



# Distribution Fitting for Arbitrarily Censored Data

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# Censored Data

- Censored data arise when the value of an observation is only partially known.
- For example, in a medical trial the survival time of a patient may only be known to greater than some value because the patient left the study.
- Or a measurement taken to study groundwater contamination may only be known to be less than some detection or quantitation limit.

# Types of Censored Data

- Right-censored – known to be greater than.

$> 45$

- Left-censored – known to be less than.

$< 7$

- Interval-censored – known to be between.

[6,10]

# Example #1: Arsenic Concentrations

- Arsenic concentrations (ug/L) in an urban stream, Manoa Stream at Kanewai Field, on Oahu, Hawaii. (Tomlinson, 2003)

0.5	0.5	0.5	0.6	0.7	0.7	<0.9	0.9
<1.0	<1.0	<1.0	<1.0	1.5	1.7	<2.0	<2.0
<2.0	<2.0	<2.0	<2.0	<2.0	<2.0	2.8	3.2

# Example #2: Breast Cancer Study

- Days between treatment and breast retraction – Finkelstein and Wolfe (1985)

	patient	days	left	right	treatment
	Numeric	Censored	Numeric	Numeric	Character
1	1	>45	45		Rad
2	2	[6,10]	6	10	Rad
3	3	<7	0	7	Rad
4	4	>46	46		Rad
5	5	>46	46		Rad
6	6	[7,16]	7	16	Rad
7	7	>17	17		Rad
8	8	[7,14]	7	14	Rad
9	9	[37,44]	37	44	Rad
10	10	<8	0	8	Rad
11	11	[4,11]	4	11	Rad
12	12	>15	15		Rad

# Statgraphics Procedures

*Option #1: Describe – Distribution Fitting – Fitting Censored Data.*

- Handles left-censored and right-censored data.
- Includes goodness-of-fit tests.

*Option #2: R Interface – Distribution Fitting (Arbitrarily Censored Data).*

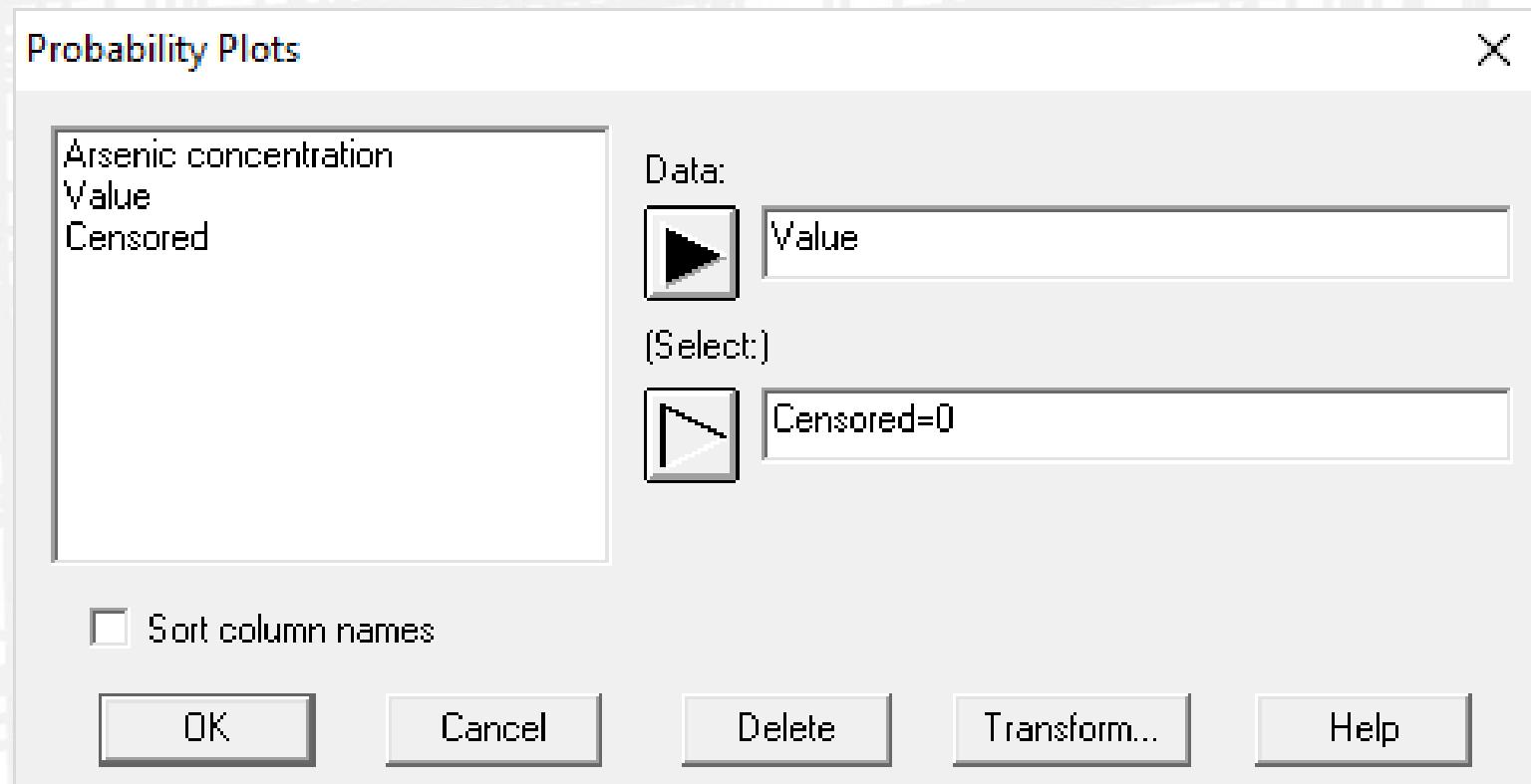
- Also handles interval-censored data.
- Does not include goodness-of-fit tests.

# Fitting the Arsenic Data

	Arsenic concentration	Value	Censored
	mu-g/L		
	Censored	Numeric	Numeric
1	.5	.5	0
2	.5	.5	0
3	.5	.5	0
4	.6	.6	0
5	.7	.7	0
6	.7	.7	0
7	<.9	.9	-1
8	.9	.9	0
9	<1	1	-1
10	<1	1	-1
11	<1	1	-1
12	<1	1	-1
13	1.5	1.5	0
14	1.7	1.7	0

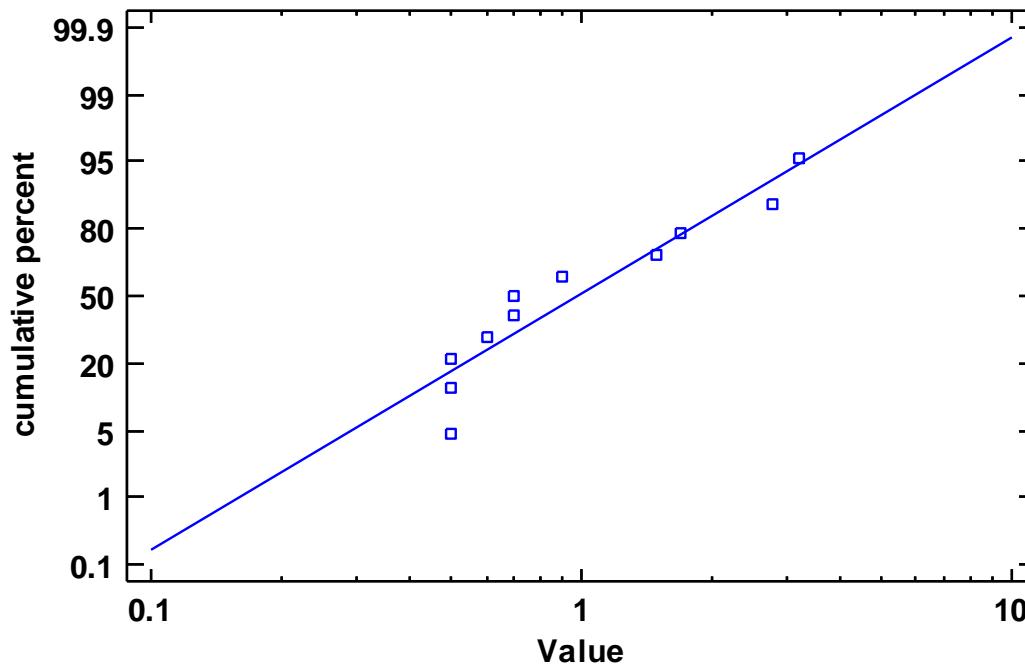
Note: may use *Edit – Replace Censored Values* to create the *Value* column.

# Probability Plots

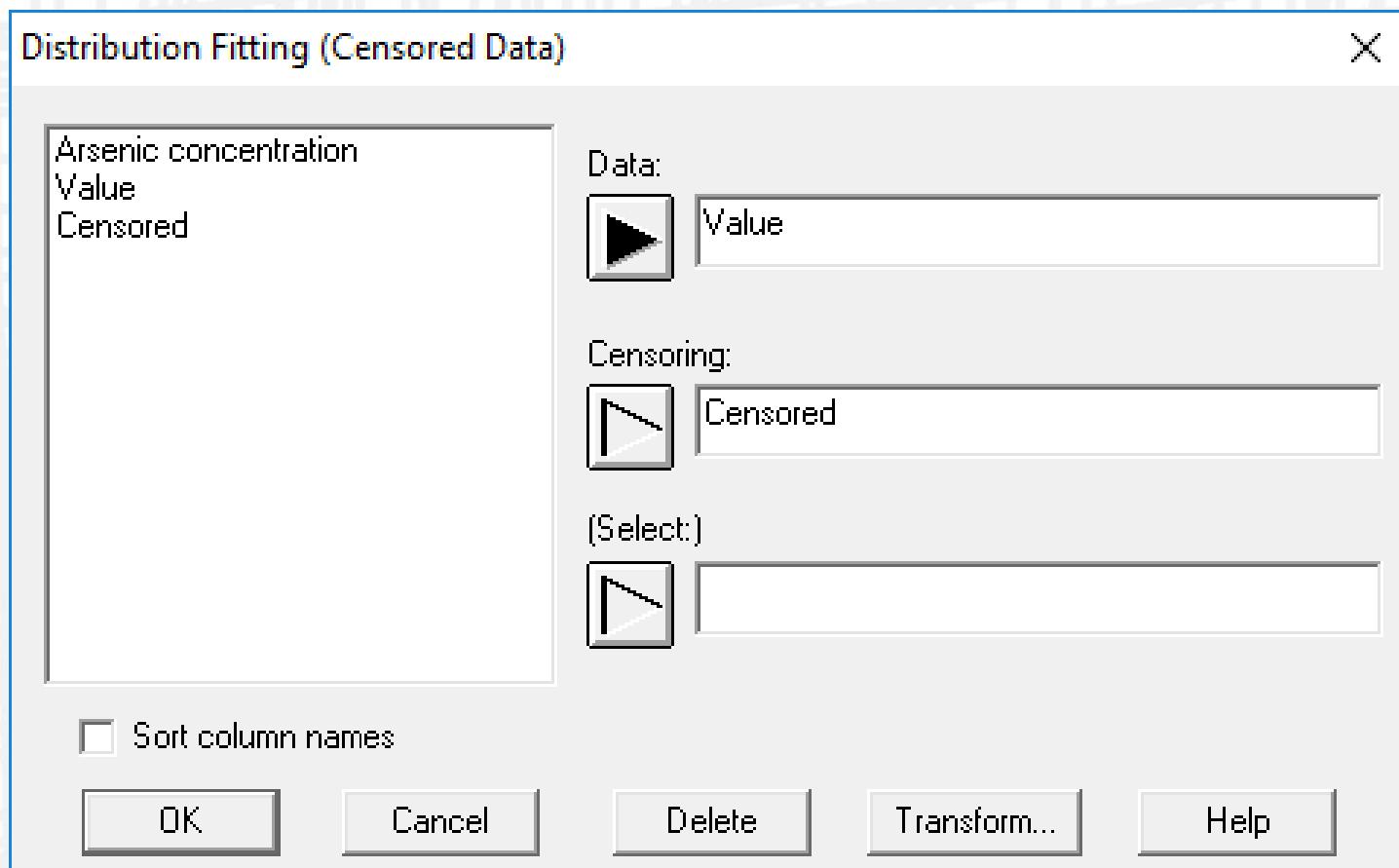


# Probability Plots

Lognormal Probability Plot



# Distribution Fitting (Censored Data)



# Analysis Options

Distribution Fitting Options X

**Distribution**

<input type="checkbox"/> Bernoulli	<input type="checkbox"/> Exponential Power	<input type="checkbox"/> Lognormal (3-parameter)
<input type="checkbox"/> Binomial	<input type="checkbox"/> F (Variance Ratio)	<input type="checkbox"/> Maxwell (2-parameter)
<input type="checkbox"/> Discrete Uniform	<input type="checkbox"/> Folded Normal	<input type="checkbox"/> Noncentral Chi-Square
<input type="checkbox"/> Geometric	<input type="checkbox"/> Gamma	<input type="checkbox"/> Noncentral F
<input type="checkbox"/> Hypergeometric	<input type="checkbox"/> Gamma (3-parameter)	<input type="checkbox"/> Noncentral t
<input type="checkbox"/> Negative Binomial	<input type="checkbox"/> Generalized Gamma	<input type="checkbox"/> Normal
<input type="checkbox"/> Poisson	<input type="checkbox"/> Generalized Logistic	<input type="checkbox"/> Pareto
<input type="checkbox"/> Beta	<input type="checkbox"/> Half Normal (2-parameter)	<input type="checkbox"/> Pareto (2-parameter)
<input type="checkbox"/> Beta (4-parameter)	<input type="checkbox"/> Inverse Gaussian	<input type="checkbox"/> Rayleigh (2-parameter)
<input type="checkbox"/> Birnbaum-Saunders	<input type="checkbox"/> Laplace	<input type="checkbox"/> Smallest Extreme Value
<input type="checkbox"/> Cauchy	<input type="checkbox"/> Largest Extreme Value	<input type="checkbox"/> Student's t
<input type="checkbox"/> Chi-Square	<input type="checkbox"/> Logistic	<input type="checkbox"/> Triangular
<input type="checkbox"/> Erlang	<input type="checkbox"/> Loglogistic	<input type="checkbox"/> Uniform
<input type="checkbox"/> Exponential	<input type="checkbox"/> Loglogistic (3-parameter)	<input type="checkbox"/> Weibull
<input type="checkbox"/> Exponential (2-parameter)	<input checked="" type="checkbox"/> Lognormal	<input type="checkbox"/> Weibull (3-parameter)

**Binomial Trials**  
Sample Size n:

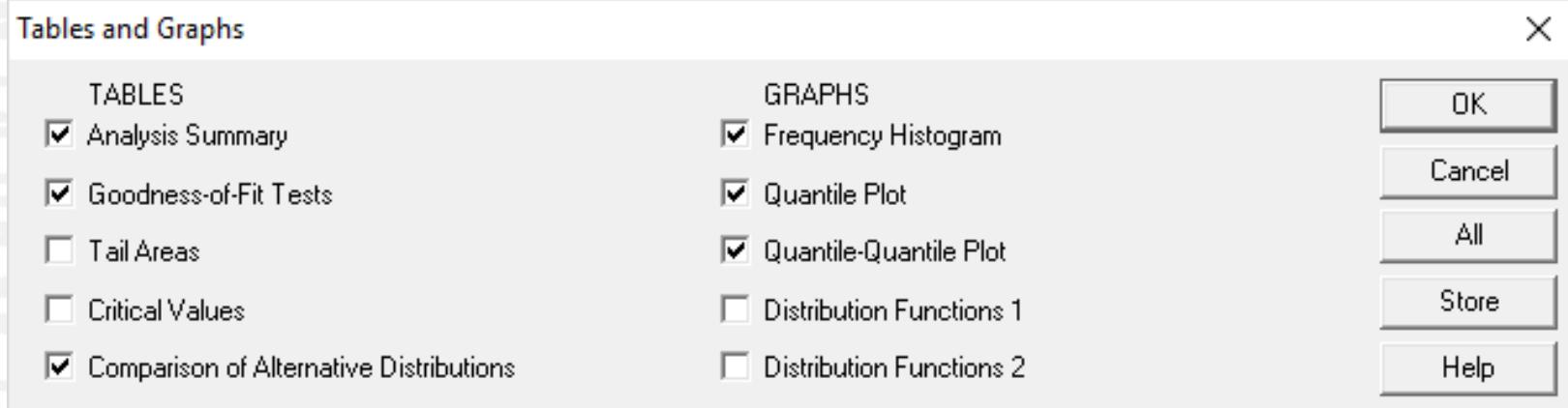
**Hypergeometric Trials**  
Sample Size n:  
  
 Estimate N  
 Specify N

**Negative Binomial Trials**  
 Estimate k  
 Specify k

**Extended Threshold Parameters**  
 Estimate  
 Specify lower/upper

**OK** **Cancel** **Help**

# Tables and Graphs



# Analysis Summary

## Distribution Fitting (Censored Data) - Value

Data variable: Value

Censoring: Censored

24 values ranging from 0.5 to 3.2

Number of left-censored observations: 13

Number of right-censored observations: 0

### Fitted Distributions

Lognormal

mean = 0.94526

standard deviation = 0.655927

Log scale: mean = -0.252828

Log scale: std. dev. = 0.626949

# Goodness-of-Fit Tests

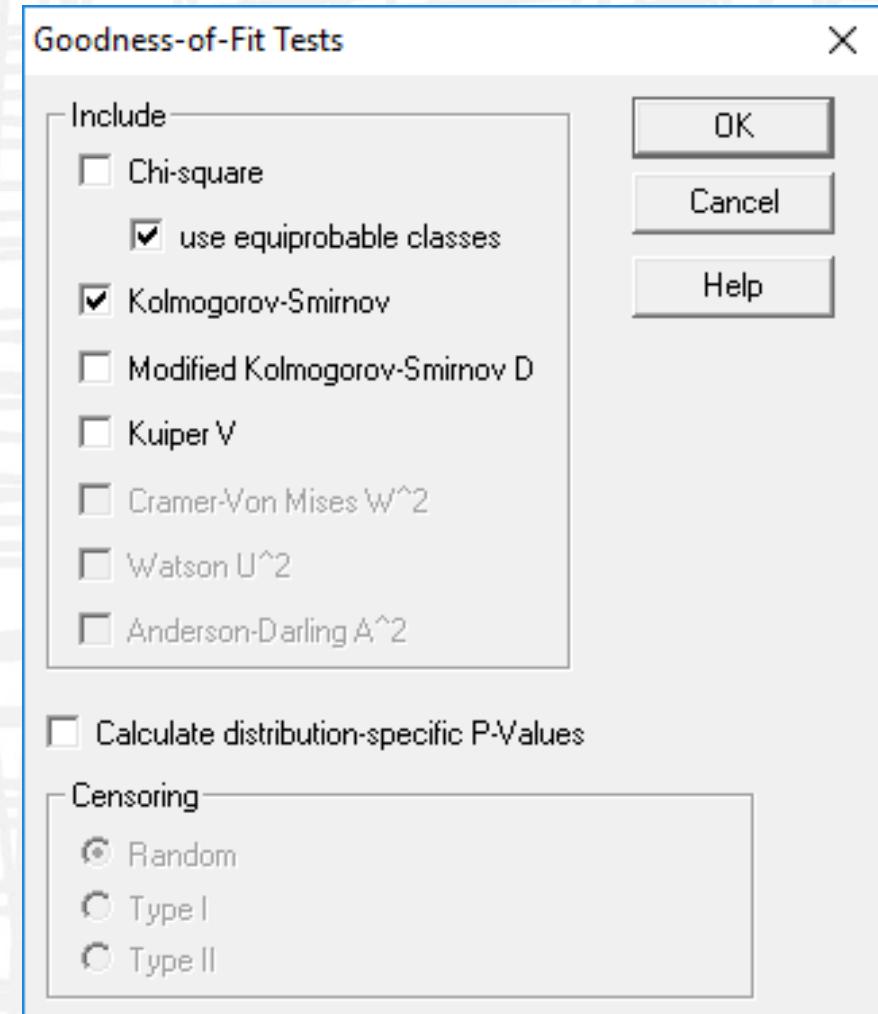
## Goodness-of-Fit Tests for Value

### Kolmogorov-Smirnov Test

	<i>Lognormal</i>
DPLUS	0.253282
DMINUS	0.168321
DN	0.253282
P-Value	0.091989

Type I censoring – items removed after prespecified times.

Type II censoring – test stopped after prespecified number of failures.



# Comparison of Alternative Distributions

Comparison of Alternative Distributions

Distribution	Est. Parameters	KS D
Loglogistic	2	0.219342
Lognormal	2	0.253282
Weibull	2	0.255712
Gamma	2	0.257573
Inverse Gaussian	2	0.260837
Birnbaum-Saunders	2	0.268475
Laplace	2	0.276909
Logistic	2	0.283439
Largest Extreme Value	2	0.285595
Normal	2	0.287314
Exponential	1	0.366133
Smallest Extreme Value	2	0.374146
Uniform	2	0.409164
Pareto	<no fit>	

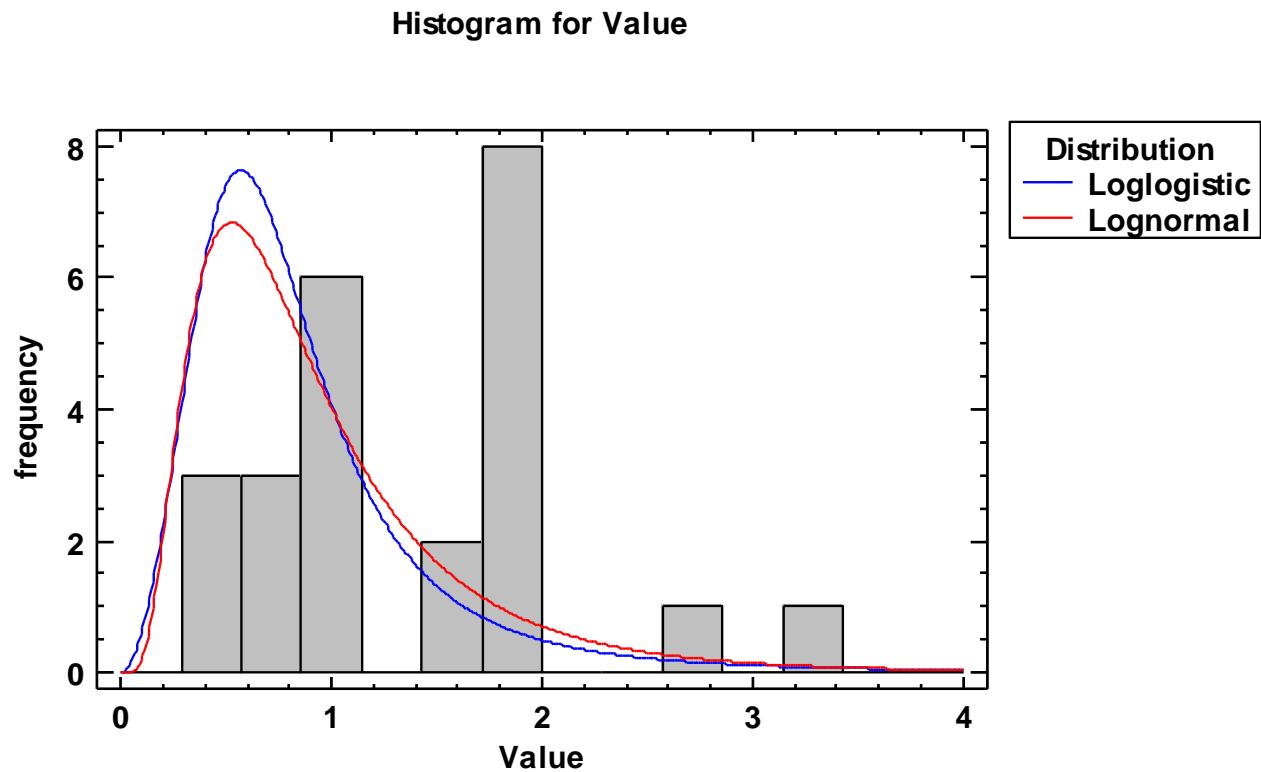
# Goodness-of-Fit Tests

## Goodness-of-Fit Tests for Value

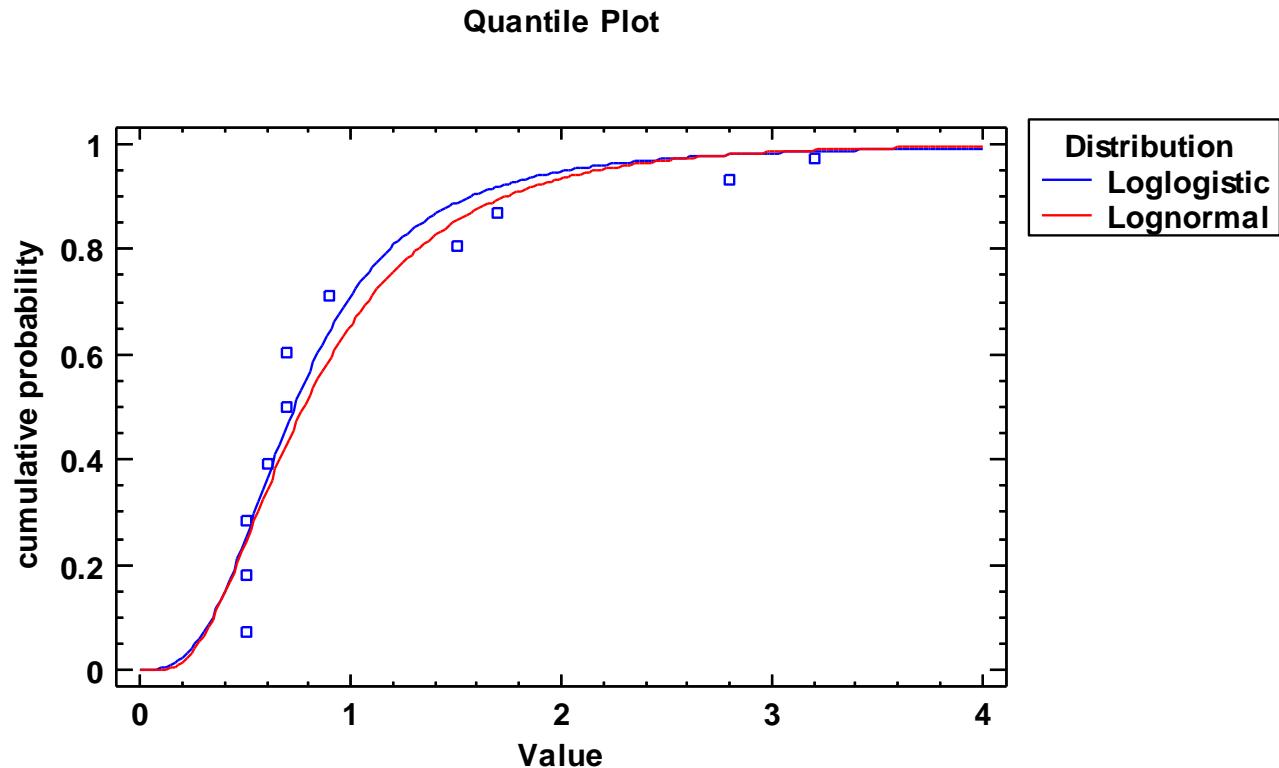
### Kolmogorov-Smirnov Test

	<i>Loglogistic</i>	<i>Lognormal</i>
DPLUS	0.219342	0.253282
DMINUS	0.177452	0.168321
DN	0.219342	0.253282
P-Value	0.198755	0.091989

# Histogram with Fits



# Quantile Plot

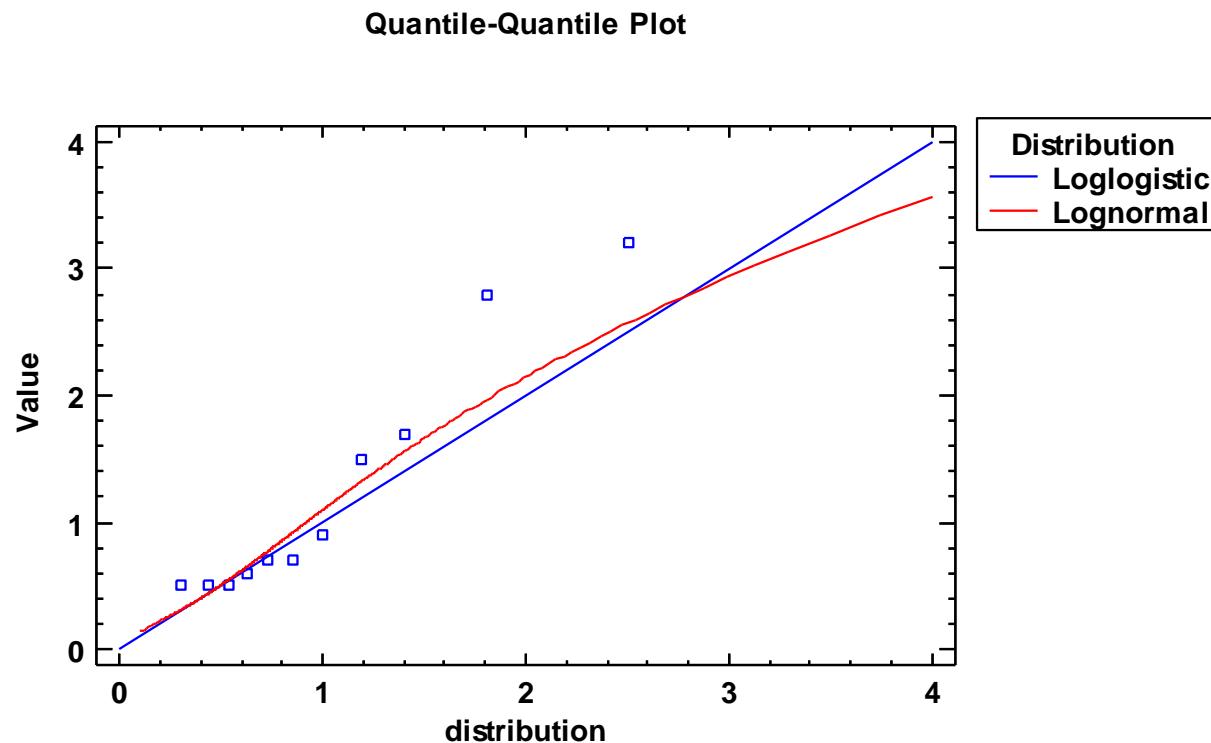


# Critical Values

## Critical Values for Value

<i>Lower Tail Area (<math>\leq</math>)</i>	<i>Loglogistic</i>	<i>Lognormal</i>
0.01	0.148405	0.180625
0.1	0.341207	0.347741
0.5	0.731694	0.776602
0.9	1.56907	1.73436
0.99	3.60754	3.33901

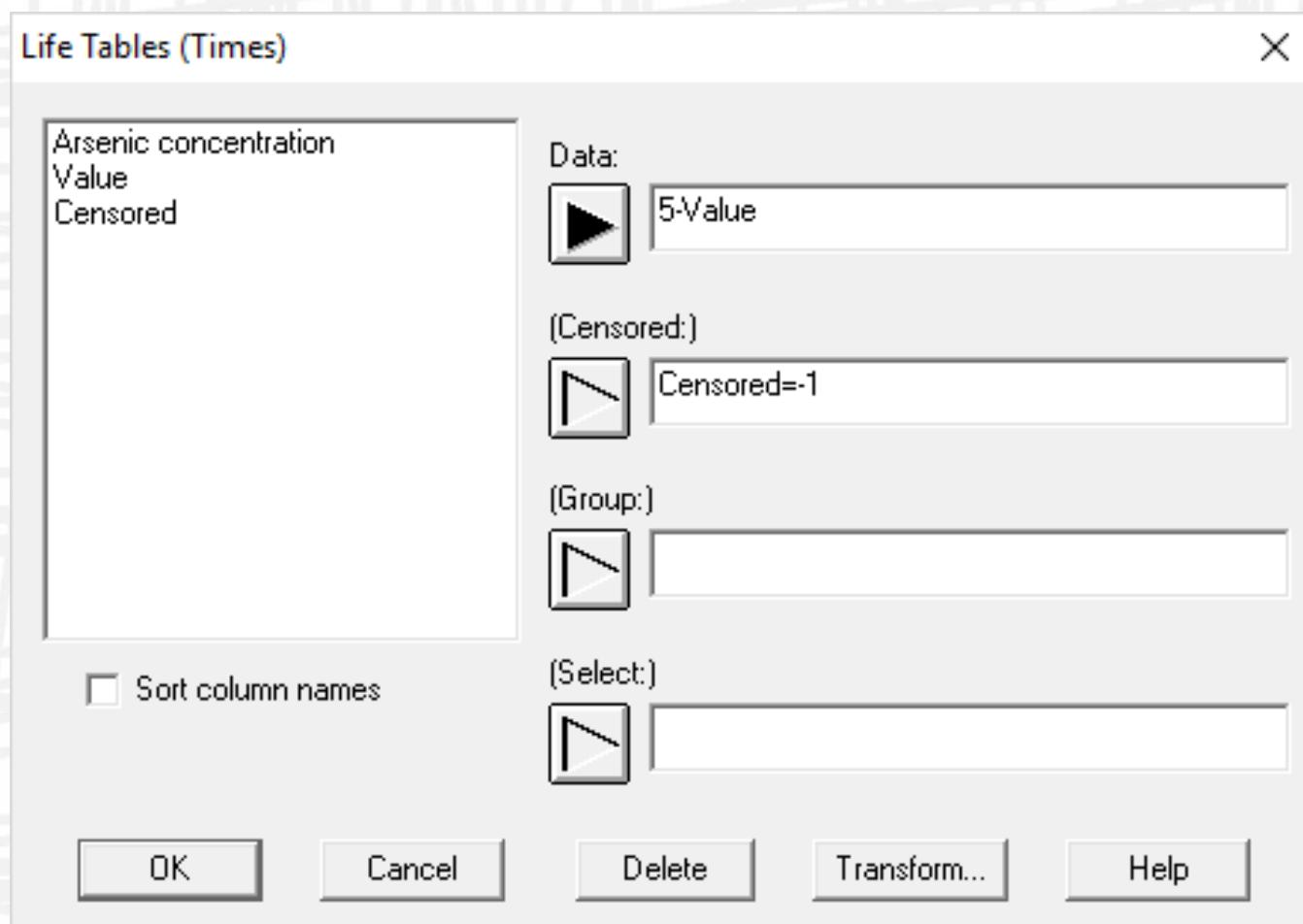
# Quantile-Quantile Plot



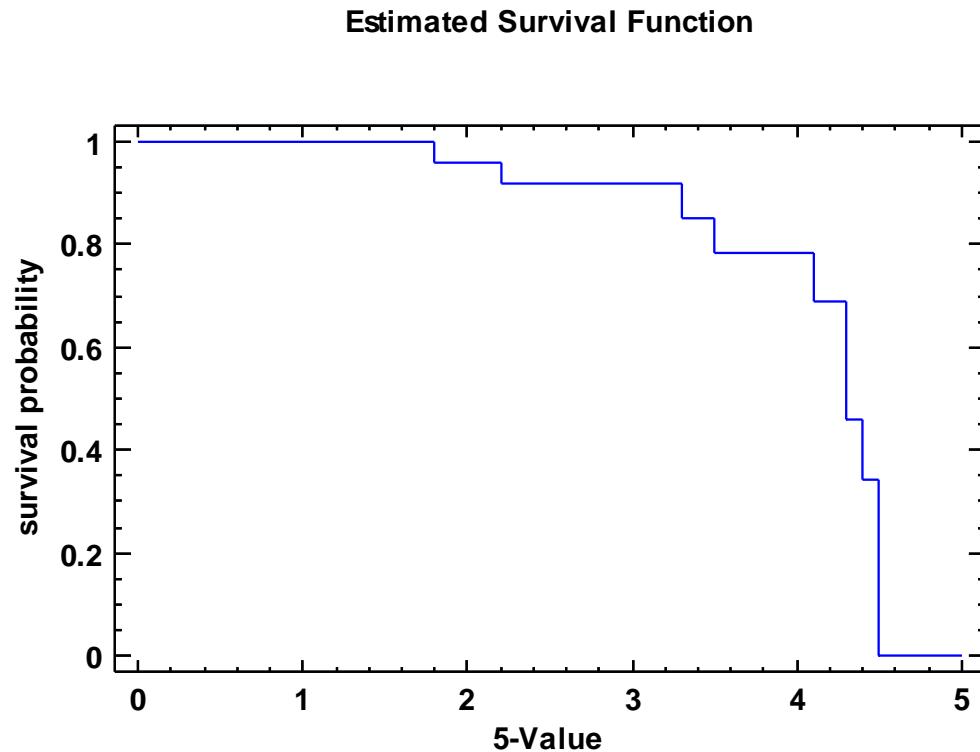
# Nonparametric Estimates

- Statgraphics contains a procedure under *Describe – Life Data – Life Tables (Times)* which will estimate a nonparametric survival function for right-censored data.
- Helsel (2012) shows how such a procedure may be used to analyze left-censored data by “flipping” the data values.

# Data Input Dialog Box



# Kaplan-Meier Estimate



# Results

- Subtract each result from 5 to get what you want.

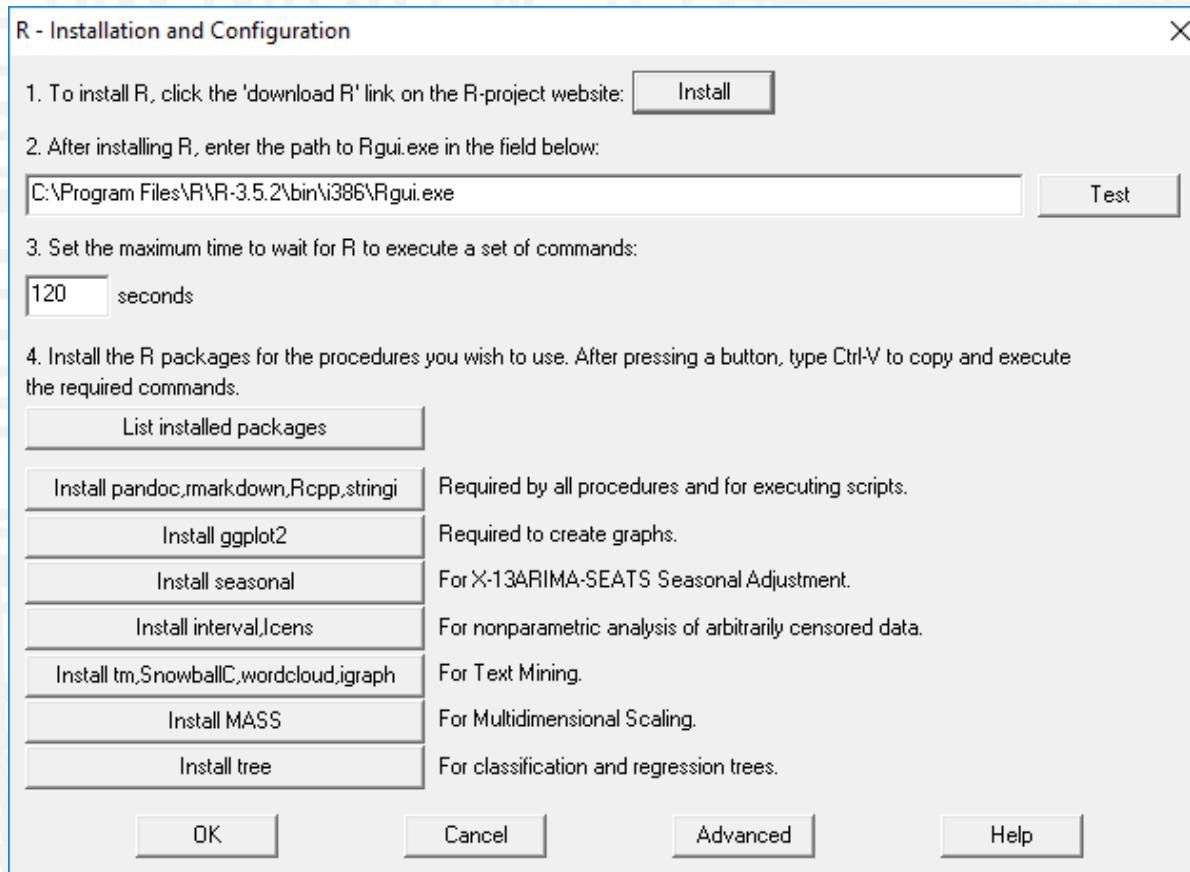
Percentile Table

		Standard
Percentile	Estimate	Error
75.0	4.1	0.615438
50.0	4.3	0.136753
25.0	4.5	

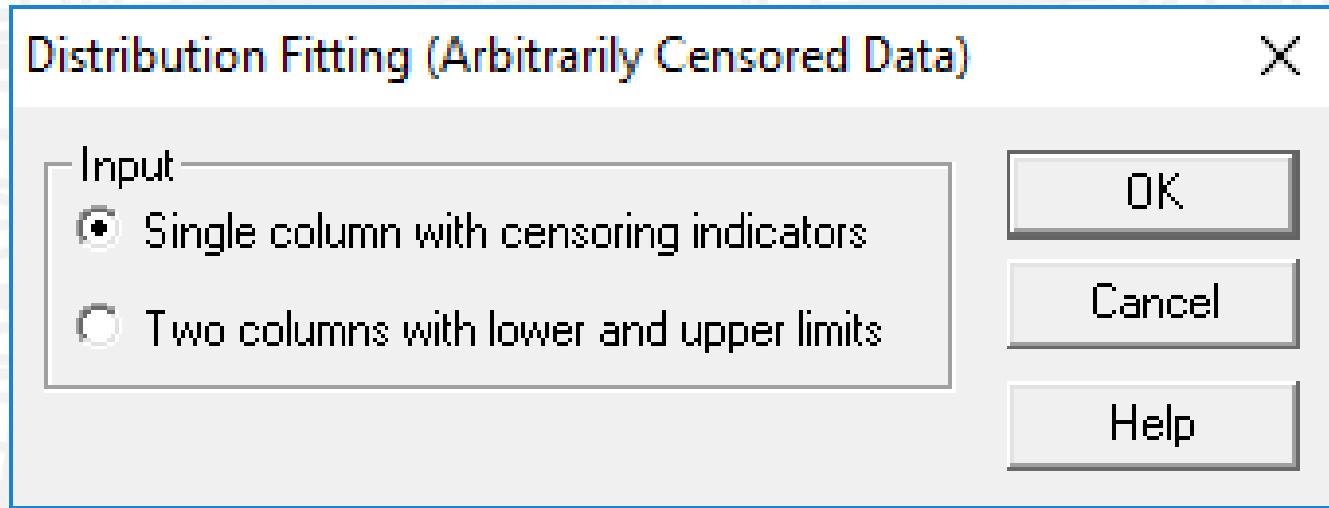
- $Q1 = 0.5$ , Median = 0.7,  $Q3 = 0.9$

# Fitting the Breast Cancer Data

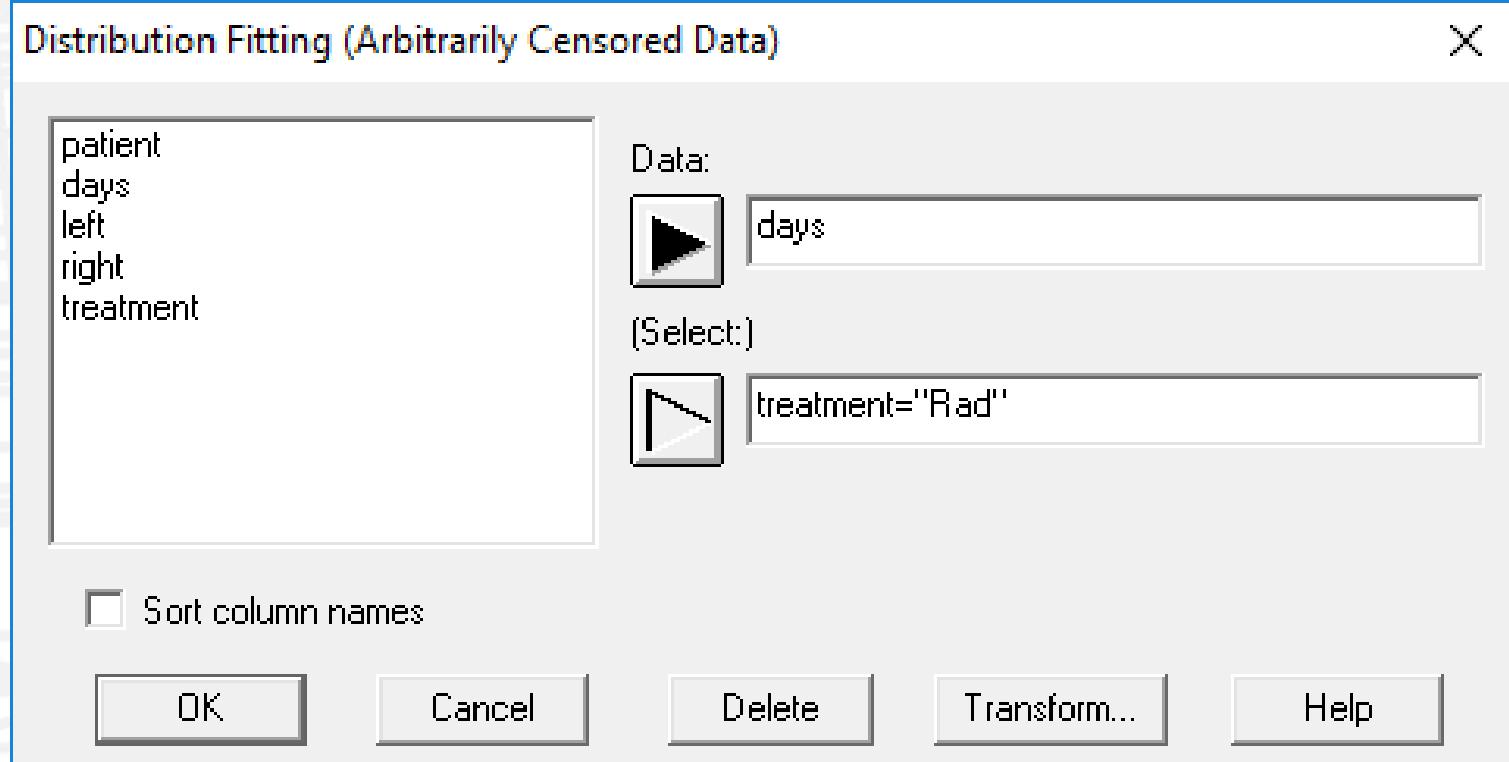
- Uses the “interval” and “icens” packages written for R.



# Data Input Dialog Boxes



# Data Input Dialog Boxes



# Analysis Options

Distribution Fitting (Arbitrarily Censored Data) Options X

Assumed Distribution

Birnbaum-Saunders     Generalized Logistic     Lognormal (3-parameter)  
 Cauchy     Half Normal (2-parameter)     Maxwell (2-parameter)  
 Exponential     Inverse Gaussian     Normal  
 Exponential (2-parameter)     Laplace     Pareto  
 Exponential Power     Largest Extreme Value     Pareto (2-parameter)  
 Folded Normal     Logistic     Rayleigh (2-parameter)  
 Gamma     Loglogistic     Smallest Extreme Value  
 Gamma (3-parameter)     Loglogistic (3-parameter)     Weibull  
 Generalized Gamma     Lognormal     Weibull (3-parameter)

Lower threshold:   Apply Efron bias correction

Confidence level for intervals:  % Number of bootstrap subsamples:

# Options

- **Assumed distribution** – will be fit to the data.
- **Lower threshold** – for distributions such as lognormal, the minimum possible value.
- **Apply Efron bias correction** – if smallest value is left-censored, sets KMT nonparametric CDF at that observation to 0 for purposes of calculating the mean and standard deviation. Otherwise, CDF is assumed to decay linearly to 0 at lower threshold.
- **Confidence level for intervals:** confidence level used to create confidence limits for distribution parameters and other quantities.
- **Number of bootstrap subsamples:** number of subsamples to be used when estimating confidence limits for the distribution parameters and other quantities.

# Analysis Summary

## Distribution Fitting (Arbitrarily Censored Data) (treatment="Rad")

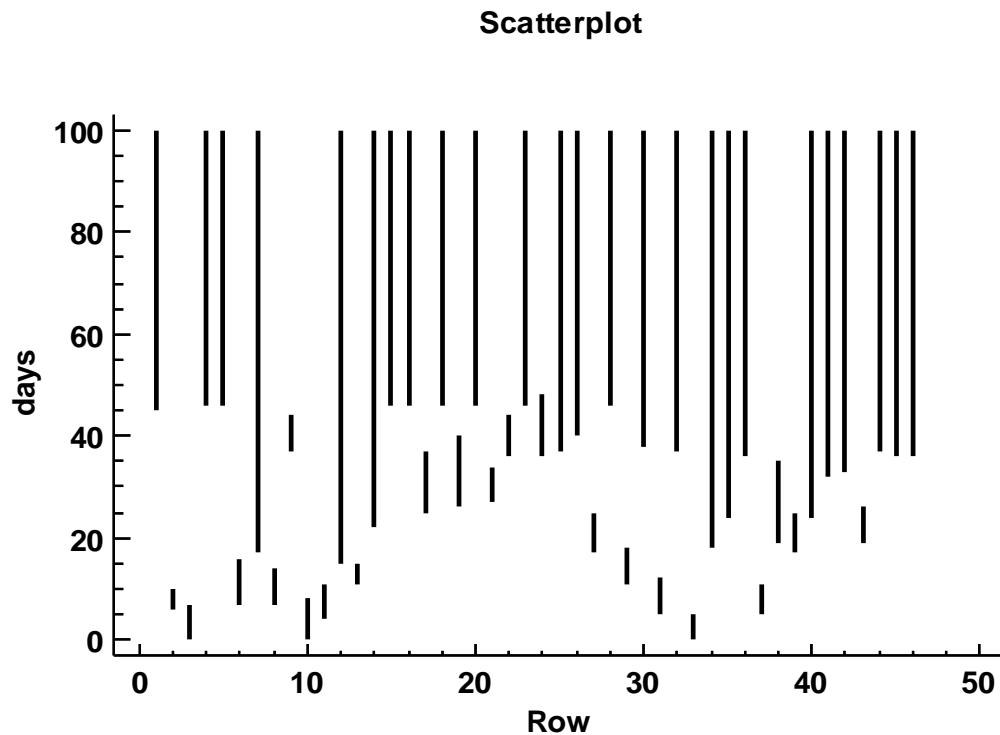
Data variable: days

Selection variable: treatment="Rad"

### Observations

Value	Frequency
Uncensored	0
Left-censored	3
Interval-censored	18
Right-censored	25

# Plotting the Data



# Distribution Fitting

Parameter estimates are obtained by maximizing the likelihood function:

$$L = \prod_{i=1}^n l(x_i)$$

where

$l(x_i) = f(x_i)$  if the observation  $x_i$  is uncensored

$l(x_i) = F(L_i)$  if the observation is left-censored at  $L_i$

$l(x_i) = 1 - F(U_i)$  if the observation is right-censored at  $U_i$

$l(x_i) = F(U_i) - F(L_i)$  if the observation is interval-censored between  $[L_i, U_i]$

# Example

## Distribution Fitting

Fitted distribution: Lognormal

Parameter	Estimate	95% LCL	95% UCL
Mean	100.514	50.0203	447.337
Std. Dev.	214.7	63.2189	2986.17

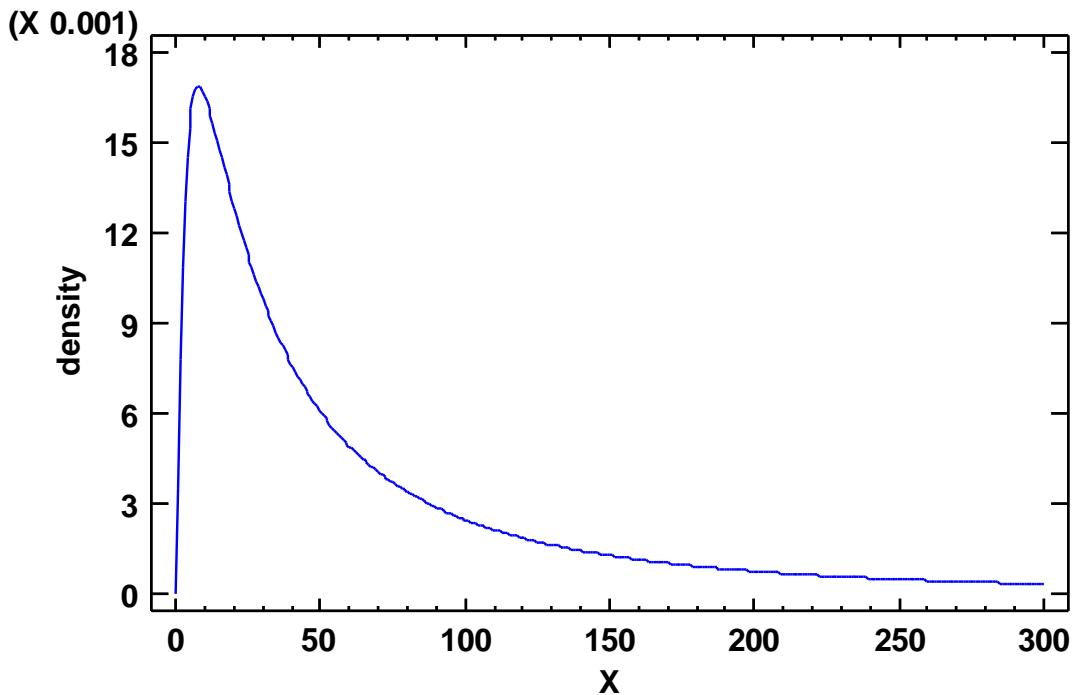
## Distribution Properties

	Estimate	95% LCL	95% UCL
Mean	100.514	50.0203	447.337
Standard deviation	214.7	63.2189	2986.17
Median	42.6171	28.1219	77.6418
Lower quartile	17.6137	11.2882	27.5537
Upper quartile	103.114	59.1971	257.675
Interquartile range	85.5001	44.1705	238.614

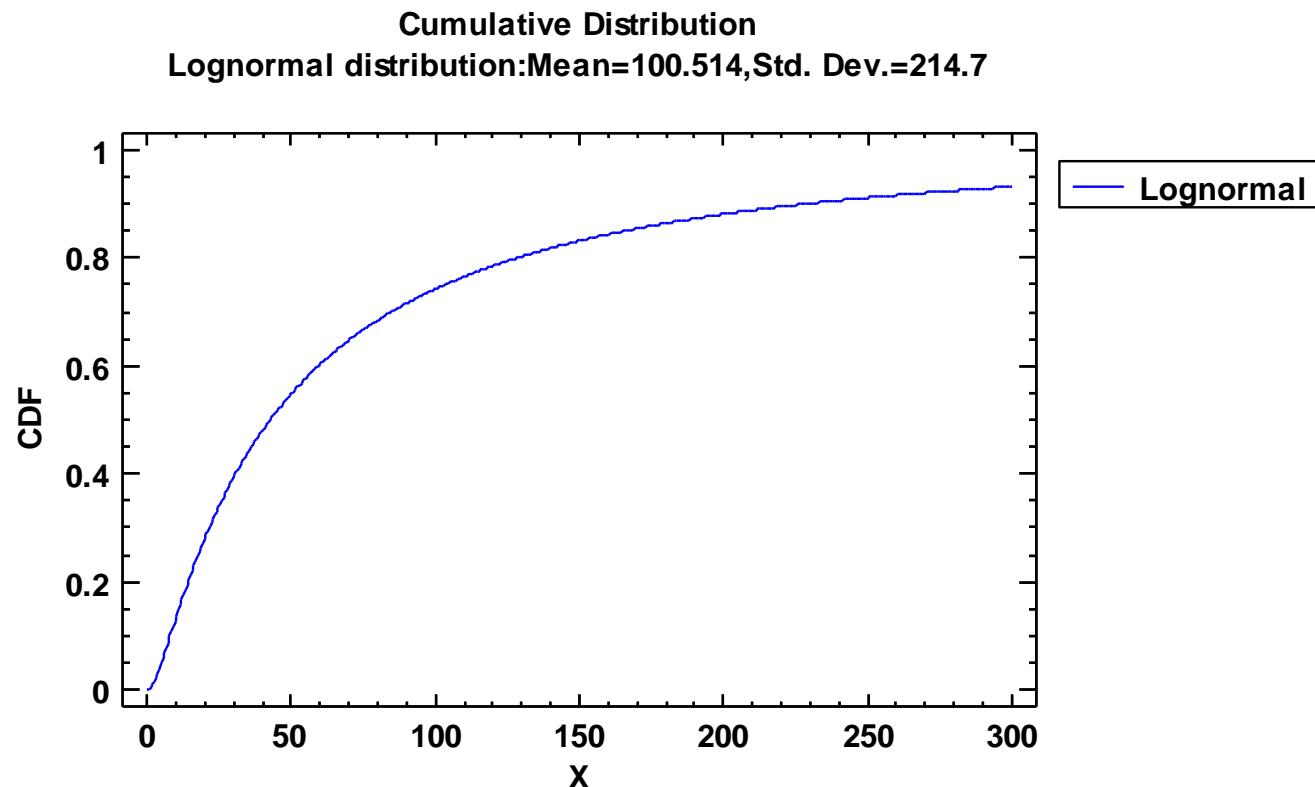
Number of bootstrap subsamples: 1000

# Plot of Fitted Distribution

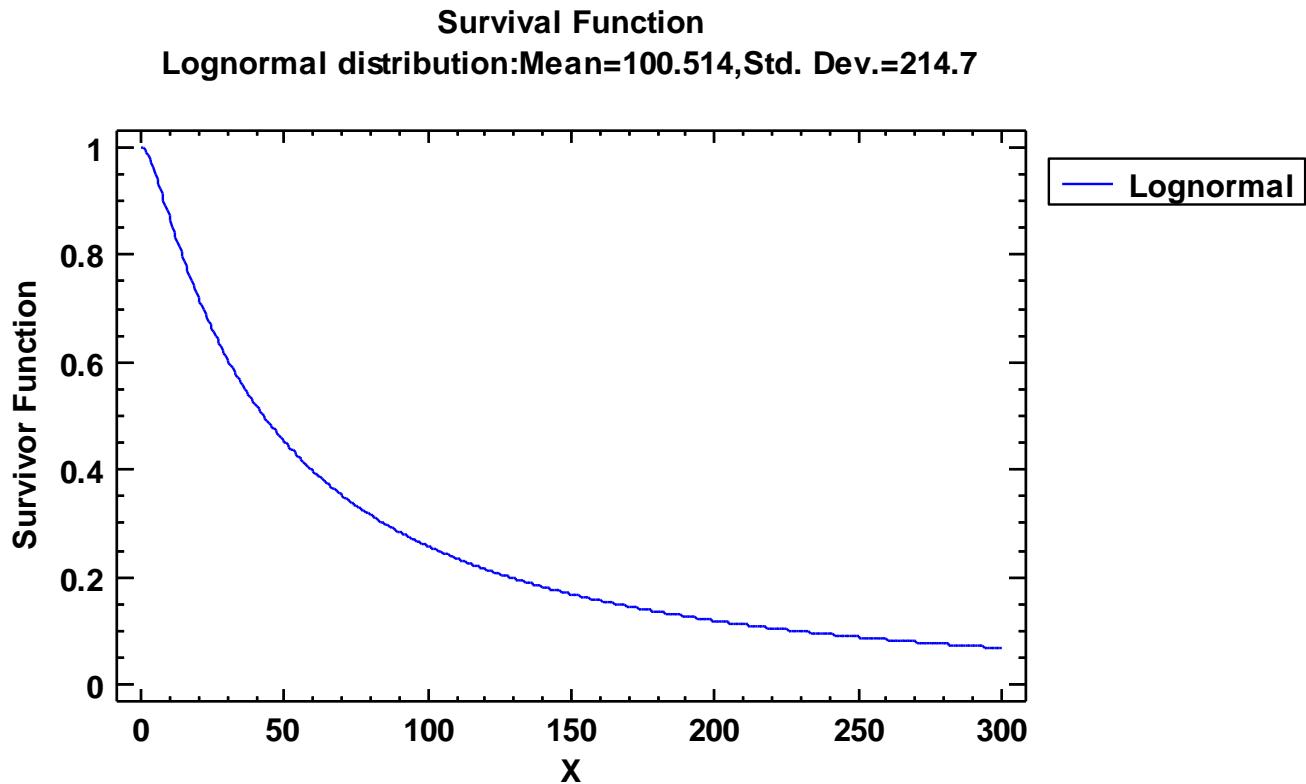
Lognormal distribution (Mean=100.514, Std. Dev.=214.7)



# Cumulative Distribution Function



# Survival Function



# Nonparametric Estimates

- A nonparametric estimate of the survival function may be obtained without assuming any particular distributional form.
- Estimated using the methods of Kaplan, Meier and Turnbull.
- Can also calculate confidence limits for the nonparametric survival function or CDF.

# Example

## Nonparametric Estimates

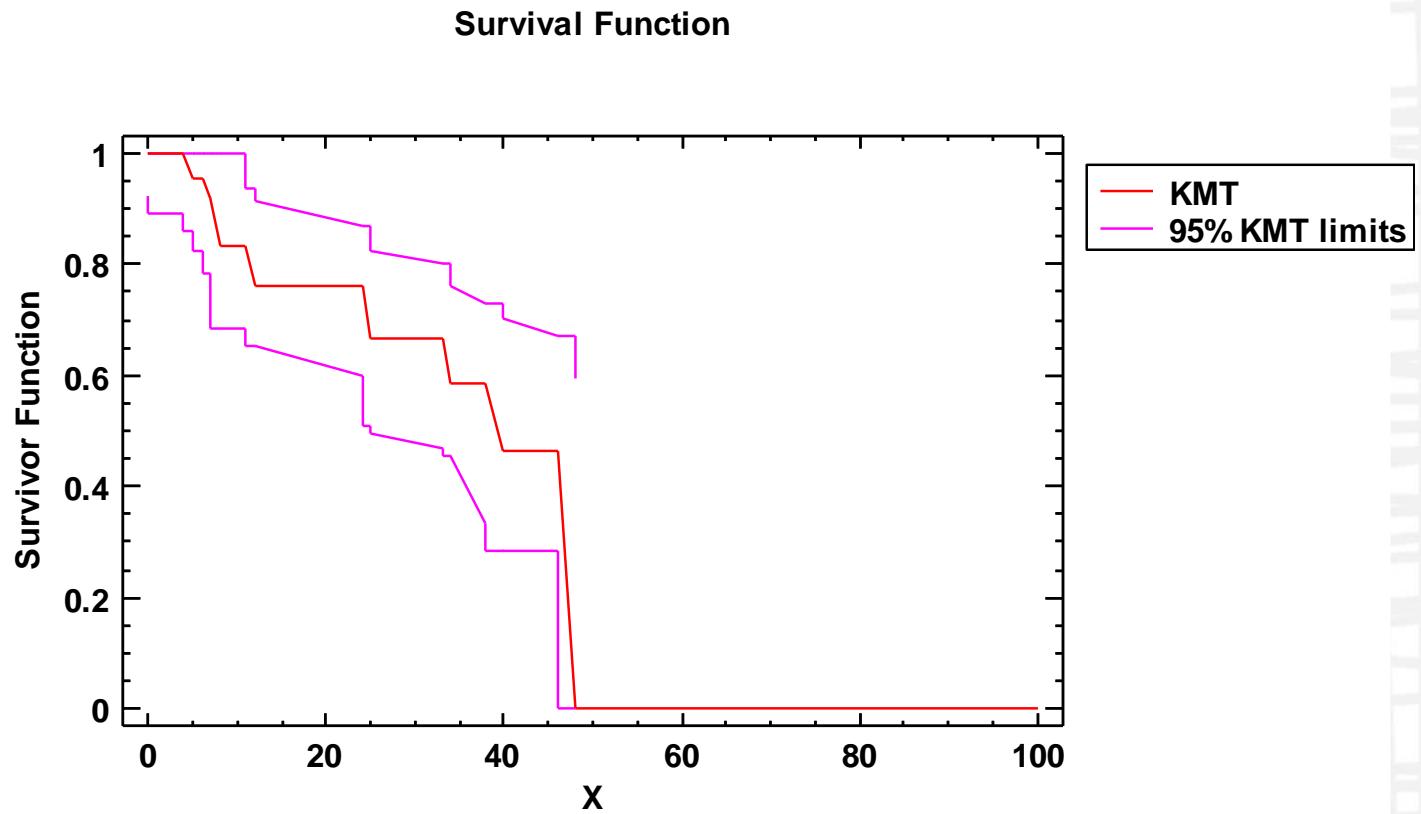
### Kaplan-Meier-Turnbull Estimates

<i>days</i>	<i>CDF</i>	<i>Survival</i>	<i>95% LCL</i>	<i>95% UCL</i>
4.0	0.0	1.0	0.891313	1.0
6.0	0.0463468	0.953653	0.826087	1.0
7.0	0.0797102	0.92029	0.782609	1.0
11.0	0.168378	0.831622	0.685386	1.0
24.0	0.23913	0.76087	0.599359	0.869565
33.0	0.331776	0.668224	0.470669	0.800272
38.0	0.413562	0.586438	0.332921	0.730274
46.0	0.534442	0.465558	0.283185	0.66923

### Statistics

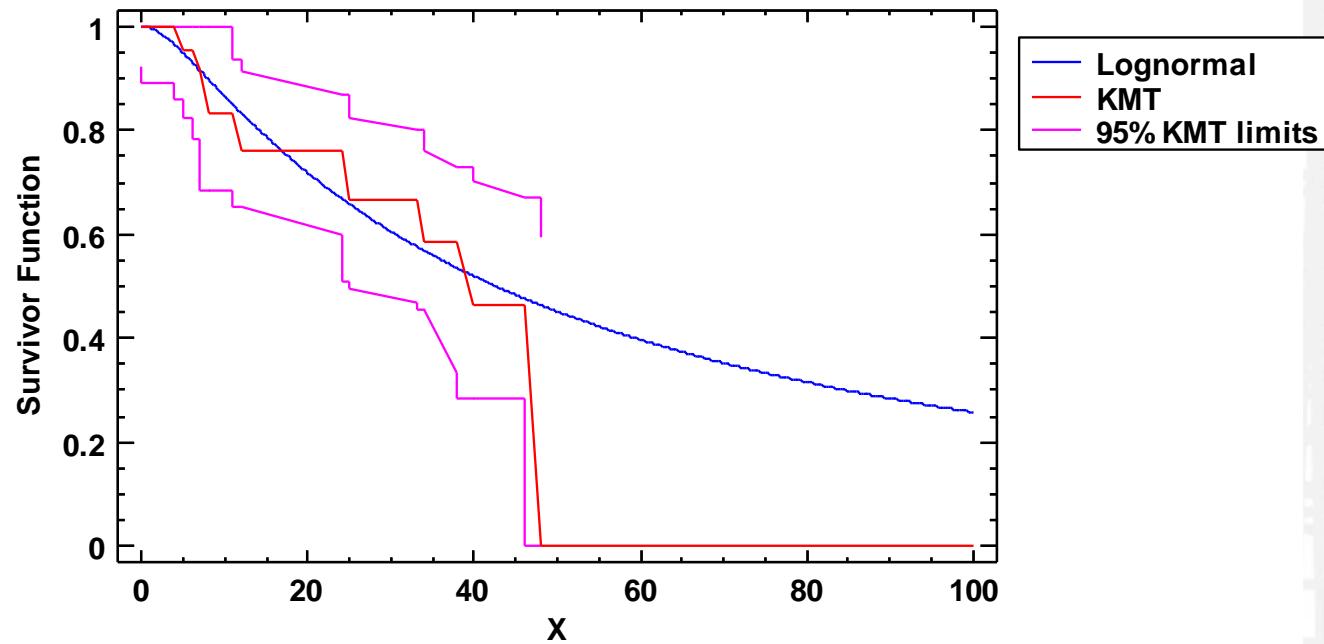
Mean	33.5093
Standard deviation	15.9287
Standard error	2.34855

# Example



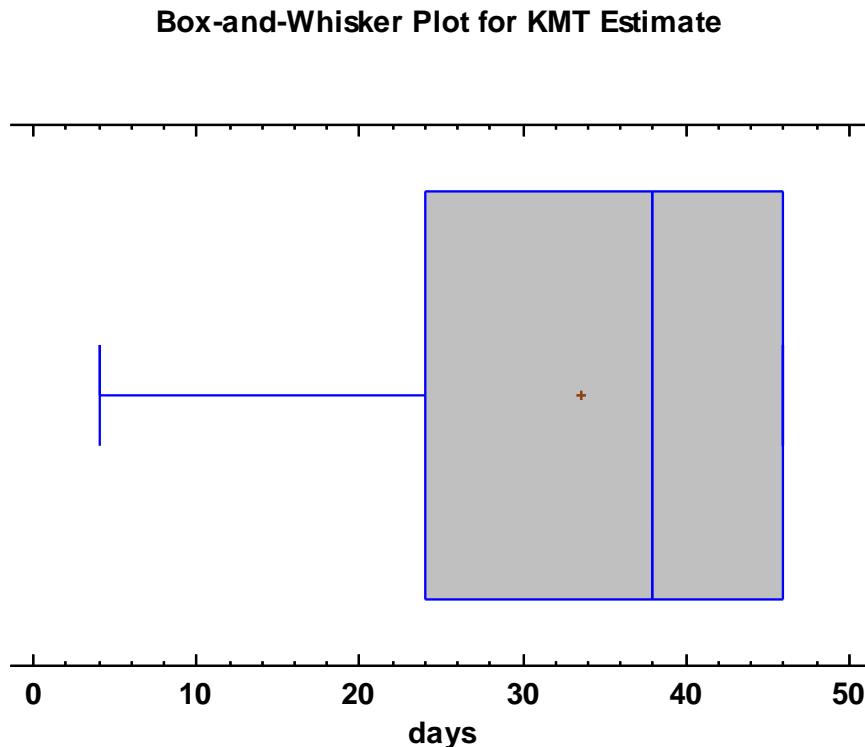
# Example

Survival Function  
Lognormal distribution: Mean=100.514, Std. Dev.=214.7

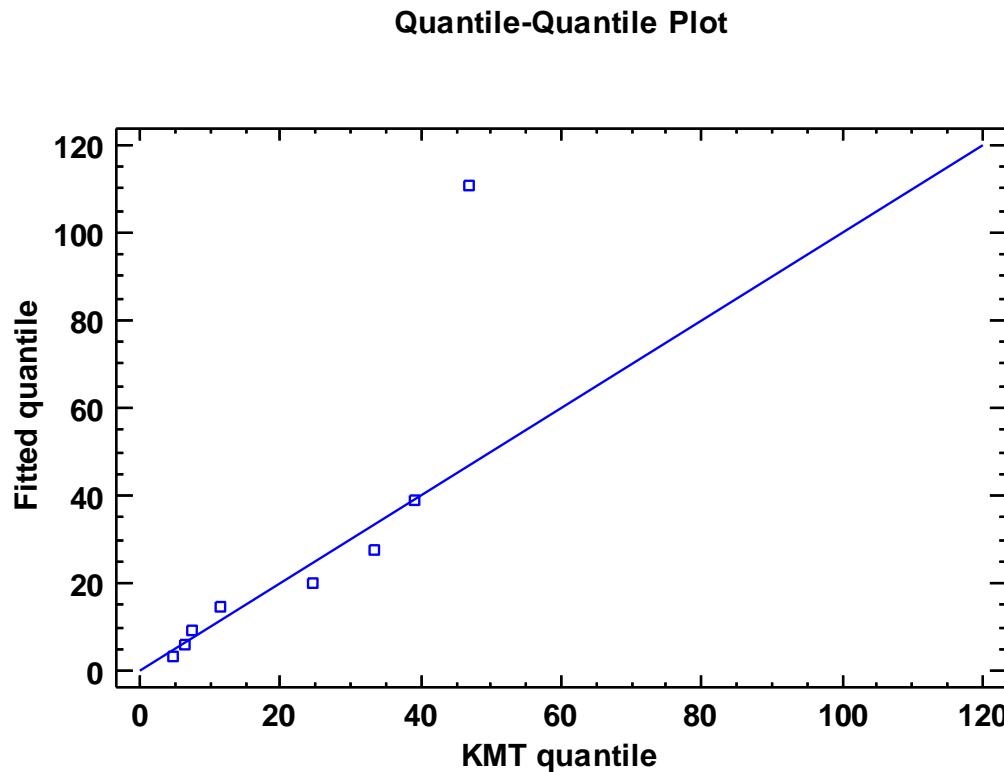


# Box-and-Whisker Plot

- Based on 1<sup>st</sup>, 25<sup>th</sup>, 50<sup>th</sup>, 75<sup>th</sup> and 99<sup>th</sup> percentiles.



# Quantile-Quantile Plot



# References

- StatFolios and data files are at: [www.statgraphics.com/webinars](http://www.statgraphics.com/webinars)
- Finkelstein, D.M. and Wolfe, R.A. (1985). “A semiparametric model for regression analysis of interval-censored failure time data.” *Biometrics* 41, 731-740.
- Gentleman R, Vandal A (2018). *Icens: NPMLE for Censored and Truncated Data*. R package version 1.54.0.
- Helsel, D.R. (2005). Nondetects and Data Analysis: Statistics for Censored Environmental Data. Wiley, New York.
- Helsel, D.R. (2012). Statistics for Censored Environmental Data using Minitab and R, second edition. Wiley, Hoboken, N.J.
- Lee, E.T. and Wang, J.W. (2003). Statistical Methods for Survival Data Analysis, 3<sup>rd</sup> edition. Wiley, New York.
- R Package “interval” - <https://cran.r-project.org/web/packages/interval/interval.pdf>
- Tomlinson, M. S. (2003). “Effects of ground-water/surface-water interactions and land use on water quality.” Written communication (draft USGS report).
- Turnbull BW (1976). “The Empirical Distribution Function with Arbitrarily Grouped, Censored and Truncated Data.” *Journal of the Royal Statistical Society. Series B*, 38(3), 290–295.