

eAppendix:

RESTRICTED QUADRATIC SPLINES

A spline is a function of a continuous regressor (“independent variable”) which allows the polynomial representing the regressor to change at specified values (knots) of the regressor. Splines may be linear or higher-order polynomials. For linear splines, the slope for the relationship between the variables of interest is allowed to change solely at the knots resulting in linear functions with inflection points at the knots. Its fit can be improved by using quadratic or cubic spline functions which allow for smooth functions at the knots rather than inflection points. A spline may be unrestricted or restricted in the tails (i.e., the function before the first knot and after the last knot) of the spline. Unrestricted splines allow the tails to be non-linear, while restricted splines are linear in the tails. Restriction can prevent poor fit in the tails due to small sample size or outliers. Here we concentrate on quadratic splines and we restrict the tails of the spline to be linear.

Splines of any order or restriction can be used when fitting various regression models including generalized linear and Cox proportional hazards models. To fit any standard regression model with a spline, one must include the original continuous variable as well as a set of “basis functions”, which are constructed variables that are added to the model. The number of basis functions is determined by the number of knots in the spline.

For unrestricted quadratic splines, if there are k knots for continuous regressor x , in relation to the dependent variable y , then there are $k + 1$ basis functions. The functional form for a quadratic spline regression model with k knots is

$$y = f(x) = \beta_0 + \beta_1 x + \gamma_1 x^2 + \gamma_2 (x - \chi_1)_+^2 + \dots + \gamma_{k+1} (x - \chi_k)_+^2 \quad (\text{Equation 1})$$

where χ_i corresponds to the location of the i th knot ($i = 1$ to k), the positive-part function $(\dots)_+$ is defined by $(x - \chi_k)_+ = (x - \chi_k)$ if $(x - \chi_k) > 0$; $(x - \chi_k)_+ = 0$ if $(x - \chi_k) \leq 0$, and x^2 as well as $(x - \chi_i)_+^2$ are the basis functions for the quadratic spline regression model.

For quadratic splines restricted in both tails, if there are k knots for continuous regressor x , in relation to the dependent variable y , then there are $k - 1$ basis functions. To restrict the quadratic spline to be linear before the first knot, the first quadratic term (i.e., x^2) in Equation 1 should be removed. To restrict the quadratic spline to be linear after the last knot, the quadratic term corresponding to the last knot (i.e., $(x - \chi_k)_+^2$) should be subtracted from all quadratic terms. Therefore, the functional form for the restricted quadratic spline regression model is

$$\begin{aligned}
 y = f(x) = & \beta_0 + \beta_1 x \\
 & + \beta_2 \left[(x - \chi_1)_+^2 - (x - \chi_k)_+^2 \right] \\
 & + \beta_3 \left[(x - \chi_2)_+^2 - (x - \chi_k)_+^2 \right] & \text{(Equation 2)} \\
 & + \dots \\
 & + \beta_k \left[(x - \chi_{k-1})_+^2 - (x - \chi_k)_+^2 \right]
 \end{aligned}$$

where $\left[(x - \chi_i)_+^2 - (x - \chi_k)_+^2 \right]$ are the basis functions for the restricted quadratic spline regression model.

The restricted quadratic spline model with k knots has the same model degrees of freedom (df) as the restricted cubic spline model with $k+1$ knots, perhaps counter to intuition. As demonstrated by the SAS output for the included example, when the model df are the same and

the knots are placed at comparable locations (e.g., cubic: 16th, 33rd, 50th, 67th and 84th; quadratic: 20th, 40th, 60th, and 80th) restricted quadratic and cubic spline models yield similar results. This equivalence between the quadratic and cubic models likely occurs given that both models are allowed to be equally flexible at similar knot locations of the continuous regressor.

DESCRIPTION OF SAS CODE FOR ESTIMATION OF RESTRICTED QUADRATIC SPLINES

To generate the restricted spline basis functions in Equation 2 the user can specify the dataset to be used (i.e., *data*). If a dataset is not specified, the provided code will automatically select the last dataset used in SAS. The user must specify the continuous variable for which the spline basis functions are to be constructed (i.e., x) as well as choose the number of knots (i.e., k) from 3, 4, 5, 6, or 7. A finite range for the potential number of knots was selected rather than allowing the user to specify any integer to keep the macro as simple as possible. The specific range of 3-7 was recommended by Harrell.¹ If the number of knots is not chosen, the provided code will automatically select 4 knots. The user must specify whether the knots should be placed at equally or unequally spaced percentile intervals. By default the macro generates knots at equally spaced percentile intervals. If the user specifies unequal intervals, then following Harrell² the knots will be placed at the symmetric, but unequally spaced percentile intervals summarized in the provided SAS macro.

The user must also specify whether the percentiles are to be based on the entire distribution of x or solely on the cases to maximize power when non-cases substantially outnumber cases.³ The default is to estimate the percentiles using the entire data. The macro allows for the exploration of various knot specifications in terms of number and location to facilitate careful consideration during model selection. Finally, the macro prints to the SAS log the values corresponding to the specified knot locations.

The spline basis functions for variable x are automatically added to the data set *data* as variables $_x$, with the number of leading underscores coincident with the number of the basis

function (i.e., if 4 unequal knots are chosen then basis function variables $_x$, $__x$, and $___x$ are

returned where $_x = \left[(x - P5)_+^2 - (x - P95)_+^2 \right] / (P95 - P5)$,

$__x = \left[(x - P35)_+^2 - (x - P95)_+^2 \right] / (P95 - P5)$, $___x = \left[(x - P65)_+^2 - (x - P95)_+^2 \right] / (P95 - P5)$,

$P5$ is the quantile for the 5th percentile of x , $P35$ is the quantile for the 35th percentile, $P65$ is the

quantile for the 65th percentile, and $P95$ is the quantile for the 95th percentile). Following

Harrell,² our macro divides each basis function in Equation 2 by the difference of the outer knots

(e.g., 5th and 95th percentiles) to ensure that the basis functions generated by the macro are in the

original units of x . To generate splines with distinct specifications (e.g., different number of

knots) or for more than one continuous variable, the user must re-call the SAS code with the

specification or regressor x changed.

To fit any standard regression model with the restricted quadratic spline, the user must

include the original continuous variable as well as the basis function variables constructed by the

macro (e.g., $_x$, $__x$, and $___x$) in the relevant regression model which can be any generalized

linear (e.g., Poisson, logistic) or Cox proportional hazards model.

STRATEGIES FOR MODEL SELECTION

The provided code is intended to aid in selecting between various model specifications of the continuous regressor (e.g., linear, categorical, spline, fractional polynomial) as well as assessing robustness of inferences. Given that these different specifications correspond to non-nested models, the Akaike information criterion (AIC)⁴ should be used to assess model fit. The AIC can also be used to inform selection regarding the number of knots (e.g., 3, 4, 5, 6, or 7) as well as knot location (e.g., equal versus unequal percentile intervals, percentiles intervals based on the entire distribution of x versus solely on the cases).

SAS (VERSION 9.2) CODE FOR ESTIMATION OF RESTRICTED QUADRATIC SPLINES (RQSMACRO)

```

*****
** 1. User names data set (e.g., data), single variable (e.g., var), outcome of interest (e.g., event) **
** number of knots (3, 4, 5, 6, or 7), knots at equal intervals (e.g., 1 for equal), and percentiles **
** based on cases (e.g., 1 for cases). **
** 2. Macro adds k-1 restricted quadratic spline basis functions to data set. **
** 3. Basis functions are named _var, __var, etc. **
** 4. Multiple macro runs are required for multiple variables. **
** 5. 3 knots at equal intervals or at: 5, 50, 95 **
** 4 knots at equal intervals or at: 5, 35, 65, 95 **
** 5 knots at equal intervals or at: 5, 28, 50, 73, 95 **
** 6 knots at equal intervals or at: 5, 23, 41, 59, 77, 95 **
** 7 knots at equal intervals or at: 3, 18, 34, 50, 66, 82, 98 **
*****

```

```

%macro rqspline(data=_last_,x=,event=,k=4,equal=1,cases=0);
  options nonotes;
  %local _p_ _z_;
  data &data;
    set &data;
    _z_=1;
  %if &equal=0 %then %do;
  %if &cases=0 %then %do;
  proc univariate data=&data noprint;
    var &x; output out=_p_ pctlpts=3 5 18 23 28 34 35 41 50 59 65 66 73 77 82 95 98 pctlpre=p;
  %end;
  %if &cases=1 %then %do;
  proc univariate data=&data noprint;
    where &event=1; var &x; output out=_p_ pctlpts=3 5 18 23 28 34 35 41 50 59 65 66 73 77 82 95 98 pctlpre=p;
  %end; %end;

```

```

%if &equal=1 %then %do;
%if &cases=0 %then %do;
proc univariate data=&data noprint;
    var &x; output out=_p_ pctlpts=12 15 16 20 24 25 29 33 37 40 43 50 57 60 63 67 71 75 76 80 84 85 88 pctlpre=p;
%end;
%if &cases=1 %then %do;
proc univariate data=&data noprint;
    where &event=1; var &x; output out=_p_ pctlpts=12 15 16 20 24 25 29 33 37 40 43 50 57 60 63 67 71 75 76 80 84 85
88 pctlpre=p;
%end; %end;
data _p_;
    set _p_;
    _z_=1;
    if &k=3 and &equal=0 then put "Knots for &x =" p5 p50 p95;
    else if &k=3 and &equal=1 then put "Knots for &x =" p25 p50 p75;
    if &k=4 and &equal=0 then put "Knots for &x =" p5 p35 p65 p95;
    else if &k=4 and &equal=1 then put "Knots for &x =" p20 p40 p60 p80;
    if &k=5 and &equal=0 then put "Knots for &x =" p5 p28 p50 p73 p95;
    else if &k=5 and &equal=1 then put "Knots for &x =" p16 p33 p50 p67 p84;
    if &k=6 and &equal=0 then put "Knots for &x =" p5 p23 p41 p59 p77 p95;
    else if &k=6 and &equal=1 then put "Knots for &x =" p15 p29 p43 p57 p71 p85;
    if &k=7 and &equal=0 then put "Knots for &x =" p3 p18 p34 p50 p66 p82 p98;
    else if &k=7 and &equal=1 then put "Knots for &x =" p12 p24 p37 p50 p63 p76 p88;

data &data;
    merge &data _p_;
    by _z_;
    drop _z_;
data &data;
    set &data;
    %if &equal=0 %then %do;

```



```

%if &k=3 %then %do;
  _&x=(max(0,&x-p5)**2-max(0,&x-p95)**2)/(p95-p5);
  __&x=(max(0,&x-p50)**2-max(0,&x-p95)**2)/(p95-p5);
%end;
%else %if &k=4 %then %do;
  _&x=(max(0,&x-p5)**2-max(0,&x-p95)**2)/(p95-p5);
  __&x=(max(0,&x-p35)**2-max(0,&x-p95)**2)/(p95-p5);
  ___&x=(max(0,&x-p65)**2-max(0,&x-p95)**2)/(p95-p5);
%end;
%else %if &k=5 %then %do;
  _&x=(max(0,&x-p5)**2-max(0,&x-p95)**2)/(p95-p5);
  __&x=(max(0,&x-p28)**2-max(0,&x-p95)**2)/(p95-p5);
  ___&x=(max(0,&x-p50)**2-max(0,&x-p95)**2)/(p95-p5);
  ____&x=(max(0,&x-p73)**2-max(0,&x-p95)**2)/(p95-p5);
%end;
%else %if &k=6 %then %do;
  _&x=(max(0,&x-p5)**2-max(0,&x-p95)**2)/(p95-p5);
  __&x=(max(0,&x-p23)**2-max(0,&x-p95)**2)/(p95-p5);
  ___&x=(max(0,&x-p41)**2-max(0,&x-p95)**2)/(p95-p5);
  ____&x=(max(0,&x-p59)**2-max(0,&x-p95)**2)/(p95-p5);
  _____&x=(max(0,&x-p77)**2-max(0,&x-p95)**2)/(p95-p5);
%end;
%else %if &k=7 %then %do;
  _&x=(max(0,&x-p3)**2-max(0,&x-p98)**2)/(p98-p3);
  __&x=(max(0,&x-p18)**2-max(0,&x-p98)**2)/(p98-p3);
  ___&x=(max(0,&x-p34)**2-max(0,&x-p98)**2)/(p98-p3);
  ____&x=(max(0,&x-p50)**2-max(0,&x-p98)**2)/(p98-p3);
  _____&x=(max(0,&x-p66)**2-max(0,&x-p98)**2)/(p98-p3);
  _____&x=(max(0,&x-p82)**2-max(0,&x-p98)**2)/(p98-p3);
%end; %end;
%if &equal=1 %then %do;

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```

%if &k=3 %then %do;
  _&x=(max(0,&x-p25)**2-max(0,&x-p75)**2)/(p75-p25);
  __&x=(max(0,&x-p50)**2-max(0,&x-p75)**2)/(p75-p25);
%end;
%else %if &k=4 %then %do;
  _&x=(max(0,&x-p20)**2-max(0,&x-p80)**2)/(p80-p20);
  __&x=(max(0,&x-p40)**2-max(0,&x-p80)**2)/(p80-p20);
  ___&x=(max(0,&x-p60)**2-max(0,&x-p80)**2)/(p80-p20);
%end;
%else %if &k=5 %then %do;
  _&x=(max(0,&x-p16)**2-max(0,&x-p84)**2)/(p84-p16);
  __&x=(max(0,&x-p33)**2-max(0,&x-p84)**2)/(p84-p16);
  ___&x=(max(0,&x-p50)**2-max(0,&x-p84)**2)/(p84-p16);
  ____&x=(max(0,&x-p67)**2-max(0,&x-p84)**2)/(p84-p16);
%end;
%else %if &k=6 %then %do;
  _&x=(max(0,&x-p15)**2-max(0,&x-p85)**2)/(p85-p15);
  __&x=(max(0,&x-p29)**2-max(0,&x-p85)**2)/(p85-p15);
  ___&x=(max(0,&x-p43)**2-max(0,&x-p85)**2)/(p85-p15);
  ____&x=(max(0,&x-p57)**2-max(0,&x-p85)**2)/(p85-p15);
  _____&x=(max(0,&x-p71)**2-max(0,&x-p85)**2)/(p85-p15);
%end;
%else %if &k=7 %then %do;
  _&x=(max(0,&x-p12)**2-max(0,&x-p88)**2)/(p88-p12);
  __&x=(max(0,&x-p24)**2-max(0,&x-p88)**2)/(p88-p12);
  ___&x=(max(0,&x-p37)**2-max(0,&x-p88)**2)/(p88-p12);
  ____&x=(max(0,&x-p50)**2-max(0,&x-p88)**2)/(p88-p12);
  _____&x=(max(0,&x-p63)**2-max(0,&x-p88)**2)/(p88-p12);
  _____&x=(max(0,&x-p76)**2-max(0,&x-p88)**2)/(p88-p12);
%end; %end;

```

```
drop p3 p5 p12 p15 p16 p18 p20 p23 p24 p25 p28 p29 p33 p34 p35 p37 p40 p41 p43 p50 p57 p59 p60 p63 p65 p66
p67 p71 p73 p75 p76 p77 p80 p82 p84 p85 p88 p95 p98;
run; quit; run;
options notes;
%mend rqspline;
```

THE UNIVERSITY OF NORTH CAROLINA CENTER FOR AIDS RESEARCH HIV CLINICAL COHORT

In 1999 the University of North Carolina Center for AIDS Research (UNC CFAR) initiated a prospective clinic cohort study of HIV-infected patients attending the UNC HIV clinic. Details regarding this cohort are provided elsewhere.⁵ The institutional review board at UNC approved all study forms. Written informed consent was obtained in English or Spanish from all enrolled participants. Here we analyze 557 male participants seen for a scheduled visit after 1 January 1999 in the clinic cohort who initiated combination antiretroviral therapy (henceforth, therapy) at or after their first visit to the UNC HIV clinic, and had CD4 and HIV-1 viral load measured at least once during the 12 months prior to therapy initiation. Patients were administratively censored at 31 May 2010. Therapy was defined as the concurrent use of three or more antiretroviral agents. Over the 2,537 person years of follow up 59 participants died. Given that therapy could have been initiated between entry into the UNC HIV clinic and cohort enrollment the person time included during this interval is immortal and may introduce a survival bias.⁶ However, because the objective of this example is to demonstrate the use of RQS this issue was ignored in all corresponding analyses.

DATA USED IN EXAMPLES

1	15	3.5740300	0.000000000	10.143737166	0
2	339	1.8286600	0.000000000	4.559206023	0
3	35	3.0345400	0.000000000	8.164271047	1
4	203	2.4417700	0.000000000	2.231348392	1
5	67	3.5355900	0.000000000	5.947296372	0
6	22	3.1953300	0.000000000	9.927446954	0
7	5	3.3090900	0.000000000	9.333333333	0
8	11	3.1473000	0.000000000	9.982203970	0
9	10	3.5740300	0.000000000	9.114305270	0
10	447	2.7224700	0.000000000	9.429158111	0
11	266	1.7430400	0.000000000	8.060232717	0
12	5	3.0267700	0.000000000	10.080766598	0
13	35	2.5740400	0.000000000	7.718001369	0
14	209	2.2560300	0.000000000	8.944558522	0
15	237	2.9435500	0.000000000	3.064339493	0
16	26	2.8577400	0.000000000	0.643394935	1
17	27	2.1945100	0.000000000	5.708418891	0
18	398	0.0000000	0.000000000	10.083504449	0
19	384	1.7893700	0.000000000	6.697467488	0
20	265	0.0000000	0.000000000	1.853524983	0
21	136	2.5708900	0.000000000	8.222450376	0
22	69	3.3670800	0.000000000	5.527720739	1
23	350	2.5837700	0.000000000	2.738535250	0
24	131	1.1147800	0.000000000	8.777549624	0
25	207	2.9345000	0.000000000	9.122518823	0
26	168	2.9599300	0.000000000	10.231348392	0
27	67	2.1253700	0.000000000	10.146475017	0
28	456	2.3927300	0.000000000	7.874743326	0
29	13	3.2819000	0.000000000	10.036960986	0
30	545	1.5651400	0.000000000	9.875427789	0
31	202	2.2403700	0.000000000	3.491444216	0
32	193	3.1034100	0.000000000	3.543463381	0
33	119	3.2897800	0.000000000	4.388774812	1
34	146	1.4336900	0.000000000	2.932922656	0
35	5	3.5267700	0.000000000	0.260095825	1
36	150	1.9794300	0.000000000	6.639288159	1
37	529	1.9001800	0.000000000	8.704312115	0
38	311	2.5174800	0.000000000	2.952087611	0
39	106	2.5934000	0.000000000	9.943874059	0
40	227	3.0341700	0.000000000	9.842573580	0
41	12	3.5559400	0.000000000	10.165639973	0
42	203	2.2388700	0.000000000	9.667351129	0
43	363	2.7130800	0.000000000	5.120465435	0

44	47	2.0776600	0.000000000	3.712525667	1
45	309	1.7328800	0.000000000	10.015058179	0
46	44	3.3866500	0.000000000	10.053388090	0
47	415	2.4649600	0.000000000	9.946611910	0
48	634	2.3848300	0.000000000	9.880903491	0
49	5	3.2086100	0.000000000	9.831622177	0
50	58	3.3576000	0.000000000	9.708418891	0
51	14	3.2583200	0.000000000	9.524982888	0
52	449	2.6216600	0.000000000	9.273100616	0
53	37	2.9230200	0.000000000	7.986310746	1
54	5	3.5740300	0.000000000	9.560574949	0
55	50	2.2120400	0.000000000	9.152635181	0
56	5	3.5401300	0.000000000	7.759069131	1
57	181	3.1254700	0.000000000	9.275838467	0
58	103	3.5740300	0.000000000	9.253935661	0
59	44	3.4374200	0.000000000	9.382614648	0
60	5	0.0000000	0.000000000	9.327857632	0
61	5	2.6242200	0.000000000	9.360711841	0
62	62	3.1646400	0.000000000	9.267624914	0
63	150	2.2233000	0.000000000	9.002053388	0
64	52	3.5740300	0.000000000	8.441478439	0
65	5	1.9868200	0.000000000	2.494866530	0
66	542	2.2809200	0.000000000	8.843258042	0
67	235	2.7004100	0.000000000	8.884325804	0
68	169	2.4567500	0.000000000	1.468172485	0
69	5	3.3521500	0.000000000	8.583846680	0
70	239	2.4201000	0.000000000	7.819986311	0
71	5	2.5740400	0.000000000	6.919233402	0
72	201	2.2238200	0.000000000	4.536618754	0
73	692	1.8257500	0.000000000	9.670088980	0
74	176	2.8057800	0.000000000	9.806981520	0
75	62	1.9439400	0.000000000	3.616700890	1
76	5	3.5296200	0.000000000	1.848049281	1
77	111	2.5740400	0.000000000	8.908966461	0
78	555	2.4894700	0.000000000	0.950034223	0
79	203	0.9604700	0.000000000	10.135523614	0
80	28	2.2001400	0.000000000	9.462012320	0
81	99	2.6214600	0.000000000	4.982888433	1
82	712	2.5069900	0.000000000	9.639972621	0
83	5	3.5740300	0.000000000	6.004106776	1
84	107	3.5740300	0.000000000	3.127310062	0
85	48	3.5740300	0.000000000	5.913757700	0
86	5	3.1186200	0.000000000	2.237508556	0
87	160	2.3566500	0.000000000	8.711841205	0
88	128	3.1120300	0.000000000	7.926078029	0

89	11	3.4327700	0.000000000	1.785763176	0
90	113	2.2696300	0.000000000	9.390828200	0
91	195	3.4835000	0.000000000	9.618069815	0
92	407	1.6902800	0.000000000	9.716632444	0
93	548	2.3244200	0.000000000	5.544832307	0
94	5	2.9425100	0.000000000	6.611909651	1
95	16	3.0475700	0.000000000	8.777549624	0
96	47	3.5740300	0.000000000	8.804928131	0
97	279	2.2707500	0.000000000	4.533880903	0
98	17	3.5740300	0.000000000	7.798083504	0
99	504	1.1931200	0.000000000	0.911704312	0
100	89	2.7920800	0.000000000	7.685831622	0
101	303	2.1453100	0.000000000	5.079397673	0
102	90	2.5740400	0.000000000	8.758384668	0
103	94	3.4935600	0.000000000	2.904859685	1
104	123	2.4456000	0.000000000	8.733744011	0
105	5	2.5740400	0.000000000	8.709103354	0
106	90	3.5740300	0.000000000	6.009582478	1
107	651	3.4392000	0.000000000	2.787816564	0
108	12	3.4715500	0.000000000	8.616016427	0
109	86	3.5740300	0.000000000	4.832991102	0
110	544	2.0140600	0.000000000	2.809719370	0
111	288	2.9519000	0.000000000	4.255304586	0
112	17	2.7821200	0.000000000	8.659822040	0
113	264	1.7391400	0.000000000	8.177960301	0
114	214	1.4616500	0.000000000	2.308008214	0
115	5	3.5740300	0.000000000	0.561259411	1
116	275	3.0491500	0.000000000	0.232717317	1
117	95	3.4023800	0.000000000	3.581793292	0
118	197	0.0000000	0.000000000	8.487337440	0
119	161	2.5740400	0.000000000	8.216290212	0
120	266	2.6365500	0.000000000	7.488021903	1
121	162	2.8020900	0.000000000	1.160848734	0
122	286	2.3316600	0.000000000	0.720739220	0
123	21	3.1838100	0.000000000	2.270362765	0
124	290	2.4332300	0.000000000	7.129363450	0
125	262	3.3371400	0.000000000	7.126625599	0
126	33	2.6933400	0.000000000	5.402464066	0
127	235	1.4609000	0.000000000	5.670773443	0
128	176	2.1049800	0.000000000	1.251882272	0
129	388	2.3927000	0.000000000	6.225872690	0
130	157	2.9433000	0.000000000	6.533196441	0
131	269	2.8870100	0.000000000	8.079397673	0
132	23	3.3961100	0.000000000	8.227241615	0
133	332	2.9086600	0.000000000	4.652292950	0

134	286	0.00000000	0.000000000	5.210130048	0
135	50	2.2577700	0.000000000	2.418206708	0
136	22	3.5740300	0.000000000	2.031485284	1
137	416	2.3179800	0.000000000	6.763175907	0
138	316	0.9867700	0.000000000	6.108145106	0
139	437	0.4337700	0.000000000	7.707049966	0
140	349	2.9458600	0.000000000	1.785763176	0
141	205	2.2019600	0.000000000	7.854893908	0
142	330	0.00000000	0.000000000	6.707734428	0
143	68	3.4721300	0.000000000	5.577686516	0
144	5	2.8882300	0.000000000	0.936344969	1
145	49	3.5740300	0.000000000	0.487337440	1
146	402	2.6989700	0.000000000	1.104038330	0
147	20	3.5740300	0.000000000	7.871321013	0
148	39	3.5740300	0.000000000	7.723477070	0
149	193	0.4321700	0.000000000	7.575633128	0
150	167	3.2056200	0.000000000	7.728952772	0
151	26	3.3865100	0.000000000	7.042436687	0
152	14	3.5328500	0.000000000	0.788501027	1
153	518	2.5498900	0.000000000	7.186858316	0
154	386	3.1440700	0.000000000	7.693360712	0
155	26	2.8724200	0.000000000	7.704312115	0
156	11	2.5159800	0.000000000	7.638603696	0
157	158	2.8572800	0.000000000	7.570157426	0
158	184	2.5256100	0.000000000	4.583846680	0
159	206	2.5868800	0.000000000	7.679671458	0
160	316	2.4133000	0.000000000	5.957563313	0
161	106	3.3362100	0.000000000	7.233401780	0
162	38	2.7136000	0.000000000	3.472279261	0
163	216	2.4788500	0.000000000	6.973305955	0
164	109	2.5911300	0.000000000	1.457221081	0
165	49	2.7042200	0.000000000	1.769336071	0
166	52	2.9641800	0.000000000	5.535934292	1
167	181	1.8952000	0.000000000	6.954140999	0
168	262	2.1653600	0.000000000	6.951403149	0
169	38	2.4397400	0.000000000	3.898699521	0
170	12	2.4625200	0.000000000	7.271731691	0
171	5	2.7976700	0.000000000	7.140314853	0
172	254	2.4756700	0.000000000	2.147159480	0
173	62	3.1430600	0.000000000	4.495550992	1
174	33	3.3878100	0.000000000	4.698836413	0
175	88	3.3936200	0.000000000	7.085557837	0
176	261	2.4281800	0.000000000	6.609171800	0
177	112	2.4460600	0.000000000	9.535934292	0
178	70	2.2236600	0.000000000	7.101984942	0

179	64	2.4607700	0.000000000	6.984257358	0
180	120	3.4852700	0.000000000	6.666666667	1
181	42	3.2538400	0.000000000	7.063655031	0
182	5	3.3509700	0.000000000	1.344969199	0
183	307	1.8208300	0.000000000	6.891170431	0
184	17	3.5740300	0.000000000	1.311430527	1
185	247	2.3566400	0.000000000	1.761122519	0
186	181	3.0758500	0.000000000	4.958247775	0
187	5	3.3025700	0.000000000	0.342231348	1
188	13	2.8688300	0.000000000	1.160848734	1
189	131	2.6550700	0.000000000	1.295687885	0
190	401	1.7323900	0.000000000	1.623545517	0
191	5	2.3255500	0.000000000	3.674195756	0
192	69	3.4977400	0.000000000	6.877481177	0
193	123	2.9220100	0.000000000	0.747433265	1
194	250	2.6117200	0.000000000	3.843942505	0
195	61	2.6181800	0.000000000	6.819986311	0
196	301	0.9894500	0.000000000	4.533880903	0
197	5	3.5740300	0.000000000	6.850102669	0
198	59	3.5740300	0.000000000	6.376454483	0
199	51	2.2290300	0.000000000	6.773442847	0
200	108	3.4936700	0.000000000	5.405201916	0
201	51	2.2579100	0.000000000	6.718685832	0
202	276	2.0273500	0.000000000	6.050650240	0
203	16	3.5740300	0.000000000	6.707734428	0
204	198	2.6989700	0.000000000	3.247091034	0
205	184	3.5740300	0.000000000	6.669404517	0
206	276	2.8558500	0.000000000	6.625598905	0
207	157	0.0000000	0.000000000	6.644763860	0
208	63	3.5740300	0.000000000	6.475017112	0
209	129	3.4855200	0.000000000	6.631074606	0
210	249	2.5932900	0.000000000	3.031485284	0
211	88	3.1546800	0.000000000	6.532511978	0
212	116	2.6989700	0.000000000	2.280629706	0
213	518	1.4224300	0.000000000	5.338809035	0
214	5	2.3335000	0.000000000	6.245037645	0
215	394	1.6190900	0.000000000	3.236824093	0
216	118	3.5740300	0.000000000	6.198494182	0
217	130	3.5740300	0.000000000	6.401095140	0
218	254	1.4409100	0.000000000	6.015742642	0
219	428	1.5969800	0.000000000	3.000684463	0
220	46	2.2228900	0.000000000	6.439425051	0
221	73	3.4548400	0.000000000	5.831622177	0
222	5	2.9956400	0.000000000	0.945242984	0
223	361	2.8388500	0.000000000	2.034907598	0

224	941	0.00000000	0.000000000	0.720054757	0
225	326	2.6989700	0.000000000	5.152635181	0
226	57	3.0700400	0.000000000	5.798767967	0
227	258	2.8059100	0.000000000	5.787816564	0
228	50	2.5194400	0.000000000	1.859685147	0
229	406	3.1089000	0.000000000	5.489390828	0
230	81	2.6989700	0.000000000	3.839151266	0
231	56	3.3128100	0.000000000	5.820670773	0
232	15	3.3453700	0.000000000	5.790554415	0
233	5	1.9754300	0.000000000	1.757700205	1
234	11	3.5740300	0.000000000	5.309377139	0
235	5	2.8162400	0.000000000	3.529774127	0
236	43	3.1258400	0.000000000	5.571526352	0
237	43	3.5740300	0.000000000	5.459274470	0
238	53	3.1687900	0.000000000	4.759069131	0
239	542	0.00000000	0.000000000	0.720054757	0
240	149	1.8388500	0.000000000	5.054072553	0
241	268	2.7708500	0.000000000	3.159479808	1
242	353	1.7593300	0.000000000	3.072553046	0
243	68	3.5740300	0.000000000	5.459274470	0
244	128	2.7520500	0.000000000	2.223819302	0
245	433	2.6811200	0.000000000	3.529089665	0
246	386	0.00000000	0.000000000	5.325119781	0
247	504	2.3819300	0.000000000	2.064339493	1
248	79	3.5740300	0.000000000	5.210130048	0
249	272	2.4690400	0.000000000	2.990417522	0
250	16	3.3032000	0.000000000	2.503080082	0
251	44	3.5740300	0.000000000	5.364134155	0
252	123	3.5740300	0.000000000	5.182751540	0
253	24	1.9684800	0.000000000	2.284052019	0
254	299	1.9545600	0.000000000	2.924709103	0
255	10	2.6369300	0.000000000	5.021218344	0
256	128	3.5450000	0.000000000	3.362080767	1
257	32	2.5247100	0.000000000	4.744695414	0
258	84	3.2625100	0.000000000	1.364134155	0
259	180	3.5740300	0.000000000	4.922655715	0
260	163	2.6190900	0.000000000	4.703627652	0
261	155	2.1222200	0.000000000	5.494866530	0
262	577	2.4848800	0.000000000	4.635181383	0
263	94	3.5740300	0.000000000	4.837782341	0
264	102	2.4649600	0.000000000	3.392881588	0
265	53	3.4511400	0.000000000	2.618069815	0
266	139	3.3774900	0.000000000	4.533880903	0
267	5	1.9365400	0.000000000	1.872689938	1
268	217	3.1268600	0.000000000	3.403832991	0

269	22	2.0022500	0.000000000	4.700889802	0
270	381	2.0673500	0.000000000	2.335386721	0
271	311	2.1673600	0.000000000	5.338809035	0
272	311	2.0719400	0.000000000	3.830253251	0
273	39	2.6989700	0.000000000	3.510609172	0
274	388	2.2384900	0.000000000	3.156741958	0
275	260	1.4672400	0.000000000	3.802874743	0
276	591	1.9916200	0.000000000	2.927446954	0
277	370	2.5414400	0.000000000	4.465434634	0
278	5	3.2464800	0.000000000	4.479123888	0
279	28	3.2530600	0.000000000	4.421629021	0
280	207	3.2712600	0.000000000	3.023271732	0
281	53	3.5740300	0.000000000	5.314852841	0
282	463	0.4005400	0.000000000	1.383299110	0
283	361	1.7991300	0.000000000	4.167008898	0
284	466	2.8461600	0.000000000	4.347707050	0
285	239	2.5481200	0.000000000	4.331279945	0
286	262	2.4730300	0.000000000	4.331279945	0
287	104	2.1486600	0.000000000	4.446269678	0
288	118	3.5740300	0.000000000	4.342231348	0
289	304	2.6836900	0.000000000	4.386036961	0
290	622	0.0000000	0.000000000	2.283367556	0
291	179	2.2599300	0.000000000	4.290212183	0
292	477	2.2467800	0.000000000	3.316221766	0
293	49	3.5104900	0.000000000	1.368925394	1
294	234	2.6419700	0.000000000	4.112251882	0
295	294	2.1717300	0.000000000	2.201232033	0
296	99	3.4175300	0.000000000	4.136892539	0
297	273	1.4800100	0.000000000	4.169746749	0
298	187	2.4158100	0.000000000	4.057494867	0
299	309	2.6662500	0.000000000	3.250513347	0
300	468	1.9826600	0.000000000	3.961670089	0
301	342	0.9588000	0.000000000	3.928815880	0
302	152	3.5428300	0.000000000	1.448323066	0
303	227	2.3239800	0.000000000	3.805612594	0
304	106	2.3184300	0.000000000	3.405886379	0
305	345	3.2459800	0.000000000	3.860369610	0
306	5	3.3058800	0.000000000	3.947980835	0
307	37	2.7318600	0.000000000	3.616700890	0
308	27	2.3554800	0.000000000	3.967145791	0
309	385	0.7937900	0.000000000	3.926078029	0
310	182	1.3555500	0.000000000	3.693360712	0
311	32	3.5740300	0.000000000	4.427104723	0
312	23	2.4438100	0.000000000	3.843942505	0
313	564	0.0000000	0.000000000	1.705681040	1

314	176	2.5005200	0.000000000	3.832991102	0
315	343	2.6000200	0.000000000	3.485284052	0
316	47	3.5740300	0.000000000	2.804243669	0
317	140	2.5332900	0.000000000	3.157426420	0
318	27	2.6192400	0.000000000	3.843942505	0
319	127	2.4357100	0.000000000	3.597535934	0
320	88	2.8479400	0.000000000	3.657768652	0
321	648	2.8508700	0.000000000	1.453798768	0
322	183	2.1666400	0.000000000	3.085557837	0
323	22	3.0636100	0.000000000	3.635865845	0
324	805	0.0000000	0.000000000	3.693360712	0
325	203	2.5924600	0.000000000	3.581108830	0
326	286	1.0993400	0.000000000	3.537303217	0
327	277	3.5740300	0.000000000	3.594798084	0
328	305	2.2627500	0.000000000	3.468856947	0
329	388	1.7075700	0.000000000	1.894592745	0
330	343	1.9469400	0.000000000	1.831622177	0
331	145	2.6509600	0.000000000	2.374401095	0
332	318	1.9216900	0.000000000	1.607118412	0
333	84	2.7823300	0.000000000	3.351129363	0
334	23	3.2686700	0.000000000	3.252566735	0
335	389	3.4694100	0.000000000	3.441478439	0
336	30	2.3858900	0.000000000	3.381245722	0
337	252	2.8895700	0.000000000	3.471594798	1
338	242	1.6758700	0.000000000	2.447638604	0
339	86	2.6078600	0.000000000	3.233401780	0
340	17	2.6110700	0.000000000	3.189596167	0
341	516	1.3580300	0.000000000	2.598220397	0
342	539	2.0320900	0.000000000	2.040383299	0
343	480	1.9258500	0.000000000	2.354551677	0
344	347	1.7157500	0.000000000	1.095824778	0
345	61	0.3909400	0.000000000	3.222450376	0
346	305	3.5484600	0.000000000	2.948665298	0
347	525	2.6455800	0.000000000	0.919917864	0
348	587	3.4748800	0.000000000	3.044490075	0
349	21	3.5740300	0.000000000	2.852840520	0
350	1012	3.5740300	0.000000000	2.981519507	0
351	5	2.3171600	0.000000000	2.770704997	0
352	696	1.4673100	0.000000000	2.872005476	0
353	363	1.8317400	0.000000000	2.702258727	0
354	272	1.9445600	0.000000000	2.447638604	0
355	5	2.1389800	0.000000000	2.485968515	0
356	225	1.8654900	0.000000000	1.402464066	0
357	30	3.5740300	0.000000000	0.796714579	1
358	414	2.3736200	0.000000000	2.464065708	0

359	9	2.5665000	0.000000000	2.642026010	0
360	5	3.2718000	0.000000000	1.969199179	0
361	5	2.9878100	0.000000000	2.587268994	0
362	403	1.7155400	0.000000000	2.644763860	0
363	257	2.2594000	0.000000000	2.639288159	0
364	189	2.3074600	0.000000000	2.428473648	0
365	78	2.6989700	0.000000000	2.524298426	0
366	235	1.9559000	0.000000000	2.469541410	0
367	310	1.2430400	0.000000000	1.598904860	0
368	286	1.4771200	0.000000000	2.502395619	0
369	420	2.7103900	0.000000000	0.843258042	0
370	5	3.5740300	0.000000000	2.179329227	0
371	25	3.4773800	0.000000000	2.294318960	0
372	33	2.8755600	0.000000000	2.272416153	0
373	56	2.5963300	0.000000000	2.228610541	0
374	241	1.5422000	0.000000000	1.278576318	0
375	165	2.4083800	0.000000000	1.724845996	0
376	14	3.5740300	0.000000000	2.067077344	0
377	440	0.5415800	0.000000000	2.061601643	0
378	29	2.5860200	0.000000000	2.223134839	0
379	18	2.9777200	0.000000000	2.105407255	0
380	58	3.5740300	0.000000000	2.086242300	0
381	158	2.5927300	0.000000000	2.028747433	0
382	108	3.5740300	0.000000000	1.820670773	0
383	61	2.4661300	0.000000000	2.086242300	0
384	801	2.7921400	0.000000000	2.058863792	0
385	540	2.5085300	0.000000000	0.355920602	0
386	478	3.2870700	0.000000000	2.042436687	0
387	430	1.3953300	0.000000000	0.268309377	0
388	254	1.8481900	0.000000000	1.932922656	0
389	5	2.8162400	0.000000000	1.713894593	0
390	97	3.5740300	0.000000000	3.986310746	0
391	290	0.8388500	0.000000000	0.785763176	0
392	466	0.0000000	0.000000000	1.779603012	0
393	345	2.1861100	0.000000000	1.752224504	0
394	301	2.9493900	0.000000000	0.421629021	0
395	142	1.9956400	0.000000000	1.530458590	0
396	668	2.2342600	0.000000000	0.265571526	0
397	17	3.5740300	0.000000000	1.620807666	0
398	300	1.8779500	0.000000000	1.601642710	0
399	272	3.5740300	0.000000000	1.541409993	0
400	557	1.6932900	0.000000000	0.262833676	0
401	458	3.5740300	0.000000000	1.103353867	0
402	49	2.8721600	0.000000000	1.393566051	0
403	509	2.2002800	0.000000000	4.709103354	1

404	444	2.7714900	0.000000000	7.030800821	1
405	16	2.5740400	0.000000000	8.323750856	0
406	90	3.1507700	0.000000000	10.825462012	0
407	195	2.3496400	0.000000000	10.694045175	0
408	268	2.2237700	0.000000000	3.026009582	0
409	502	2.4839100	0.000000000	10.480492813	0
410	5	2.9825100	0.000000000	6.844626968	1
411	576	1.3821100	0.000000000	9.100616016	0
412	20	2.7103500	0.000000000	2.209445585	1
413	353	2.5644400	0.000000000	10.291581109	0
414	9	3.4201000	0.000000000	11.011635866	0
415	81	3.2821300	0.000000000	10.721423682	0
416	410	1.8750000	0.000000000	7.696098563	0
417	402	0.5769200	0.000000000	3.069815195	0
418	57	2.6169400	0.000000000	9.900068446	0
419	305	0.0000000	0.000000000	10.209445585	0
420	137	2.7217700	0.000000000	4.603011636	0
421	19	3.5740300	0.000000000	10.138261465	0
422	292	2.2230300	0.000000000	0.673511294	1
423	241	2.4826800	0.000000000	1.109514031	0
424	402	2.0839100	0.000000000	11.052703628	0
425	278	2.6677800	0.000000000	10.039698836	0
426	12	3.5740300	0.000000000	10.733059548	0
427	459	2.7143700	0.000000000	6.623545517	0
428	9	2.2825200	0.000000000	1.629705681	0
429	150	2.1359300	0.000000000	2.582477755	0
430	1174	1.1166100	0.000000000	0.865845311	0
431	56	3.4375100	0.000000000	1.769336071	0
432	108	3.3067300	0.000000000	10.948665298	1
433	95	2.5972000	0.000000000	11.148528405	0
434	339	3.0099100	0.000000000	11.148528405	0
435	119	3.2549400	0.000000000	0.736481862	1
436	131	3.5618000	0.000000000	3.332648871	0
437	140	1.4277300	0.000000000	1.492813142	0
438	351	0.0000000	0.000000000	1.440793977	0
439	9	2.3773300	0.000000000	0.492813142	1
440	252	1.9168000	0.000000000	10.858316222	0
441	306	1.8980400	0.000000000	6.644763860	1
442	9	3.0093700	0.000000000	2.277891855	1
443	499	1.4488600	0.000000000	8.137577002	0
444	219	2.2976600	0.000000000	10.647501711	0
445	518	2.5289700	0.000000000	7.839151266	0
446	95	3.5295600	0.000000000	0.950034223	0
447	40	3.5740300	0.000000000	10.798083504	0
448	99	2.7834000	0.000000000	4.339493498	1

449	182	3.5127900	0.000000000	5.292950034	0
450	447	3.5740300	0.000000000	3.061601643	0
451	84	3.4975200	0.000000000	4.457905544	0
452	271	2.2673200	0.000000000	7.775496235	1
453	9	1.0119900	0.000000000	10.787132101	0
454	218	2.6224700	0.000000000	2.894592745	0
455	12	3.0411100	0.000000000	7.679671458	1
456	68	3.1845700	0.000000000	10.663928816	0
457	548	0.0000000	0.000000000	10.683093771	0
458	113	3.5328800	0.000000000	3.948665298	0
459	591	2.0835600	0.000000000	9.618754278	0
460	2	3.4170000	0.000000000	7.075290897	0
461	18	2.7531100	0.000000000	0.503764545	1
462	67	2.5801300	0.000000000	10.543463381	0
463	462	2.4493500	0.000000000	2.686516085	0
464	413	3.0808100	0.000000000	10.387405886	0
465	306	2.5179200	0.000000000	7.748802190	0
466	266	2.5750500	0.000000000	10.516084873	0
467	22	2.5964000	0.000000000	10.496919918	0
468	150	3.3344900	0.000000000	9.418206708	1
469	29	3.1536600	0.000000000	10.039698836	0
470	13	3.4885100	0.000000000	4.017111567	0
471	14	2.4983800	0.000000000	0.057494867	1
472	1005	0.0000000	0.000000000	8.958247775	0
473	165	3.0583900	0.000000000	2.538672142	0
474	72	2.2789500	0.000000000	10.968514716	0
475	356	1.5262700	0.000000000	8.062970568	0
476	234	2.6479600	0.000000000	1.000000000	0
477	32	3.5740300	0.000000000	0.646132786	0
478	265	1.7611000	0.000000000	0.885010267	0
479	89	2.4528200	0.000000000	7.225188227	0
480	155	2.9211000	0.000000000	0.501026694	1
481	366	2.5403000	0.000000000	3.794661191	0
482	313	2.5949400	0.000000000	5.639972621	1
483	103	2.5118800	0.000000000	5.971252567	0
484	5	3.3215500	0.000000000	3.075290897	0
485	5	3.0395200	0.000000000	4.054757016	0
486	552	1.0683700	0.000000000	2.798083504	0
487	5	2.6989700	0.000000000	3.173169062	0
488	13	3.4811900	0.000000000	3.104722793	0
489	569	2.2749800	0.000000000	2.685831622	0
490	275	2.7181900	0.000000000	2.718685832	0
491	631	2.4043900	0.000000000	4.191649555	0
492	63	3.5740300	0.000000000	2.579055441	0
493	679	2.2928900	0.000000000	2.064339493	0

494	630	0.00000000	0.000000000	2.242299795	0
495	387	2.6478700	0.000000000	2.154688569	0
496	322	2.4432600	0.000000000	2.154688569	0
497	5	3.3404400	0.000000000	2.020533881	0
498	542	2.7422800	0.000000000	2.028747433	0
499	39	3.5740300	0.000000000	1.848049281	0
500	98	3.5037900	0.000000000	1.837097878	0
501	973	2.0889300	0.000000000	1.812457221	0
502	145	2.6989700	0.000000000	3.230663929	0
503	183	1.5544900	0.000000000	1.661875428	0
504	638	2.9455500	0.000000000	1.763175907	0
505	337	2.4742200	0.000000000	1.618069815	0
506	46	3.0354300	0.000000000	1.645448323	0
507	417	2.6833700	0.000000000	0.525667351	0
508	33	3.2304500	0.000000000	1.284736482	0
509	169	2.7242800	0.000000000	1.533196441	0
510	65	2.6803400	0.000000000	1.552361396	0
511	715	2.3019400	0.000000000	1.396303901	0
512	683	1.9542400	0.000000000	1.549623546	0
513	354	2.2405500	0.000000000	1.429158111	0
514	133	2.0827900	0.000000000	1.379876797	0
515	839	0.6725600	0.000000000	1.000000000	0
516	202	2.2576800	0.000000000	1.278576318	0
517	54	3.5071800	0.000000000	1.352498289	0
518	278	2.1174200	0.000000000	1.215605749	0
519	79	2.8095600	0.000000000	1.399041752	0
520	366	2.6491200	0.000000000	0.947296372	0
521	60	2.9637900	0.000000000	1.368925394	0
522	289	2.5854600	0.000000000	1.232032854	0
523	144	2.9754300	0.000000000	0.610540726	0
524	313	2.5763400	0.000000000	0.824093087	1
525	494	1.1861100	0.000000000	1.084188912	0
526	418	2.6725600	0.000000000	1.205338809	0
527	25	2.6906400	0.000000000	1.316906229	0
528	212	2.7118100	0.000000000	1.152635181	0
529	408	1.6941200	0.000000000	0.947296372	0
530	561	2.3317100	0.000000000	0.744695414	0
531	754	2.0393500	0.000000000	1.199178645	0
532	209	2.5165400	0.000000000	1.032169747	0
533	72	2.2977600	0.000000000	1.051334702	0
534	960	1.5409500	0.000000000	0.492813142	0
535	24	3.1361900	0.000000000	1.097878166	0
536	412	1.9150600	0.000000000	0.955509925	0
537	202	1.5938400	0.000000000	1.051334702	0
538	209	3.3854300	0.000000000	0.988364134	0

539	426	2.2821700	0.000000000	0.977412731	0
540	324	1.4143000	0.000000000	0.900752909	0
541	336	1.9622000	0.000000000	0.958247775	0
542	621	2.8673100	0.000000000	0.793976728	0
543	603	0.9003700	0.000000000	0.758384668	0
544	196	3.5740300	0.000000000	0.791238877	0
545	461	2.9420100	0.000000000	0.829568789	0
546	380	1.9867700	0.000000000	0.878850103	0
547	253	3.4674700	0.000000000	0.914442163	0
548	374	1.6345300	0.000000000	0.783025325	0
549	307	1.9092100	0.000000000	0.372347707	0
550	840	2.6666000	0.000000000	0.783025325	0
551	5	2.1861100	0.000000000	0.610540726	0
552	1175	1.4970000	0.000000000	0.651608487	0
553	25	2.7993400	0.000000000	0.555783710	0
554	11	2.9138100	0.000000000	0.624229979	0
555	411	3.5072700	0.000000000	0.476386037	0
556	417	3.4741700	0.000000000	0.476386037	0
557	575	2.1748700	0.000000000	0.454483231	0

SAS (VERSION 9.2) CODE USED IN EXAMPLES

****NOTE:** Before running the below SAS code the user must save the restricted quadratic spline macro (i.e., rqsmacro) in the same location as the SAS library name. To compare results from the restricted quadratic spline macro to Harrell's daspline macro, the user must download both the daspline and dshide macros from the referenced website and save these macros in the same location as the SAS library name. The user may also have to tailor graphing 'goptions' for their particular SAS version and computing system.**

*Automatically clearing the log and output windows;
dm log "clear;" continue; dm out "clear;" continue;

*Setting the following SAS options: page size "ps", line size "ls", turn-off date and start each run of output with page 1;
options ps=60 ls=90 nodate pageno=1;

*Specifying SAS libray name;
libname spline "C:\";

*Reading in UNC CFAR data on 557 male participants and calling temporary data file, 'a';
*Only first 5 records are displayed, however all 557 records must be included after the 'datalines' statement;
data a;

```
input id cd4 log10rnac in out dead;
label id="Participant ID";
label cd4="CD4 cell count (cells/mm^3) at therapy initiation";
label log10rnac="Centered HIV-1 viral load (log10 copies/ml) at therapy initiation";
label in="Years since therapy initiation at start of follow up in years";
label out="Years since therapy initiation at end of follow up in years";
label dead="Indicator of death";
datalines;
```

```
1 15 3.5740300 0.000000000 10.143737166 0
2 339 1.8286600 0.000000000 4.559206023 0
3 35 3.0345400 0.000000000 8.164271047 1
```

```
4 203 2.4417700 0.000000000 2.231348392 1
5 67 3.5355900 0.000000000 5.947296372 0
run;
```

```
*Printing contents of 'a' data;
proc contents data=a; run;
```

```
*Creating indicators for HIV-1 viral load and CD4 cell count;
data a;
```

```
    set a;
    highlog10nac=0;
    if log10nac>1 then highlog10nac=1;
```

```
    log10nac1i=0; log10nac2i=0; log10nac3i=0;
    if log10nac>2.17487 and log10nac<=2.61172 then log10nac1i=1;
    if log10nac>2.61172 and log10nac<=3.20562 then log10nac2i=1;
    if log10nac>3.20562 then log10nac3i=1;
```

```
    log10nacd1i=0; log10nacd2i=0; log10nacd3i=0; log10nacd4i=0;
    log10nacd5i=0;    log10nacd6i=0; log10nacd7i=0;    log10nacd8i=0;
    log10nacd9i=0;
    if log10nac>1.47712 and log10nac<=1.98677 then log10nacd1i=1;
    if log10nac>1.98677 and log10nac<=2.26732 then log10nacd2i=1;
    if log10nac>2.26732 and log10nac<=2.46904 then log10nacd3i=1;
    if log10nac>2.46904 and log10nac<=2.61172 then log10nacd4i=1;
    if log10nac>2.61172 and log10nac<=2.78233 then log10nacd5i=1;
    if log10nac>2.78233 and log10nac<=3.03952 then log10nacd6i=1;
    if log10nac>3.03952 and log10nac<=3.34537 then log10nacd7i=1;
    if log10nac>3.34537 and log10nac<=3.5618 then log10nacd8i=1;
    if log10nac>3.5618 then log10nacd9i=1;
```

```
cd4bin=0;
if cd4<=350 then cd4bin=1;
```

```
label highlog10rnac="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation >1 ";
```

```
label log10rnac1i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >2.17487 to 2.61172";
label log10rnac2i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >2.61172 to 3.20562";
label log10rnac3i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >3.20562 to 3.57403";
```

```
label log10rnacd1i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >1.47712 to 1.98677";
label log10rnacd2i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >1.98677 to 2.26732";
label log10rnacd3i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >2.26732 to 2.46904";
label log10rnacd4i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >2.46904 to 2.61172";
label log10rnacd5i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >2.61172 to 2.78233";
label log10rnacd6i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >2.78233 to 3.03952";
label log10rnacd7i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >3.03952 to 3.34537";
label log10rnacd8i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >3.34537 to 3.5618";
label log10rnacd9i="Indicator of centered HIV-1 viral load (log10 copies/ml) at therapy initiation from >3.5618 to 3.57403";
```

```
label cd4bin="Indicator of CD4 cell count (cells/mm3) at therapy initiation <=350";
```

```
run;
```

```
data a;
```

```
retain id cd4 cd4bin log10rnac highlog10rnac log10rnac1i log10rnac2i log10rnac3i log10rnacd1i log10rnacd2i log10rnacd3i
log10rnacd4i log10rnacd5i log10rnacd6i log10rnacd7i log10rnacd8i log10rnacd9i in out dead;
```

```
set a;
```

```
keep id cd4 cd4bin log10rnac highlog10rnac log10rnac1i log10rnac2i log10rnac3i log10rnacd1i log10rnacd2i log10rnacd3i
log10rnacd4i log10rnacd5i log10rnacd6i log10rnacd7i log10rnacd8i log10rnacd9i in out dead;
```

```
run;
```

*Examine distribution of the data;

```
proc means data=a n min mean p25 median p75 max sum; var cd4 log10rnac in out;
proc freq data=a; tables cd4bin highlog10rnac log10rnac1i log10rnac2i log10rnac3i log10rnacd1i log10rnacd2i log10rnacd3i
log10rnacd4i log10rnacd5i log10rnacd6i log10rnacd7i log10rnacd8i log10rnacd9i dead;
run;
```

Example 1: Use of a spline when estimating the association between centered log10 HIV-1 viral load centered at 2.3010 log10 copies/ml and hazard of death

*****;

*Generating scatterplot of centered log10 HIV-1 viral load by CD4 count;

```
proc sort data=a; by cd4;
```

```
options nogstyle FONTRENDERING=HOST_PIXELS;
```

```
goptions reset=all;
```

```
filename grafout "C:\figure1.png";
```

```
goptions device=zpng target=png gsfname=grafout gsfmode=replace xmax=6 ymax=6 xpixels=3600 ypixels=3600;
```

```
axis1 color=black
```

```
label=(angle=90 font=swiss h=2 justify=center "HIV-1, log10 copies/ml") w=2
```

```
major=(h=1 w=2) minor=none value=(font=swiss h=2) order=(-1 to 4 by 1) offset=(0,0);
```

```
axis2 color=black
```

```
label=(font=swiss h=2 justify=center "CD4, cells/mm^3") w=2
```

```
major=(h=1 w=2) minor=none value=(font=swiss h=2) order=(0 to 1500 by 500) offset=(0,0);
```

```
symbol1 c=black l=1 w=3 h=.8 v=dot i=none;
```

```
proc gplot data=a;
```

```
plot log10rnac*cd4/overlay vaxis=axis1 haxis=axis2 noframe nolegend; run; quit; run;
```

Assumption of log-linear association between centered log10 HIV-1 viral load and hazard of death;

*Fitting Cox model;

```
proc phreg data=a;
```

```
model out*dead(0)= log10rnac /entry=in ties=efron rl;
```

```
output out=xbeta (keep=xbeta xbetase log10rnac) xbeta=xbeta STDXBETA=xbetase; run;
```

```
proc sort data=xbeta; by log10rnac;
```

```

data log10rnacrplot;
  set xbeta;
  xbetaul=xbeta+1.96*xbetase;
  xbetall=xbeta-1.96*xbetase;
  rh=exp(xbeta);
  rhul=exp(xbetaul);
  rhll=exp(xbetall);
proc means data=log10rnacrplot min max; var rh rhul rhll log10rnac; run;
*Plotting relative hazard of death as a function of centered log10 HIV-1 viral load;
options nogstyle FONTRENDERING=HOST_PIXELS;
goptions reset=all;
filename grafout "C:\figure2a.png";
goptions device=zpng target=png gsfname=grafout gsfmode=replace xmax=6 ymax=6 xpixels=3600 ypixels=3600;
axis1 color=black logbase=e logstyle=expand interval=uneven order=(.1 .2 .5 1 2 5 10 100)
      label=(angle=90 font=swiss h=2 "Relative hazard of death") w=2
      major=(h=1 w=2) minor=none value=(font=swiss h=2 tick=1 '0.1' tick=2 '0.2' tick=3 '0.5' tick=4 '1.0' tick=5 '2.0' tick=6
'5.0' tick=7 '10.0' tick=8 '100.0') offset=(0,0);
axis2 color=black
      label=(font=swiss h=2 justify=center "HIV-1, log10 copies/ml") w=2
      major=(h=1 w=2) minor=none value=(font=swiss h=2) offset=(0,0);
symbol1 c=black l=1 w=3 h=.4 v=none i=join;
symbol2 c=black l=2 w=2 h=.4 v=none i=join;
symbol3 c=black l=2 w=2 h=.4 v=none i=join;
footnote move=(25,42) font=swiss color=black h=2 "AIC=655.91";
proc gplot data=log10rnacrplot;
  plot (rh rhul rhll)*log10rnac/overlay vaxis=axis1 haxis=axis2 noframe nolegend; run; quit; run;
footnote " "; run;

```

Using indicators corresponding to quartiles of centered log10 viral load with categories: >2.17487 to 2.61172, >2.61172 to 3.20562, >3.20562 to 3.57403 log10 copies/ml to model association between centered log10 HIV-1 viral load and hazard of death;

```

*Fitting Cox model;
proc phreg data=a;
    model out*dead(0)= log10rnac1i log10rnac2i log10rnac3i /entry=in ties=efron rl;
    output out=xbeta (keep=xbeta xbetase log10rnac log10rnac1i log10rnac2i log10rnac3i) xbeta=xbeta STDXBETA=xbetase;
run;
proc sort data=xbeta; by log10rnac;
data log10rnacrplot;
    set xbeta;
    xbetaul=xbeta+1.96*xbetase;
    xbetall=xbeta-1.96*xbetase;
    rh=exp(xbeta);
    rhul=exp(xbetaul);
    rhll=exp(xbetall);
proc means data=log10rnacrplot min max; var rh rhul rhll log10rnac; run;
*Plotting relative hazard of death as a function of centered log10 HIV-1 viral load;
options nogstyle FONTRENDERING=HOST_PIXELS;
goptions reset=all;
filename grafout "C:\figure2b.png";
goptions device=zpng target=png gsfname=grafout gsfmode=replace xmax=6 ymax=6 xpixels=3600 ypixels=3600;
axis1 color=black logbase=e logstyle=expand interval=uneven order=(.1 .2 .5 1 2 5 10 100)
    label=(angle=90 font=swiss h=2 "Relative hazard of death") w=2
    major=(h=1 w=2) minor=none value=(font=swiss h=2 tick=1 '0.1' tick=2 '0.2' tick=3 '0.5' tick=4 '1.0' tick=5 '2.0' tick=6
'5.0' tick=7 '10.0' tick=8 '100.0') offset=(0,0);
axis2 color=black
    label=(font=swiss h=2 justify=center "HIV-1, log10 copies/ml") w=2
    major=(h=1 w=2) minor=none value=(font=swiss h=2) offset=(0,0);
symbol1 c=black l=1 w=3 h=.4 v=none i=stepjs;
symbol2 c=black l=2 w=2 h=.4 v=none i=stepjs;
symbol3 c=black l=2 w=2 h=.4 v=none i=stepjs;
footnote move=(25,42) font=swiss color=black h=2 "AIC=656.47";
proc gplot data=log10rnacrplot;

```

```
plot (rh rhul rhll)*log10rnac/overlay vaxis=axis1 haxis=axis2 noframe nolegend; run; quit; run;
footnote " "; run;
```

```
***Using restricted quadratic spline with 4 equal knots at the 20th, 40th, 60th, and 80th percentiles
based on the case distribution to model association between centered log10 HIV-1 viral load and hazard of death***;
```

```
*Running rqs macro;
```

```
%include "C:\rqsmacro.sas";
```

```
%rqspline(data=a,x=log10rnac,event=dead,k=4,equal=1,cases=1);
```

```
*Fitting Cox model;
```

```
proc phreg data=a;
```

```
model out*dead(0)= log10rnac _log10rnac __log10rnac ___log10rnac /entry=in ties=efron rl;
```

```
test_nonlinear: test _log10rnac, __log10rnac, ___log10rnac;
```

```
output out=xbeta (keep=xbeta xbetase log10rnac) xbeta=xbeta STDXBETA=xbetase; run;
```

```
proc sort data=xbeta; by log10rnac;
```

```
data log10rnacrplot;
```

```
set xbeta;
```

```
xbetaul=xbeta+1.96*xbetase;
```

```
xbetall=xbeta-1.96*xbetase;
```

```
rh=exp(xbeta);
```

```
rhul=exp(xbetaul);
```

```
rhll=exp(xbetall);
```

```
proc means data=log10rnacrplot min max; var rh rhul rhll log10rnac; run;
```

```
*Plotting relative hazard of death as a function of centered log10 HIV-1 viral load;
```

```
options nogstyle FONTRENDERING=HOST_PIXELS;
```

```
goptions reset=all;
```

```
filename grafout "C:\figure2c.png";
```

```
goptions device=zpng target=png gsfname=grafout gsfmode=replace xmax=6 ymax=6 xpixels=3600 ypixels=3600;
```

```
axis1 color=black logbase=e logstyle=expand interval=uneven order=(.1 .2 .5 1 2 5 10 100)
```

```
label=(angle=90 font=swiss h=2 "Relative hazard of death") w=2
```

```
major=(h=1 w=2) minor=none value=(font=swiss h=2 tick=1 '0.1' tick=2 '0.2' tick=3 '0.5' tick=4 '1.0' tick=5 '2.0' tick=6
'5.0' tick=7 '10.0' tick=8 '100.0') offset=(0,0);
```



```

axis2 color=black
      label=(font=swiss h=2 justify=center "HIV-1, log10 copies/ml") w=2
      major=(h=1 w=2) minor=none value=(font=swiss h=2) offset=(0,0);
symbol1 c=black l=1 w=3 h=.4 v=none i=join;
symbol2 c=black l=2 w=2 h=.4 v=none i=join;
symbol3 c=black l=2 w=2 h=.4 v=none i=join;
footnote move=(25,42) font=swiss color=black h=2 "AIC=651.71";
proc gplot data=log10rnacrhplot;
      plot (rh rhul rhll)*log10rnac/overlay vaxis=axis1 haxis=axis2 noframe nolegend; run; quit; run;
footnote " "; run;

```

Using restricted cubic spline with 5 equal knots at the 16th, 33rd, 50th, 67th, and 84th percentiles based on the case distribution to model association between centered log10 HIV-1 viral load and hazard of death;

```

*Running daspline macro;
%include "C:\dshide.sas";
%include "C:\daspline.sas";
%daspline(x=log10rnac,nk=5,knot1=2.26732 2.75311 2.92302 3.28978 3.53285,data=a);
data a; set a; &_log10rnac; run;
*Fitting Cox model;
proc phreg data=a;
      model out*dead(0)= log10rnac log10rnac1 log10rnac2 log10rnac3 /entry=in ties=efron rl;
      test_nonlinear: test log10rnac1, log10rnac2, log10rnac3;
      output out=xbeta (keep=xbeta xbetase log10rnac) xbeta=xbeta STDXBETA=xbetase; run;
proc sort data=xbeta; by log10rnac;
data log10rnacrhplot;
      set xbeta;
      xbetaul=xbeta+1.96*xbetase;
      xbetall=xbeta-1.96*xbetase;
      rh=exp(xbeta);
      rhul=exp(xbetaul);
      rhll=exp(xbetall);

```

```

proc means data=log10rnacrplot min max; var rh rhul rhll log10rnac; run;
*Plotting relative hazard of death as a function of centered log10 HIV-1 viral load;
options nogstyle FONTRENDERING=HOST_PIXELS;
goptions reset=all;
filename grafout "C:\figure2d.png";
goptions device=zpng target=png gsfname=grafout gsfmode=replace xmax=6 ymax=6 xpixels=3600 ypixels=3600;
axis1 color=black logbase=e logstyle=expand interval=uneven order=(.1 .2 .5 1 2 5 10 100)
      label=(angle=90 font=swiss h=2 "Relative hazard of death") w=2
      major=(h=1 w=2) minor=none value=(font=swiss h=2 tick=1 '0.1' tick=2 '0.2' tick=3 '0.5' tick=4 '1.0' tick=5 '2.0' tick=6
'5.0' tick=7 '10.0' tick=8 '100.0') offset=(0,0);
axis2 color=black
      label=(font=swiss h=2 justify=center "HIV-1, log10 copies/ml") w=2
      major=(h=1 w=2) minor=none value=(font=swiss h=2) offset=(0,0);
symbol1 c=black l=1 w=3 h=.4 v=none i=join;
symbol2 c=black l=2 w=2 h=.4 v=none i=join;
symbol3 c=black l=2 w=2 h=.4 v=none i=join;
footnote move=(25,42) font=swiss color=black h=2 "AIC=652.67";
proc gplot data=log10rnacrplot;
      plot (rh rhul rhll)*log10rnac/overlay vaxis=axis1 haxis=axis2 noframe nolegend; run; quit; run;
footnote " "; run;

```

Example 2: Use of a spline when controlling for centered log10 HIV-1 viral load
(i.e., centered viral load > 1 log10 copies/ml) as a confounder of the association
between an indicator of CD4 <= 350 cells/mm³ high at therapy initiation and hazard of death

*****;

*Unadjusted for centered log10 HIV-1 viral load;

```

proc phreg data=a;
      model out*dead(0)= cd4bin /entry=in ties=efron rl; run;

```

*Adjusted for centered log10 HIV-1 viral load using binary indicator for centered HIV-1 viral load >1 log10 copies/ml;

```

proc phreg data=a;

```

```

    model out*dead(0)= cd4bin highlog10rnac/entry=in ties=efron rl; run;
*Adjusted for centered log10 HIV-1 viral load using 9 indicators for centered log10 HIV-1 viral load categories:
>1.47712 to 1.98677, >1.98677 to 2.26732, >2.26732 to 2.46904, >2.46904 to 2.61172, >2.61172 to 2.78233,
>2.78233 to 3.03952,>3.03952 to 3.34537, >3.34537 to 3.5618, and >3.5618 to 3.57403 cells/mm^3;
proc phreg data=a;
    model out*dead(0)= cd4bin log10rnacd1i log10rnacd2i log10rnacd3i log10rnacd4i log10rnacd5i log10rnacd6i log10rnacd7i
log10rnacd8i log10rnacd9i /entry=in ties=efron rl; run;
*Adjusted for centered log10 HIV-1 viral load assuming log-linear association between centered log10 HIV-1 viral load;
*and hazard of death;
proc phreg data=a;
    model out*dead(0)= cd4bin log10rnac /entry=in ties=efron rl; run;
*Adjusted for centered log10 HIV-1 viral load using restricted quadratic spline with with 4 equal knots at
the 20th, 40th, 60th, and 80th percentiles based on the case distribution;
proc phreg data=a;
    model out*dead(0)= cd4bin log10rnac _log10rnac __log10rnac ___log10rnac /entry=in ties=efron rl;
run;
*Adjusted for centered log10 HIV-1 viral load using restricted cubic spline with with 5 equal knots at
the 16th, 33rd, 50th, 67th, and 84th percentiles based on the case distribution;
proc phreg data=a;
    model out*dead(0)= cd4bin log10rnac log10rnac1 log10rnac2 log10rnac3 /entry=in ties=efron rl;
run;

```

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