

Interpreting Heterogeneity in Meta Analyses of Clinical Trials

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Introduction

- Outline how to investigate heterogeneity
- Give statistical test
- Highlight graphical methods
- Use worked examples throughout

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Assessing Statistical Assumptions

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Statistical Tests

- The test of homogeneity of the treatment effect across studies is based on the statistic

$$Q = \sum_{i=1}^k (\hat{\theta}_i - \hat{\theta})^2 w_i$$

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The Residuals from a Meta Analysis

- The weighted residuals from fitting a fixed-effects meta-analysis are given by

$$q_i = (\hat{\theta}_i - \hat{\theta})\sqrt{w_i}$$

- Under the standard fixed-effects meta-analysis the standardised weighted residuals are

$$q'_i = \frac{(\hat{\theta}_i - \hat{\theta})\sqrt{w_i}}{\sqrt{1 - w_i / \sum_{i=1}^k w_i}}$$

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Assessing the Residuals

- The standardised residuals should follow a standard Normal distribution
- Under the standard random-effects meta-analysis the weighted residuals and standardised weighted residuals would be calculated from same equations by replacing w_i and $\hat{\theta}$ with w_i^* and $\hat{\theta}^*$

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Assessing the Residuals

- A Normal probability plot of the q_i or q'_i may be used to check the distributional assumptions of the meta-analysis model

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Multi-Centre Trials v Meta Analysis

- A multi-centre trial may be analysed in the same way as a meta-analysis
 - That is by treating centres as studies.
- Multi-centre trials can be analysed using an overall model which includes the terms centre and treatment.
- Homogeneity of treatment effects across centres can be undertaken by fitting a treatment by centre interaction to obtain an estimate of treatment effect with its variance for each centre.
- The standardised residual can be calculated replacing the w 's with the inverse of the variance.

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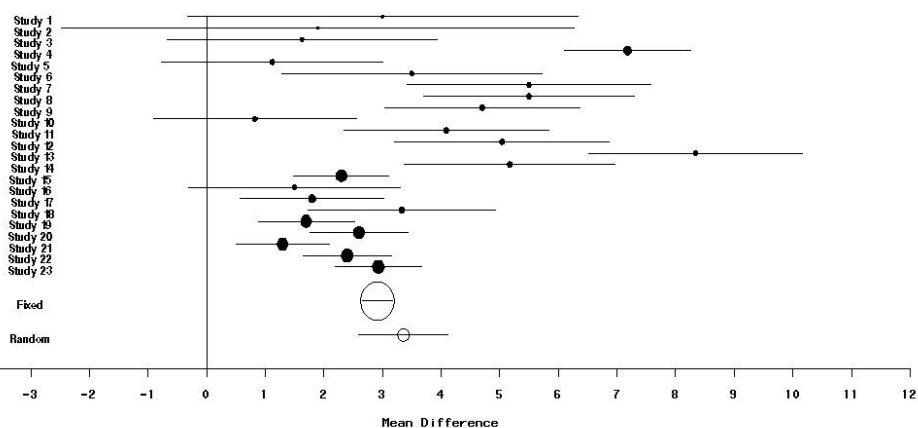
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Worked Example Paroxetine in Adults

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Meta analysis of 23 Paroxetine Trials



Paroxetine Trials

- The treatment effect is measured by the difference in mean HAMD between Paroxetine and placebo.
- There seems to be some evidence of heterogeneity in treatment effect though most of the heterogeneity seems to be in the smaller trials
- The fixed effect meta-analysis gave an overall mean difference of 2.91 with a 95% confidence interval of (2.66 to 3.17).
 - The test of homogeneity gives a P-value of <0.001 (chi-squared of 170.23 on 22 degrees of freedom).
- The overall mean difference from the random effects meta-analysis is 3.36 with 95% confidence interval of (2.59 to 4.12)

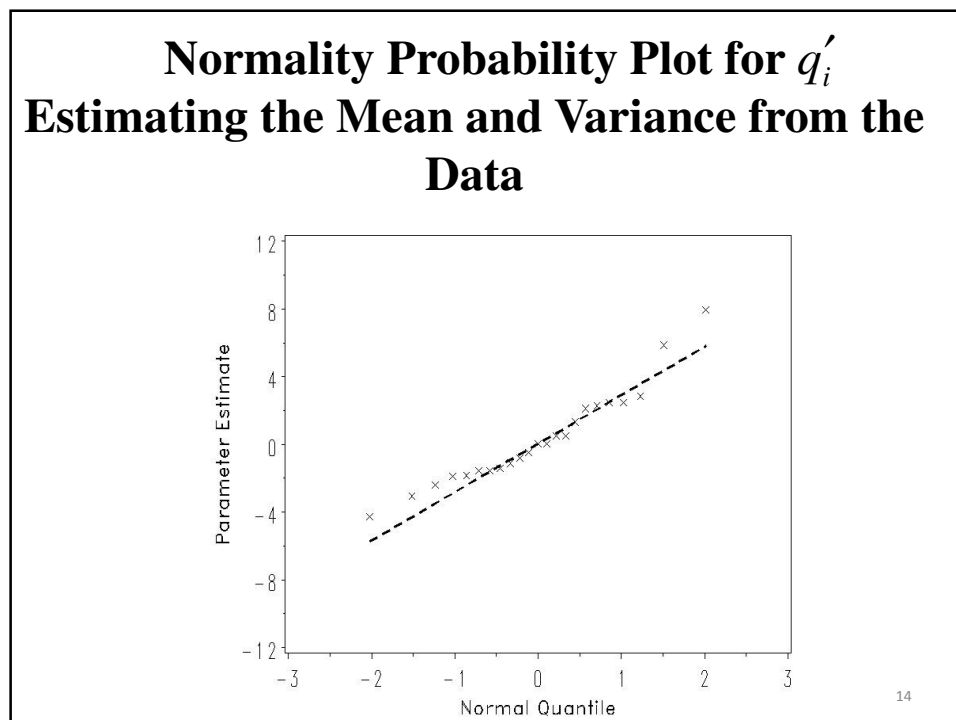
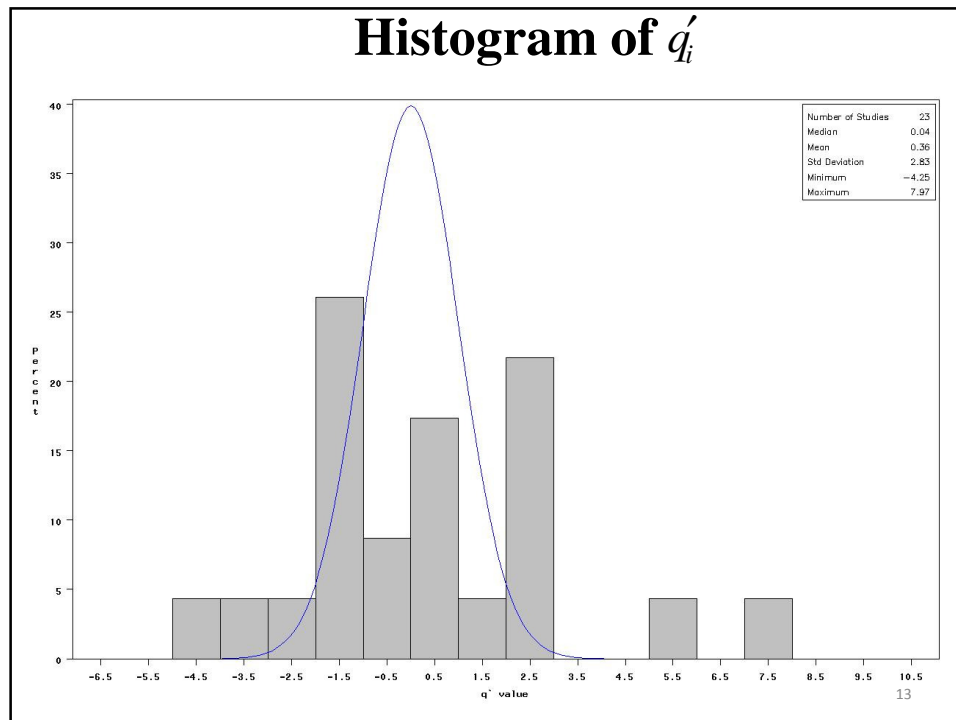
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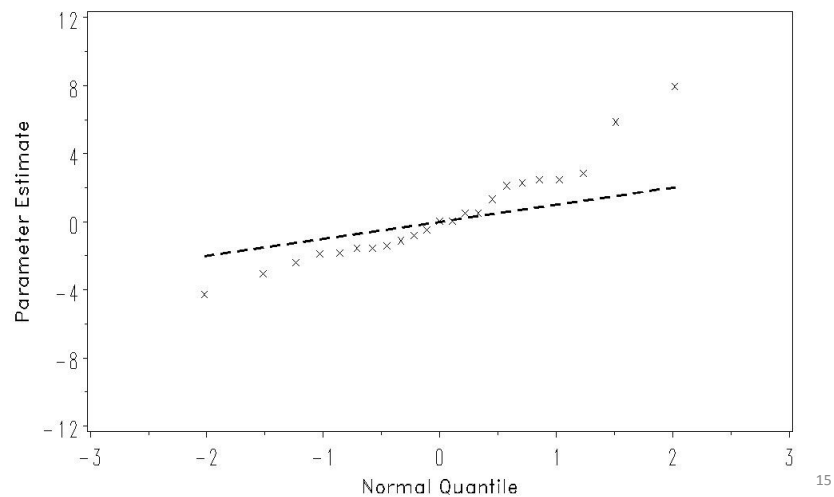
Fixed Effects

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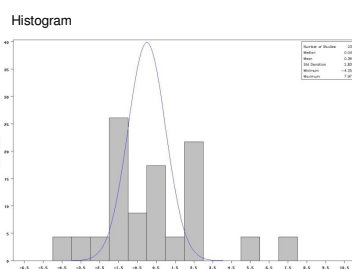
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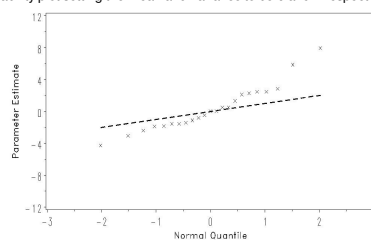
Normal Probability Plot for q_i' Setting the Mean and Variance to be 0 and 1 Respectively



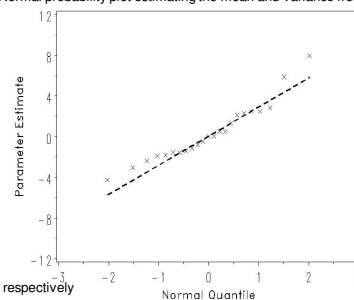
Assessment of Homogeneity of Treatment Effects with Normality Assumption of Fixed Effects Meta-analysis



Normal probability plot setting the mean and variance to be 0 and 1 respectively



Normal probability plot estimating the mean and variance from the data

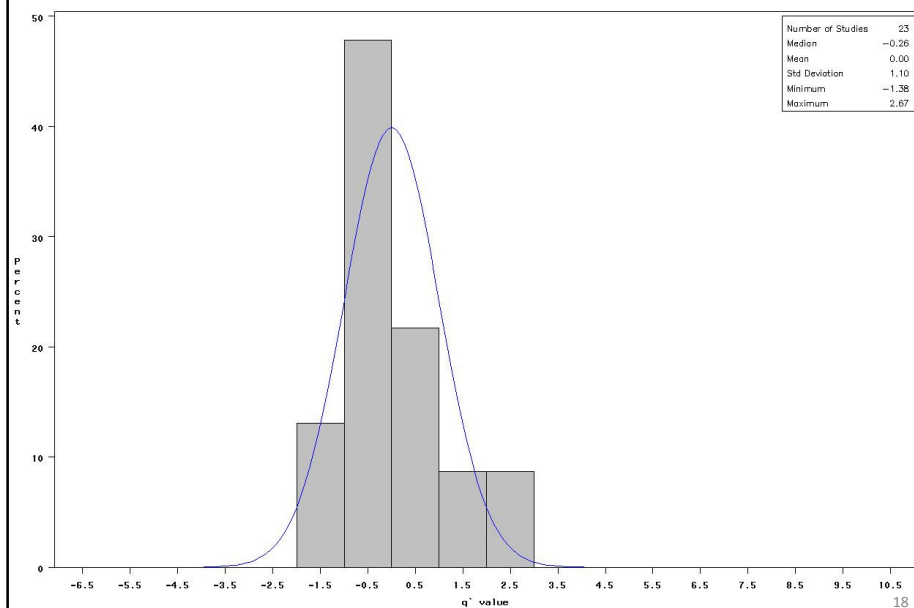


Random Effects

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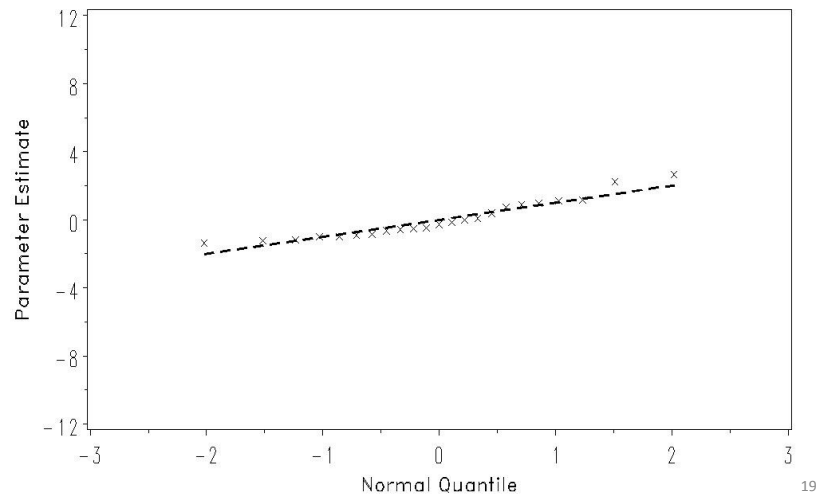
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Histogram of q'_i



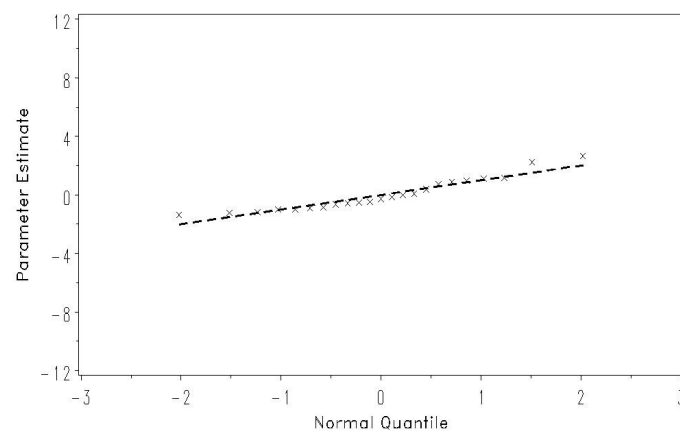
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Normality Probability Plot for q'_i Estimating the Mean and Variance from the Data



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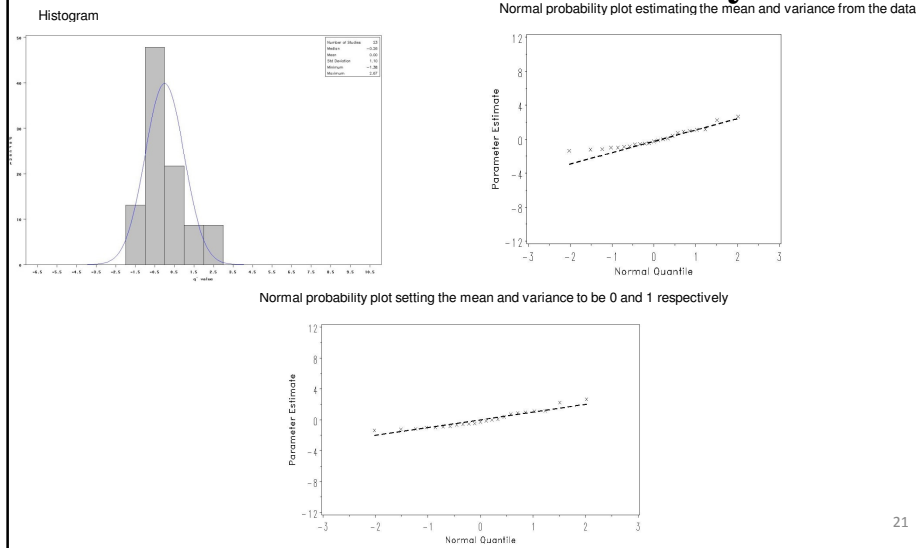
Normal Probability Plot for q'_i setting the Mean and Variance to be 0 and 1 Respectively



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Assessment of Homogeneity of Treatment Effects with Normality Assumption of Random Effects meta-Analysis



Summary of Results so Far

- There seems to be deviation from the assumptions for a fixed effects meta analysis although the data do seem to take a Normal form
- The assumptions seem to hold better for a random effects meta analysis
- If a fixed effects analysis was the primary analysis then the results would need to be interpreted with care
- The diagnostics so far although easy to produce in most statistics packages but have limitations
- Could an assessment of outliers better be made?

Confidence Bands for the Normal Probability Plots

- Confidence bands for the Normal probability plots can be constructed to assist in interpretation
- For the Normal probability line confidence bands can be calculate
 - Using Normal approximation (using the Friendly macro)
 - Using simulation
- For the observed data a confidence band can be calculated
 - Using bootstrapping

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Simulation Envelopes

1. Simulate a value for the estimated treatment effect from study i from a $N(0, w_i^{-1})$, for $i = 1, \dots, k$.
2. Perform the fixed-effects meta-analysis and calculate the q'_i
3. Order the q'_i from smallest to largest
4. Repeat 1. to 3. a number of times, for example 1,000
5. For each i , calculate the 2.5th and 97.5 percentile of the 1,000 values, to give the 95% lower and upper bounds for all points

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Bootstrapping

1. From the observed data pairs (θ_i, w_i^{-1}) , $i = 1, \dots, k$, take a random sample with replacement of size k
2. Perform the fixed-effects meta-analysis and calculate the q'_i
3. Order the q'_i from smallest to largest
4. Repeat 1. to 3. a number of times, for example 1,000.
5. For each i , calculate the 2.5th and 97.5 percentile of the 1,000 values, to give the 95% lower and upper bounds for all points

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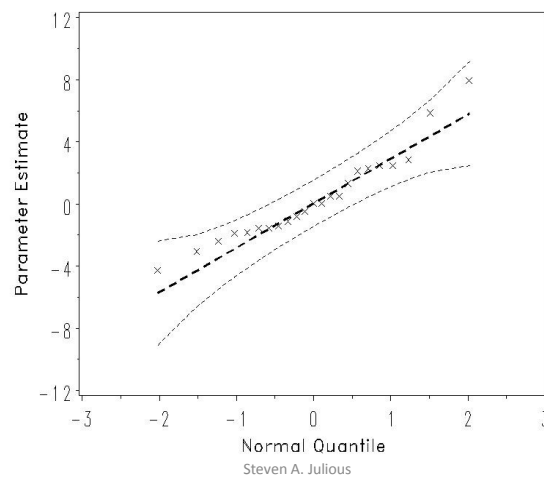
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Fixed Effects

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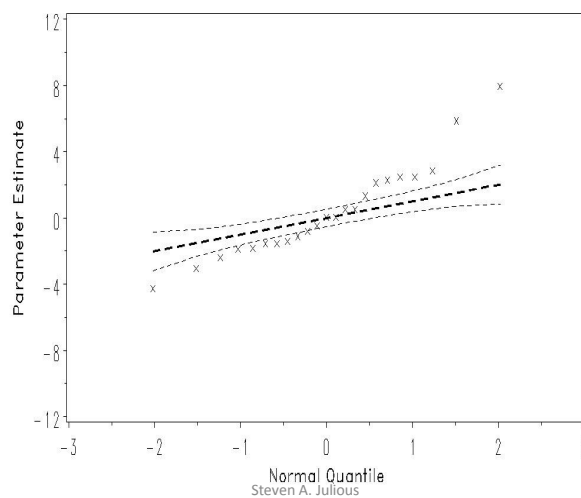
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Normal Approximation, Estimating the Expected Values with the Mean and Variance from the Data



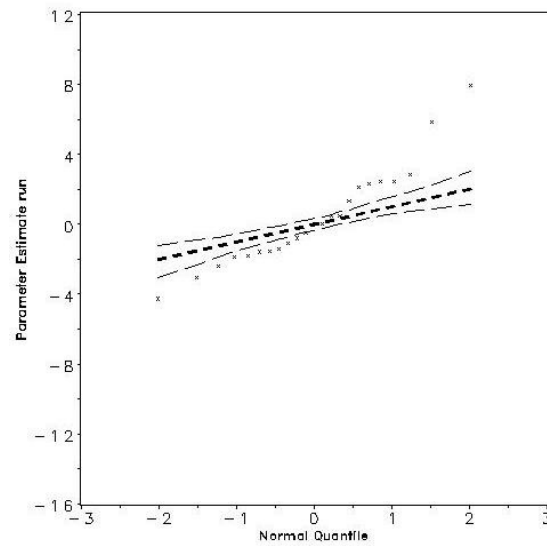
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Normal Approximation, Estimating the Expected Values from $N(0,1)$



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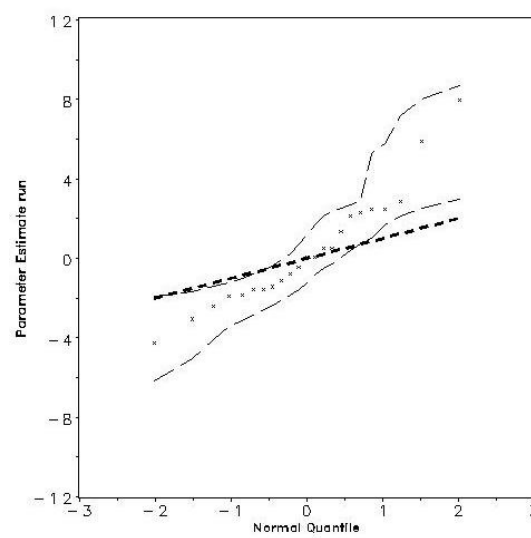
Simulated Envelopes



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Bootstrap Envelopes

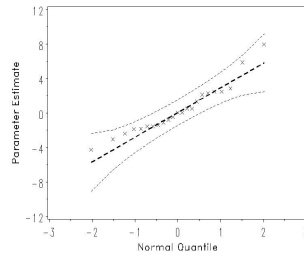


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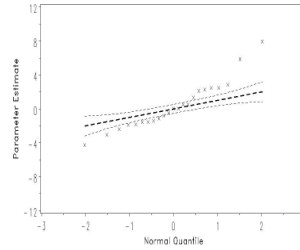
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Comparison of Plots

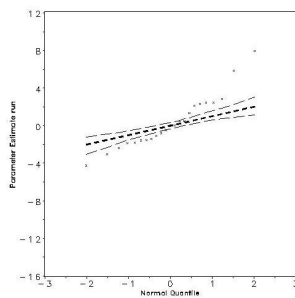
Normal using data



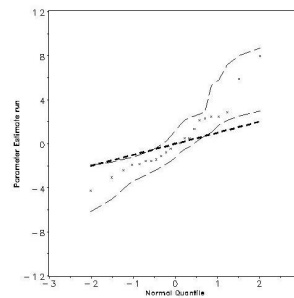
Normal using $N(0,1)$



Simulation using $N(0,1)$



Simulation



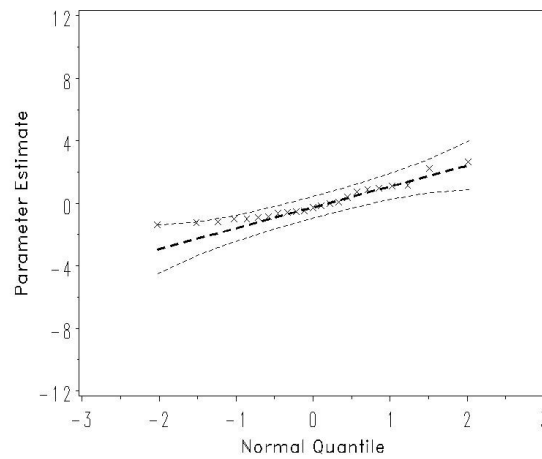
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Random Effects

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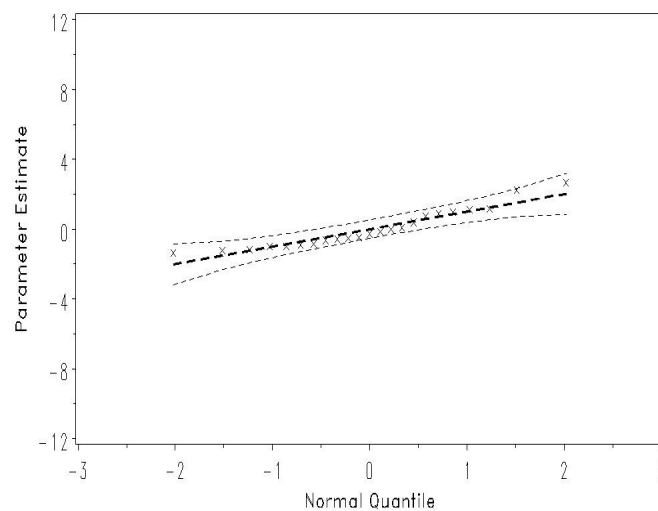
Normal Approximation, Estimating the Expected Values with the Mean and Variance from the Data



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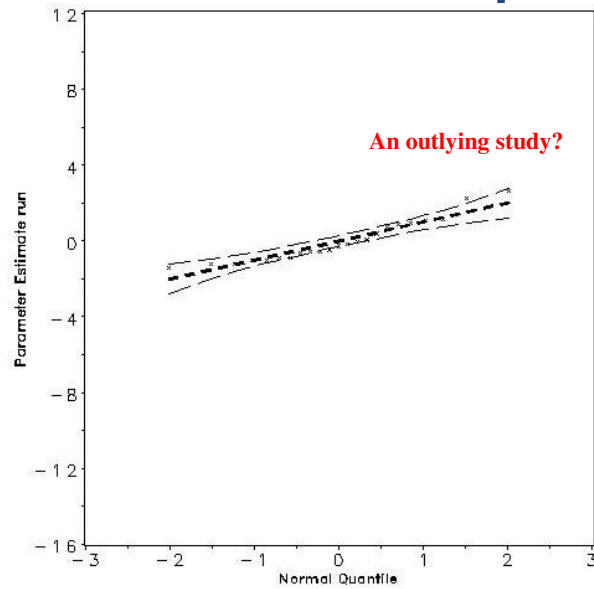
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Normal Approximation, Estimating the Expected Values from $N(0,1)$



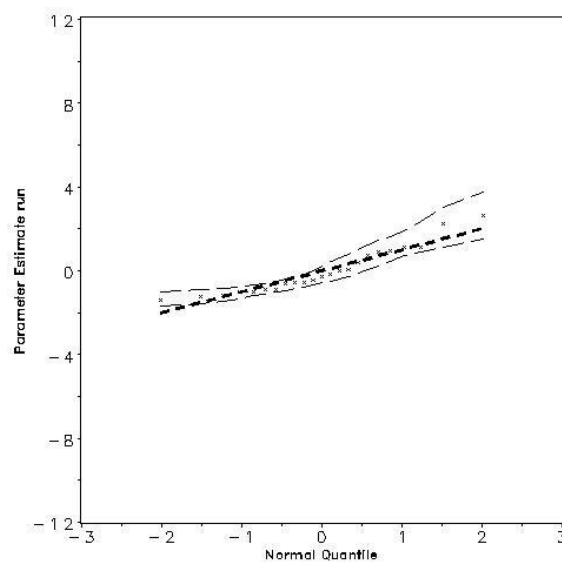
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Simulated Envelopes



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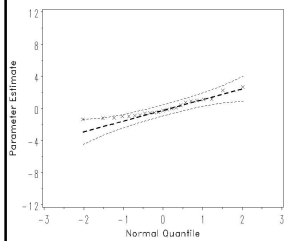
Bootstrap Envelopes



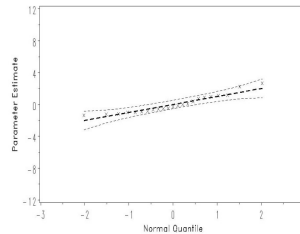
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Comparison of Plots

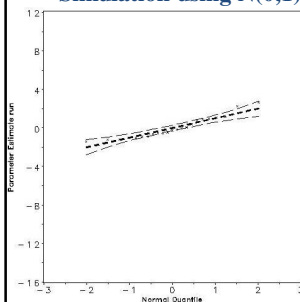
Normal using data



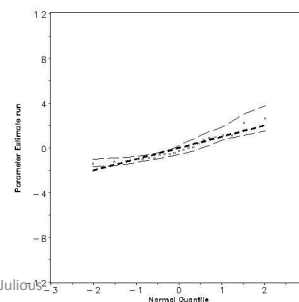
Normal using $N(0,1)$



Simulation using $N(0,1)$



Bootstrapping



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Summary of Results so Far (Again)

- The additional analyses confirm the original analysis
- The bands are wider at the extremes of the range
- For this example bootstrapping has wide bands at one end
- Allows for an assessment of outliers

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What if the Treatment Effects Differ Study to Study?

- Find explanation
 - Investigate baseline imbalances
 - Look at subgroups
- Are there any possible explanations?

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Summary of Meta Analysis

- Gave overview of how to investigate heterogeneity in a meta analysis
- Recommended not to rely on statistical tests to assess heterogeneity
- Described graphical approaches to assess heterogeneity

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