

1. (a) The randomization matrix for treatments A and B along with the means and mean difference for each of the 20 unique possible designs

	1	2	3	4	5	6	MeanA	MeanB	MeanDiff
Yield	3	5	4	2	7	4	0.00	0.00	0.00
Design 1	A	B	B	A	B	A	3.00	5.33	2.33
Design 2	A	B	B	A	A	B	4.00	4.33	0.33
Design 3	A	B	A	B	A	B	4.67	3.67	-1.00
Design 4	A	A	B	B	A	B	5.00	3.33	-1.67
Design 5	A	A	B	B	B	A	4.00	4.33	0.33
Design 6	A	B	A	B	B	A	3.67	4.67	1.00
Design 7	A	B	B	B	A	A	4.67	3.67	-1.00
Design 8	B	A	B	B	A	A	5.33	3.00	-2.33
Design 9	B	A	B	A	B	A	3.67	4.67	1.00
Design 10	B	A	A	B	B	A	4.33	4.00	-0.33
Design 11	B	A	A	B	A	B	5.33	3.00	-2.33
Design 12	B	A	B	A	A	B	4.67	3.67	-1.00
Design 13	A	B	A	A	B	B	3.00	5.33	2.33
Design 14	A	A	B	A	B	B	3.33	5.00	1.67
Design 15	A	A	A	B	B	B	4.00	4.33	0.33
Design 16	B	A	A	A	B	B	3.67	4.67	1.00
Design 17	B	B	A	A	B	A	3.33	5.00	1.67
Design 18	B	B	A	B	A	A	5.00	3.33	-1.67
Design 19	B	B	A	A	A	B	4.33	4.00	-0.33
Design 20	B	B	B	A	A	A	4.33	4.00	-0.33

Table 1: Randomization Matrix

- (b) The mean difference between treatments in Design 1, the actual experiment, is 2.33. There are a total of 4 experiments out of twenty that have a mean absolute difference greater than or equal to 2.33. So the p-value, calculated as $Pr(|Difference| \geq 2.33 | randomization) = \frac{4}{20} = .20$.
- (c) For the two-sample t-test with $H_0 : \mu_a = \mu_b$ vs. $H_a : \mu_b \neq \mu_a$, the t statistic is -2.214 and the p-value is 0.102
- (d) The two-sided two-sample t-test gives a smaller p-value than the randomization test, most likely because of the small sample size for each treatment.
- 2.15 (a) We need to test the hypothesis that the population mean μ is greater than $\mu_0 = 120$. The Hypotheses are $H_0 : \mu \leq 120$ vs. $H_a : \mu > 120$. We will use the t-test, rejecting the null if

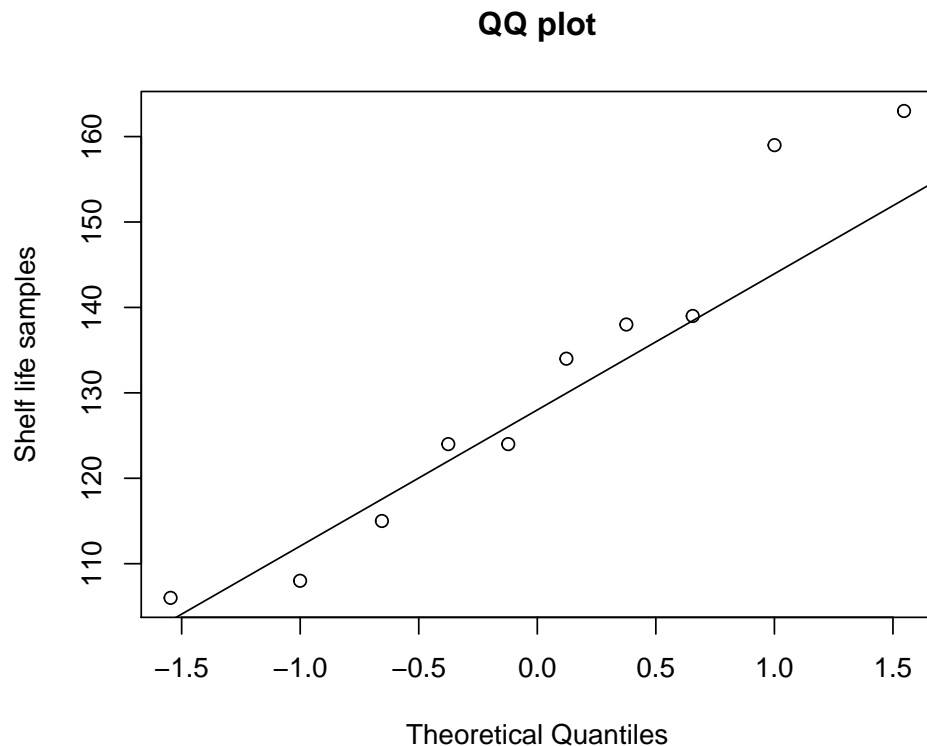
$$t_0 = \frac{\bar{y} - \mu_0}{S/\sqrt{n}} > t_{\alpha, n-1}$$

- (b) Here $\bar{y} = 131$, $S = 19.545$, and $n = 10$. Therefore $t_0 = 1.78$. According to the table in the appendix of Montgomery, $t_{0.01, 9} = 2.821$. Since t_0 is less than $t_{0.01, 9}$ we can not reject the null hypothesis at the 1% level.
- (c) $t_{0.05, 9} = 1.833$ and $t_{0.10, 9} = 1.833$ so the actual p-value lies between .05 and .10, but much closer to .05. According to R it is 0.054.

(d) The 99% confidence interval is given by formula 2.38 in Montgomery

$$\begin{aligned} P(\bar{y} - t_{.01/2, n-1} * S/\sqrt{n} &\leq \mu \leq \bar{y} + t_{.01/2, n-1} * S/\sqrt{n}) = .01 \\ P(131 - 3.25 * 19.545/\sqrt{10} &\leq \mu \leq 131 + 3.25 * 19.545/\sqrt{10}) = .01 \\ P(131 - 20.087 &\leq \mu \leq 131 + 20.087) = .01 \\ P(110.93 &\leq \mu \leq 151.087) = .01 \end{aligned}$$

2.16 We can evaluate the normality assumption by constructing a QQ plot.

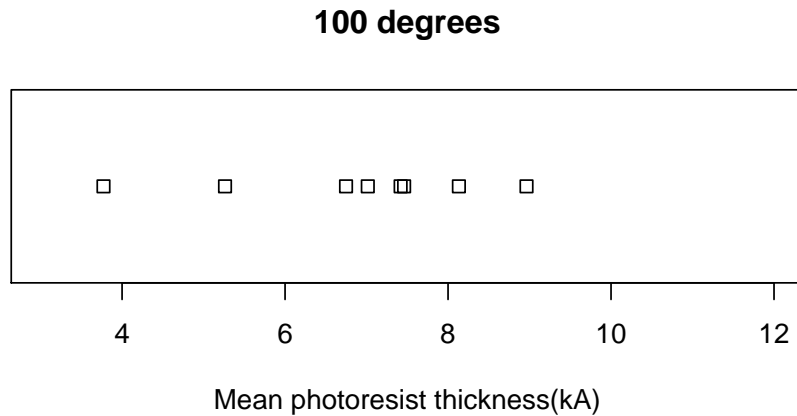
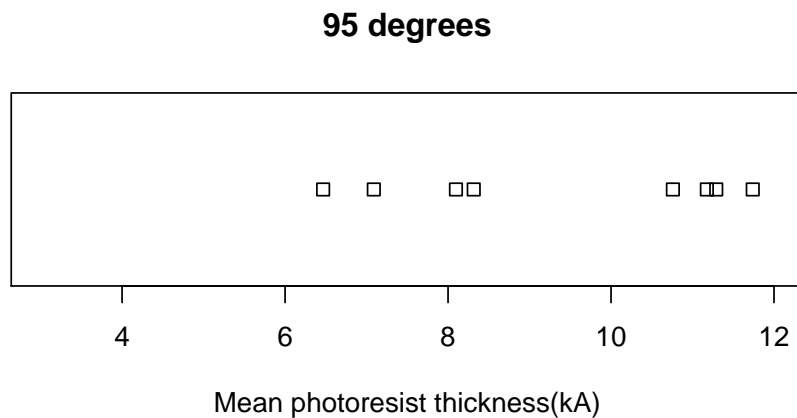


Subjectively, based on the small sample provided, it appears reasonable to conclude that the shelf life of the population is normally distributed. If this assumption were substantially violated, the t-test would not be valid, and another way would have to be found to estimate the mean shelf life.

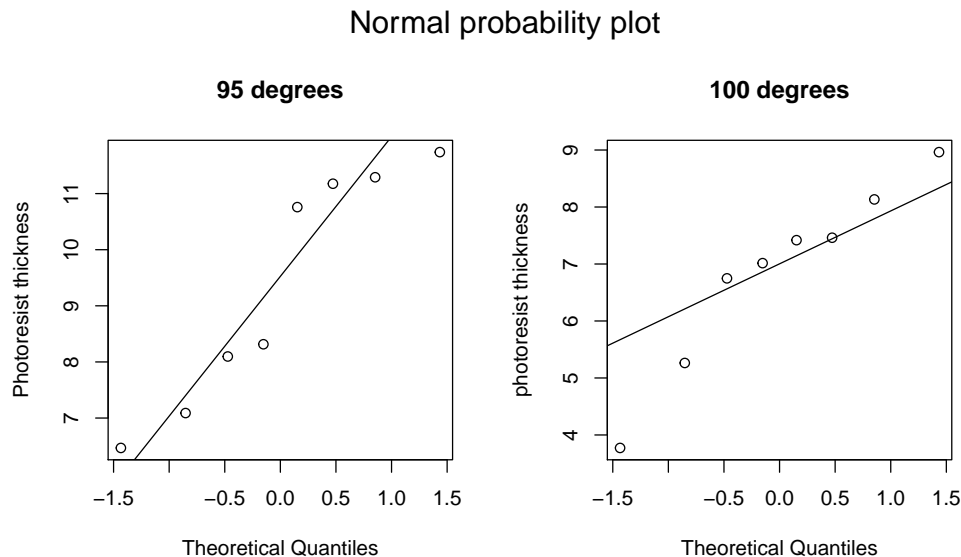
2.24 I will use the R function `t.test` to generate numbers in this question (a)-(e).

- (a) The two-sample t-statistic for this sample, testing the hypothesis $H_0 : \mu_{95} > \mu_{100}$ vs. $H_a : \mu_{95} \leq \mu_{100}$ is 2.675 while $t_{0.05, 14}$ is 1.761. Thus we have can not reject the null, that higher temperatures provide lower mean photoresist thickness, at the 5% level.
- (b) The actual p-value for this two-sample t-test is 0.009
- (c) $\Pr(0.854 \leq \mu_{100} - \mu_{95} \leq \inf) = .05$. There is a 95% probability that the difference between the mean photoresist thickness is greater than 0.854 kÅ.

- (d) The figures below show that the mean photoresist thickness at 95 degrees is generally greater than that for 100 degrees, visually verifying our previous result.



- (e) We will test the normality assumption using a QQ plot as in 2.16.



The normality assumption is justified by the data, although the two low values at 100 degrees seem to diverge from normal.

- (f) Using the pooled sample variance $S_p^2 = \frac{(n_1-1)S_1^2 + (n_2-1)S_2^2}{n_1+n_2-2} = 3.55$, the effect size for a 2.5kA mean difference is $d = \frac{2.5}{2\sqrt{3.55}} = 0.663$. Using $n = 16$ the R function `pwr.t.test` gives $\text{power} = \beta = 0.443$. With the current sample size of 16 there is a 44.3% chance of rejecting $H_0 : \mu_{95} \neq \mu_{100}$ when it is false.
- (g) Using the same pooled variance as above, $d = \frac{1.5}{2\sqrt{3.55}} = 0.398$. To detect an effect that size with a power of 90% you need a total sample size of $n^* = 134$, so each treatment will need to be sampled $\frac{n^*+1}{2} \approx 67.5$ times, which we round up to $n_1 = n_2 = 68$.

2.27 I will use the R function `t.test` to generate numbers in this problem.

- (a) There is not a significant difference in means between the populations the samples were drawn from at the $\alpha = 0.05$ level.
- (b) The two sample t-test for $H_0 : \mu_1 = \mu_2$ has a p-value of 0.522 at the $\alpha = 0.05$ level.
- (c) The 95% confidence interval for the mean difference in mean diameter measurements is $-0.00092 < \mu_1 - \mu_2 < 0.00175$.