

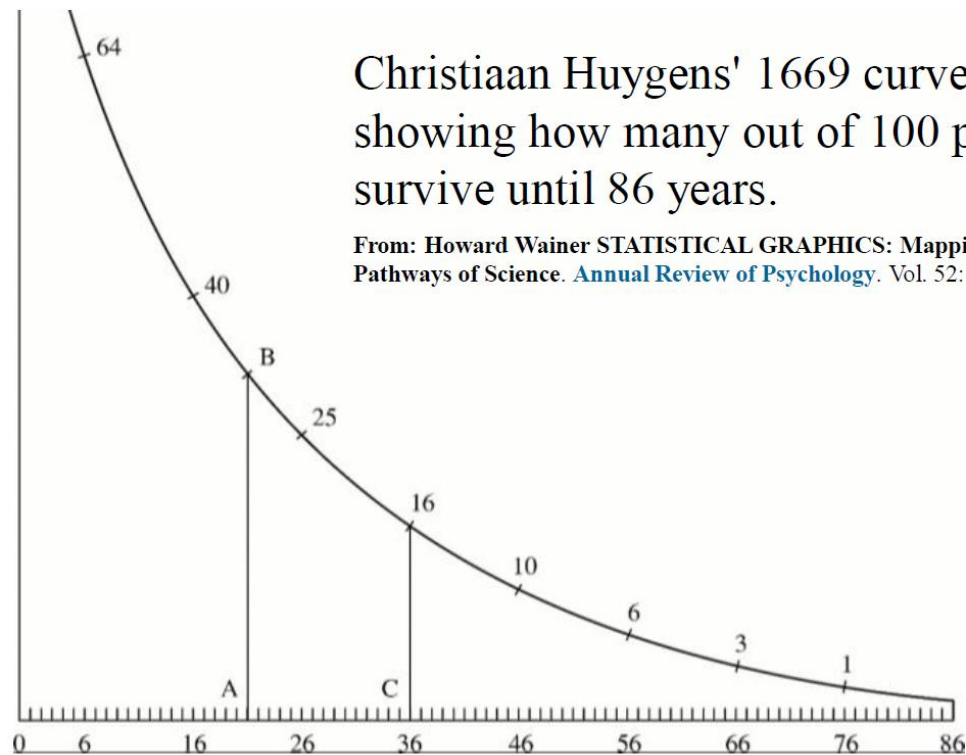
# Survival Analysis

*STATS – Modelação Estatística*

*PhD Programme in Health Data Science*

**Cristina Costa Santos & Andreia Teixeira**

# One of the first examples of survival analysis 1669



# What is survival analysis?

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- Statistical methods for analyzing **longitudinal time to event** data.
- **Events** may include: death, injury, onset of illness, recovery from illness or transition above or below the clinical threshold of a meaningful continuous variable.



# Objectives of survival analysis

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- **Estimate time-to-event for a group of individuals**, such as time until death for a group of cancer patients.
- **To compare time-to-event between two or more groups**, such as treated vs placebo cancer patients in a randomized controlled trial.
- **To assess the relationship of co-variables to time-to-event**, such as: does age, a certain biomarker, or tumor size influence survival time of cancer patients?

# Censoring

- The patients who did not have the event are considered censored.
  - We know that they survived a specific amount of time, but do not know the exact time of the event.
  - We believe that the event would have happened if we observed them long enough.
- These patients provide some information, but not complete information.

- **How could we account for censoring?**

- **Ignore it and say event occurred at time of censoring**
  - Incorrect because this is almost certainly not true
- **Remove patient from analysis**
  - Potential bias and loss of power
- **Survival analysis**

- **Our objective is to estimate the survival distribution of patients in the presence of censoring.**



**Censoring is taken into account in the analysis.**

# Why use survival analysis?



- **Why not compare mean time-to-event between groups using a t-test or linear regression?**
- **Why not compare proportion of events in groups using risk/odds ratios or logistic regression?**

# Why use survival analysis?



- **Why not compare mean time-to-event between groups using a t-test or linear regression?**
  - Ignores censoring
  
- **Why not compare proportion of events in groups using risk/odds ratios or logistic regression?**
  - Ignores time



# Why use survival analysis?



- In survival analysis the time until the occurrence of a well-defined event is recorded (time-to-event data).
  - Survival time
- If everyone had an event, some of the methods we have already learned could be applied.
- Often, not everyone has event - censoring
  - Loss to follow-up
  - End of study

# Survival analysis: Terms

- **Time-to-event:** the time from entry into a study until a subject has a particular outcome
- **Censoring:** subjects are said to be censored if they are lost to follow up or drop out of the study, or if they die of unrelated causes or the study ends before they have the event of interest. They are counted as alive or disease-free for the time they were enrolled in the study.

# Data Structure: survival analysis



## Two-variable outcome:

- **Time variable (T):**  $t_i$  = time at last event-free observation or time at event
- **Censoring variable (C):**  $c_i = 1$  if had the event;  $c_i = 0$  no event by time  $t_i$

# Introduction to survival distributions

- $T_i$  the event time for an individual, is a random variable having a probability distribution.
- Different models for survival data are distinguished by different choice of distribution for  $T_i$ .

- The probability (density) of the event time occurring at exactly time  $t$ :

$$f(t) = \lim_{\Delta t \rightarrow 0} \frac{P(t \leq T < t + \Delta t)}{\Delta t}$$

- The cumulative probability (density) of the event time occurring before time  $t$ :

$$F(t) = P(T \leq t) = \int_{-\infty}^t f(t) dt$$

# Survival function: $1-F(t)$

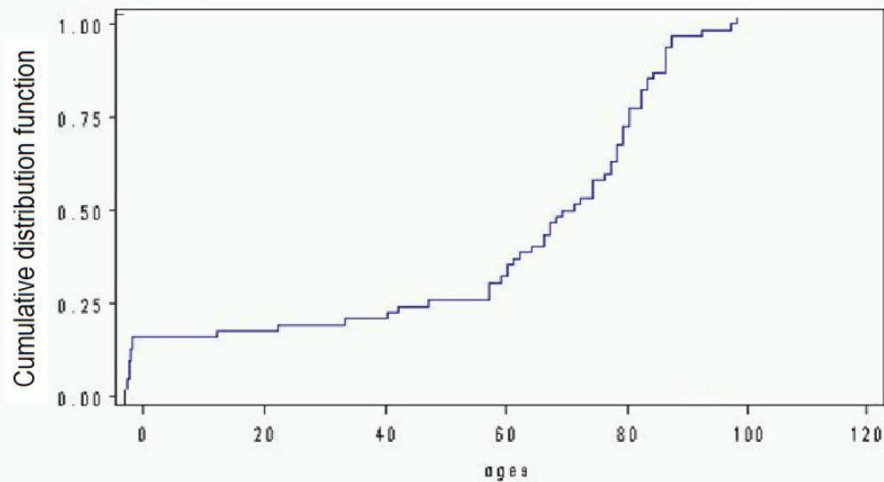
The goal of survival analysis is to estimate and compare survival experiences of different groups.

Survival experience is described by the cumulative survival function:

$$S(t) = 1 - P(T \leq t) = 1 - F(t)$$

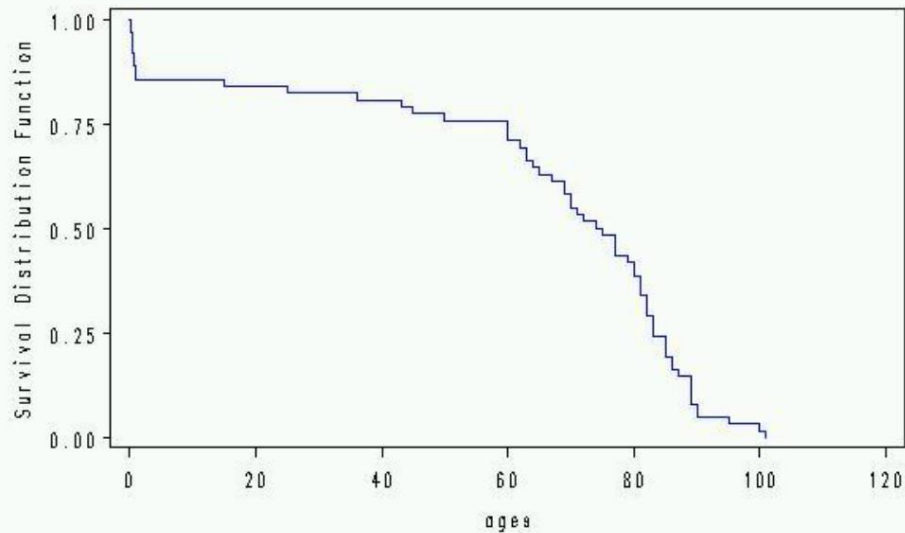
**Example:** If  $t = 100$  years,  $S(t = 100)$  = probability of surviving beyond 100 years.

Cumulative distribution function

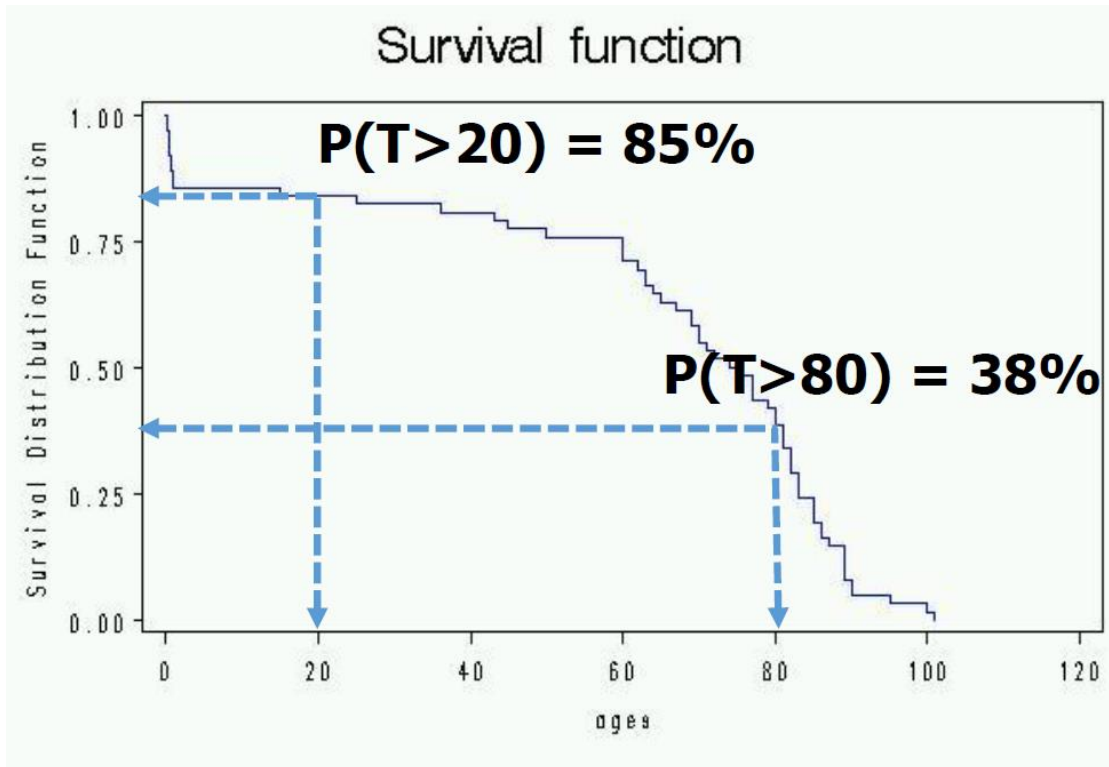


$F(t)$

Survival function



$S(t)$



# Hazard Function: new concept

- The probability that if you survive to  $t$ , you will succumb to the event in the next instant:

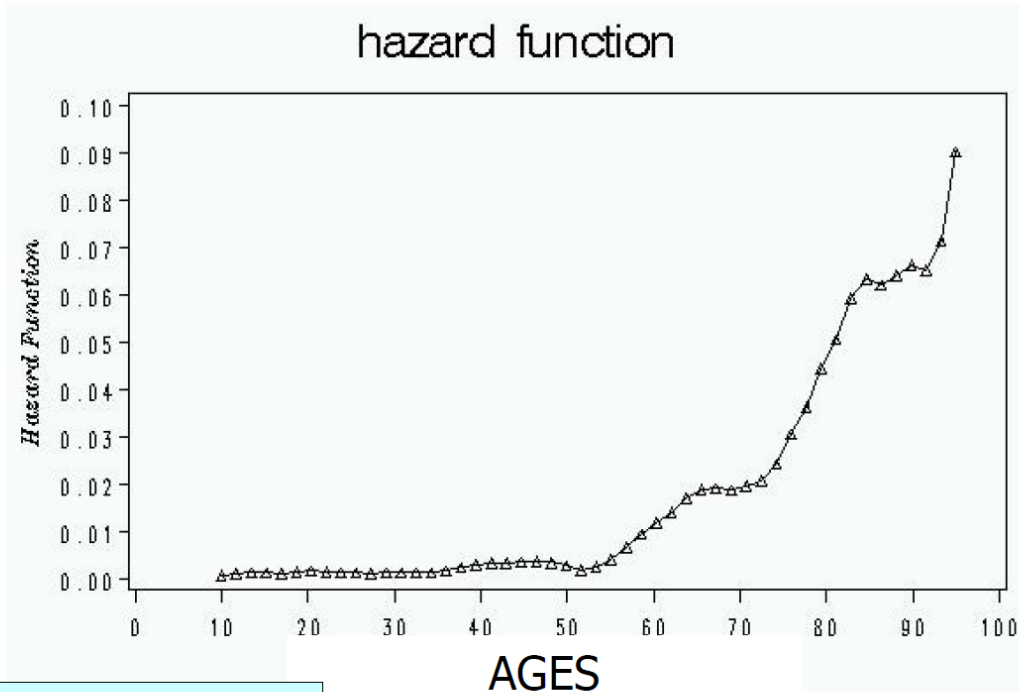
$$h(t) = \lim_{\Delta t \rightarrow 0} \frac{P(t \leq T < t + \Delta t | T \geq t)}{\Delta t}$$

$$h(t)dt = P(t \leq T < t + dt | T \geq t) = \frac{P(t \leq T < t + dt \wedge T \geq t)}{P(T \geq t)} = \frac{P(t \leq T < t + dt)}{P(T \geq t)} = \frac{f(t)dt}{S(t)} = \frac{-S'(t)dt}{S(t)}$$

$$h(t) = \frac{-S'(t)}{S(t)}$$



# Hazard Function: new concept



Hazard rate is an instantaneous incidence rate.

# Hazard vs Density example

- When you are born, you have a certain probability of dying at any age; that's the probability density

*(Example: a woman born today has, say, a 1% chance of dying at 80 years).*

- However, as you survive for awhile, your probabilities keep changing; that's the hazard rate (think: conditional probability)

*(Example: a woman who is 79 today has, say, a 5% chance of dying at 80 years).*

# Practical example

## Basics of Survival Analysis



In **R** (as with other packages) we require the following **two variables when dealing with survival time data**:

- A **continuous time variable** that measures the time until either the event or the individuals withdrawal (censoring).
- A **categorical variable that acts as an indicator for whether the subject experienced the event of interest or whether they did not and were censored.**

# Kaplan-Meier Analysis Example



Rather than categorising, we can estimate the survival function directly from the continuous survival times.

**File:** data2.csv

# Kaplan-Meier Analysis Example

The **variables** included in the data are:

- id: identification of the patient
- fu: survival time in months
- event: event observed [0 = censored; 1 = death]
- eventHD: event observed [0 = censored; 1 = transfer to haemodialysis]
- eventTR: event observed [0 = censored; 1 = renal transplant]
- eventCR: event observed considering a competing risks approach [0 = censored; 1 = death; 2 = transfer to haemodialysis; 3 = renal transplant]
- peritonitis: occurrence of a peritonites [0 = no; 1 = yes]
- sex [0 = female; 1 = male]
- age: in years
- diab: diabetes [0 = no; 1 = yes]
- first: prior treatment [0 = no; 1 = yes]

# Kaplan-Meier Analysis Example



#Load the library required for a survival analysis

```
library(splines)  
library(survival)
```

#Read data

```
data=read.csv("data2.csv", sep=";", dec=",")
```

# Kaplan-Meier Analysis Example

```
#Read data
```

```
data=read.csv("data2.csv", sep=";", dec=",")
```

```
> head(data)
```

	id	fu	event	event_HD	event_RT	event_CR	peritonitis	sex	age	diab	first
1	1	7	1	0	0	1	0	1	60	1	0
2	3	22	0	0	1	3	1	1	41	1	0
3	7	13	0	0	1	3	1	0	38	0	1
4	9	42	1	0	0	1	1	0	46	0	0
5	11	5	0	1	0	2	1	0	51	0	0
6	13	13	1	0	0	1	0	0	36	0	1

# Kaplan-Meier Analysis Example

#Provides the names of variables included in the database

```
> names(data)
[1] "id"      "fu"      "event"   "event_HD" "event_RT" "event_CR" "peritonitis"
[8] "sex"     "age"     "diab"    "first"
```

#Provides dimension of the database

```
> dim(data)
[1] 274 11
```



## Kaplan-Meier Analysis Example

The first step is to create a survival object, considering the variables follow-up and event. Note that a “+” after the time indicates censoring.

### #Create a survival object

```
> Surv(data$fu,data$event)
 [1] 7.00 22.00+ 13.00+ 42.00 5.00+ 13.00 70.00+ 51.00 25.00+ 5.00+ 17.00 42.00+ 6.00
[14] 10.00+ 49.00+ 55.00+ 18.00+ 7.00+ 2.00+ 22.00 24.00 2.00+ 1.00+ 10.00 17.00+ 35.00
[27] 32.00+ 30.00+ 2.00 5.00 6.00+ 78.00+ 4.00+ 63.00+ 6.00+ 9.00 22.00+ 6.00+ 33.00+
[40] 1.00+ 9.00 27.00+ 2.00+ 27.00 21.00+ 14.00+ 2.00 53.00+ 38.00 62.00+ 44.00+ 58.00+
[53] 33.00+ 23.00 22.00+ 26.00+ 11.00+ 2.00 28.00+ 74.00+ 28.00+ 13.00 17.00+ 30.00+ 24.00+
[66] 109.00+ 105.00+ 36.00+ 14.00+ 7.00+ 43.00+ 6.57+ 37.00 22.00+ 1.00 22.00+ 84.00+ 37.00
[79] 14.00 60.00 8.00 35.00 23.00+ 21.00+ 92.00+ 36.00 31.00+ 2.00+ 28.00+ 13.00 12.00
[92] 8.00+ 7.00+ 20.00+ 18.00 40.00+ 78.00+ 61.00+ 10.00+ 101.00+ 2.00+ 9.00+ 4.00+ 83.00+
[105] 8.00+ 9.00+ 1.00+ 34.00+ 33.00+ 11.00+ 32.00+ 31.00+ 29.00+ 5.00+ 21.00+ 26.00+ 26.00+
[118] 24.00+ 24.00+ 8.00+ 14.00+ 19.00+ 8.00+ 18.00+ 16.00+ 16.00+ 14.00+ 15.00+ 10.00+ 10.00+
[131] 22.00+ 8.00+ 7.00+ 6.00+ 5.00+ 3.00+ 2.00+ 7.00 22.00+ 13.00+ 42.00 5.00+ 13.00
[144] 70.00+ 51.00 25.00+ 5.00+ 17.00 42.00+ 6.00 10.00+ 49.00+ 55.00+ 18.00+ 7.00+ 2.00+
[157] 22.00 24.00 2.00+ 1.00+ 10.00 17.00+ 35.00 32.00+ 30.00+ 2.00 5.00 6.00+ 78.00+
[170] 4.00+ 63.00+ 6.00+ 9.00 22.00+ 6.00+ 33.00+ 1.00+ 9.00 27.00+ 2.00+ 27.00 21.00+
[183] 14.00+ 2.00 53.00+ 38.00 62.00+ 44.00+ 58.00+ 33.00+ 23.00 22.00+ 26.00+ 11.00+ 2.00
[196] 28.00+ 74.00+ 28.00+ 13.00 17.00+ 30.00+ 24.00+ 109.00+ 105.00+ 36.00+ 14.00+ 7.00+ 43.00+
[209] 6.57+ 37.00 22.00+ 1.00 22.00+ 84.00+ 37.00 14.00 60.00 8.00 35.00 23.00+ 21.00+
[222] 92.00+ 36.00 31.00+ 2.00+ 28.00+ 13.00 12.00 8.00+ 7.00+ 20.00+ 18.00 40.00+ 78.00+
[235] 61.00+ 10.00+ 101.00+ 2.00+ 9.00+ 4.00+ 83.00+ 8.00+ 9.00+ 1.00+ 34.00+ 33.00+ 11.00+
[248] 32.00+ 31.00+ 29.00+ 5.00+ 21.00+ 26.00+ 26.00+ 24.00+ 24.00+ 8.00+ 14.00+ 19.00+ 8.00+
[261] 18.00+ 16.00+ 16.00+ 14.00+ 15.00+ 10.00+ 10.00+ 22.00+ 8.00+ 7.00+ 6.00+ 5.00+ 3.00+
[274] 2.00+
```

# Kaplan-Meier Analysis Example

To produce the Kaplan-Meier estimates of the probability of survival over time. The median of survival and the survival in several points of time could be requested.

## #Kaplan-Meier survival curve

```
> KM_1 <- survfit(Surv(data$fu,data$event)~1)
> KM_1
Call: survfit(formula = Surv(data$fu, data$event) ~ 1)

      n  events  median 0.95LCL 0.95UCL
274      62      NA      51      NA
> summary(KM_1)
Call: survfit(formula = Surv(data$fu, data$event) ~ 1)

   time n.risk n.event survival std.err lower 95% CI upper 95% CI
1      1    274      2    0.993 0.00514    0.983    1.000
2      2    266      6    0.970 0.01034    0.950    0.991
5      5    242      2    0.962 0.01171    0.940    0.986
6      6    232      2    0.954 0.01299    0.929    0.980
7      7    220      2    0.945 0.01425    0.918    0.974
8      8    210      2    0.936 0.01547    0.906    0.967
9      9    198      4    0.917 0.01782    0.883    0.953
10     10    190      2    0.908 0.01889    0.871    0.946
12     12    176      2    0.897 0.02004    0.859    0.938
13     13    174      6    0.866 0.02299    0.823    0.913
14     14    166      2    0.856 0.02386    0.811    0.904
17     17    150      2    0.845 0.02487    0.797    0.895
18     18    144      2    0.833 0.02587    0.784    0.885
22     22    128      2    0.820 0.02706    0.769    0.875
23     23    114      2    0.806 0.02843    0.752    0.863
24     24    110      2    0.791 0.02974    0.735    0.851
27     27     94      2    0.774 0.03140    0.715    0.838
35     35     62      4    0.724 0.03802    0.653    0.803
36     36     58      2    0.699 0.04061    0.624    0.783
37     37     54      4    0.647 0.04510    0.565    0.742
38     38     50      2    0.621 0.04687    0.536    0.720
42     42     46      2    0.594 0.04857    0.506    0.698
51     51     36      2    0.561 0.05118    0.470    0.671
60     60     28      2    0.521 0.05482    0.424    0.641
```

# Kaplan-Meier Analysis Example

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To produce the Kaplan-Meier estimates of the probability of survival over time. The median of survival and the survival in several points of time could be requested.

### #Kaplan-Meier survival curve

The **median survival** is the smallest time at which the survival probability drops to 0.5 (50%) or below. If the survival curve does not drop to 0.5 or below then the median time cannot be computed.

```
> KM_1 <- survfit(Surv(data$fu, data$event)~1)
> KM_1
Call: survfit(formula = Surv(data$fu, data$event) ~ 1)

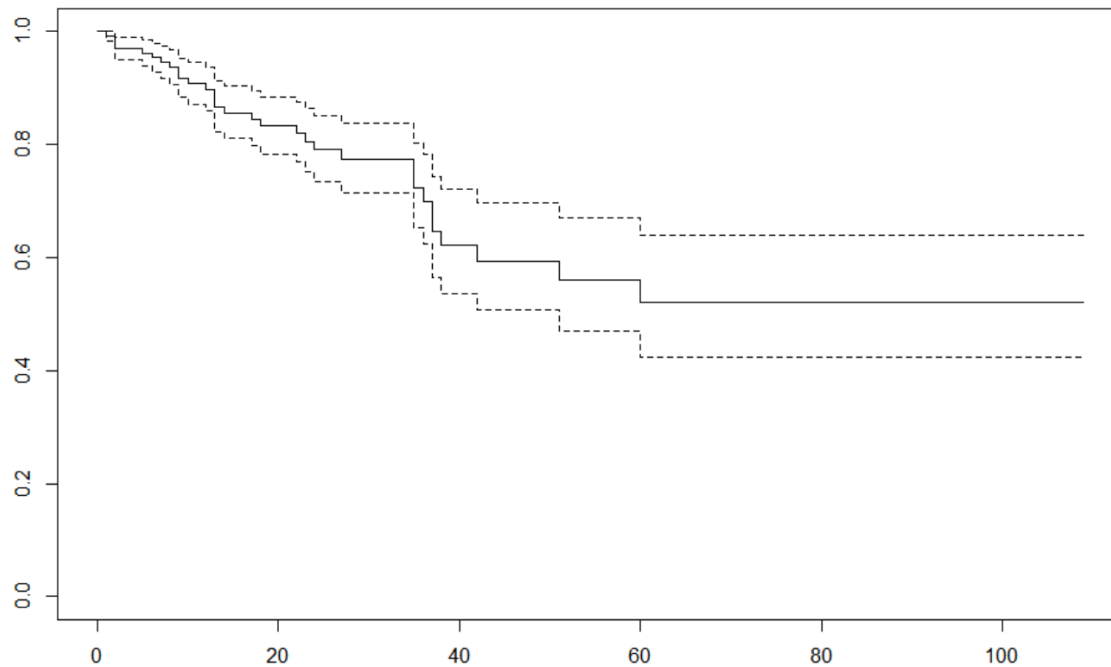
      n  events median 0.95LCL 0.95UCL
    274      62      NA       51      NA
> summary(KM_1)
Call: survfit(formula = Surv(data$fu, data$event) ~ 1)
```

time	n.risk	n.event	survival	std.err	lower 95% CI	upper 95% CI
1	274	2	0.993	0.00514	0.983	1.000
2	266	6	0.970	0.01034	0.950	0.991
5	242	2	0.962	0.01171	0.940	0.986
6	232	2	0.954	0.01299	0.929	0.980
7	220	2	0.945	0.01425	0.918	0.974
8	210	2	0.936	0.01547	0.906	0.967
9	198	4	0.917	0.01782	0.883	0.953
10	190	2	0.908	0.01889	0.871	0.946
12	176	2	0.897	0.02004	0.859	0.938
13	174	6	0.866	0.02299	0.823	0.913
14	166	2	0.856	0.02386	0.811	0.904
17	150	2	0.845	0.02487	0.797	0.895
18	144	2	0.833	0.02587	0.784	0.885
22	128	2	0.820	0.02706	0.769	0.875
23	114	2	0.806	0.02843	0.752	0.863
24	110	2	0.791	0.02974	0.735	0.851
27	94	2	0.774	0.03140	0.715	0.838
35	62	4	0.724	0.03802	0.653	0.803
36	58	2	0.699	0.04061	0.624	0.783
37	54	4	0.647	0.04510	0.565	0.742
38	50	2	0.621	0.04687	0.536	0.720
42	46	2	0.594	0.04857	0.506	0.698
51	36	2	0.561	0.05118	0.470	0.671
60	28	2	0.521	0.05482	0.424	0.641

# Kaplan-Meier Analysis Example

#Plot survival curve with confidence intervals

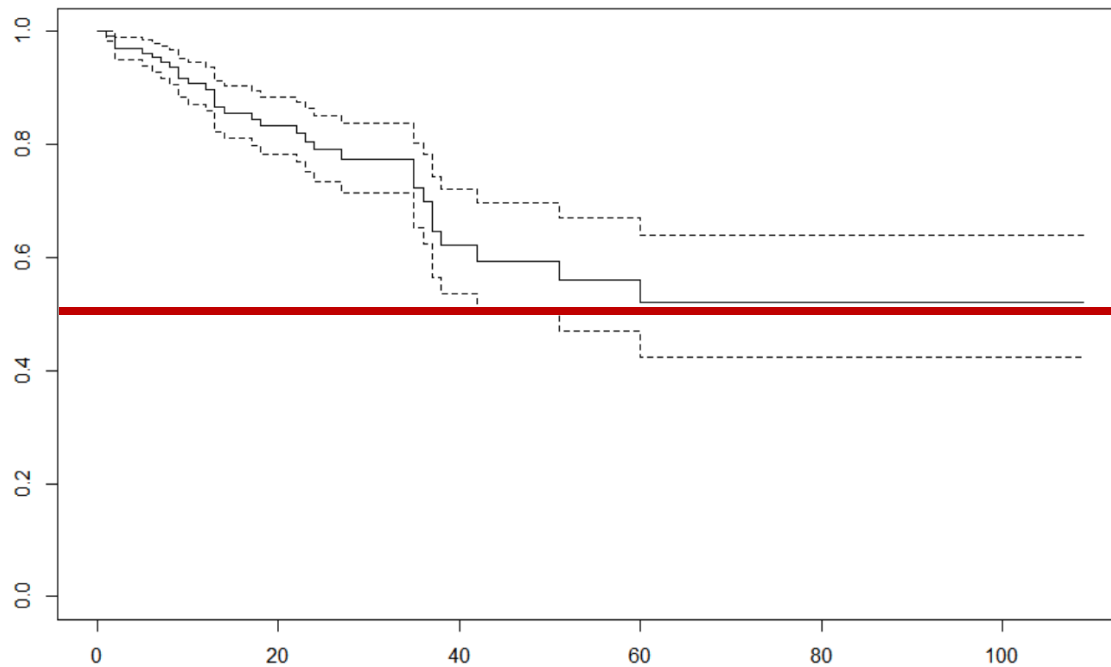
```
> plot(KM_1, mark.time=F)
```



# Kaplan-Meier Analysis Example

#Plot survival curve with confidence intervals

```
> plot(KM_1, mark.time=F)
```



# Kaplan-Meier Analysis Example

**To compare curves for different groups of subjects: Kaplan Meier Analysis according to one factor**

To produce the Kaplan-Meier curves according to one factor. To compare survival curves according to a factor, log-rank test could be used. For the log-rank test “rho=0”.

#Kaplan-Meier survival curves according to diabetes

```
> KM.diab <- survfit(Surv(data$fu,data$event)~ 1+data$diab)
> KM.diab
Call: survfit(formula = Surv(data$fu, data$event) ~ 1 + data$diab)
```

	n	events	median	0.95LCL	0.95UCL
data\$diab=0	206	38	NA	NA	NA
data\$diab=1	68	24	37	36	NA

# Kaplan-Meier Analysis Example

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```
> KM.diab <- survfit(Surv(data$fu,data$event)~ 1+data$diab)
> KM.diab
Call: survfit(formula = Surv(data$fu, data$event) ~ 1 + data$diab)
```

	n	events	median	0.95LCL	0.95UCL
data\$diab=0	206	38	NA	NA	NA
data\$diab=1	68	24	37	36	NA

				data\$diab=0				
time	n.risk	n.event	survival	std.err	lower	95% CI	upper	95% CI
1	206	2	0.990	0.00683		0.977		1.000
2	202	4	0.971	0.01179		0.948		0.994
6	178	2	0.960	0.01396		0.933		0.988
8	160	2	0.948	0.01616		0.917		0.980
9	148	2	0.935	0.01830		0.900		0.972
10	146	2	0.922	0.02017		0.883		0.963
12	132	2	0.908	0.02215		0.866		0.953
13	130	6	0.866	0.02694		0.815		0.921
17	110	2	0.851	0.02866		0.796		0.909
23	88	2	0.831	0.03110		0.772		0.894
27	72	2	0.808	0.03425		0.744		0.878
35	48	4	0.741	0.04500		0.658		0.834
37	42	2	0.705	0.04929		0.615		0.809
38	40	2	0.670	0.05276		0.574		0.782
42	36	2	0.633	0.05601		0.532		0.753

				data\$diab=1				
time	n.risk	n.event	survival	std.err	lower	95% CI	upper	95% CI
2	64	2	0.969	0.0217		0.9270		1.000
5	56	2	0.934	0.0319		0.8737		0.999
7	52	2	0.898	0.0395		0.8240		0.979
9	50	2	0.862	0.0454		0.7778		0.956
14	44	2	0.823	0.0511		0.7288		0.930
18	38	2	0.780	0.0568		0.6760		0.900
22	34	2	0.734	0.0621		0.6218		0.866
24	24	2	0.673	0.0704		0.5481		0.826
36	14	2	0.577	0.0872		0.4288		0.775
37	12	2	0.481	0.0955		0.3255		0.709
51	6	2	0.320	0.1123		0.1612		0.637
60	4	2	0.160	0.0978		0.0484		0.530

# Kaplan-Meier Analysis Example

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```
> KM.diab <- survfit(Surv(data$fu,data$event)~ 1+data$diab)
> KM.diab
Call: survfit(formula = Surv(data$fu, data$event) ~ 1 + data$diab)
```

	n	events	median	0.95LCL	0.95UCL
data\$diab=0	206	38	NA	NA	NA
data\$diab=1	68	24	37	36	NA

data\$diab=0									
time	n.risk	n.event	survival	std.err	lower	95% CI	upper	95% CI	
1	206	2	0.990	0.00683		0.977		1.000	
2	202	4	0.971	0.01179		0.948		0.994	
6	178	2	0.960	0.01396		0.933		0.988	
8	160	2	0.948	0.01616		0.917		0.980	
9	148	2	0.935	0.01830		0.900		0.972	
10	146	2	0.922	0.02017		0.883		0.963	
12	132	2	0.908	0.02215		0.866		0.953	
13	130	6	0.866	0.02694		0.815		0.921	
17	110	2	0.851	0.02866		0.796		0.909	
23	88	2	0.831	0.03110		0.772		0.894	
27	72	2	0.808	0.03425		0.744		0.878	
35	48	4	0.741	0.04500		0.658		0.834	
37	42	2	0.705	0.04929		0.615		0.809	
38	40	2	0.670	0.05276		0.574		0.782	
42	36	2	0.633	0.05601		0.532		0.753	

Smallest time at which the survival probability drops to 0.5 (50%) or below.

**Median survival time = 37.**

data\$diab=1									
time	n.risk	n.event	survival	std.err	lower	95% CI	upper	95% CI	
2	64	2	0.969	0.0217		0.9270		1.000	
5	56	2	0.934	0.0319		0.8737		0.999	
7	52	2	0.898	0.0395		0.8240		0.979	
9	50	2	0.862	0.0454		0.7778		0.956	
14	44	2	0.823	0.0511		0.7288		0.930	
18	38	2	0.780	0.0568		0.6760		0.900	
22	34	2	0.734	0.0621		0.6218		0.866	
24	24	2	0.673	0.0704		0.5481		0.826	
36	14	2	0.577	0.0872		0.4288		0.775	
37	12	2	0.481	0.0955		0.3255		0.709	
51	6	2	0.320	0.1123		0.1612		0.637	
60	4	2	0.160	0.0978		0.0484		0.530	

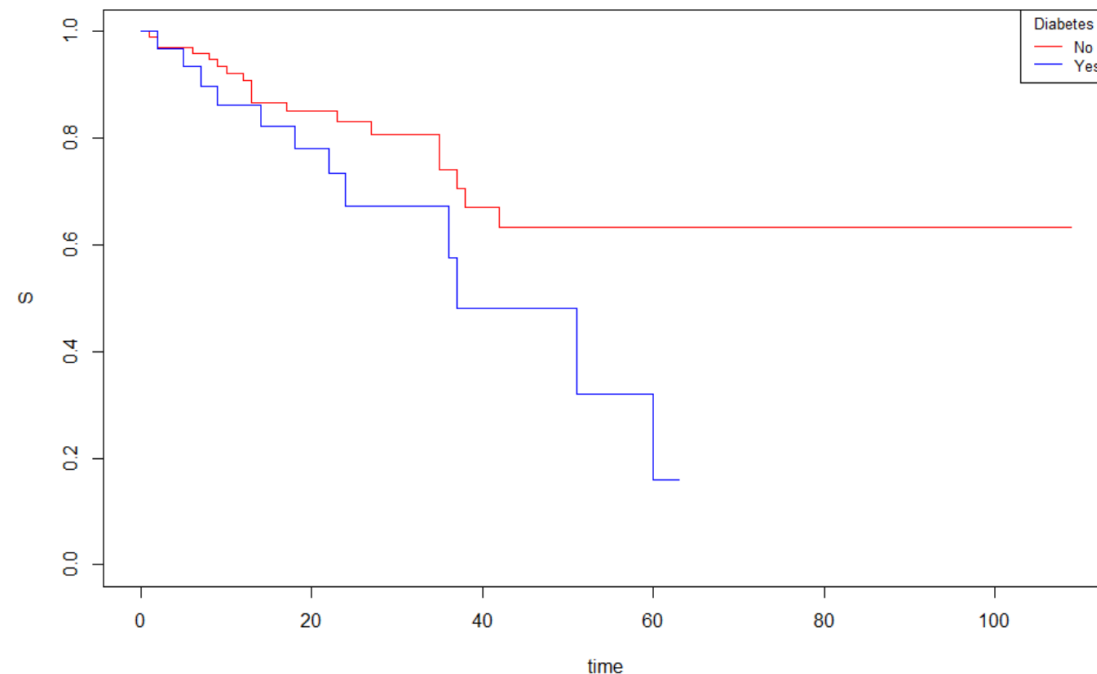




# Kaplan-Meier Analysis Example

#Plot survival curve according to diabetes

```
> plot(KM.diab,col=c("red","blue"),mark.time=F,ylim=c(0,1),xlab="time",ylab="s")  
> legend("topright", title="Diabetes", legend=c("No", "Yes"), col=c("red", "blue"),  
+       lty=1:1, cex=0.8)
```

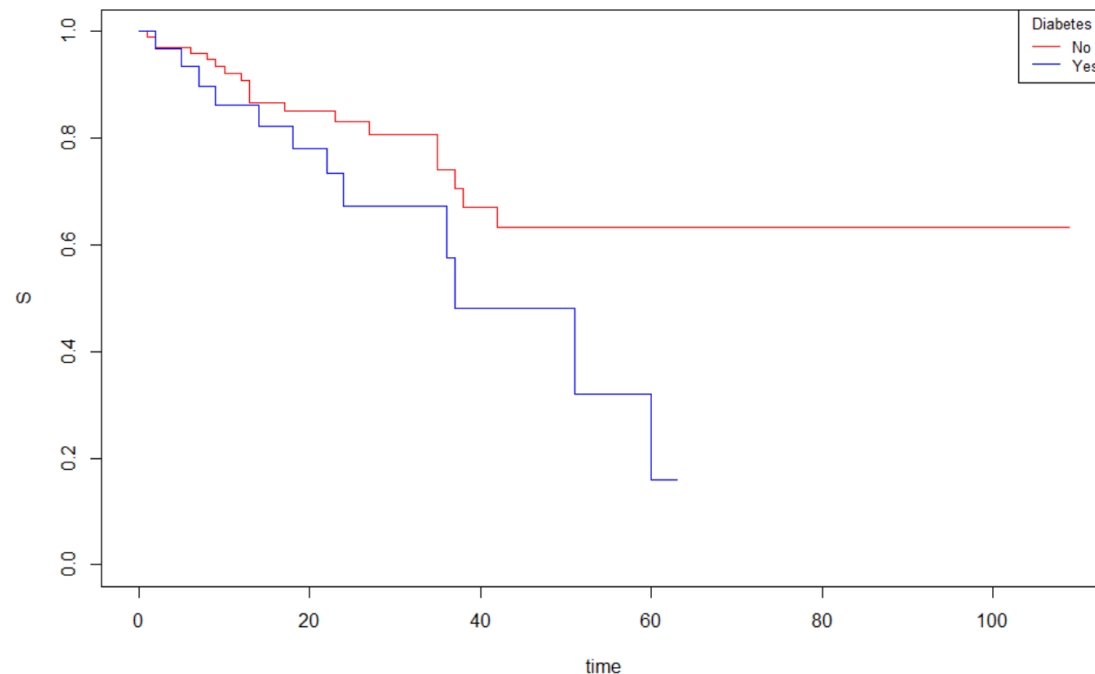


# Kaplan-Meier Analysis Example

#Plot survival curve according to diabetes

```
> plot(KM.diab,col=c("red","blue"),mark.time=F,ylim=c(0,1),xlab="time",ylab="s")  
> legend("topright", title="Diabetes", legend=c("No", "Yes"), col=c("red", "blue"),  
+       lty=1:1, cex=0.8)
```

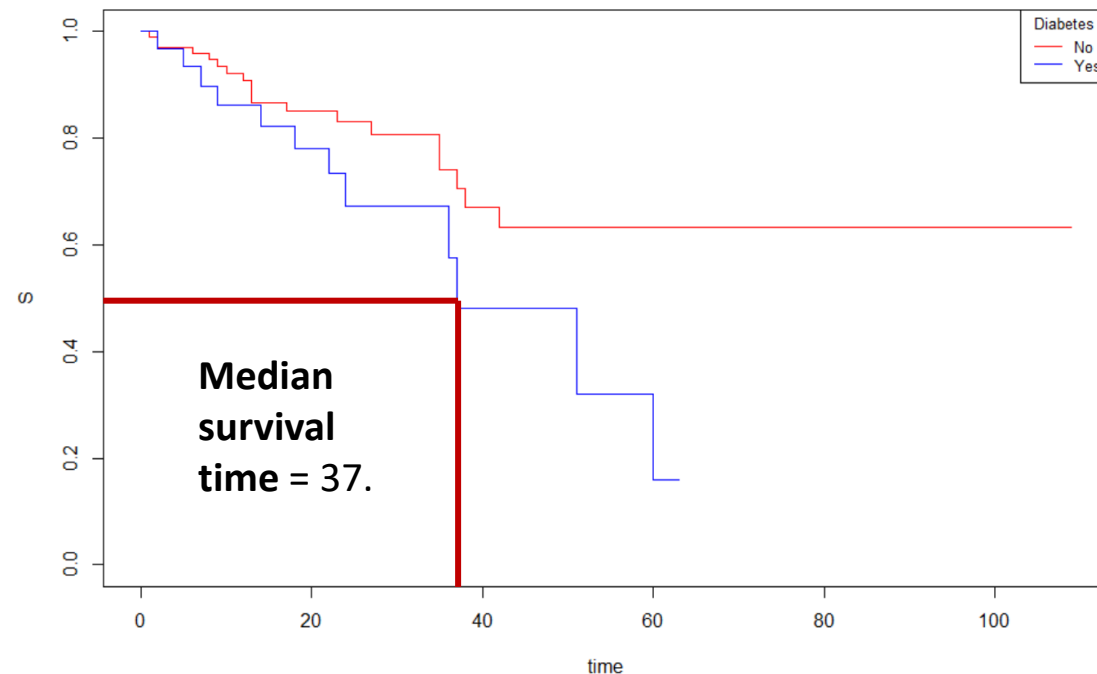
The Kaplan-Meier plot is now split into each of the levels of the categorical variable (2 groups in this case).



# Kaplan-Meier Analysis Example

#Plot survival curve according to diabetes

```
> plot(KM.diab,col=c("red","blue"),mark.time=F,ylim=c(0,1),xlab="time",ylab="s")  
> legend("topright", title="Diabetes", legend=c("No", "Yes"), col=c("red", "blue"),  
+       lty=1:1, cex=0.8)
```



# Kaplan-Meier Analysis Example



## #Log-rank test

- Allows for comparison between groups.
- Possible to compute hand (based on Chi-square).

$H_0$ : No difference between the groups.

$H_1$ : The groups are different.

# Kaplan-Meier Analysis Example

## #Log-rank test

```
> plot(KM.diab,col=c("red","blue"),mark.time=F,ylim=c(0,1),xlab="time",ylab="s")
> legend("topright", title="Diabetes", legend=c("No", "Yes"), col=c("red", "blue"),
+       lty=1:1, cex=0.8)
> survdiff(Surv(data$fu,data$event)~ 1+data$diab,rho=0)
```

```
Call:
survdiff(formula = Surv(data$fu, data$event) ~ 1 + data$diab,
        rho = 0)
```

	N	Observed	Expected	(O-E) <sup>2</sup> /E	(O-E) <sup>2</sup> /V
data\$diab=0	206	38	47.5	1.89	8.27
data\$diab=1	68	24	14.5	6.19	8.27

Chisq= 8.3 on 1 degrees of freedom, p= 0.004

# Kaplan-Meier Analysis Example

## #Log-rank test

```
> plot(KM.diab,col=c("red","blue"),mark.time=F,ylim=c(0,1),xlab="time",ylab="s")
> legend("topright", title="Diabetes", legend=c("No", "Yes"), col=c("red", "blue"),
+       lty=1:1, cex=0.8)
> survdiff(Surv(data$fu,data$event)~ 1+data$diab,rho=0)
```

```
Call:
survdiff(formula = Surv(data$fu, data$event) ~ 1 + data$diab,
         rho = 0)
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data\$diab=0	206	38	47.5	1.89	8.27
data\$diab=1	68	24	14.5	6.19	8.27

chisq= 8.3 on 1 degrees of freedom, **p= 0.004**



The log rank test here shows significant difference between the groups ( $p < 0.05$ ).

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- Kalbfleisch, J. and Prentice, R. (2002). **The Statistical Analysis of Failure Time Data, 2<sup>nd</sup> Edition**. New York: Wiley.
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