

# **Prognosis of survival for breast cancer patients**

**Ken Ryder**

**Breast Cancer Unit Data Section**

**Guy's Hospital**

**Patrick Royston,**

**MRC Clinical Trials Unit**

**London**

# Outline

- **Introduce the data and outcomes requested by the medics.**
- **Work through the stages of the analyses.**

**Imputation**

**Fractional Polynomials**

**Producing a simple score**

**Fitting Splines**

- **Illustrate the menu call developed for everyday use.**

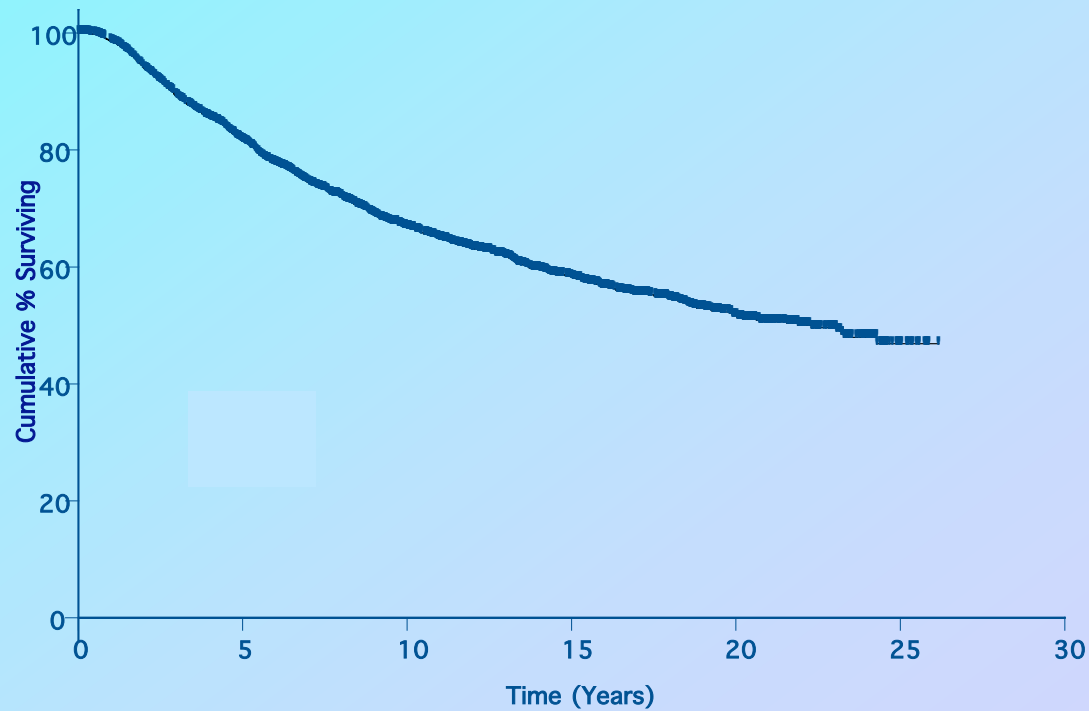
# The Problem

- **Over 8000 patients with invasive Breast Cancer with 3083 operable stage 1 and 2 disease.**
- **Time of relapse-free survival**
- **Time to a distant relapse**
- **Overall survival**
  - **Death from breast cancer**
  - **All-cause mortality**

# The Problem

- **Kaplan-Meier Curves**

Overall Survival for Stage 1 and 2 patients



# The Problem

## Factors affecting outcome

- **Histological type ( $x_4$ )**  
**Ductal ( Grade I, II & III), Lobular, others (Mucoid, Medullary, Tubular)**
- **Tumour Size ( $x_3$ )**  
**( $\leq 2.0$  cms, 2.0 - 5.0 cms,  $>5.0$  cms)**
- **Axilla Nodal status ( $x_5$ )**  
**(All negative, 1-3, 4 - 9,  $\geq 10$  positive)**
- **Age ( $x_1$ )**  
**(Under 40, 41-50, 51-60, 61-70, Over 70)**

# The Problem

## Factors affecting outcome

- Postmenopausal status ( $x_2$ )
- Type of surgery ( $x_{11}$ )
- Progesterone receptor status ( $x_{6a}$ )
- Oestrogen receptor status ( $x_{7a}$ )
- Adjuvant treatment  
Chemotherapy ( $x_8$ )
- Hormonal therapy ( $x_9$ )
- Radiotherapy ( $x_{10}$ )

## The Task

- **Given the information on the prognostic factors for a new patient**
- **To provide estimates of the probabilities of being relapse-free or surviving after 5, 10, 15, 20 years.**

# Prognostic Factors

	Code	Mean	% missing
Age (years)	$x_1$	55	0
Postmenopausal status	$x_2$	0.55	2.0
Clinical tumour size (mm)	$x_3$	28	1.1
Histology Ductal Grade 2	$x_{4a}$	0.39	0
Histology Ductal Grade 3	$x_{4b}$	0.34	0
Histology Lobular	$x_{4x}$	0.13	0
Histology Other	$x_{4y}$	0.03	0
No of positive lymph nodes	$x_5$	2.8	0
Progesterone receptor positive	$x_{6a}$	0.52	13.7
Oestrogen receptor positive	$x_{7a}$	0.70	11.5
Chemotherapy (any type)	$x_8$	0.20	0
Hormonal therapy	$x_9$	0.35	0
Radiotherapy	$x_{10}$	0.42	0
Impalable tumour	$x_{3x}$	0.07	1.1



# Imputation

**van Buuren S., H. C. Boshuizen and D. L. Knook. 1999. Multiple imputation of missing blood pressure covariates in survival analysis. *Statistics in Medicine* 18:681-694.**

**mvisamp imputes missing values using "switching regression", an iterative multivariable regression technique.**

**mvisamp varlist [if exp] [in range] [weight], gen(name)  
[ cc(ccvarlist) cmd(cmdlist) seed(#) ]**

# Imputation

**Impute x6a and x7a on complete variables and transformed year and survival time**

```
mvisamp x6a x7a x1 x4a x4b x4x x4y x5 yr_1 yr_2  
yr_3 lnt _d, gen(z1) seed(101)
```

**Impute x3 on these variables and completed x6a, x7a**

```
mvisamp x3 z1_1 z1_2 x1 x4a x4b x4x x4y x5 yr_1  
yr_2 yr_3 lnt _d if x3>0, gen(z2) seed(103)
```

# Cox's Proportional Hazards model

## Fractional Polynomials

- For the continuous variables, the Fractional Polynomials, based on  $X^{-2}$ ,  $X^{-1}$ ,  $X^{-0.5}$ ,  $X^{0.5}$ ,  $X^2$ ,  $X^3$ , and  $\log_e(X)$  are compared with the linear polynomial to determine if one or more of these polynomials provides a better fit.
- All other factors are (0,1) variables, including the interaction of Hormone Therapy with ER status.

# Cox's Proportional Hazards model

## Fractional Polynomials

**mfracpol** selects the fractional polynomial (FP) model which best predicts the outcome variable, **yvar**, from the RHS variables, **xvarlist**. After execution, **mfracpol** leaves variables in the data named **Ivar1\_\_1**, **Ivar2\_\_2**, ...**Ivar2\_1**, **Ivar2\_2** etc. The new variables contain the best-fitting fractional polynomial powers of **var1**, **var2**, .

```
mfracpol regression_cmd yvar xvarlist [in range] [if exp]
[weight]      [, alpha(alpha_list) df(df_list)
select(select_list)      regression_cmd_options ]
```

# Cox's Proportional Hazards model

## Fractional Polynomials

\* Force in ER and hormon, since know ER x hormon is significant.

```
mfracpol cox _t x1 ix2 ix3 ix3x x4a x4b x4x x4y x5 ix6a ix7a  
x7ah hormon chemo rt yr, dead(_d) sel(.05, ix7a hormon  
x7ah:0.9999) df(ix3 x5:2)
```

# Cox's Proportional Hazards model

## Fractional Polynomials

-> gen double lx1\_\_1 =  $X^{-2-.0327}$  if e(sample)

(where:  $X = x1/10$ )

-> gen double lix3\_\_1 =  $\ln(X)+1.243$  if e(sample)

(where:  $X = (ix3+1)/100$ )

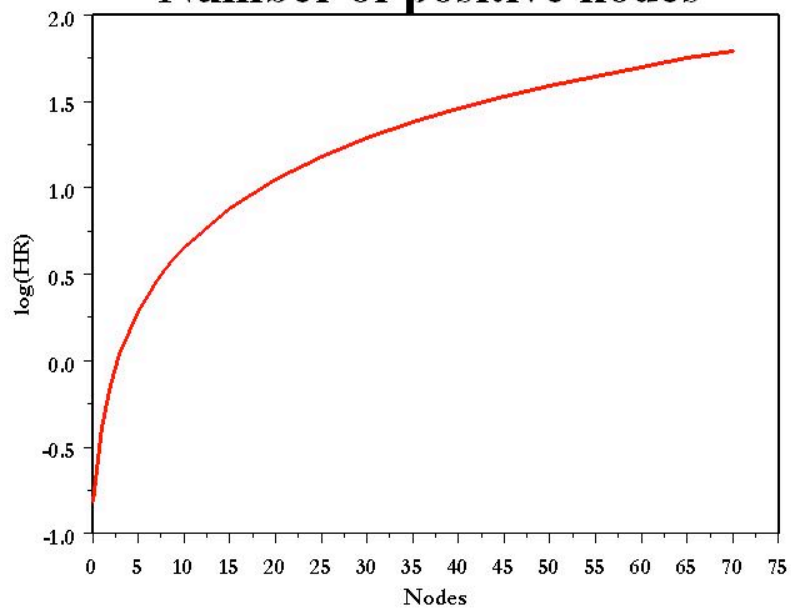
-> gen double lx5\_\_1 =  $\ln(X)+.974$  if e(sample)

(where:  $X = (x5+1)/10$ )

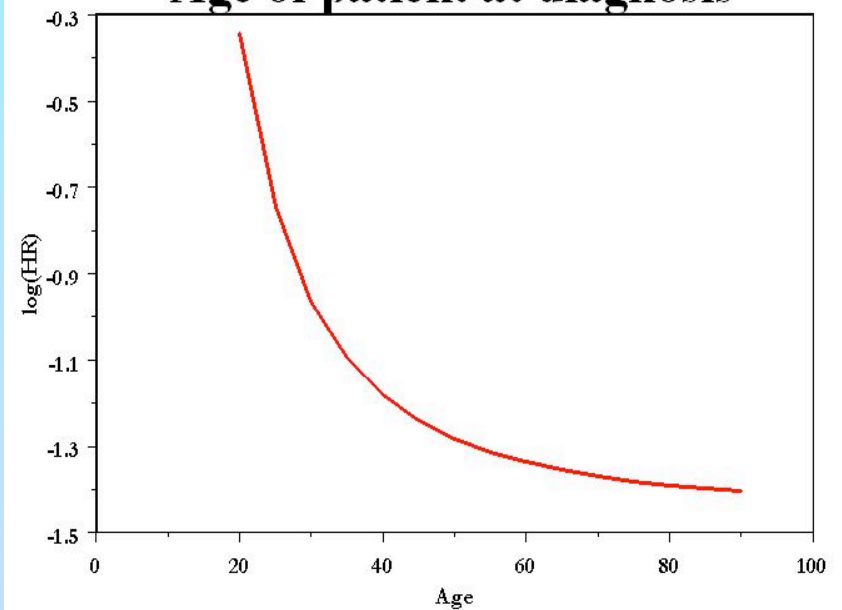
# Cox's Proportional Hazards model

## Fractional Polynomials

### Number of positive nodes



### Age of patient at diagnosis



# Cox's Proportional Hazards model

## Fractional Polynomials

Final multivariable fractional polynomial model for \_t

Variable	-----Initial-----			-----Final-----		
	df	Select	Alpha	Status	df	Powers
x1	4	0.0500	0.0500	in	2	-2
ix2	1	0.0500	0.0500	out	0	
ix3	2	0.0500	0.0500	in	2	0
ix3x	1	0.0500	0.0500	in	1	1
x4a	1	0.0500	0.0500	in	1	1
x4b	1	0.0500	0.0500	in	1	1
x4x	1	0.0500	0.0500	in	1	1
x4y	1	0.0500	0.0500	out	0	
x5	2	0.0500	0.0500	in	2	0
ix6a	1	0.0500	0.0500	in	1	1
ix7a	1	0.9999	0.0500	in	1	1
x7ah	1	0.9999	0.0500	in	1	1
hormon	1	0.9999	0.0500	in	1	1
chemo	1	0.0500	0.0500	in	1	1
rt	1	0.0500	0.0500	out	0	
yr	4	0.0500	0.0500	out	0	



# Cox's Proportional Hazards model

## Fractional Polynomials

_t   _d	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
Ix1__1	4.707014	1.439364	3.27	0.001	1.885911	7.528116
Iix3__1	.5143961	.0655541	7.85	0.000	.3859123	.6428798
ix3x	1.320571	.2790599	4.73	0.000	.7736239	1.867518
x4a	1.122357	.1557526	7.21	0.000	.8170873	1.427626
x4b	1.420355	.1588664	8.94	0.000	1.108983	1.731728
x4x	1.028325	.1704158	6.03	0.000	.6943161	1.362334
Ix5__1	.6070695	.0299867	20.24	0.000	.5482966	.6658423
ix6a	-.1314727	.066974	-1.96	0.050	-.2627393	-.000206
ix7a	.1902246	.0781319	2.43	0.015	.037089	.3433603
x7ah	-.708684	.1504531	-4.71	0.000	-1.003567	-.4138013
hormon	-.0381773	.1254731	-0.30	0.761	-.2841	.2077454
chemo	-.3198936	.0782234	-4.09	0.000	-.4732086	-.1665787

Deviance:16524.958.

# Cox's Proportional Hazards model

## Fractional Polynomials

- . global model `e(fp\_fvl) `
- . stcox \$model, nolog sca(sca\*) sch(sch\*)
- . stphtest, rank detail

Test of proportional hazards assumption				
	rho	chi2	df	Prob>chi2
Ix1__1	-0.08739	10.11	1	0.0015
Ix3_1__1	0.03250	1.23	1	0.2666
x4a	-0.06038	4.17	1	0.0412
x4b	-0.11912	16.27	1	0.0001
x4x	-0.01873	0.40	1	0.5268
Ix5__1	-0.14572	23.72	1	0.0000
ix6a	0.06161	4.59	1	0.0321
ix7a	0.08467	8.35	1	0.0039
x7ah	0.03161	1.14	1	0.2847
hormon	-0.01300	0.19	1	0.6609
chemo	0.11522	16.23	1	0.0001
global test		145.80	11	0.0000

## Simple scoring system

- A scoring system was developed so that a unit increase in the score was approximately an increase of 0.1 in the  $\log_e(\text{HR})$ .
- For the (0,1) binary variables the scores were taken to the nearest 0.1 of the  $\log_e(\text{HR})$ , but for the continuous variables the cut-off points were selected to have an increment of approximately 0.2 in the  $\log_e(\text{HR})$ .

# Simple scoring system

Node status	Score	Tumour size	Score
0	0	$\leq 11\text{mm}$	0
1	4	12 – 19	2
2	6	20 – 27	4
3 – 4	8	28 – 38	6
5 – 6	10	$\geq 39$	8
7 – 9	12		
10 – 14	14		
15 – 20	16	ER <sup>+</sup> and HT	0
21 – 29	18	ER <sup>+</sup> and no HT	8
30 – 42	20	ER <sup>-</sup>	7
43 – 52	22		
$\geq 53$	24		

## Simple scoring system

### Fitting splines

**stpm [varlist] [if exp] [in range] , model\_complexity  
scale(hazardlnormalodds) stratify(strat\_varlist)**

**stpm fits spline-based distributional models to right- or  
interval-censored survival-time data.**

**varlist is a set of covariates.**

**model\_complexity given by defining the number of degrees of  
freedom to use or by defining the knot positions**

**scale specifies the scale of model**

**stratify**

**stpm guysrisk, st(guysrisk) sc(h) knots(0.643 1.614)  
nolog**

## Simple scoring system

<b>Score</b>	<b>1</b>	<b>2</b>	<b>5</b>	<b>10</b>	<b>20</b>
<b>10</b>	<b>0.989</b>	<b>0.973</b>	<b>0.942</b>	<b>0.914</b>	<b>0.881</b>
<b>14</b>	<b>0.984</b>	<b>0.960</b>	<b>0.914</b>	<b>0.873</b>	<b>0.827</b>
<b>18</b>	<b>0.976</b>	<b>0.940</b>	<b>0.873</b>	<b>0.815</b>	<b>0.751</b>
<b>22</b>	<b>0.964</b>	<b>0.911</b>	<b>0.815</b>	<b>0.735</b>	<b>0.650</b>
<b>26</b>	<b>0.947</b>	<b>0.869</b>	<b>0.735</b>	<b>0.628</b>	<b>0.522</b>
<b>30</b>	<b>0.921</b>	<b>0.809</b>	<b>0.629</b>	<b>0.497</b>	<b>0.376</b>
<b>34</b>	<b>0.884</b>	<b>0.727</b>	<b>0.498</b>	<b>0.348</b>	<b>0.229</b>
<b>38</b>	<b>0.830</b>	<b>0.619</b>	<b>0.350</b>	<b>0.204</b>	<b>0.109</b>
<b>42</b>	<b>0.755</b>	<b>0.485</b>	<b>0.205</b>	<b>0.092</b>	<b>0.035</b>
<b>46</b>	<b>0.656</b>	<b>0.337</b>	<b>0.092</b>	<b>0.027</b>	<b>0.006</b>

# Menu system

