

## nag\_prod\_limit\_surviv\_fn (g12aac)

### 1. Purpose

**nag\_prod\_limit\_surviv\_fn (g12aac)** computes the Kaplan-Meier, (or product-limit), estimates of survival probabilities for a sample of failure times.

### 2. Specification

```
#include <nag.h>
#include <nagg12.h>

void nag_prod_limit_surviv_fn(Integer n, double t[], Integer ic[],
                             Integer freq[], Integer *nd, double tp[], double p[],
                             double psig[], NagError *fail)
```

### 3. Description

A survivor function,  $S(t)$ , is the probability of surviving to at least time  $t$  with  $S(t) = 1 - F(t)$ , where  $F(t)$  is the cumulative distribution function of the failure times. The Kaplan-Meier or product limit estimator provides an estimate of  $S(t)$ ,  $\hat{S}(t)$ , from sample of failure times which may be progressively right-censored.

Let  $t_i$ ,  $i = 1, 2, \dots, n_d$ , be the ordered distinct failure times for the sample of observed failure/censored times, and let the number of observations in the sample that have not failed by time  $t_i$  be  $n_i$ . If a failure and a loss (censored observation) occur at the same time  $t_i$ , then the failure is treated as if it had occurred slightly before time  $t_i$  and the loss as if it had occurred slightly after  $t_i$ .

The Kaplan-Meier estimate of the survival probabilities is a step function which in the interval  $t_i$  to  $t_{i+1}$  is given by

$$\hat{S}(t) = \prod_{j=1}^i \left( \frac{n_j - d_j}{n_j} \right)$$

where  $d_j$  is the number of failures occurring at time  $t_j$ .

**nag\_prod\_limit\_surviv\_fn** computes the Kaplan-Meier estimates and the corresponding estimates of the variances,  $\text{var}(\hat{S}(t))$ , using Greenwood's formula,

$$\text{var}(\hat{S}(t)) = \hat{S}(t)^2 \sum_{j=1}^i \frac{d_j}{n_j(n_j - d_j)}.$$

### 4. Parameters

**n**

Input: the number of failure and censored times given in **t**.

Constraint: **n**  $\geq$  2.

**t[n]**

Input: the failure and censored times; these need not be ordered.

**ic[n]**

Input: **ic**[ $i - 1$ ] contains the censoring code of the  $i$ th observation, for  $i = 1, 2, \dots, \mathbf{n}$ .

If **ic**[ $i - 1$ ] = 0 the  $i$ th observation is a failure time.

If **ic**[ $i - 1$ ] = 1 the  $i$ th observation is right-censored.

Constraint: **ic**[ $i - 1$ ] = 0 or 1 for  $i = 1, 2, \dots, \mathbf{n}$ .

**freq[n]**

Input: indicates whether frequencies are provided for each failure and censored time point. If frequencies are provided then **freq** must be dimensioned at least **n**. If the failure and censored times are to be considered as single observations, i.e., a frequency of 1 is to be assumed then **freq** must be set to NULL. Constraints: Either **freq** = (Integer \*)0 or **freq**[ $i - 1$ ]  $\geq 0$ , for  $i = 1, 2, \dots, \mathbf{n}$ .

**nd**

Output: the number of distinct failure times,  $n_d$ .

**tp[n]**

Output: **tp**[ $i - 1$ ] contains the  $i$ th ordered distinct failure time,  $t_i$ , for  $i = 1, 2, \dots, n_d$ .

**p[n]**

Output: **p**[ $i - 1$ ] contains the Kaplan-Meier estimate of the survival probability,  $\hat{S}(t)$ , for time **tp**[ $i - 1$ ], for  $i = 1, 2, \dots, n_d$ .

**psig[n]**

Output: **psig**[ $i - 1$ ] contains an estimate of the standard deviation of **p**[ $i - 1$ ], for  $i = 1, 2, \dots, n_d$ .

**fail**

The NAG error parameter, see the Essential Introduction to the NAG C Library.

**5. Error Indications and Warnings****NE\_INT\_ARG\_LT**

On entry, **n** must not be less than 2: **n**= $\langle \text{value} \rangle$ .

**NE\_INVALID\_CENSOR\_CODE**

On entry, **ic**[ $\langle \text{value} \rangle$ ]= $\langle \text{value} \rangle$ . The censor code for an observation must be either 0 or 1.

**NE\_INVALID\_FREQ**

On entry, **freq**[ $\langle \text{value} \rangle$ ]= $\langle \text{value} \rangle$ . The value of frequency for an observation must be  $\geq 0$ .

**NE\_ALLOC\_FAIL**

Memory allocation failed.

**NE\_INTERNAL\_ERROR**

An internal error has occurred in this function. Check the function call and any array sizes. If the call is correct then please consult NAG for assistance.

**6. Further Comments**

If there are no censored observations,  $\hat{S}(t)$ , reduces to the ordinary binomial estimate of the probability of survival at time  $t$ .

**6.1. Accuracy**

The computations are believed to be stable.

**6.2. References**

Gross A J and Clark V A (1975) *Survival Distributions: Reliability Applications in the Biomedical Sciences*. Wiley.

Kalbfleisch J D and Prentice R L (1980) *The Statistical Analysis of Failure Time Data*. Wiley.

**7. See Also**

None

**8. Example**

The remission times for a set of 21 leukemia patients at 18 distinct time points are read in and the Kaplan-Meier estimate computed and printed. For further details see Gross and Clark (1975), page 242.

## 8.1. Program Text

```

/* nag_prod_limit_surviv_fn(g12aac) Example Program.
 *
 * Copyright 1996 Numerical Algorithms Group.
 *
 * Mark 4, 1996.
 *
 */

#include <nag.h>
#include <stdio.h>
#include <nag_stdlib.h>
#include <nagg12.h>

#define NMAX 18

main()
{
    double psig[NMAX];
    double p[NMAX];
    double t[NMAX];
    double tp[NMAX];

    Integer i, n;
    Integer ifreq[NMAX], ic[NMAX], nd;

    Vprintf("g12aac Example Program Results\n");

    /*      Skip heading in data file */
    Vscanf("%*[^\\n] ");

    Vscanf("%ld ", &n);
    if (n <= NMAX)
    {
        for (i = 0; i < n; ++i)
            Vscanf("%lf %ld %ld ", &t[i], &ic[i], &ifreq[i]);

        g12aac(n, t, ic, ifreq, &nd, tp, p, psig, NAGERR_DEFAULT);

        Vprintf("\n   Time   Survival   Standard\n");
        Vprintf("           probability deviation\n\n");
        for (i = 0; i < nd; ++i)
            Vprintf(" %6.1f%10.3f %10.3f\n", tp[i], p[i], psig[i]);
    }
    exit(EXIT_SUCCESS);
}

```

## 8.2. Program Data

```

g12aac Example Program Data
18
6.0 1 1 6.0 0 3 7.0 0 1 9.0 1 1 10.0 0 1 10.0 1 1
11.0 1 1 13.0 0 1 16.0 0 1 17.0 1 1 19.0 1 1 20.0 1 1
22.0 0 1 23.0 0 1 25.0 1 1 32.0 1 2 34.0 1 1 35.0 1 1

```

**8.3. Program Results**

g12aac Example Program Results

Time	Survival probability	Standard deviation
6.0	0.857	0.076
7.0	0.807	0.087
10.0	0.753	0.096
13.0	0.690	0.107
16.0	0.627	0.114
22.0	0.538	0.128
23.0	0.448	0.135

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