

HW_3

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#####

6.2. For the covariates with complete data (in Columns 1-22), use stepwise regression with AIC to identify the best model for (a) overall survival, and (b) recurrence-free survival.

#####

```
library(survival)
library(asauro)
library(MASS)
attach (hepatoCellular)
head(hepatoCellular[, 1:8])
```

##	Number	Age	Gender	HBsAg	Cirrhosis	ALT	AST	AFP
## 1	1	57	0	1	1	1	2	2
## 2	2	58	1	0	0	1	1	2
## 3	3	65	1	0	0	1	1	2
## 4	4	54	1	1	0	2	1	2
## 5	5	71	1	1	0	2	2	2
## 6	6	32	1	0	0	2	2	2

```
head(hepatoCellular[, 9:13])
```

##	Tumorsize	Tumordifferentiation	Vascularinvasion	Tumormultiplicity	Capsulation
## 1	2	1	0	1	0
## 2	1	1	0	1	1
## 3	2	2	1	1	1
## 4	2	2	0	1	1
## 5	2	2	1	2	0
## 6	2	2	0	1	0

```
head(hepatoCellular[, 14:22])
```

##	TNM	BCLC	OS	Death	RFS	Recurrence	CXCL17T	CXCL17P	CXCL17N
## 1	2	1	83	0	13	1	113.947238	299.325404	138.247177
## 2	1	1	81	0	81	0	54.071542	63.468462	6.214912
## 3	2	2	79	0	79	0	22.188831	34.470336	22.131934
## 4	1	1	76	0	76	0	8.442809	16.002843	11.146906
## 5	2	2	7	1	3	1	8.271131	22.034540	32.271196
## 6	1	1	13	1	3	1	13.708073	6.423604	0.000000

(a) overall survival

With RFS Included

```
modelAll1.coxph <- coxph(Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT +AST + AFP
```

```

+ Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity + Capsula
tion + TNM + BCLC + RFS + Recurrence + CXCL17T + CXCL17P + CXCL17N)

## Warning in coxph.fit(X, Y, istrat, offset, init, control, weights = weights, :
## Loglik converged before variable 15 ; coefficient may be infinite.

result.step1 <-suppressWarnings(step(modelAll1.coxph, scope=list(upper=~ Age + HBs
Ag + Cirrhosis + ALT +AST + AFP+ Tumorsize + Tumordifferentiation + Vascularinvasio
n + Tumormultiplicity + Capsulation + TNM + BCLC + RFS + Recurrence + CXCL17T + CX
CL17P + CXCL17N, lower=~Age)))

## Start: AIC=818.36
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
## Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
## Capsulation + TNM + BCLC + RFS + Recurrence + CXCL17T + CXCL17P +
## CXCL17N
##
##
## Df AIC
## - AST 1 816.36
## - Tumormultiplicity 1 816.40
## - ALT 1 816.49
## - CXCL17T 1 816.59
## - CXCL17N 1 816.64
## - Capsulation 1 816.88
## - TNM 1 816.95
## - BCLC 1 817.26
## - Cirrhosis 1 817.49
## <none> 818.36
## - AFP 1 819.21
## - CXCL17P 1 819.51
## - HBsAg 1 819.57
## - Tumordifferentiation 1 819.98
## - Tumorsize 1 820.63
## - Vascularinvasion 1 821.04
## - Recurrence 1 829.41
## - RFS 1 841.35
##
## Step: AIC=816.36
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AFP + Tumorsize +
## Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
## Capsulation + TNM + BCLC + RFS + Recurrence + CXCL17T + CXCL17P +
## CXCL17N
##
##
## Df AIC
## - Tumormultiplicity 1 814.40
## - ALT 1 814.51
## - CXCL17T 1 814.60
## - CXCL17N 1 814.65
## - Capsulation 1 814.89
## - TNM 1 814.95

```

```

## - BCLC 1 815.26
## - Cirrhosis 1 815.49
## <none> 816.36
## - AFP 1 817.24
## - CXCL17P 1 817.55
## - HBsAg 1 817.57
## - Tumordifferentiation 1 818.05
## + AST 1 818.36
## - Tumorsize 1 818.72
## - Vascularinvasion 1 819.06
## - Recurrence 1 827.47
## - RFS 1 839.39
##
## Step: AIC=814.4
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AFP + Tumorsize +
## Tumordifferentiation + Vascularinvasion + Capsulation + TNM +
## BCLC + RFS + Recurrence + CXCL17T + CXCL17P + CXCL17N
##
## Df AIC
## - ALT 1 812.53
## - CXCL17T 1 812.62
## - CXCL17N 1 812.71
## - Capsulation 1 812.97
## - TNM 1 812.99
## - Cirrhosis 1 813.50
## - BCLC 1 814.20
## <none> 814.40
## - HBsAg 1 815.57
## - CXCL17P 1 815.61
## - AFP 1 815.62
## - Tumordifferentiation 1 816.05
## + Tumormultiplicity 1 816.36
## + AST 1 816.40
## - Tumorsize 1 816.80
## - Vascularinvasion 1 817.48
## - Recurrence 1 825.48
## - RFS 1 837.99
##
## Step: AIC=812.53
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + AFP + Tumorsize +
## Tumordifferentiation + Vascularinvasion + Capsulation + TNM +
## BCLC + RFS + Recurrence + CXCL17T + CXCL17P + CXCL17N
##
## Df AIC
## - CXCL17T 1 810.76
## - CXCL17N 1 810.81
## - TNM 1 811.05
## - Capsulation 1 811.07
## - Cirrhosis 1 811.69
## - BCLC 1 812.53

```

```

## <none> 812.53
## - CXCL17P 1 813.70
## - HBsAg 1 813.83
## - AFP 1 813.87
## - Tumordifferentiation 1 814.22
## + ALT 1 814.40
## + Tumormultiplicity 1 814.51
## + AST 1 814.51
## - Tumorsize 1 814.82
## - Vascularinvasion 1 815.49
## - Recurrence 1 823.73
## - RFS 1 836.20
##
## Step: AIC=810.76
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + AFP + Tumorsize +
## Tumordifferentiation + Vascularinvasion + Capsulation + TNM +
## BCLC + RFS + Recurrence + CXCL17P + CXCL17N
##
## Df AIC
## - CXCL17N 1 809.26
## - Capsulation 1 809.33
## - TNM 1 809.34
## - Cirrhosis 1 809.95
## <none> 810.76
## - BCLC 1 811.14
## - CXCL17P 1 811.73
## - HBsAg 1 811.89
## + CXCL17T 1 812.53
## - Tumordifferentiation 1 812.55
## + ALT 1 812.62
## - AFP 1 812.64
## + Tumormultiplicity 1 812.75
## + AST 1 812.76
## - Tumorsize 1 813.07
## - Vascularinvasion 1 813.54
## - Recurrence 1 822.05
## - RFS 1 834.45
##
## Step: AIC=809.26
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + AFP + Tumorsize +
## Tumordifferentiation + Vascularinvasion + Capsulation + TNM +
## BCLC + RFS + Recurrence + CXCL17P
##
## Df AIC
## - TNM 1 807.72
## - Capsulation 1 807.87
## - Cirrhosis 1 808.30
## <none> 809.26
## - BCLC 1 809.46
## - CXCL17P 1 810.11

```

```

## - HBsAg          1 810.32
## - AFP            1 810.73
## + CXCL17N        1 810.76
## + CXCL17T        1 810.81
## - Tumordifferentiation 1 810.95
## + ALT            1 811.18
## + Tumormultiplicity 1 811.22
## + AST            1 811.23
## - Tumorsize      1 811.26
## - Vascularinvasion 1 812.13
## - Recurrence     1 820.98
## - RFS            1 832.49
##
## Step: AIC=807.72
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + AFP + Tumorsize +
##      Tumordifferentiation + Vascularinvasion + Capsulation + BCLC +
##      RFS + Recurrence + CXCL17P
##
##              Df      AIC
## - Capsulation      1 806.37
## - Cirrhosis        1 806.61
## - BCLC              1 807.46
## <none>              807.72
## - CXCL17P          1 808.16
## - AFP              1 808.87
## - HBsAg            1 809.14
## + CXCL17T          1 809.23
## + TNM              1 809.26
## - Tumorsize        1 809.29
## + CXCL17N          1 809.34
## - Tumordifferentiation 1 809.61
## + Tumormultiplicity 1 809.68
## + ALT              1 809.69
## + AST              1 809.71
## - Vascularinvasion 1 810.77
## - Recurrence       1 820.39
## - RFS              1 831.37
##
## Step: AIC=806.37
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + AFP + Tumorsize +
##      Tumordifferentiation + Vascularinvasion + BCLC + RFS + Recurrence +
##      CXCL17P
##
##              Df      AIC
## - Cirrhosis        1 805.16
## - BCLC              1 806.18
## <none>              806.37
## - CXCL17P          1 806.96
## + Capsulation      1 807.72
## - Tumordifferentiation 1 807.73

```

```

## - Tumorsize          1 807.78
## + CXCL17T            1 807.80
## - AFP                1 807.86
## + TNM                1 807.87
## + CXCL17N            1 807.95
## + Tumormultiplicity  1 808.27
## - HBsAg              1 808.33
## + AST                1 808.34
## + ALT                1 808.36
## - Vascularinvasion   1 808.94
## - Recurrence         1 818.65
## - RFS                1 832.12
##
## Step: AIC=805.16
## Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize + Tumordifferentiation +
##      Vascularinvasion + BCLC + RFS + Recurrence + CXCL17P
##
##              Df      AIC
## - BCLC          1 804.95
## <none>           805.16
## - CXCL17P       1 805.60
## - Tumorsize     1 806.14
## + Cirrhosis     1 806.37
## - AFP           1 806.38
## - HBsAg         1 806.45
## + Capsulation   1 806.61
## + CXCL17T       1 806.63
## + TNM           1 806.82
## + CXCL17N       1 806.83
## - Tumordifferentiation 1 806.95
## + Tumormultiplicity 1 807.12
## + ALT           1 807.13
## + AST           1 807.13
## - Vascularinvasion 1 807.65
## - Recurrence    1 817.79
## - RFS           1 830.23
##
## Step: AIC=804.95
## Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize + Tumordifferentiation +
##      Vascularinvasion + RFS + Recurrence + CXCL17P
##
##              Df      AIC
## - CXCL17P       1 804.72
## <none>           804.95
## + BCLC          1 805.16
## + Tumormultiplicity 1 805.91
## - HBsAg         1 805.98
## + CXCL17T       1 806.03
## - Tumordifferentiation 1 806.08
## + Cirrhosis     1 806.18

```

```
## + Capsulation          1 806.32
## + CXCL17N              1 806.65
## + ALT                  1 806.71
## + AST                  1 806.84
## - AFP                  1 806.90
## + TNM                  1 806.94
## - Tumorsize            1 807.28
## - Vascularinvasion     1 813.51
## - Recurrence           1 816.81
## - RFS                  1 835.43
##
```

```
## Step: AIC=804.72
```

```
## Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize + Tumordifferentiation +
## Vascularinvasion + RFS + Recurrence
```

```
##
##              Df      AIC
## <none>          804.72
## + CXCL17P        1 804.95
## - HBsAg          1 805.46
## + BCLC           1 805.60
## - Tumordifferentiation 1 805.66
## + Capsulation    1 805.95
## + Tumormultiplicity 1 806.00
## + Cirrhosis      1 806.08
## - AFP            1 806.18
## + ALT            1 806.42
## + CXCL17N        1 806.45
## + AST            1 806.57
## + TNM            1 806.62
## + CXCL17T        1 806.68
## - Tumorsize      1 806.93
## - Vascularinvasion 1 814.13
## - Recurrence     1 817.44
## - RFS            1 835.80
```

Without RFS Included

```
modelAll1n.coxph <- coxph(Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
P+ Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P + CXCL17N)
```

```
## Warning in coxph.fit(X, Y, istrat, offset, init, control, weights = weights, :
## Loglik converged before variable 14 ; coefficient may be infinite.
```

```
result.step1n <- suppressWarnings(step(modelAll1n.coxph, scope=list(upper=~ Age + HBsAg + Cirrhosis + ALT + AST + AFP + Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P + CXCL17N, lower=~Age)))
```

```
## Start: AIC=841.35
```

```
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
```

```
## Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
```

```

##      Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P +
##      CXCL17N
##
##              Df      AIC
## - AST              1 839.39
## - CXCL17N          1 839.41
## - ALT              1 839.47
## - CXCL17T          1 839.52
## - Cirrhosis        1 839.62
## - TNM              1 839.88
## - Tumormultiplicity 1 839.92
## - BCLC             1 840.21
## - CXCL17P          1 840.52
## - Tumorsize        1 840.74
## - HBsAg            1 840.77
## - AFP              1 840.87
## <none>              841.35
## - Capsulation      1 841.57
## - Vascularinvasion 1 843.68
## - Tumordifferentiation 1 846.16
## - Recurrence       1 952.96
##
## Step:  AIC=839.39
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AFP + Tumorsize +
##      Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##      Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P +
##      CXCL17N
##
##              Df      AIC
## - CXCL17N          1 837.47
## - ALT              1 837.58
## - CXCL17T          1 837.59
## - Cirrhosis        1 837.67
## - TNM              1 837.94
## - Tumormultiplicity 1 837.99
## - BCLC             1 838.25
## - CXCL17P          1 838.61
## - Tumorsize        1 838.80
## - HBsAg            1 838.82
## - AFP              1 838.96
## <none>              839.39
## - Capsulation      1 839.83
## + AST              1 841.35
## - Vascularinvasion 1 841.80
## - Tumordifferentiation 1 844.44
## - Recurrence       1 951.50
##
## Step:  AIC=837.47
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AFP + Tumorsize +
##      Tumordifferentiation + Vascularinvasion + Tumormultiplicity +

```



```

##      Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P
##
##              Df      AIC
## - CXCL17T      1 835.61
## - ALT          1 835.63
## - Cirrhosis    1 835.74
## - TNM          1 836.09
## - Tumormultiplicity 1 836.13
## - BCLC         1 836.27
## - Tumorsize    1 836.82
## - CXCL17P      1 836.84
## - HBsAg        1 836.91
## - AFP          1 836.96
## <none>          837.47
## - Capsulation  1 837.91
## + CXCL17N      1 839.39
## + AST          1 839.41
## - Vascularinvasion 1 840.01
## - Tumordifferentiation 1 842.46
## - Recurrence   1 950.66
##
## Step: AIC=835.61
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AFP + Tumorsize +
##      Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##      Capsulation + TNM + BCLC + Recurrence + CXCL17P
##
##              Df      AIC
## - ALT          1 833.78
## - Cirrhosis    1 833.86
## - TNM          1 834.31
## - Tumormultiplicity 1 834.31
## - BCLC         1 834.35
## - Tumorsize    1 834.93
## - AFP          1 834.97
## - HBsAg        1 835.25
## <none>          835.61
## - CXCL17P      1 835.93
## - Capsulation  1 836.09
## + CXCL17T      1 837.47
## + AST          1 837.53
## + CXCL17N      1 837.59
## - Vascularinvasion 1 838.36
## - Tumordifferentiation 1 840.64
## - Recurrence   1 951.13
##
## Step: AIC=833.78
## Surv(OS, Death) ~ Age + HBsAg + Cirrhosis + AFP + Tumorsize +
##      Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##      Capsulation + TNM + BCLC + Recurrence + CXCL17P
##

```

```

##          Df    AIC
## - Cirrhosis      1 832.05
## - Tumormultiplicity 1 832.49
## - TNM             1 832.58
## - BCLC            1 832.61
## - Tumorsize       1 832.97
## - AFP             1 833.21
## - HBsAg           1 833.41
## <none>            833.78
## - CXCL17P         1 834.16
## - Capsulation     1 834.22
## + AST             1 835.61
## + ALT             1 835.61
## + CXCL17T         1 835.63
## + CXCL17N         1 835.77
## - Vascularinvasion 1 836.36
## - Tumordifferentiation 1 838.76
## - Recurrence      1 949.31
##
## Step: AIC=832.05
## Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize + Tumordifferentiation +
##   Vascularinvasion + Tumormultiplicity + Capsulation + TNM +
##   BCLC + Recurrence + CXCL17P
##
##          Df    AIC
## - Tumormultiplicity 1 830.70
## - TNM               1 830.85
## - BCLC              1 830.92
## - Tumorsize         1 831.10
## - AFP               1 831.45
## - HBsAg             1 831.48
## <none>              832.05
## - Capsulation       1 832.39
## - CXCL17P           1 832.45
## + Cirrhosis         1 833.78
## + AST               1 833.85
## + ALT               1 833.86
## + CXCL17T           1 833.92
## + CXCL17N           1 834.04
## - Vascularinvasion  1 834.51
## - Tumordifferentiation 1 837.22
## - Recurrence        1 948.72
##
## Step: AIC=830.7
## Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize + Tumordifferentiation +
##   Vascularinvasion + Capsulation + TNM + BCLC + Recurrence +
##   CXCL17P
##
##          Df    AIC
## - TNM              1 829.58

```

```

## - Tumorsize          1 829.74
## - HBsAg              1 830.07
## <none>                830.70
## - AFP                1 830.81
## - CXCL17P            1 831.14
## - Capsulation        1 831.40
## + Tumormultiplicity  1 832.05
## - BCLC                1 832.15
## + AST                 1 832.40
## + Cirrhosis          1 832.49
## + ALT                 1 832.51
## - Vascularinvasion   1 832.52
## + CXCL17T            1 832.53
## + CXCL17N            1 832.67
## - Tumordifferentiation 1 835.44
## - Recurrence         1 947.37
##
## Step: AIC=829.58
## Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize + Tumordifferentiation +
##      Vascularinvasion + Capsulation + BCLC + Recurrence + CXCL17P
##
##              Df      AIC
## - HBsAg          1 828.63
## <none>            829.58
## - Tumorsize      1 829.66
## - AFP            1 829.84
## - Capsulation    1 830.23
## + TNM            1 830.70
## + Tumormultiplicity 1 830.85
## - Vascularinvasion 1 830.90
## - CXCL17P        1 830.96
## + AST            1 831.13
## + ALT            1 831.29
## + CXCL17T        1 831.32
## + Cirrhosis      1 831.37
## + CXCL17N        1 831.52
## - BCLC           1 833.63
## - Tumordifferentiation 1 833.74
## - Recurrence     1 945.73
##
## Step: AIC=828.63
## Surv(OS, Death) ~ Age + AFP + Tumorsize + Tumordifferentiation +
##      Vascularinvasion + Capsulation + BCLC + Recurrence + CXCL17P
##
##              Df      AIC
## - Tumorsize      1 828.51
## - AFP            1 828.57
## <none>            828.63
## + HBsAg          1 829.58
## - CXCL17P        1 829.71

```

```
## + Tumormultiplicity      1 829.98
## + TNM                    1 830.07
## - Capsulation            1 830.10
## + AST                    1 830.19
## + CXCL17T                1 830.19
## - Vascularinvasion       1 830.23
## + ALT                    1 830.39
## + Cirrhosis              1 830.58
## + CXCL17N                1 830.58
## - BCLC                   1 832.53
## - Tumordifferentiation    1 832.56
## - Recurrence             1 943.97
##
## Step: AIC=828.51
## Surv(OS, Death) ~ Age + AFP + Tumordifferentiation + Vascularinvasion +
##   Capsulation + BCLC + Recurrence + CXCL17P
##
##              Df      AIC
## <none>                828.51
## + Tumorsize           1 828.63
## - AFP                 1 828.73
## + TNM                 1 829.10
## - CXCL17P             1 829.57
## - Capsulation         1 829.62
## - Vascularinvasion    1 829.64
## + HBsAg               1 829.66
## + Tumormultiplicity   1 829.86
## + CXCL17T             1 830.08
## + AST                 1 830.11
## + ALT                 1 830.45
## + CXCL17N             1 830.49
## + Cirrhosis           1 830.51
## - Tumordifferentiation 1 831.95
## - BCLC                1 833.80
## - Recurrence          1 946.00
```

With RFS Included

```
summary(result.step1)
```

```
## Call:
## coxph(formula = Surv(OS, Death) ~ Age + HBsAg + AFP + Tumorsize +
##   Tumordifferentiation + Vascularinvasion + RFS + Recurrence)
##
## n= 227, number of events= 97
##
##              coef exp(coef) se(coef)      z Pr(>|z|)
## Age             1.220e-02  1.012e+00  8.200e-03  1.488 0.136687
## HBsAg           -7.472e-01  4.737e-01  4.133e-01 -1.808 0.070611 .
## AFP             4.526e-01  1.572e+00  2.513e-01  1.801 0.071724 .
```

```
## Tumorsize          4.432e-01  1.558e+00  2.203e-01  2.011 0.044296 *
## Tumordifferentiation 3.615e-01  1.436e+00  2.097e-01  1.724 0.084642 .
## Vascularinvasion    9.035e-01  2.468e+00  2.504e-01  3.608 0.000309 ***
## RFS                 -6.237e-02  9.395e-01  1.252e-02 -4.981 6.32e-07 ***
## Recurrence          1.804e+01  6.835e+07  2.393e+03  0.008 0.993985
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##              exp(coef) exp(-coef) lower .95 upper .95
## Age          1.012e+00  9.879e-01    0.9961    1.0287
## HBsAg         4.737e-01  2.111e+00    0.2107    1.0648
## AFP           1.572e+00  6.360e-01    0.9608    2.5732
## Tumorsize     1.558e+00  6.420e-01    1.0114    2.3990
## Tumordifferentiation 1.436e+00  6.966e-01    0.9518    2.1650
## Vascularinvasion 2.468e+00  4.052e-01    1.5108    4.0322
## RFS           9.395e-01  1.064e+00    0.9168    0.9629
## Recurrence    6.835e+07  1.463e-08    0.0000      Inf
##
## Concordance= 0.872 (se = 0.014 )
## Likelihood ratio test= 191.6 on 8 df,  p=<2e-16
## Wald test              = 52.85 on 8 df,  p=1e-08
## Score (logrank) test = 143.9 on 8 df,  p=<2e-16
```

Without RFS Included

```
result.step1n
```

```
## Call:
## coxph(formula = Surv(OS, Death) ~ Age + AFP + Tumordifferentiation +
##       Vascularinvasion + Capsulation + BCLC + Recurrence + CXCL17P)
##
##              coef exp(coef) se(coef)      z      p
## Age          3.140e-03  1.003e+00  8.147e-03  0.385 0.69991
## AFP          3.672e-01  1.444e+00  2.531e-01  1.451 0.14682
## Tumordifferentiation 5.082e-01  1.662e+00  2.161e-01  2.351 0.01870
## Vascularinvasion  5.353e-01  1.708e+00  2.997e-01  1.786 0.07412
## Capsulation    -4.471e-01  6.395e-01  2.468e-01 -1.811 0.07010
## BCLC           7.099e-01  2.034e+00  2.551e-01  2.783 0.00538
## Recurrence     2.019e+01  5.879e+08  2.454e+03  0.008 0.99344
## CXCL17P        1.185e-03  1.001e+00  6.293e-04  1.882 0.05981
##
## Likelihood ratio test=167.8 on 8 df, p=< 2.2e-16
## n= 227, number of events= 97
```

(b) recurrence-free survival.

With OS Included

```
modelAll2.coxph <- suppressWarnings(coxph(Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP + Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormult
```

```
iplicity + Capsulation + TNM + BCLC + OS + Recurrence + CXCL17T + CXCL17P + CXCL17N))
```

```
result.step2 <-suppressWarnings(step(modelAll2.coxph, scope=list(upper=~ Age + HBsAg + Cirrhosis + ALT +AST + AFP+ Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity + Capsulation + TNM + BCLC + OS + Recurrence + CXCL17T + CXCL17P + CXCL17N, lower=~Age)))
```

```
## Start: AIC=691.43
```

```
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +  
## Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +  
## Capsulation + TNM + BCLC + OS + Recurrence + CXCL17T + CXCL17P +  
## CXCL17N
```

```
##  
## Df AIC  
## - TNM 1 689.43  
## - Capsulation 1 689.43  
## - Tumorsize 1 689.45  
## - Vascularinvasion 1 689.48  
## - ALT 1 689.58  
## - AST 1 689.68  
## - Cirrhosis 1 689.80  
## - AFP 1 689.92  
## - BCLC 1 689.94  
## - Tumormultiplicity 1 690.25  
## - CXCL17T 1 690.28  
## - HBsAg 1 690.63  
## - Tumordifferentiation 1 690.65  
## - CXCL17P 1 691.41  
## <none> 691.43  
## - CXCL17N 1 691.60  
## - Recurrence 1 773.38  
## - OS 1 785.80  
##
```

```
## Step: AIC=689.43
```

```
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +  
## Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +  
## Capsulation + BCLC + OS + Recurrence + CXCL17T + CXCL17P +  
## CXCL17N
```

```
##  
## Df AIC  
## - Capsulation 1 687.43  
## - Tumorsize 1 687.47  
## - Vascularinvasion 1 687.48  
## - ALT 1 687.58  
## - AST 1 687.68  
## - Cirrhosis 1 687.80  
## - AFP 1 687.92  
## - BCLC 1 688.08  
## - Tumormultiplicity 1 688.25
```

```

## - CXCL17T          1 688.28
## - HBsAg            1 688.65
## - Tumordifferentiation 1 688.66
## <none>              689.43
## - CXCL17P          1 689.55
## - CXCL17N          1 689.61
## + TNM              1 691.43
## - Recurrence       1 773.99
## - OS               1 786.31
##
## Step: AIC=687.43
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
##   Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##   BCLC + OS + Recurrence + CXCL17T + CXCL17P + CXCL17N
##
##              Df    AIC
## - Tumorsize      1 685.48
## - Vascularinvasion 1 685.48
## - ALT            1 685.58
## - AST            1 685.68
## - Cirrhosis      1 685.82
## - AFP            1 685.93
## - BCLC           1 686.08
## - Tumormultiplicity 1 686.25
## - CXCL17T        1 686.30
## - Tumordifferentiation 1 686.69
## - HBsAg          1 686.75
## <none>            687.43
## - CXCL17P        1 687.57
## - CXCL17N        1 687.65
## + Capsulation    1 689.43
## + TNM            1 689.43
## - Recurrence     1 772.03
## - OS             1 785.86
##
## Step: AIC=685.48
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
##   Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##   BCLC + OS + Recurrence + CXCL17T + CXCL17P + CXCL17N
##
##              Df    AIC
## - Vascularinvasion 1 683.54
## - ALT              1 683.62
## - AST              1 683.74
## - Cirrhosis        1 683.83
## - AFP              1 684.00
## - BCLC             1 684.09
## - CXCL17T          1 684.30
## - Tumormultiplicity 1 684.34
## - Tumordifferentiation 1 684.69

```

```

## - HBsAg                1 684.95
## <none>                  685.48
## - CXCL17P              1 685.59
## - CXCL17N              1 685.74
## + Tumorsize            1 687.43
## + TNM                  1 687.46
## + Capsulation          1 687.47
## - Recurrence           1 771.09
## - OS                   1 784.43
##
## Step: AIC=683.54
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
##      Tumordifferentiation + Tumormultiplicity + BCLC + OS + Recurrence +
##      CXCL17T + CXCL17P + CXCL17N
##
##              Df      AIC
## - ALT                1 681.68
## - AST                1 681.81
## - Cirrhosis          1 681.96
## - AFP                1 682.25
## - Tumormultiplicity  1 682.36
## - CXCL17T            1 682.47
## - Tumordifferentiation 1 682.70
## - BCLC               1 682.76
## - HBsAg              1 683.01
## <none>               683.54
## - CXCL17P            1 683.60
## - CXCL17N            1 683.74
## + Vascularinvasion   1 685.48
## + Tumorsize          1 685.48
## + TNM                1 685.53
## + Capsulation        1 685.53
## - Recurrence         1 771.17
## - OS                 1 783.67
##
## Step: AIC=681.68
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + AST + AFP + Tumordifferentiation +
##      Tumormultiplicity + BCLC + OS + Recurrence + CXCL17T + CXCL17P +
##      CXCL17N
##
##              Df      AIC
## - Cirrhosis          1 680.03
## - AST                1 680.11
## - AFP                1 680.37
## - Tumormultiplicity  1 680.55
## - CXCL17T            1 680.56
## - BCLC               1 680.79
## - Tumordifferentiation 1 680.83
## - HBsAg              1 681.13
## <none>               681.68

```



```

## - CXCL17P          1 681.68
## - CXCL17N          1 681.83
## + ALT              1 683.54
## + Vascularinvasion 1 683.62
## + Tumorsize        1 683.64
## + Capsulation      1 683.67
## + TNM              1 683.67
## - Recurrence       1 769.43
## - OS               1 781.68
##
## Step: AIC=680.03
## Surv(RFS, Death) ~ Age + HBsAg + AST + AFP + Tumordifferentiation +
##   Tumormultiplicity + BCLC + OS + Recurrence + CXCL17T + CXCL17P +
##   CXCL17N
##
##              Df    AIC
## - AST          1 678.43
## - AFP          1 678.74
## - Tumormultiplicity 1 678.83
## - CXCL17T      1 678.88
## - Tumordifferentiation 1 679.16
## - BCLC         1 679.26
## - HBsAg        1 679.58
## - CXCL17P      1 679.73
## - CXCL17N      1 680.03
## <none>         680.03
## + Cirrhosis    1 681.68
## + Vascularinvasion 1 681.90
## + ALT          1 681.96
## + Capsulation  1 682.01
## + Tumorsize    1 682.03
## + TNM          1 682.03
## - Recurrence   1 767.93
## - OS           1 780.07
##
## Step: AIC=678.43
## Surv(RFS, Death) ~ Age + HBsAg + AFP + Tumordifferentiation +
##   Tumormultiplicity + BCLC + OS + Recurrence + CXCL17T + CXCL17P +
##   CXCL17N
##
##              Df    AIC
## - Tumormultiplicity 1 677.05
## - CXCL17T           1 677.09
## - AFP               1 677.18
## - Tumordifferentiation 1 677.57
## - CXCL17P          1 677.82
## - HBsAg             1 677.94
## - CXCL17N           1 678.27
## <none>              678.43
## - BCLC              1 678.53

```

```

## + AST 1 680.03
## + Cirrhosis 1 680.11
## + ALT 1 680.23
## + Vascularinvasion 1 680.28
## + Tumorsize 1 680.42
## + Capsulation 1 680.42
## + TNM 1 680.42
## - Recurrence 1 766.15
## - OS 1 779.46
##
## Step: AIC=677.05
## Surv(RFS, Death) ~ Age + HBsAg + AFP + Tumordifferentiation +
## BCLC + OS + Recurrence + CXCL17T + CXCL17P + CXCL17N
##
## Df AIC
## - CXCL17T 1 675.60
## - Tumordifferentiation 1 675.95
## - AFP 1 676.05
## - CXCL17P 1 676.14
## - HBsAg 1 676.38
## - CXCL17N 1 676.74
## <none> 677.05
## + Tumormultiplicity 1 678.43
## + Cirrhosis 1 678.79
## + AST 1 678.83
## + ALT 1 678.84
## + Tumorsize 1 679.04
## + Capsulation 1 679.05
## + TNM 1 679.05
## + Vascularinvasion 1 679.05
## - BCLC 1 679.47
## - Recurrence 1 764.18
## - OS 1 779.03
##
## Step: AIC=675.6
## Surv(RFS, Death) ~ Age + HBsAg + AFP + Tumordifferentiation +
## BCLC + OS + Recurrence + CXCL17P + CXCL17N
##
## Df AIC
## - CXCL17P 1 674.28
## - Tumordifferentiation 1 674.34
## - AFP 1 674.38
## - HBsAg 1 674.80
## - CXCL17N 1 675.34
## <none> 675.60
## + CXCL17T 1 677.05
## + Tumormultiplicity 1 677.09
## + Cirrhosis 1 677.34
## + ALT 1 677.48
## + AST 1 677.50

```

```

## + Vascularinvasion      1 677.58
## + TNM                    1 677.60
## + Capsulation            1 677.60
## + Tumorsize              1 677.60
## - BCLC                   1 678.24
## - Recurrence             1 762.84
## - OS                      1 779.34
##
## Step: AIC=674.28
## Surv(RFS, Death) ~ Age + HBsAg + AFP + Tumordifferentiation +
##      BCLC + OS + Recurrence + CXCL17N
##
##              Df      AIC
## - Tumordifferentiation  1 673.21
## - HBsAg                 1 673.34
## - AFP                   1 673.34
## - CXCL17N               1 673.34
## <none>                   674.28
## + CXCL17P               1 675.60
## + Tumormultiplicity     1 675.97
## + CXCL17T               1 676.14
## + ALT                   1 676.19
## + Cirrhosis             1 676.21
## + TNM                   1 676.23
## + AST                   1 676.25
## + Capsulation           1 676.28
## + Tumorsize             1 676.28
## + Vascularinvasion      1 676.28
## - BCLC                   1 677.85
## - Recurrence            1 761.68
## - OS                    1 778.64
##
## Step: AIC=673.21
## Surv(RFS, Death) ~ Age + HBsAg + AFP + BCLC + OS + Recurrence +
##      CXCL17N
##
##              Df      AIC
## - CXCL17N               1 671.99
## - AFP                   1 672.13
## - HBsAg                 1 672.56
## <none>                   673.21
## + Tumordifferentiation  1 674.28
## + CXCL17P               1 674.34
## + Tumormultiplicity     1 675.09
## + ALT                   1 675.12
## + TNM                   1 675.14
## + Cirrhosis             1 675.16
## + CXCL17T               1 675.17
## + AST                   1 675.17
## + Tumorsize             1 675.19

```

```

## + Vascularinvasion      1 675.21
## + Capsulation           1 675.21
## - BCLC                  1 675.86
## - Recurrence            1 759.68
## - OS                    1 782.36
##
## Step: AIC=671.99
## Surv(RFS, Death) ~ Age + HBsAg + AFP + BCLC + OS + Recurrence
##
##              Df    AIC
## - AFP          1 670.93
## - HBsAg        1 671.40
## <none>          671.99
## + CXCL17N      1 673.21
## + Tumordifferentiation 1 673.34
## + CXCL17T      1 673.73
## + Tumormultiplicity 1 673.83
## + Cirrhosis    1 673.88
## + ALT          1 673.93
## + AST          1 673.94
## + CXCL17P      1 673.97
## + TNM          1 673.98
## + Tumorsize    1 673.98
## + Capsulation  1 673.99
## + Vascularinvasion 1 673.99
## - BCLC         1 674.33
## - Recurrence   1 758.80
## - OS           1 783.07
##
## Step: AIC=670.93
## Surv(RFS, Death) ~ Age + HBsAg + BCLC + OS + Recurrence
##
##              Df    AIC
## - HBsAg        1 670.16
## <none>          670.93
## + AFP          1 671.99
## + CXCL17N      1 672.13
## + Tumordifferentiation 1 672.39
## + Tumormultiplicity 1 672.61
## + CXCL17T      1 672.80
## + Cirrhosis    1 672.84
## + AST          1 672.87
## + ALT          1 672.88
## + CXCL17P      1 672.88
## + Vascularinvasion 1 672.90
## + TNM          1 672.92
## + Capsulation  1 672.93
## + Tumorsize    1 672.93
## - BCLC         1 672.98
## - Recurrence   1 760.49

```

```
## - OS 1 784.73
##
## Step: AIC=670.16
## Surv(RFS, Death) ~ Age + BCLC + OS + Recurrence
##
## Df AIC
## <none> 670.16
## + HBsAg 1 670.93
## + CXCL17N 1 671.30
## + Tumordifferentiation 1 671.40
## + AFP 1 671.40
## - BCLC 1 671.69
## + Tumormultiplicity 1 671.94
## + Cirrhosis 1 671.97
## + CXCL17T 1 672.04
## + AST 1 672.07
## + Capsulation 1 672.11
## + ALT 1 672.12
## + Vascularinvasion 1 672.12
## + TNM 1 672.14
## + CXCL17P 1 672.14
## + Tumorsize 1 672.14
## - Recurrence 1 758.53
## - OS 1 782.89
```

Without OS Included

```
modelAll2n.coxph <- suppressWarnings(coxph(Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP + Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P + CXCL17N))
```

```
result.step2n <- suppressWarnings(step(modelAll2n.coxph, scope=list(upper=~ Age + HBsAg + Cirrhosis + ALT + AST + AFP + Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P + CXCL17N, lower=~Age)))
```

```
## Start: AIC=785.8
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
## Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
## Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17P +
## CXCL17N
##
## Df AIC
## - CXCL17P 1 783.80
## - Tumorsize 1 783.83
## - HBsAg 1 783.88
## - Cirrhosis 1 783.88
## - ALT 1 783.92
## - AST 1 784.24
```

```

## - CXCL17N          1 784.39
## - AFP              1 785.30
## - BCLC             1 785.30
## - CXCL17T          1 785.38
## - Capsulation      1 785.56
## - Vascularinvasion 1 785.64
## - Tumormultiplicity 1 785.75
## <none>              785.80
## - TNM              1 786.31
## - Tumordifferentiation 1 794.11
## - Recurrence       1 941.17
##
## Step: AIC=783.8
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
##      Tumorsize + Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##      Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17N
##
##              Df      AIC
## - Tumorsize      1 781.83
## - HBsAg          1 781.88
## - Cirrhosis      1 781.88
## - ALT            1 781.93
## - AST            1 782.25
## - CXCL17N        1 782.71
## - AFP            1 783.32
## - BCLC           1 783.40
## - Capsulation    1 783.56
## - CXCL17T        1 783.59
## - Vascularinvasion 1 783.70
## <none>            783.80
## - Tumormultiplicity 1 783.85
## - TNM            1 784.47
## + CXCL17P        1 785.80
## - Tumordifferentiation 1 792.11
## - Recurrence     1 946.79
##
## Step: AIC=781.83
## Surv(RFS, Death) ~ Age + HBsAg + Cirrhosis + ALT + AST + AFP +
##      Tumordifferentiation + Vascularinvasion + Tumormultiplicity +
##      Capsulation + TNM + BCLC + Recurrence + CXCL17T + CXCL17N
##
##              Df      AIC
## - HBsAg          1 779.92
## - Cirrhosis      1 779.93
## - ALT            1 779.95
## - AST            1 780.28
## - CXCL17N        1 780.77
## - AFP            1 781.39
## - BCLC           1 781.42
## - Capsulation    1 781.56

```

```

## - CXCL17T          1 781.59
## - Vascularinvasion 1 781.74
## <none>              781.83
## - Tumormultiplicity 1 781.87
## - TNM              1 783.32
## + Tumorsize        1 783.80
## + CXCL17P          1 783.83
## - Tumordifferentiation 1 790.12
## - Recurrence       1 947.82
##
## Step: AIC=779.92
## Surv(RFS, Death) ~ Age + Cirrhosis + ALT + AST + AFP + Tumordifferentiation +
##      Vascularinvasion + Tumormultiplicity + Capsulation + TNM +
##      BCLC + Recurrence + CXCL17T + CXCL17N
##
##
##      Df      AIC
## - Cirrhosis      1 778.01
## - ALT            1 778.06
## - AST            1 778.38
## - CXCL17N        1 778.90
## - AFP            1 779.47
## - BCLC           1 779.48
## - Capsulation    1 779.57
## - CXCL17T        1 779.60
## - Vascularinvasion 1 779.83
## - Tumormultiplicity 1 779.91
## <none>            779.92
## + HBsAg          1 781.83
## + Tumorsize      1 781.88
## - TNM            1 781.89
## + CXCL17P        1 781.92
## - Tumordifferentiation 1 788.38
## - Recurrence     1 946.27
##
## Step: AIC=778.01
## Surv(RFS, Death) ~ Age + ALT + AST + AFP + Tumordifferentiation +
##      Vascularinvasion + Tumormultiplicity + Capsulation + TNM +
##      BCLC + Recurrence + CXCL17T + CXCL17N
##
##
##      Df      AIC
## - ALT            1 776.16
## - AST            1 776.42
## - CXCL17N        1 776.92
## - AFP            1 777.61
## - BCLC           1 777.64
## - CXCL17T        1 777.73
## - Vascularinvasion 1 777.85
## - Capsulation    1 777.85
## - Tumormultiplicity 1 777.95
## <none>            778.01

```

```

## + Cirrhosis          1 779.92
## + HBsAg              1 779.93
## + Tumorsize          1 779.95
## + CXCL17P            1 780.01
## - TNM                1 780.14
## - Tumordifferentiation 1 786.42
## - Recurrence         1 944.35
##
## Step: AIC=776.16
## Surv(RFS, Death) ~ Age + AST + AFP + Tumordifferentiation + Vascularinvasion +
##   Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence +
##   CXCL17T + CXCL17N
##
##              Df    AIC
## - AST          1 774.87
## - CXCL17N       1 775.21
## - AFP          1 775.71
## - BCLC          1 775.72
## - CXCL17T       1 775.81
## - Vascularinvasion 1 775.95
## - Capsulation   1 775.96
## <none>          776.16
## - Tumormultiplicity 1 776.28
## + ALT          1 778.01
## + HBsAg        1 778.06
## + Cirrhosis    1 778.06
## + Tumorsize    1 778.11
## + CXCL17P      1 778.16
## - TNM          1 778.25
## - Tumordifferentiation 1 784.44
## - Recurrence   1 942.35
##
## Step: AIC=774.87
## Surv(RFS, Death) ~ Age + AFP + Tumordifferentiation + Vascularinvasion +
##   Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence +
##   CXCL17T + CXCL17N
##
##              Df    AIC
## - CXCL17N       1 773.77
## - AFP          1 774.44
## - BCLC          1 774.49
## - CXCL17T       1 774.65
## - Vascularinvasion 1 774.74
## <none>          774.87
## - Tumormultiplicity 1 775.05
## - Capsulation   1 775.29
## + AST          1 776.16
## + ALT          1 776.42
## + HBsAg        1 776.74
## + Cirrhosis    1 776.82

```



```

## + Tumorsize          1 776.83
## + CXCL17P            1 776.85
## - TNM                1 777.15
## - Tumordifferentiation 1 783.30
## - Recurrence         1 943.67
##
## Step: AIC=773.77
## Surv(RFS, Death) ~ Age + AFP + Tumordifferentiation + Vascularinvasion +
##   Tumormultiplicity + Capsulation + TNM + BCLC + Recurrence +
##   CXCL17T
##
##              Df    AIC
## - BCLC          1 773.26
## - Vascularinvasion 1 773.58
## - AFP           1 773.64
## <none>           773.77
## - Tumormultiplicity 1 773.85
## - Capsulation      1 774.21
## + CXCL17N         1 774.87
## - CXCL17T         1 775.16
## + ALT             1 775.18
## + AST             1 775.21
## + CXCL17P         1 775.38
## + HBsAg           1 775.59
## + Tumorsize       1 775.69
## + Cirrhosis       1 775.76
## - TNM             1 776.26
## - Tumordifferentiation 1 782.08
## - Recurrence      1 945.30
##
## Step: AIC=773.26
## Surv(RFS, Death) ~ Age + AFP + Tumordifferentiation + Vascularinvasion +
##   Tumormultiplicity + Capsulation + TNM + Recurrence + CXCL17T
##
##              Df    AIC
## - AFP           1 772.64
## <none>           773.26
## + BCLC          1 773.77
## - CXCL17T       1 774.08
## - Capsulation   1 774.21
## + CXCL17N       1 774.49
## + AST           1 774.65
## + ALT           1 774.80
## + HBsAg         1 775.14
## + Tumorsize     1 775.18
## + CXCL17P       1 775.20
## + Cirrhosis     1 775.23
## - TNM           1 777.93
## - Tumormultiplicity 1 778.04
## - Vascularinvasion 1 778.31

```

```

## - Tumordifferentiation 1 781.87
## - Recurrence          1 946.01
##
## Step: AIC=772.64
## Surv(RFS, Death) ~ Age + Tumordifferentiation + Vascularinvasion +
##      Tumormultiplicity + Capsulation + TNM + Recurrence + CXCL17T
##
##              Df    AIC
## - CXCL17T      1 772.53
## <none>          772.64
## + AFP          1 773.26
## - Capsulation  1 773.60
## + CXCL17N      1 773.60
## + BCLC         1 773.64
## + AST          1 774.07
## + ALT          1 774.22
## + Tumorsize    1 774.48
## + HBsAg        1 774.53
## + CXCL17P      1 774.58
## + Cirrhosis    1 774.60
## - TNM          1 777.45
## - Vascularinvasion 1 778.95
## - Tumormultiplicity 1 779.29
## - Tumordifferentiation 1 782.19
## - Recurrence   1 953.48
##
## Step: AIC=772.53
## Surv(RFS, Death) ~ Age + Tumordifferentiation + Vascularinvasion +
##      Tumormultiplicity + Capsulation + TNM + Recurrence
##
##              Df    AIC
## + CXCL17N      1 772.33
## <none>          772.53
## + CXCL17T      1 772.64
## - Capsulation  1 773.47
## + CXCL17P      1 773.65
## + BCLC         1 773.77
## + AST          1 773.88
## + AFP          1 774.08
## + ALT          1 774.16
## + Cirrhosis    1 774.50
## + Tumorsize    1 774.51
## + HBsAg        1 774.52
## - TNM          1 777.80
## - Tumormultiplicity 1 778.73
## - Vascularinvasion 1 779.56
## - Tumordifferentiation 1 781.56
## - Recurrence   1 959.33
##
## Step: AIC=772.33

```

```
## Surv(RFS, Death) ~ Age + Tumordifferentiation + Vascularinvasion +
##   Tumormultiplicity + Capsulation + TNM + Recurrence + CXCL17N
##
##               Df      AIC
## <none>                772.33
## - CXCL17N             1 772.53
## - Capsulation         1 773.24
## + BCLC                 1 773.30
## + AST                  1 773.44
## + CXCL17T             1 773.60
## + AFP                  1 773.88
## + ALT                  1 774.08
## + Cirrhosis           1 774.18
## + Tumorsize           1 774.31
## + HBsAg               1 774.32
## + CXCL17P             1 774.33
## - TNM                  1 776.95
## - Tumormultiplicity   1 779.17
## - Vascularinvasion    1 779.47
## - Tumordifferentiation 1 781.66
## - Recurrence          1 953.67
```

With OS Included

```
summary(result.step2)
```

```
## Call:
## coxph(formula = Surv(RFS, Death) ~ Age + BCLC + OS + Recurrence)
##
##   n= 227, number of events= 97
##
##               coef exp(coef) se(coef)      z Pr(>|z|)
## Age           2.410e-03 1.002e+00 8.389e-03 0.287  0.774
## BCLC           4.237e-01 1.528e+00 2.262e-01 1.873  0.061 .
## OS            -9.582e-02 9.086e-01 1.203e-02 -7.962 1.69e-15 ***
## Recurrence    2.027e+01 6.375e+08 2.400e+03 0.008  0.993
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##               exp(coef) exp(-coef) lower .95 upper .95
## Age           1.002e+00 9.976e-01 0.9861 1.0190
## BCLC           1.528e+00 6.546e-01 0.9806 2.3797
## OS            9.086e-01 1.101e+00 0.8874 0.9303
## Recurrence    6.375e+08 1.569e-09 0.0000 Inf
##
## Concordance= 0.923 (se = 0.011 )
## Likelihood ratio test= 314.5 on 4 df, p=<2e-16
## Wald test              = 76.38 on 4 df, p=1e-15
## Score (logrank) test = 239 on 4 df, p=<2e-16
```

Without OS Included

result.step2n

Call:

coxph(formula = Surv(RFS, Death) ~ Age + Tumordifferentiation +
Vascularinvasion + Tumormultiplicity + Capsulation + TNM +
Recurrence + CXCL17N)

	coef	exp(coef)	se(coef)	z	p
## Age	-1.082e-02	9.892e-01	8.320e-03	-1.301	0.193380
## Tumordifferentiation	7.655e-01	2.150e+00	2.261e-01	3.385	0.000711
## Vascularinvasion	7.959e-01	2.216e+00	2.481e-01	3.208	0.001338
## Tumormultiplicity	7.570e-01	2.132e+00	2.498e-01	3.030	0.002446
## Capsulation	-4.383e-01	6.451e-01	2.508e-01	-1.748	0.080533
## TNM	6.127e-01	1.845e+00	2.385e-01	2.568	0.010216
## Recurrence	2.147e+01	2.109e+09	2.392e+03	0.009	0.992838
## CXCL17N	7.706e-04	1.001e+00	4.861e-04	1.585	0.112888

##

Likelihood ratio test=220.3 on 8 df, p=< 2.2e-16

n= 227, number of events= 97

#####

7.1. Consider the case deletion and dfbeta residuals discussed in Sect. 7.1.2. For each of the covariates in the final pharmacoSmoking model (grp, employment Level s 2 and 3 vs. 1, and age), plot the case deletion residuals versus the dfbeta residuals. Also plot the “dfbeta” residuals versus the “dfbetas” residuals. Do you see any differences?

#####

library(survival)

library(asauro)

library(MASS)

library(survminer) *# Load the package*

Warning: package 'survminer' was built under R version 4.2.3

Loading required package: ggplot2

Loading required package: ggpubr

##

Attaching package: 'survminer'

The following object is masked from 'package:survival':

##

myeloma

library(ggfortify)

attach(pharmacoSmoking)

```

library(ggplot2)
head(pharmacoSmoking)

##      id ttr relapse      grp age gender      race employment yearsSmoking
## 1  21 182      0  patchOnly 36  Male    white      ft          26
## 2 113  14      1  patchOnly 41  Male    white    other          27
## 3  39   5      1 combination 25 Female  white    other          12
## 4  80  16      1 combination 54  Male    white      ft          39
## 5  87   0      1 combination 45  Male    white    other          30
## 6  29 182      0 combination 43  Male  hispanic      ft          30
##      levelSmoking ageGroup2 ageGroup4 priorAttempts longestNoSmoke
## 1      heavy      21-49      35-49          0          0
## 2      heavy      21-49      35-49          3          90
## 3      heavy      21-49      21-34          3          21
## 4      heavy      50+      50-64          0          0
## 5      heavy      21-49      35-49          0          0
## 6      heavy      21-49      35-49          2        1825

result.coxph <- coxph(Surv(ttr, relapse) ~ grp + employment
+ age)
result.coxph

## Call:
## coxph(formula = Surv(ttr, relapse) ~ grp + employment + age)
##
##              coef exp(coef) se(coef)      z      p
## grppatchOnly    0.60788   1.83654  0.21837  2.784 0.00537
## employmentother 0.70348   2.02077  0.26929  2.612 0.00899
## employmentpt    0.65369   1.92262  0.32732  1.997 0.04581
## age            -0.03529   0.96533  0.01075 -3.282 0.00103
##
## Likelihood ratio test=22.03 on 4 df, p=0.0001979
## n= 125, number of events= 89

coef.all_1 <- result.coxph$coef[1]
coef.all_1

## grppatchOnly
##      0.607884

n.obs1 <- length(ttr)
jkbeta.vec1 <- rep(NA, n.obs1)
for (i in 1:n.obs1) {
  tt.i <- ttr[-i]
  delta.i <- relapse[-i]
  grp.i <- grp[-i]
  employment.i <- employment[-i]
  age.i <- age[-i]
  result.coxph.i <- coxph(Surv(tt.i, delta.i) ~ grp.i +
employment.i + age.i)
  coef.i <- result.coxph.i$coef[1]

```

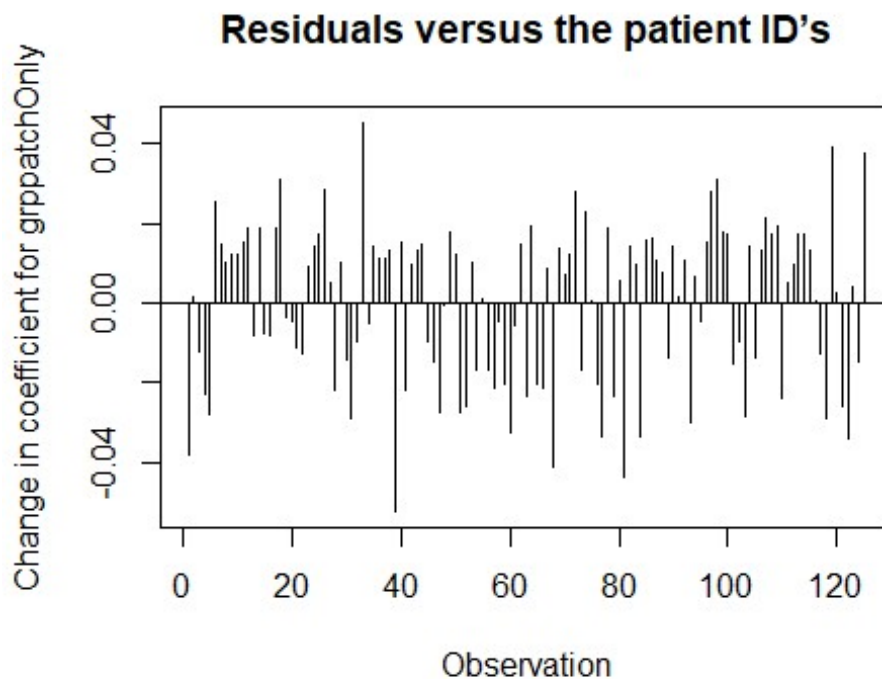
```

jkbeta.vec1[i] <- (coef.all_1 - coef.i)
}

index.obs1 <- 1:n.obs1
plot(jkbeta.vec1 ~ index.obs1, type="h", main="Residuals versus the patient ID's",
xlab="Observation", ylab="Change in coefficient for grppatchOnly")
abline(h=0)

identify(jkbeta.vec1 ~ index.obs1)

```

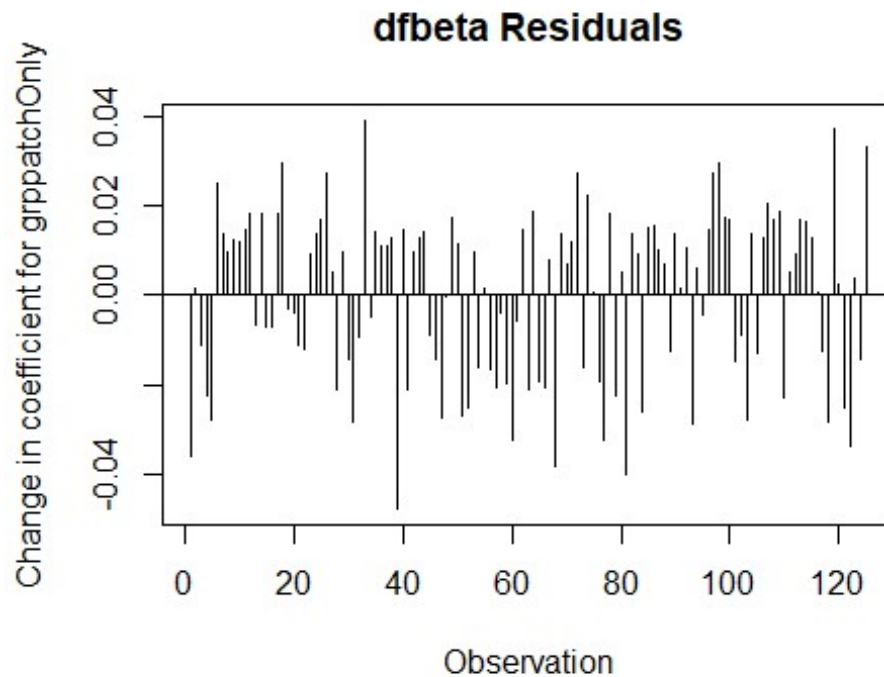


```

## integer(0)

resid.dfbeta1 <- residuals(result.coxph, type="dfbeta")
n.obs <- length(ttr)
index.obs1 <- 1:n.obs1
plot(resid.dfbeta1[,1] ~ index.obs1, type="h", main= "dfbeta Residuals",
xlab="Observation", ylab="Change in coefficient for grppatchOnly")
abline(h=0)
identify(resid.dfbeta1[,1] ~ index.obs1)

```



```
## integer(0)

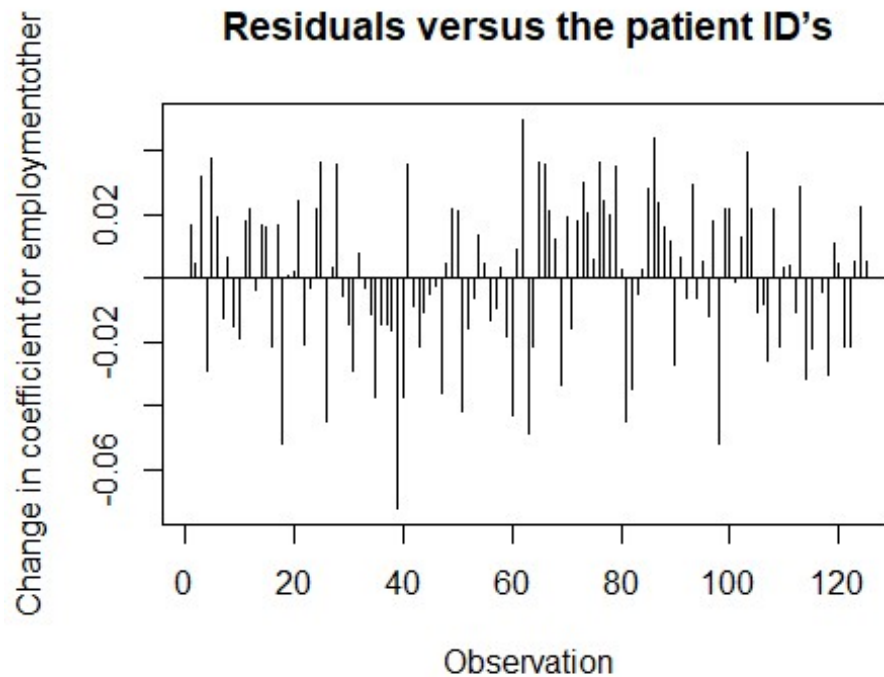
coef.all_2 <- result.coxph$coef[2]
coef.all_2

## employmenttother
##      0.7034766

n.obs2 <- length(ttr)
jkbeta.vec2 <- rep(NA, n.obs2)
for (i in 1:n.obs2) {
  tt.i <- ttr[-i]
  delta.i <- relapse[-i]
  grp.i <- grp[-i]
  employment.i <- employment[-i]
  age.i <- age[-i]
  result.coxph.i <- coxph(Surv(tt.i, delta.i) ~ grp.i +
    employment.i + age.i)
  coef.i <- result.coxph.i$coef[2]
  jkbeta.vec2[i] <- (coef.all_2 - coef.i)
}

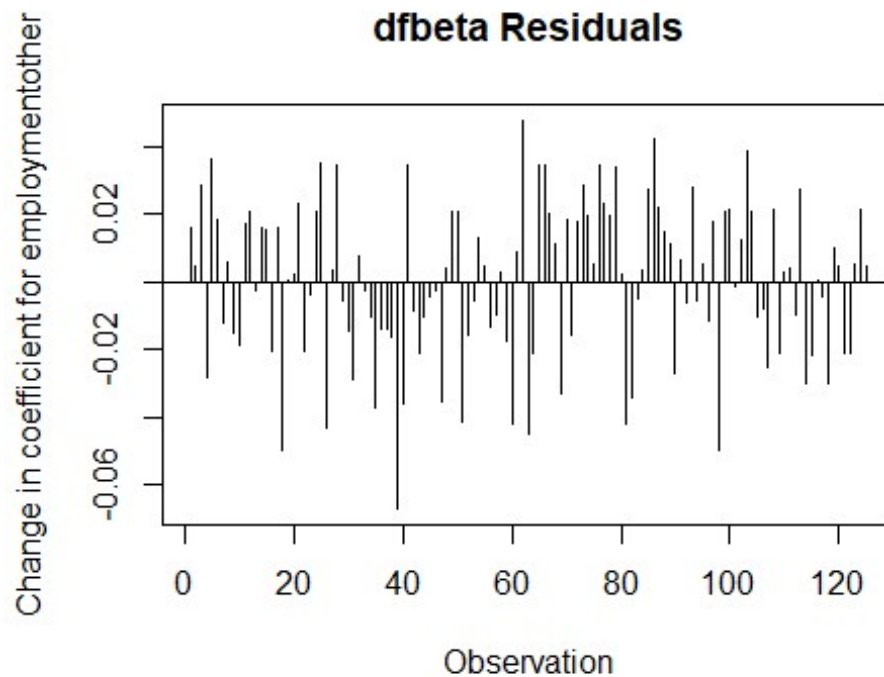
index.obs2 <- 1:n.obs2
plot(jkbeta.vec2 ~ index.obs2, type="h",
  xlab="Observation", ylab="Change in coefficient for employmenttother",
  main="Residuals versus the patient ID's")
abline(h=0)
```

```
identify(jkbeta.vec2 ~ index.obs2)
```



```
## integer(0)

resid.dfbeta2 <- residuals(result.coxph, type="dfbeta")
n.obs <- length(ttr)
index.obs2 <- 1:n.obs2
plot(resid.dfbeta2[,2] ~ index.obs2, type="h", main= "dfbeta Residuals",
xlab="Observation", ylab="Change in coefficient for employmenttother")
abline(h=0)
identify(resid.dfbeta2[,2] ~ index.obs2)
```

```
## integer(0)

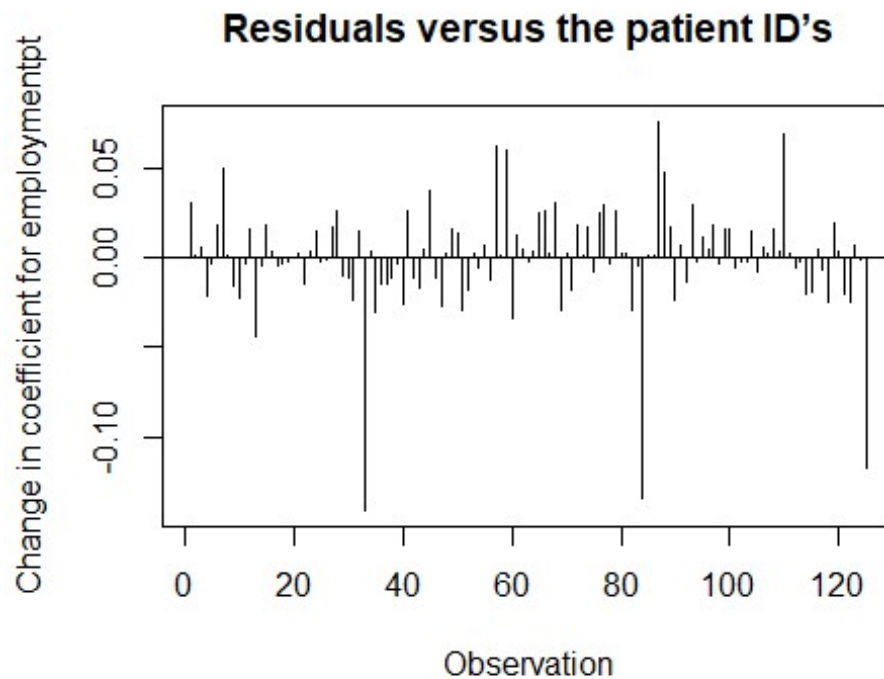
coef.all_3 <- result.coxph$coef[3]
coef.all_3

## employmentpt
##      0.6536902

n.obs3 <- length(ttr)
jkbeta.vec3 <- rep(NA, n.obs3)
for (i in 1:n.obs3) {
  tt.i <- ttr[-i]
  delta.i <- relapse[-i]
  grp.i <- grp[-i]
  employment.i <- employment[-i]
  age.i <- age[-i]
  result.coxph.i <- coxph(Surv(tt.i, delta.i) ~ grp.i +
    employment.i + age.i)
  coef.i <- result.coxph.i$coef[3]
  jkbeta.vec3[i] <- (coef.all_3 - coef.i)
}

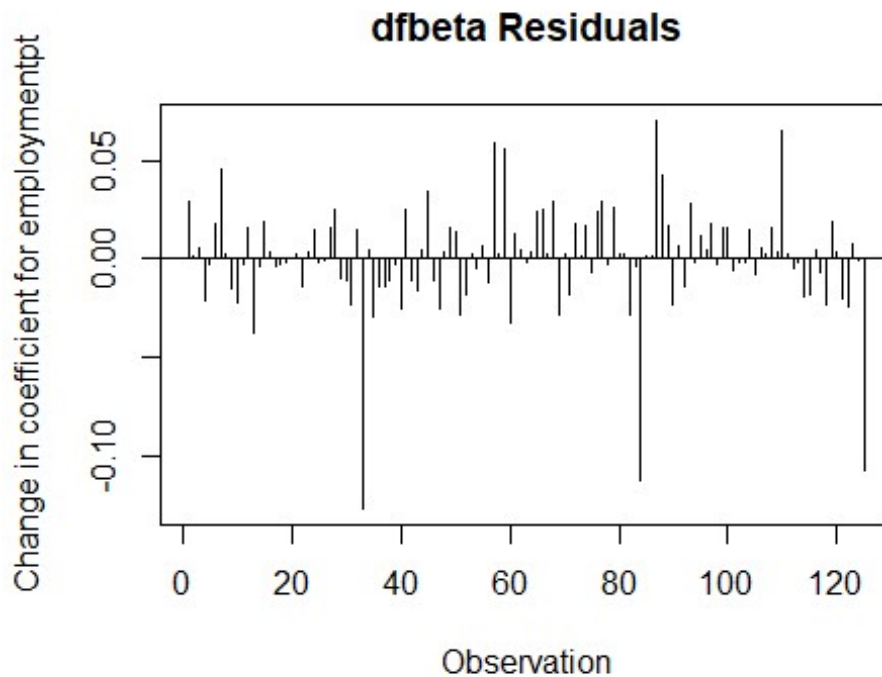
index.obs3 <- 1:n.obs3
plot(jkbeta.vec3 ~ index.obs3, type="h",
  xlab="Observation", ylab="Change in coefficient for employmentpt",
  main="Residuals versus the patient ID's")
abline(h=0)
```

```
identify(jkbeta.vec3 ~ index.obs3)
```



```
## integer(0)

resid.dfbeta3 <- residuals(result.coxph, type="dfbeta")
n.obs <- length(ttr)
index.obs3 <- 1:n.obs3
plot(resid.dfbeta3[,3] ~ index.obs3, type="h", main= "dfbeta Residuals",
xlab="Observation", ylab="Change in coefficient for employmentpt")
abline(h=0)
identify(resid.dfbeta3[,3] ~ index.obs3)
```



```
## integer(0)

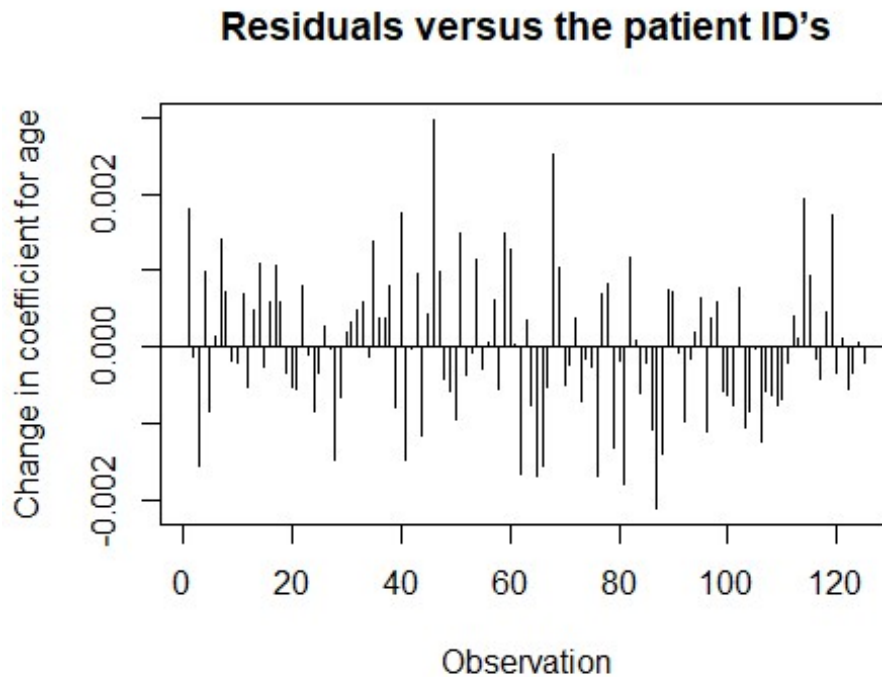
coef.all_4 <- result.coxph$coef[4]
coef.all_4

##          age
## -0.03528934

n.obs4 <- length(ttr)
jkbeta.vec4 <- rep(NA, n.obs4)
for (i in 1:n.obs4) {
  tt.i <- ttr[-i]
  delta.i <- relapse[-i]
  grp.i <- grp[-i]
  employment.i <- employment[-i]
  age.i <- age[-i]
  result.coxph.i <- coxph(Surv(tt.i, delta.i) ~ grp.i +
    employment.i + age.i)
  coef.i <- result.coxph.i$coef[4]
  jkbeta.vec4[i] <- (coef.all_4 - coef.i)
}

index.obs4 <- 1:n.obs4
plot(jkbeta.vec4 ~ index.obs4, type="h", main="Residuals versus the patient ID's",
  xlab="Observation", ylab="Change in coefficient for age"
)
abline(h=0)
```

```
identify(jkbeta.vec4 ~ index.obs4)
```



```
## integer(0)
```

```
resid.dfbeta4 <- residuals(result.coxph, type="dfbeta")
```

```
n.obs <- length(ttr)
```

```
index.obs4 <- 1:n.obs4
```

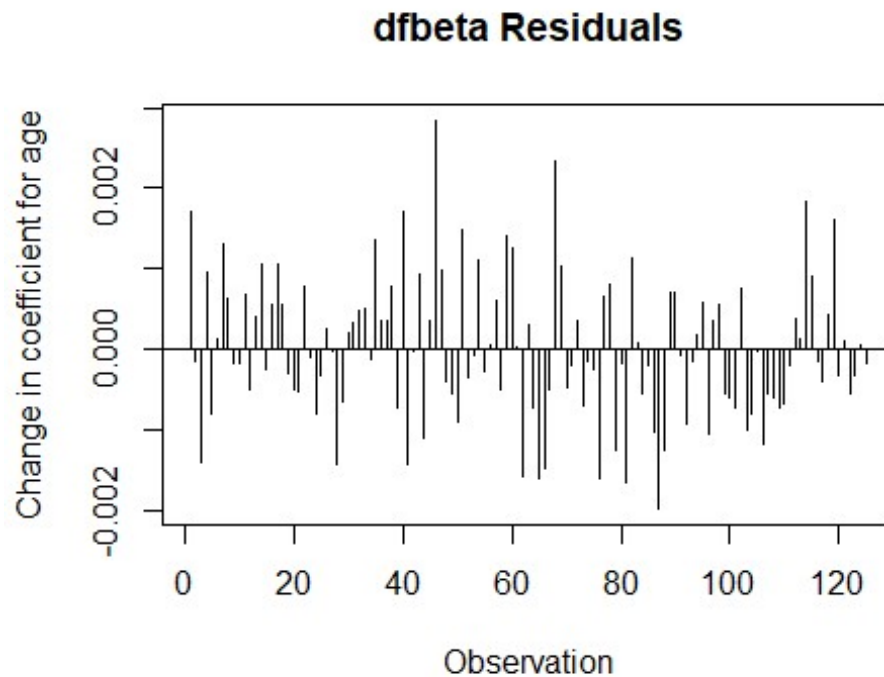
```
head(resid.dfbeta4)
```

```
##           [,1]      [,2]      [,3]      [,4]
## 1 -0.035992243  0.016039067  0.029442996  0.0017000609
## 2  0.001489819  0.004893592  0.001746782 -0.0001361549
## 3 -0.011033222  0.028769501  0.005825861 -0.0014034177
## 4 -0.022264005 -0.028321158 -0.021135909  0.0009633467
## 5 -0.027593571  0.036400215 -0.002712807 -0.0007943478
## 6  0.025089714  0.018742984  0.017774738  0.0001408364
```

```
plot(resid.dfbeta4[,4] ~ index.obs4, type="h", main= "dfbeta Residuals",
xlab="Observation", ylab="Change in coefficient for age")
```

```
abline(h=0)
```

```
identify(resid.dfbeta4[,4] ~ index.obs4)
```

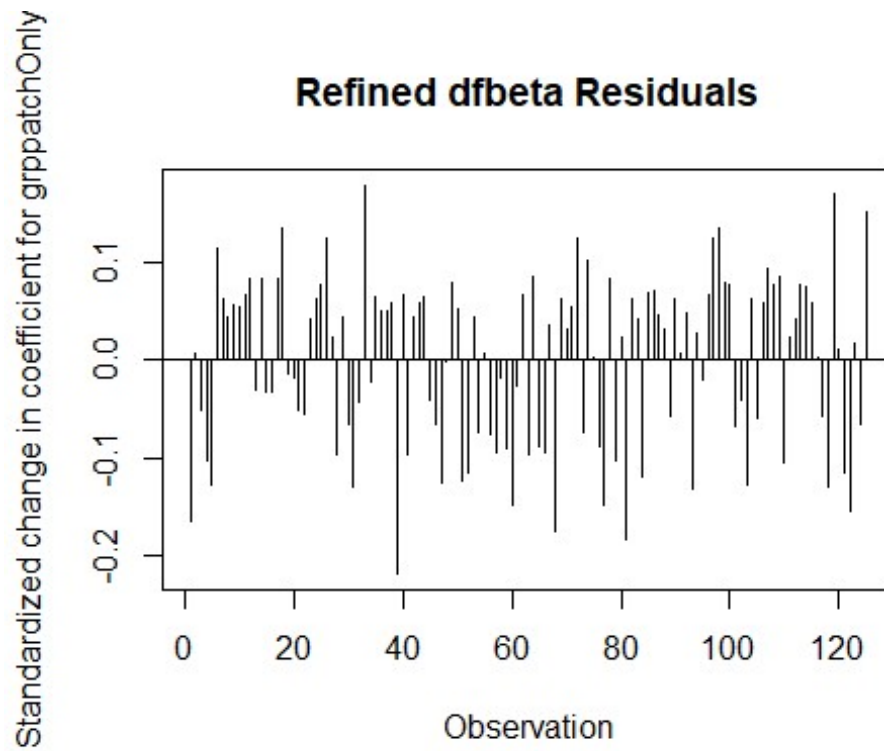


```
## integer(0)

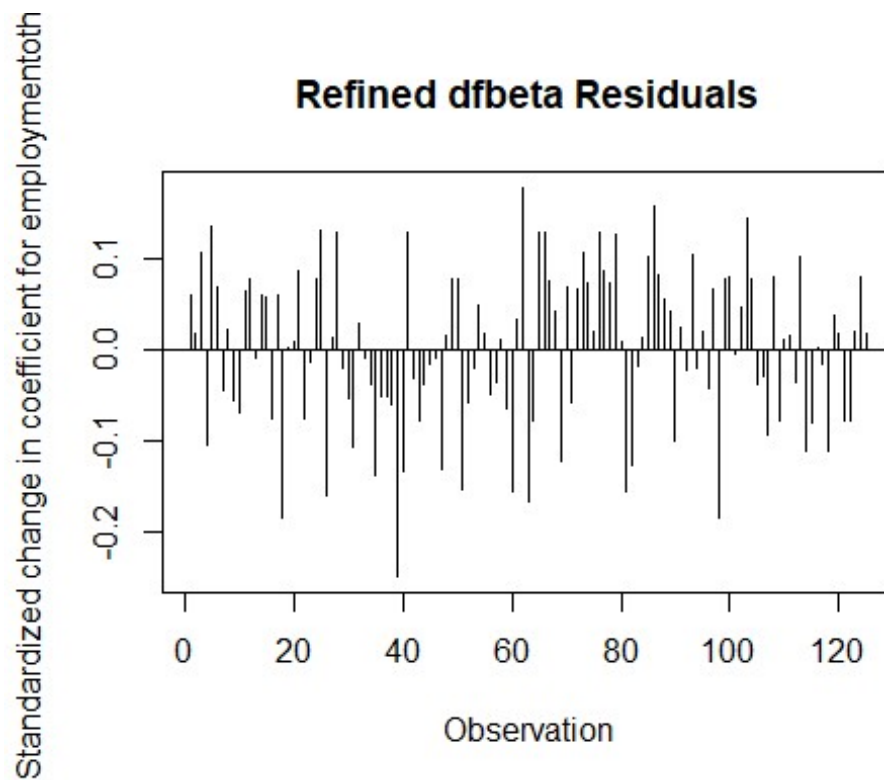
resid.dfbeta <- residuals(result.coxph, type="dfbetas")
n.obs <- length(ttr)
index.obs <- 1:n.obs
head(resid.dfbeta)

##           [,1]      [,2]      [,3]      [,4]
## 1 -0.164823287  0.05955969  0.089952431  0.15809557
## 2  0.006822493  0.01817193  0.005336660 -0.01266160
## 3 -0.050525661  0.10683305  0.017798812 -0.13050951
## 4 -0.101956037 -0.10516817 -0.064573127  0.08958552
## 5 -0.126362316  0.13516905 -0.008288001 -0.07386963
## 6  0.114896122  0.06960045  0.054304287  0.01309695

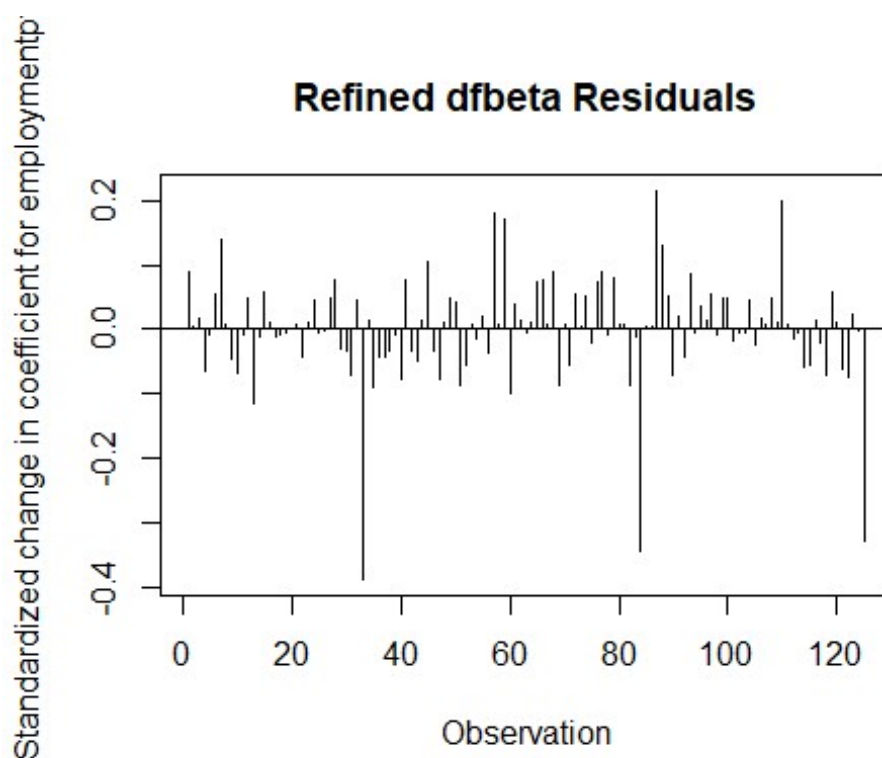
plot(resid.dfbeta[,1] ~ index.obs, type="h", main= "Refined dfbeta Residuals",
xlab="Observation", ylab="Standardized change in coefficient for grppatchOnly")
abline(h=0)
```



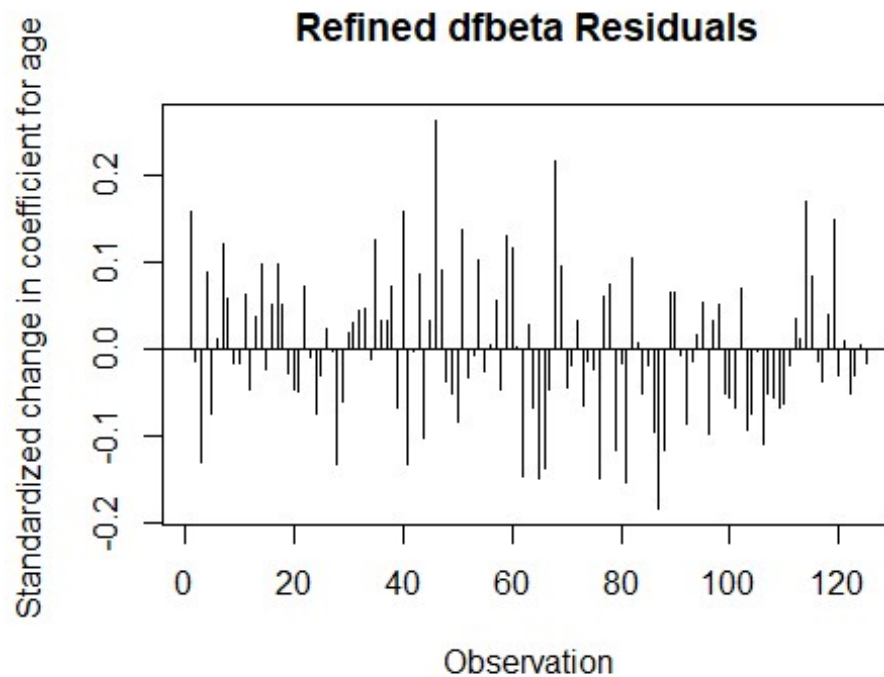
```
plot(resid.dfbeta[,2] ~ index.obs, type="h", main= "Refined dfbeta Residuals",
xlab="Observation", ylab="Standardized change in coefficient for employmentother")
abline(h=0)
```



```
plot(resid.dfbeta[,3] ~ index.obs, type="h", main= "Refined dfbeta Residuals",
xlab="Observation", ylab="Standardized change in coefficient for employmentpt")
abline(h=0)
```



```
plot(resid.dfbeta[,4] ~ index.obs, type="h", main= "Refined dfbeta Residuals",
xlab="Observation", ylab="Standardized change in coefficient for age ")
abline(h=0)
```



The dfbeta residuals plots (not shown) are nearly identical to those plots of residuals versus the patient id's.

The refined dfbeta residuals scale the dfbeta residuals by an estimate of their standard error, making it easier to compare the magnitudes of the dfbeta residuals for different predictor variables.