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Question 1

```
> d = sum(c(1, 0, 1, 1, 1, 0, 1, 1, 0, 0, 1, 1))
> H = 0.2 * sum(c(12, 15, 17, 17, 18, 19, 20, 20, 20, 21, 24, 27))
> (ts = (d - H) ^ 2 / H)
[1] 31.3913
> (cv = round(qchisq(0.95, 1), 4))
[1] 3.8415
```

```
H_0: h(t) = 0.2 \text{ vs } H_1: h(t) \neq 0.2 \forall t \in (0, 27)
```

With $\sum d_i=8$ and $\sum H_0(T_i)=0.2\sum T_i=46$, we have a test statistic $Z(\tau)=\frac{(8-46)^2}{46}\approx 31.3913$ which is greater than the critical value $\chi^2_{0.95}(1)=3.8415$, thus, we reject H_0 at 5% level of significance.

```
> time1 = c(2, 2, 3, 3, 4, 4, 5, 5, 6)
> time2 = c(2, 3, 3, 3, 4, 4, 4, 5, 5, 5, 6, 7)
> cen1 = c(1, 1, 0, 1, 1, 0, 1, 1, 0)
> cen2 = c(1, 1, 1, 0, 1, 1, 0, 1, 0, 0, 1, 1)
> data = data.frame(time = c(time1, time2), censor = c(cen1, cen2), group = c(rep(1, length(time1)), rep(2, length(time2))))
> survGroup = survdiff(Surv(data$time, data$censor) ~ data$group)
> (ts = round(survGroup$chisq, 4))
[1] 0.3357
> (cv = round(qchisq(0.95, 1), 4))
[1] 3.8415
```

$$H_0: h_1(t) = h_2(t) \ vs \ H_1: h_1(t) \neq h_2(t) \ \forall \ t$$

From the result given by above code, we notice the test statistic (0.3357) which is less than critical value (3.8415), thus, we do not reject H_0 at 5% level of significance.

```
> time = c(18, 9, 28, 31, 39, 19, 45, 6, 8, 15, 23, 28, 7, 12, 9, 8, 2, 26, 10,
4, 3, 4, 18, 8, 3, 14, 3, 13, 13, 35)
> censor = c(1, 1, 0, 1, 0, 0, 0, 1, 1, 1, 1, 0, 1, 1, 1, 1, 0, 1, 1, 1,
1, 1, 1, 1, 1, 1, 0)
> group = c(1, 3, 1, 3, 3, 3, 3, 3, 3, 3, 1, 1, 3, 2, 2, 2, 4, 2, 4, 2, 2, 2,
4, 4, 4, 4, 2, 4, 4, 2)
> data = data.frame(time, censor, group)
> 
> survGroup = survdiff(Surv(data$time, data$censor) ~ data$group)
> (ts = round(survGroup$chisq, 4))
[1] 5.6891
> (cv = round(qchisq(0.95, 3), 4))
[1] 7.8147
```

There are 4 combinations of prognostic factor in total, hence, we refine individuals into their corresponding group as

```
Group 1: x_1 = 0; x_2 = 0

Group 2: x_1 = 1; x_2 = 0

Group 3: x_1 = 0; x_2 = 1

Group 4: x_1 = 1; x_2 = 1
```

 H_0 : $h_i(t) = h_i(t)$ vs H_1 : $h_i(t) \neq h_j(t) \forall t$ and $i \neq j$

From the result given by above code, we notice the test statistic (5.6891) which is less than critical value (7.8147), thus, we do not reject H_0 at 5% level of significance.

```
> library("readxl")
> data = as.data.frame(read_excel('ass3data.xls'))
```

>> survGroup = survdiff(Surv(data\$Time) ~ data\$Group)
>> (ts = round(survGroup\$chisq, 4))
[1] 6.4561
>> (cv = round(qchisq(0.95, length(unique(data\$Group)) - 1), 4))
[1] 5.9915

```
H_0: h_i(t) = h_i(t) vs H_1: h_i(t) \neq h_i(t) \forall t and i \neq j
```

From the result given by above code, we notice the test statistic (6.4561) which is greater than critical value (5.9915), thus, we reject H_0 at 5% level of significance.

```
b)

> survTreat = survdiff(Surv(data$Time) ~ data$Treatment)
> (ts = round(survTreat$chisq, 4))
[1] 0.7369
> (cv = round(qchisq(0.95, length(unique(data$Treatment)) - 1), 4))
[1] 5.9915
```

```
H_0: h_i(t) = h_i(t) vs H_1: h_i(t) \neq h_i(t) \forall t and i \neq j
```

From the result given by above code, we notice the test statistic (0.7369) which is less than critical value (5.9915), thus, we do not reject H_0 at 5% level of significance.

```
> data = read.csv('ass3q5.csv')
> set.seed(123457)
> data = data[sample(nrow(data), 100), ]
>
> survStatus = survdiff(Surv(data$Time, data$Status) ~ data$Smoking.Status)
> (ts = round(survStatus$chisq, 4))
[1] 4.2712
> (cv = round(qchisq(0.95, length(unique(data$Smoking.Status)) - 1), 4))
[1] 5.9915
```

```
H_0: h_i(t) = h_j(t) vs H_1: h_i(t) \neq h_j(t) \forall t and i \neq j
```

From the result given by above code, we notice the test statistic (4.2712) which is less than critical value (5.9915), thus, we do not reject H_0 at 5% level of significance.