39
40
41
42
43
44
45
46
47

```

Appendix D: SAS Code for Estimating Power for an Observed F-Statistic

```

48      /* Function: optimal lower critical region *****/
49      /*
50      /* Finds the minimum confidence interval width using an
51      /* implementation of the Golden Section Search method by
52      /* Kincaid and Cheney 2009, and Jan Verschelde
53      /* http://www.math.uic.edu/~jan/mcs471/Lec9/gss.pdf
54      /*
55      /* Note(s):
56      /* The Golden Section Search method finds a value which minimizes
57      /* a function. The function must be one dimensional and unimodal.
58      /*
59      /* Imagine a 'typical' pdf of an F distribution. Say we want to
60      /* setup an interval such that the area covered by the sum of the two
61      /* tails is alpha. For each alpha we can find the
62      /* associated quantile using the SAS function 'quantile'.
63      /* Since alpha is specified, knowing the lower area easily leads to
64      /* the upper area. Here we consider only the lower
65      /* area. As such we can relate interval width as a function of
66      /* lower alpha size. As we increase or decrease the size of the
67      /* lower area we increase or decrease the width of the interval.
68      /* Assuming a minimum interval width exists, we optimize over the
69      /* values [0.00, 0.05].
70      /*
71      ****
72      start alphalU(ndf1,ddf2,ncp,s1Alpha,tol);
73      /*
74      alphaA = 0;
75      /* Initial lower bracket on optimal region */
76      alphaB = s1Alpha;
77      /* Initial upper bracket on optimal region */
78      /*
79      /* Note 1 ****
80      /* For each iteration an interval
81      /* [alphaA, alphaB] is retained
82      /*
83      /* ****
84      /*
85      phi = (-1+sqrt(5))/2;
86      a1 = alphaA+phi*(alphaB-alphaA);
87      wa1 = w(a1,ndf1,ddf2,ncp,s1Alpha);
88      a2 = alphaA+phi*(alphaB-alphaA);
89      wa2 = w(a2,ndf1,ddf2,ncp,s1Alpha);
90      /*
91      /*
92      /*
93      /*
94      /*

```

Appendix D: SAS Code for Estimating Power for an Observed *F*-Statistic

```

95      /* Note 2 *****/
96      /*
97      /* The initial setup invokes two function */
98      /* calls. The following loop uses one */
99      /* function call per iteration. */
100     /*
101     /*
102     /*
103     do while (abs(alphaB-alphaA)>tol);
104     /*
105     /*
106     /* We continue until the absolute difference */
107     /* between an upper and a lower bracket is */
108     /* negligible. Tolerance is set to 1e-4. */
109     /*
110     /*
111     /*
112     /*
113     /*
114     /* Four brackets are maintained at any */
115     /* given time. */
116     /*
117     /*
118     /* If wa1 > wa2 then minimum width is between */
119     /* alphaA and a1 */
120     /*
121     /*
122     /*
123     /*
124     /* Within the interval [alphaA, a1] we have */
125     /* the width wa2. a2 is situated by the */
126     /* expression alphaA+phi*(a1-alphaA). */
127     /*
128     /*
129     /* Updated alphaB */
130     /* Save a2 */
131     /* Save wa2 */
132     /* Set new test bracket a2 */
133     /* Calculate interval width for new a2 */
134     /*
135     /*
136     /* If wa1 < wa2 then minimum width is between */
137     /* a2 and alphaB */
138     /*
139     /*
140     /*
141

```

Appendix D: SAS Code for Estimating Power for an Observed F -Statistic

```

142      /* Note 6 *****/
143      /*
144      /* Within the interval [a2, alphaB] we have */
145      /* the width wa1. a1 is situated by the */
146      /* expression alphaA+phi*phi*(alphaB-a2). */
147      /*
148      /*
149      /*
150      alphaA = a2;
151      /*
152      /* Updated alphaA */
153      a2 = a1;
154      /*
155      /* Save a1 */
156      wa2 = wa1;
157      /*
158      /* Save wa1 */
159      a1 = alphaA+phi*(alphaB-alphaA);
160      /*
161      /* Set new test bracket a1 */
162      wa1 = w(a1,ndf1,ddf2,ncp,s1Alpha);
163      /*
164      /*
165      /* Calculations */
166      /*
167      eps = 1e-4;
168      /*
169      fcrit = quantile('F', 1-&alpha, &df1, &df2);
170      /*
171      pValue = 1-cdf('F', &F, &df1, &df2);
172      /*
173      estLambda = &F*&df1;
174      /*
175      adjEstLambda = (estLambda*(&df2-2)/&df2)-&df1;
176      /*
177      if adjEstLambda < 0 then adjEstLambda = 0;
178      /*
179      /*
180      /*
181      /*
182      /*
183      /*
184      /*
185      /*
186      /*
187      /*
188      /*

```

Appendix D: SAS Code for Estimating Power for an Observed *F*-Statistic

```

189
190   alphaLU = alphaLU(&df1,&df2,adjEstLambda,&alpha,eps); /* Lower significance */
191   alphaU = &alpha-alphaL; /* Upper significance */
192
193   adjAlphaL = 1-alphaL;
194   probCenF = cdf('F',&F,&qf1,&df2);
195
196   adjAlphaL = 1-adjAlphaL;
197
198   /* fnonct reports an error for a supplied */
199   /* probability greater than the */
200   /* probability from the cdf of a central */
201   /* F distribution. */
202
203   /* As the probability supplied gets large */
204   /* fnonct tends toward zero. probCenF */
205   /* with fnonct is small around zero. */
206   /* Using adjAlphaL would then be smaller */
207   /* about zero. Set to zero. */
208
209   if adjAlphaL > probCenF then lambdaL = 0; /* See note 8 */
210   else lambdaL = fnonct(&F,&df1,&df2,adjAlphaL);
211
212   if alphaU > probCenF then lambdaU = 0; /* See note 8 */
213   else lambdaU = fnonct(&F,&df1,&df2,alphaU);
214
215   estPower = 1-cdf('F',fcrit,&df1,&df2,adjEstLambda); /* Estimated power */
216   powerL = 1-cdf('F',fcrit,&df1,&df2,lambdaL); /* Lower limit for power */
217   powerU = 1-cdf('F',fcrit,&df1,&df2,lambdaU); /* Upper limit for power */
218
219   print estPower;
220   print powerL, powerU, pValue;
221
222   quit;
223
224 %mend application1;
225
226 %application1(F=1.5,df1=4,df2=10,alpha=0.05);
227

```

Appendix D: Section 5 Output for Example 1

```
1      The SAS System          15:16 Thursday, November 10, 2005  80
2
3      ESTPOWER
4
5      0.0975334
6
7      POWERL
8      0.05
9      POWERU
10     0.8017428
11     PVALUE
12
13     0.2278353
14
15
16
17
```

Appendix D: Section 5 Output for Example 2

```
1      The SAS System          15:16 Thursday, November 10, 2005  81
2
3      ESTPOWER
4
5      0.0777412
6
7      POWERL
8      0.05
9      POWERU
10     0.6713596
11     PVALUE
12
13     0.2741814
14
15
16
17
```
