Test 2

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This Exam is Individual Work. No Collaboration is Allowed

1 (15 points).

Consider the probability paper given in Figure 1. Do the following:

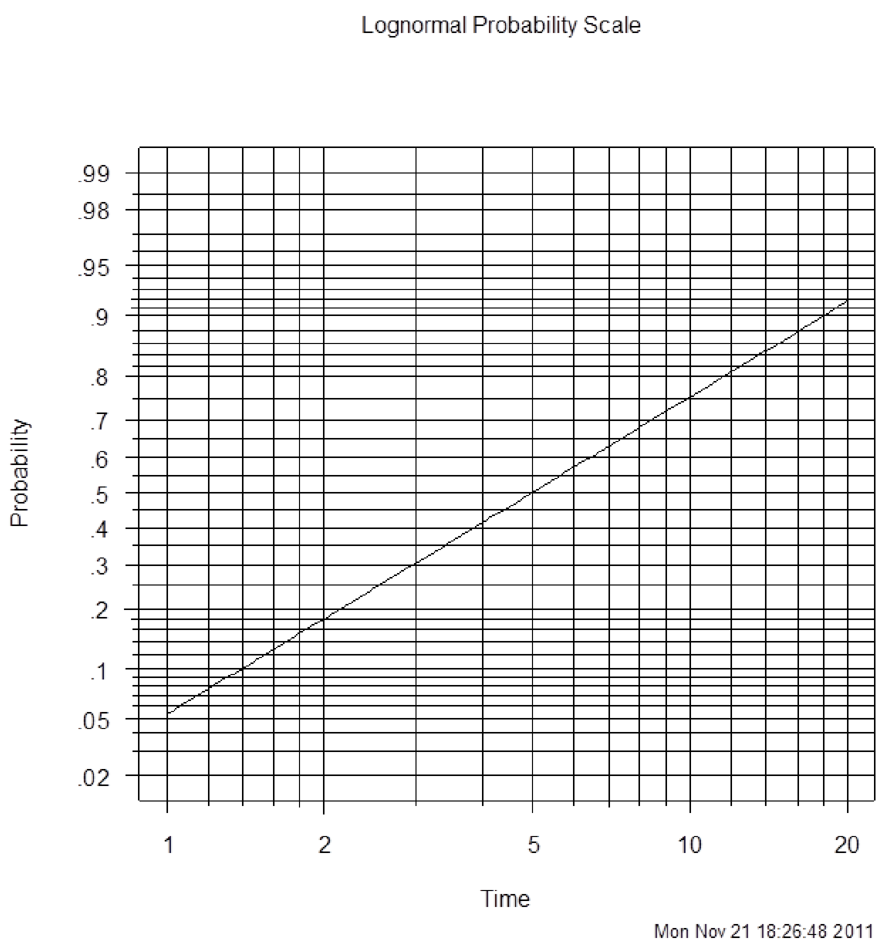

Figure 1: Lognormal Probability Plot

1.1

Plot on the paper the LOGNOR($exp(\mu) = 5, \sigma = 1$). Explain clearly the process to plot the line.

A probability plot linearizes t, some variable, against the CDF of that variable by transforming both variables such that the relationship between the two is linear. First we start with the quantile function of the CDF. In the case of the lognormal, this is $t_p = exp(\mu + \phi_{nor}^{-1}\sigma)$, where ϕ_{nor}^{-1} is the p quantile of the standard normal distribution. By taking the log of both sides we get $log(t_p) = \mu + \phi_{nor}^{-1}\sigma$. This relationship plots as a straight line. The slope of the line, given that the respective quantile function is the inverse of ϕ_{nor} is $1/\sigma$.

Therefore the steps use to generate the plot above are as follows:

a. Create a sequence of numbers between 0 and 20.

b. Log these numbers and calculate respective PDF outputs for the vector based on the normal distribution. Since the numbers have been logged this is the equivalent to the lognormal distribution.

c. Use the plotprob function in Rsplida, which creates probability paper by plotting each respective axis in log-scale (in this case).

d. Use the lines() function in Rsplida which plots t:time against y:quantiles of time.

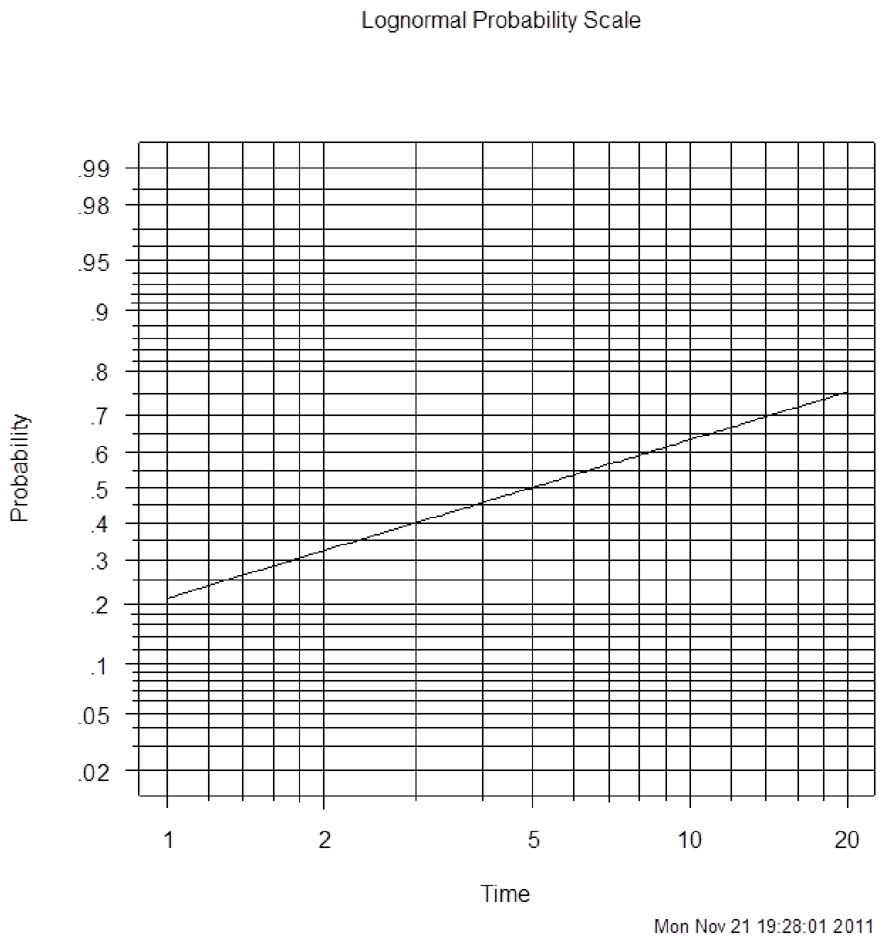

Figure 2: Lognormal Probability Plot

1.2

Plot on the paper the LOGNOR($exp(\mu) = 5, \sigma = 2$). Explain clearly the process to plot the line.

A probability plot linearizes t, some variable, against the CDF of that variable by transforming both variables such that the relationship between the two is linear. First we start with the quantile function of the CDF. In the case of the lognormal, this is $t_p = exp(\mu + \phi_{nor}^{-1}\sigma)$, where ϕ_{nor}^{-1} is the p quantile of the standard normal distribution. By taking the log of both sides we get $log(t_p) = \mu + \phi_{nor}^{-1}\sigma$. This relationship plots as a straight line. The slope of the line, given that the respective quantile function is the inverse of ϕ_{nor} is $1/\sigma$.

Therefore the steps use to generate the plot above are as follows:

a. Create a sequence of numbers between 0 and 20.

b. Log these numbers and calculate respective PDF outputs for the vector based on the normal distribution. Since the numbers have been logged this is the equivalent to the lognormal distribution.

c. Use the plotprob function in Rsplida, which creates probability paper by plotting each respective axis in log-scale (in this case).

d. Use the lines() function in Rsplida which plots t:time against y:quantiles of time.

2 (15 points).

Consider the probability paper given in Figure 1. Do the following:

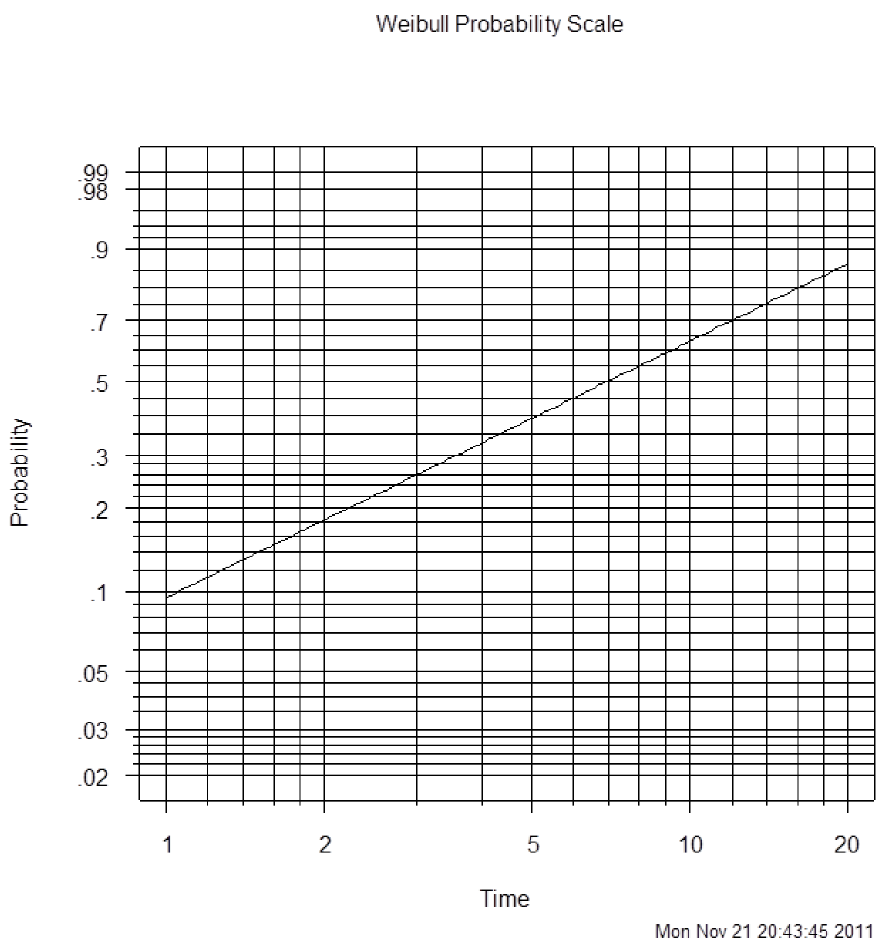

Figure 3: Weibull Probability Plot

2.1

Plot on the paper the WEIB($exp(\eta = 10, \beta = 2)$. Explain clearly the process to plot the line.

A probability plot linearizes t, some variable, against the CDF of that variable by transforming both variables such that the relationship between the two is linear. First we start with the quantile function of the CDF. In the case of the Weibull, this is $t_p = exp(\mu + \phi_{sev}^{-1}\sigma)$, where ϕ_{sev}^{-1} is the p quantile of the standard smallest extreme value distribution. By taking the log of both sides we get $log(t_p) = \mu + \phi_{sev}^{-1}\sigma$. This relationship plots as a straight line. The slope of the line, given that the respective quantile function is the inverse of ϕ_{sev} is $1/\sigma$.

Therefore the steps use to generate the plot above are as follows:

a. Create a sequence of numbers between 0 and 20.

b. Log these numbers and calculate respective PDF outputs for the vector based on the smallest extreme value distribution. Since the numbers have been logged this is the equivalent to the Weibull distribution.

c. Use the plotprob function in Rsplida, which creates probability paper by plotting each respective axis in log-scale (in this case).

d. Use the lines() function in Rsplida which plots t:time against y:quantiles of time.

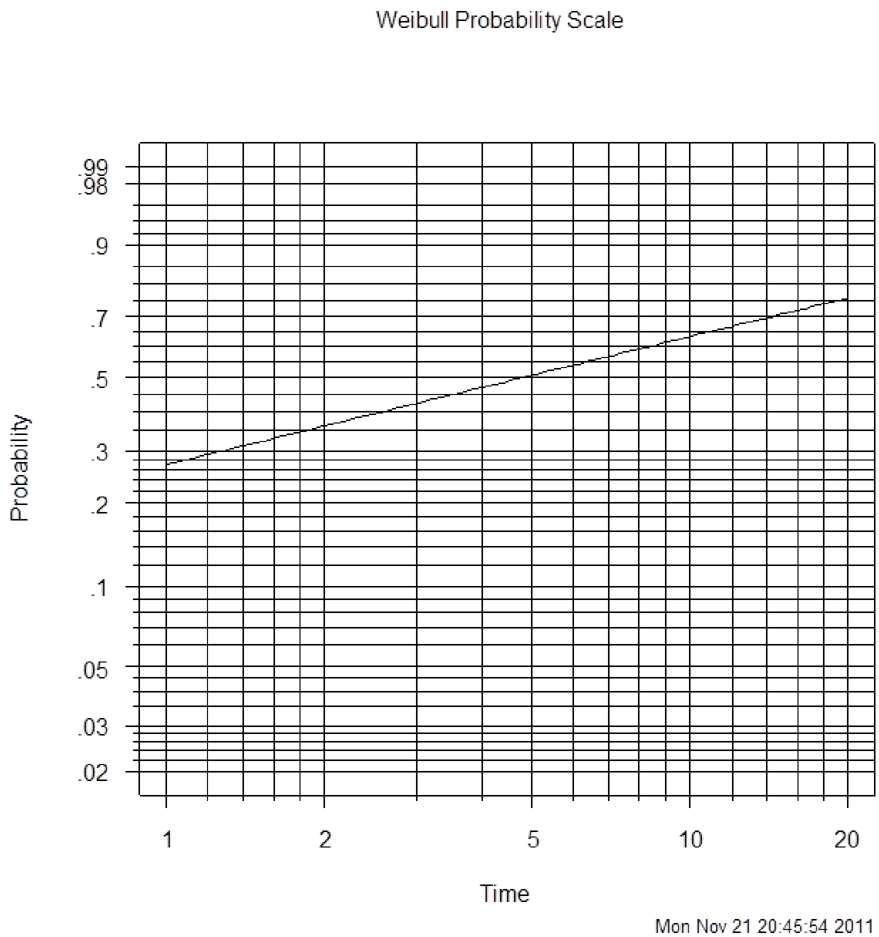

Figure 4: Weibull Probability Plot

2.2

Plot on the paper the WEIB($exp(\eta = 10, \beta = 2)$. Explain clearly the process to plot the line.

A probability plot linearizes t, some variable, against the CDF of that variable by transforming both variables such that the relationship between the two is linear. First we start with the quantile function of the CDF. In the case of the Weibull, this is $t_p = exp(\mu + \phi_{sev}^{-1}\sigma)$, where ϕ_{sev}^{-1} is the p quantile of the standard smallest extreme value distribution. By taking the log of both sides we get $log(t_p) = \mu + \phi_{sev}^{-1}\sigma$. This relationship plots as a straight line. The slope of the line, given that the respective quantile function is the inverse of ϕ_{sev} is $1/\sigma$.

Therefore the steps use to generate the plot above are as follows:

a. Create a sequence of numbers between 0 and 20.

b. Log these numbers and calculate respective PDF outputs for the vector based on the smallest extreme value distribution. Since the numbers have been logged this is the equivalent to the Weibull distribution.

c. Use the plotprob function in Rsplida, which creates probability paper by plotting each respective axis in log-scale (in this case).

d. Use the lines() function in Rsplida which plots t:time against y:quantiles of time.

3 (20 points).

Consider a data set with two observations.

Time	Status
1	Fail
2	Censored

(a) Write the likelihood of the data for a lognormal model.
The likelihood is as follows:

$$(L(\mu, \sigma) = \prod_{i=1}^{2} \left\{ \Phi_{nor} \left[\frac{log(t_i) - \mu}{\sigma} \right] \right\}^{\delta_i} \left\{ 1 - \Phi_{nor} \left[\frac{log(t_i) - \mu}{\sigma} \right] \right\}^{1-\delta_i}$$

where δ_i takes the value of 1 where an obsevation is left-censored, and 0 where an observation is right-censored.

(b) Write the likelihood of the data for a Weibull model.

The likelihood is as follows:

$$(L(\mu, \sigma) = \prod_{i=1}^{2} \left\{ \Phi_{sev} \left[\frac{log(t_i) - \mu}{\sigma} \right] \right\}^{\delta_i} \left\{ 1 - \Phi_{sev} \left[\frac{log(t_i) - \mu}{\sigma} \right] \right\}^{1-\delta_i}$$

where δ_i takes the value of 1 where an obsevation is left-censored, and 0 where an observation is right-censored.

(c) Use JMP to fit the data. Provide parameter estimates and a plot of the likelihood contour surface.

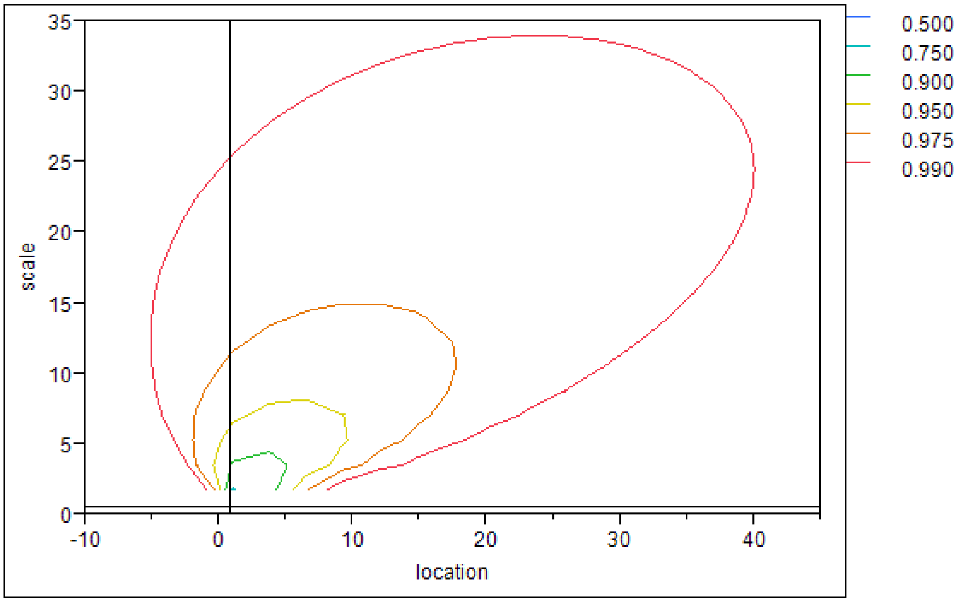

Figure 5: Weibull Likelihood Contour Surface

emg

4 (30 points).

A component has two independent failure modes, say Mode 1 and Mode 2, respectively. The independent failure modes can be modeled using the Weibull distributions WEIB(η_1, β_1) and WEIB(η_2, β_2), respectively.

Define by T the life of the component when the two failure modes are active. From a set of failure time data, the following estimates were obtained from JMP.

$$\hat{\eta_1} = 30, \hat{\beta_1} = 1$$
$$\hat{\eta_1} = 40, \hat{\beta_1} = 2$$

Using the estimates above, do the following:

(a) Provide an estimate of the survival function $S_T(t)$ of the component.

$$\hat{S_T}(t) = \left\{1 - \hat{F_1}(t)\right\}\left\{1 - \hat{F_2}(t)\right\}$$

where $\hat{F_1}$ is WEIB(30, 1) and $\hat{F_2}$ is WEIB(40, 2)

(b) Provide an estimate of the cdf $F_T(t)$ of the component.

$$\hat{F_T}(t) = \left\{1 - \left\{1 - \hat{F_1}(t)\right\}\left\{1 - \hat{F_2}(t)\right\}\right\}$$

where $\hat{F_1}$ is WEIB(30, 1) and $\hat{F_2}$ is WEIB(40, 2)

(c) Obtain an expression for the hazard estimate of T. Your answer must be in function of hazard functions ($h_1(t)$ and $h_2(t)$ for failures from Mode 1 and Mode 2, respectively.

$$h(t) = \frac{f(t)}{1-F(t)}$$

$$\frac{f_1(t)f_2(t)}{1-\{1-[1-F_1(t)][1-F_2(t)]\}}$$

$$\frac{f_1(t)f_2(t)}{[1-F_1(t)][1-F_2(t)]} = h_1(t)h_2(t)$$