

# NAG Library Function Document

## nag\_mv\_gaussian\_mixture (g03gac)

### 1 Purpose

nag\_mv\_gaussian\_mixture (g03gac) performs a mixture of Normals (Gaussians) for a given (co) variance structure.

### 2 Specification

```
#include <nag.h>
#include <nagg03.h>
void nag_mv_gaussian_mixture (Integer n, Integer m, const double x[],
    Integer pdx, const Integer isx[], Integer nvar, Integer ng,
    Nag_Boolean popt, double prob[], Integer tdprob, Integer *niter,
    Integer riter, double w[], double g[], Nag_VarCovar sopt, double s[],
    double f[], double tol, double *loglik, NagError *fail)
```

### 3 Description

A Normal (Gaussian) mixture model is a weighted sum of  $k$  group Normal densities given by,

$$p(x | w, \mu, \Sigma) = \sum_{j=1}^k w_j g(x | \mu_j, \Sigma_j), \quad x \in \mathbb{R}^p$$

where:

$x$  is a  $p$ -dimensional object of interest;

$w_j$  is the mixture weight for the  $j$ th group and  $\sum_{j=1}^k w_j = 1$ ;

$\mu_j$  is a  $p$ -dimensional vector of means for the  $j$ th group;

$\Sigma_j$  is the covariance structure for the  $j$ th group;

$g(\cdot)$  is the  $p$ -variate Normal density:

$$g(x | \mu_j, \Sigma_j) = \frac{1}{(2\pi)^{p/2} |\Sigma_j|^{1/2}} \exp \left[ -\frac{1}{2} (x - \mu_j) \Sigma_j^{-1} (x - \mu_j)^T \right].$$

Optionally, the (co)variance structure may be pooled (common to all groups) or calculated for each group, and may be full or diagonal.

### 4 References

Hartigan J A (1975) *Clustering Algorithms* Wiley

### 5 Arguments

1: **n** – Integer *Input*

*On entry:*  $n$ , the number of objects. There must be more objects than parameters in the model.

*Constraints:*

if **sopt** = Nag\_GroupCovar, **n** > **ng** × (**nvar** × **nvar** + **nvar**);  
 if **sopt** = Nag\_PooledCovar, **n** > **nvar** × (**ng** + **nvar**);

```

if supt = Nag_GroupVar, n > 2 × ng × nvar;
if supt = Nag_PooledVar, n > nvar × (ng + 1);
if supt = Nag_OverallVar, n > nvar × ng + 1.
```

2: **m** – Integer *Input*

*On entry:* the total number of variables in array **x**.

*Constraint:* **m** ≥ 1.

3: **x[n × pdx]** – const double *Input*

*On entry:* **x**[(*i* − 1) × **pdx** + *j* − 1] must contain the value of the *j*th variable for the *i*th object, for *i* = 1, 2, …, **n** and *j* = 1, 2, …, **m**.

4: **pdx** – Integer *Input*

*On entry:* the stride separating matrix column elements in the array **x**.

*Constraint:* **pdx** ≥ **m**.

5: **isx[m]** – const Integer *Input*

*On entry:* if **nvar** = **m** all available variables are included in the model and **isx** is not referenced; otherwise the *j*th variable will be included in the analysis if **isx**[*j* − 1] = 1 and excluded if **isx**[*j* − 1] = 0, for *j* = 1, 2, …, **m**.

*Constraint:* if **nvar** ≠ **m**, **isx**[*j* − 1] = 1 for **nvar** values of *j* and **isx**[*j* − 1] = 0 for the remaining **m** − **nvar** values of *j*, for *j* = 1, 2, …, **m**.

6: **nvar** – Integer *Input*

*On entry:* *p*, the number of variables included in the calculations.

*Constraint:* 1 ≤ **nvar** ≤ **m**.

7: **ng** – Integer *Input*

*On entry:* *k*, the number of groups in the mixture model.

*Constraint:* **ng** ≥ 1.

8: **popt** – Nag\_Boolean *Input*

*On entry:* if **popt** = Nag\_TRUE, the initial membership probabilities in **prob** are set internally; otherwise these probabilities must be supplied.

9: **prob[n × tdpob]** – double *Input/Output*

*On entry:* if **popt** ≠ Nag\_TRUE, **prob**[(*i* − 1) × **tdprob** + *j* − 1] is the probability that the *i*th object belongs to the *j*th group. (These probabilities are normalised internally.)

*On exit:* **prob**[(*i* − 1) × **tdprob** + *j* − 1] is the probability of membership of the *i*th object to the *j*th group for the fitted model.

10: **tdprob** – Integer *Input*

*On entry:* the stride separating matrix column elements in the array **prob**.

*Constraint:* **tdprob** ≥ **ng**.

11: **niter** – Integer \* *Input/Output*

*On entry:* the maximum number of iterations.

*Suggested value:* 15

*On exit:* the number of completed iterations.

*Constraint:* **niter**  $\geq 1$ .

12: **riter** – Integer *Input*

*On entry:* if **riter**  $> 0$ , membership probabilities are rounded to 0.0 or 1.0 after the completion of every **riter** iterations.

*Suggested value:* 5

13: **w[ng]** – double *Output*

*On exit:*  $w_j$ , the mixing probability for the  $j$ th group.

14: **g[nvar × ng]** – double *Output*

*On exit:*  $g[(i - 1) \times \text{ng} + j - 1]$  gives the estimated mean of the  $i$ th variable in the  $j$ th group.

15: **sopt** – Nag\_VarCovar *Input*

*On entry:* determines the (co)variance structure:

**sopt** = Nag\_GroupCovar  
Groupwise covariance matrices.

**sopt** = Nag\_PooledCovar  
Pooled covariance matrix.

**sopt** = Nag\_GroupVar  
Groupwise variances.

**sopt** = Nag\_PooledVar  
Pooled variances.

**sopt** = Nag\_OverallVar  
Overall variance.

*Constraint:* **sopt** = Nag\_GroupCovar, Nag\_PooledCovar, Nag\_GroupVar, Nag\_PooledVar or Nag\_OverallVar.

16: **s[dim]** – double *Output*

**Note:** the dimension,  $dim$ , of the array **s** must be at least  $a \times b \times c$ .

Where  $S(i, j, k)$  appears in this document, it refers to the array element  $s[(k - 1) \times a \times b + (j - 1) \times a + i - 1]$ .

*On exit:* if **sopt** = Nag\_GroupCovar,  $S(i, j, k)$  gives the  $(i, j)$ th element of the  $k$ th group, with  $a = b = \text{nvar}$  and  $c = \text{ng}$ .

If **sopt** = Nag\_PooledCovar,  $S(i, j, 1)$  gives the  $(i, j)$ th element of the pooled covariance, with  $a = b = \text{nvar}$  and  $c = 1$ .

If **sopt** = Nag\_GroupVar,  $S(j, k, 1)$  gives the  $j$ th variance in the  $k$ th group, with  $a = \text{nvar}$ ,  $b = \text{ng}$  and  $c = 1$ .

If **sopt** = Nag\_PooledVar,  $S(j, 1, 1)$  gives the  $j$ th pooled variance., with  $a = \text{nvar}$  and  $b = c = 1$

If **sopt** = Nag\_OverallVar,  $S(1, 1, 1)$  gives the overall variance, with  $a = b = c = 1$ .

17: **f[n × ng]** – double *Output*

*On exit:*  $f[(i - 1) \times \text{ng} + j - 1]$  gives the  $p$ -variate Normal (Gaussian) density of the  $i$ th object in the  $j$ th group.

18:	<b>tol</b> – double	<i>Input</i>
<i>On entry:</i> iterations cease the first time an improvement in log-likelihood is less than <b>tol</b> . If <b>tol</b> $\leq 0$ a value of $10^{-3}$ is used.		
19:	<b>loglik</b> – double *	<i>Output</i>
<i>On exit:</i> the log-likelihood for the fitted mixture model.		
20:	<b>fail</b> – NagError *	<i>Input/Output</i>
The NAG error argument (see Section 2.7 in How to Use the NAG Library and its Documentation).		

## 6 Error Indicators and Warnings

### NE\_ALLOC\_FAIL

Dynamic memory allocation failed.

See Section 2.3.1.2 in How to Use the NAG Library and its Documentation for further information.

### NE\_ARRAY\_SIZE

On entry, **pdx** =  $\langle\text{value}\rangle$  and **n** =  $\langle\text{value}\rangle$ .

Constraint: **pdx**  $\geq \mathbf{n}$ .

On entry, **tdprob** =  $\langle\text{value}\rangle$  and **n** =  $\langle\text{value}\rangle$ .

Constraint: **tdprob**  $\geq \mathbf{n}$ .

### NE\_BAD\_PARAM

On entry, argument  $\langle\text{value}\rangle$  had an illegal value.

### NE\_CLUSTER\_EMPTY

An iteration cannot continue due to an empty group, try a different initial allocation.

### NE\_INT

On entry, **m** =  $\langle\text{value}\rangle$ .

Constraint: **m**  $\geq 1$ .

On entry, **ng** =  $\langle\text{value}\rangle$ .

Constraint: **ng**  $\geq 1$ .

On entry, **niter** =  $\langle\text{value}\rangle$ .

Constraint: **niter**  $\geq 1$ .

### NE\_INT\_2

On entry, **nvar** =  $\langle\text{value}\rangle$  and **m** =  $\langle\text{value}\rangle$ .

Constraint:  $1 \leq \mathbf{nvar} \leq \mathbf{m}$ .

### 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 contact NAG for assistance.

An unexpected error has been triggered by this function. Please contact NAG.

See Section 2.7.6 in How to Use the NAG Library and its Documentation for further information.

### NE\_MAT\_NOT\_POS\_DEF

A covariance matrix is not positive definite, try a different initial allocation.

**NE\_NO\_LICENCE**

Your licence key may have expired or may not have been installed correctly.

See Section 2.7.5 in How to Use the NAG Library and its Documentation for further information.

**NE\_OBSERVATIONS**

On entry,  $\mathbf{n} = \langle \text{value} \rangle$  and  $p = \langle \text{value} \rangle$ .

Constraint:  $\mathbf{n} > p$ , the number of parameters, i.e., too few objects have been supplied for the model.

**NE\_PROBABILITY**

On entry, row  $\langle \text{value} \rangle$  of supplied **prob** does not sum to 1.

**NE\_VAR\_INCL\_INDICATED**

On entry,  $\mathbf{nvar} \neq \mathbf{m}$  and **isx** is invalid.

**7 Accuracy**

Not applicable.

**8 Parallelism and Performance**

`nag_mv_gaussian_mixture` (g03gac) is threaded by NAG for parallel execution in multithreaded implementations of the NAG Library.

`nag_mv_gaussian_mixture` (g03gac) makes calls to BLAS and/or LAPACK routines, which may be threaded within the vendor library used by this implementation. Consult the documentation for the vendor library for further information.

Please consult the x06 Chapter Introduction for information on how to control and interrogate the OpenMP environment used within this function. Please also consult the Users' Note for your implementation for any additional implementation-specific information.

**9 Further Comments**

None.

**10 Example**

This example fits a Gaussian mixture model with pooled covariance structure to New Haven schools test data, see Table 5.1 (p. 118) in Hartigan (1975).

**10.1 Program Text**

```
/* nag_mv_gaussian_mixture (g03gac) Example Program.
*
* NAGPRODCODE Version.
*
* Copyright 2016 Numerical Algorithms Group.
*
* Mark 26, 2016.
*/
#include <stdio.h>
#include <string.h>
#include <math.h>
#include <nag.h>
#include <nag_stdlib.h>
#include <nagg03.h>
#include <nagx04.h>

#define S(I,J,K) s[I-1 + (J-1)*(sopt==Nag_GroupVar ?ng:nvar) + (K-1)*nvar*nvar]
```

```

#define X(I,J) x[(I-1)*tdx + J-1]
#define PROB(I,J) prob[(I-1)*tdprob + J-1]

int main(void)
{
    /* Integer scalar and array declarations */
    Integer exit_status = 0, i, j, lens, m, n, ng, niter, nvar, riter,
           tdpred, tdx;
    Integer *isx = 0;

    /* Double scalar and array declarations */
    double loglik, tol;
    double *f = 0, *g = 0, *prob = 0, *s = 0, *w = 0, *x = 0;

    /* NAG structures */
    Nag_Boolean popt;
    Nag_VarCovar sopt;
    NagError fail;

    /* Character scalar and array declarations */
    char nag_enum_popt[30 + 1], nag_enum_sopt[30 + 1];

    printf("nag_mv_gaussian_mixture (g03gac) Example Program Results\n\n");
    fflush(stdout);

    /* Skip heading in data file */
#ifndef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

    /* Problem size */
#ifndef _WIN32
    scanf_s("%" NAG_IFMT "", &n);
#else
    scanf("%" NAG_IFMT "", &n);
#endif
#ifndef _WIN32
    scanf_s("%" NAG_IFMT "", &m);
#else
    scanf("%" NAG_IFMT "", &m);
#endif
#ifndef _WIN32
    scanf_s("%" NAG_IFMT "", &nvar);
#else
    scanf("%" NAG_IFMT "", &nvar);
#endif
#ifndef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

    /* Number of groups */
#ifndef _WIN32
    scanf_s("%" NAG_IFMT "", &ng);
#else
    scanf("%" NAG_IFMT "", &ng);
#endif
#ifndef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

    /* Scaling option */
#ifndef _WIN32
    scanf_s("%30s", nag_enum_sopt, (unsigned)_countof(nag_enum_sopt));
#else
    scanf("%30s", nag_enum_sopt);

```

```

#endif
#ifndef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

/* Initial probabilities option */
#ifndef _WIN32
    scanf_s("%30s", nag_enum_popt, (unsigned)_countof(nag_enum_popt));
#else
    scanf("%30s", nag_enum_popt);
#endif
#ifndef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

/* Maximum number of iterations */
#ifndef _WIN32
    scanf_s("%" NAG_IFMT "", &niter);
#else
    scanf("%" NAG_IFMT "", &niter);
#endif
#ifndef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

/* Principal dimensions */
tdx = nvar;
tdprob = ng;

/* nag_enum_name_to_value (x04nac).
 * Converts NAG enum member name to value
 */
popt = (Nag_Boolean) nag_enum_name_to_value(nag_enum_popt);
sopt = (Nag_VarCovar) nag_enum_name_to_value(nag_enum_sopt);

/* Variance/covariance array */
switch (sopt) {
case Nag_GroupCovar:
    lens = nvar * nvar * ng;
    break;
case Nag_PooledCovar:
    lens = nvar * nvar;
    break;
case Nag_GroupVar:
    lens = nvar * ng;
    break;
case Nag_PooledVar:
    lens = nvar;
    break;
case Nag_OverallVar:
    lens = 1;
    break;
}

if (!(x = NAG_ALLOC(n * tdx, double)) ||
    !(prob = NAG_ALLOC(n * tdprob, double)) ||
    !(g = NAG_ALLOC(ng * nvar, double)) ||
    !(w = NAG_ALLOC(ng, double)) ||
    !(isx = NAG_ALLOC(m, Integer)) ||
    !(f = NAG_ALLOC(ng * n, double)) || !(s = NAG_ALLOC(lens, double)))
{
    printf("Allocation failure\n");
    exit_status = -1;
    goto END;
}

```

```

/* Data matrix X */
for (i = 1; i <= n; i++)
    for (j = 1; j <= m; j++)
#ifdef _WIN32
    scanf_s("%lf", &X(i, j));
#else
    scanf("%lf", &X(i, j));
#endif
#ifdef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif

/* Included variables */
if (nvar != m) {
    for (j = 1; j <= m; j++)
#ifdef _WIN32
    scanf_s("%" NAG_IFMT "", &isx[j - 1]);
#else
    scanf("%" NAG_IFMT "", &isx[j - 1]);
#endif
#ifdef _WIN32
    scanf_s("%*[^\n] ");
#else
    scanf("%*[^\n] ");
#endif
}
}

/* Optionally read initial probabilities of group membership */
if (popt == Nag_FALSE) {
    for (i = 1; i <= n; i++)
        for (j = 1; j <= ng; j++)
#ifdef _WIN32
        scanf_s("%lf", &PROB(i, j));
#else
        scanf("%lf", &PROB(i, j));
#endif
#ifdef _WIN32
        scanf_s("%*[^\n] ");
#else
        scanf("%*[^\n] ");
#endif
}
}

/* Optimization parameters */
tol = 0.0;
riter = 5;

/* Fit the model */
/* nag_mv_gaussian_mixture (g03gac).
 * Computes a Gaussian mixture model
 */
INIT_FAIL(fail);
nag_mv_gaussian_mixture(n, m, x, tdx, isx, nvar, ng, popt, prob, tdprob,
    &niter, riter, w, g, sopt, s, f, tol, &loglik,
    &fail);

if (fail.code != NE_NOERROR) {
    printf("nag_mv_gaussian_mixture (g03gac) failed.\n%s\n", fail.message);
    exit_status = 1;
    goto END;
}

/* Results */
/* nag_gen_real_mat_print (x04cac).
 * Print real general matrix (easy-to-use)
 */
nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag,
    1, ng, w, ng, "Mixing proportions", NULL, &fail);

```

```

nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag,
                        nvar, ng, g, ng, "\n Group means", NULL, &fail);

/* Variance/Covariance */
switch (sopt) {
    case Nag_GroupCovar:
        for (i = 1; i <= ng; i++) {
            nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix,
                                   Nag_NonUnitDiag, nvar, nvar, &S(1, 1, i), nvar,
                                   "\n Variance-covariance matrix", NULL, &fail);
        }
        break;
    case Nag_PooledCovar:
        nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag,
                               nvar, nvar, s, nvar,
                               "\n Pooled Variance-covariance matrix", NULL,
                               &fail);
        break;
    case Nag_GroupVar:
        nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag,
                               nvar, ng, s, ng, "\n Groupwise Variance", NULL,
                               &fail);
        break;
    case Nag_PooledVar:
        nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag,
                               nvar, 1, s, 1, "\n Pooled Variance", NULL, &fail);
        break;
    case Nag_OverallVar:
        printf("\n Overall Variance = %g\n", S(1, 1, 1));
        break;
}
nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag, n,
                        ng, f, ng, "\n Densities", NULL, &fail);

nag_gen_real_mat_print(Nag_RowMajor, Nag_GeneralMatrix, Nag_NonUnitDiag, n,
                        ng, prob, ng, "\n Membership probabilities", NULL,
                        &fail);

printf("\nNo. iterations: %" NAG_IFMT "\n", niter);
printf("Log-likelihood: %g\n\n", loglik);

END:
NAG_FREE(f);
NAG_FREE(g);
NAG_FREE(prob);
NAG_FREE(s);
NAG_FREE(w);
NAG_FREE(x);
NAG_FREE(isx);

return exit_status;
}

```

## 10.2 Program Data

```

nag_mv_gaussian_mixture (g03gac) Example Program Data
25 4 4      : n m ip
2          : ng
Nag_PooledCovar : sopt
Nag_FALSE       : popt
15          : niter
2.7 3.2 4.5 4.8
3.9 3.8 5.9 6.2
4.8 4.1 6.8 5.5
3.1 3.5 4.3 4.6
3.4 3.7 5.1 5.6
3.1 3.4 4.1 4.7
4.6 4.4 6.6 6.1

```

### 10.3 Program Results

## naq mv gaussian mixture (q03qac) Example Program Results

Mixing proportions		
	1	2
1	0.4798	0.5202

  

Group means		
	1	2
1	4.0041	3.3350
2	3.9949	3.4434
3	5.5894	4.9870
4	5.4432	5.3602

Pooled Variance-covariance matrix				
	1	2	3	4
1	0.4539	0.2891	0.6075	0.3413
2	0.2891	0.2048	0.4101	0.2490
3	0.6075	0.4101	1.0648	0.6011
4	0.3413	0.2490	0.6011	0.3759

```
Densities
      1          2
 1  2.5836e-01  1.1853e-02
 2  3.7065e-07  1.1241e-01
```

3	5.3069e-03	1.8080e-06
4	4.2461e-01	2.8584e-05
5	5.0387e-02	1.1544e+00
6	1.1260e+00	7.2224e-02
7	2.0911e+00	2.1224e-02
8	5.7856e-03	1.3227e+00
9	1.1609e+00	2.9411e-02
10	8.9826e-02	2.4260e-05
11	3.0170e-01	1.0106e+00
12	1.2930e+00	3.5422e-01
13	2.8644e-02	6.7851e-07
14	2.0759e-02	3.1690e+00
15	7.6461e-02	1.5231e+00
16	3.0279e-04	8.4017e-01
17	5.6101e-01	4.6699e-05
18	2.6573e-05	6.4442e-01
19	2.1250e+00	5.1006e-02
20	8.6822e-04	2.7626e+00
21	1.9223e-01	2.3971e+00
22	1.2469e-02	2.8179e+00
23	1.8389e-02	5.3572e-01
24	1.2409e+00	9.6489e-03
25	2.1037e-05	4.8674e-02

## Membership probabilities

	1	2
1	9.5018e-01	4.9823e-02
2	3.3259e-06	1.0000e+00
3	9.9961e-01	3.8664e-04
4	9.9992e-01	7.9913e-05
5	3.8999e-02	9.6100e-01
6	9.3270e-01	6.7295e-02
7	9.8881e-01	1.1190e-02
8	4.1252e-03	9.9587e-01
9	9.7252e-01	2.7479e-02
10	9.9969e-01	3.0805e-04
11	2.1722e-01	7.8278e-01
12	7.6938e-01	2.3062e-01
13	9.9997e-01	2.6937e-05
14	6.1133e-03	9.9389e-01
15	4.4189e-02	9.5581e-01
16	3.5006e-04	9.9965e-01
17	9.9990e-01	9.7029e-05
18	4.0270e-05	9.9996e-01
19	9.7380e-01	2.6202e-02
20	3.0204e-04	9.9970e-01
21	6.9471e-02	9.3053e-01
22	4.1603e-03	9.9584e-01
23	3.0839e-02	9.6916e-01
24	9.9116e-01	8.8421e-03
25	4.1534e-04	9.9958e-01

No. iterations: 14

Log-likelihood: -29.6831