

Orthogonal polynomials play an important role in numerical analysis and applied mathematics. They satisfy orthogonality relations²³ of the form

$$\int_A^B dx w(x) p_n(x) p_m(x) = \delta_{nm} \equiv \begin{cases} 1, & m=n \\ 0, & m \neq n \end{cases} \quad (40)$$

where the weight function $w(x)$ is positive.

For a given $w(x)$ and interval $[A,B]$, we can construct orthogonal polynomials using the Gram-Schmidt orthogonalization process

Denote the integral in Eq. 40 by (p_n, p_m) to save having to write it many times. We start with

$$p_{-1} = 0, \\ p_0(x) = \left(\int_A^B dx w(x) \right)^{-1/2} = \text{const.},$$

and assume the polynomials satisfy the 2-term upward recursion relation

$$p_{n+1}(x) = (a_n + xb_n) p_n(x) + c_n p_{n-1}(x) \quad (41)$$

Now apply Eq. 41: assume we have calculated p_n and p_{n-1} and want to calculate p_{n+1} . Clearly, the orthogonality property gives

$$(p_{n+1}, p_n) = (p_{n+1}, p_{n-1}) = (p_n, p_{n-1}) = 0,$$

and the assumed normalization gives

$$(p_n, p_n) = 1.$$

These relations yields two equations for the three unknowns, a_n , b_n and c_n :

23. Polynomials can be thought of as vectors in a space of infinitely many dimensions ("Hilbert" space). Certain polynomials are like the vectors that point in the (mutually orthogonal) directions in ordinary 3-dimensional space, and so are called orthogonal by analogy.

$$a_n + b_n (p_n, x p_n) = 0$$

$$c_n + b_n (p_n, x p_{n-1}) = 0$$

We express a_n and c_n in terms of b_n to get

$$\begin{aligned} p_{n+1}(x) = b_n \Big[& (x - (p_n, x p_n)) p_n(x) \\ & - (p_n, x p_{n-1}) p_{n-1}(x) \Big] \end{aligned} \quad (42)$$

We determine the remaining parameter b_n by again using the normalization condition:

$$(p_{n+1}, p_{n+1}) = 1.$$

In practice, we pretend $b_n = 1$ and evaluate Eq. 42; then we calculate

$$b_n = (\bar{p}_{n+1}, \bar{p}_{n+1})^{-1/2}, \quad (43)$$

multiply the (un-normalized) \bar{p}_{n+1} by b_n , and continue.

The process of successive orthogonalization guarantees that p_n is orthogonal to all polynomials of lesser degree in the set. Why is this so? By construction, $p_{n+1} \perp p_n$ and $p_{n+1} \perp p_{n-1}$. Is it $\perp p_{n-2}$? We need to ask whether

$$(p_n, (x - \alpha_n) p_{n-2}) = 0.$$

But we know that any polynomial of degree $N-1$ can be expressed as a linear combination of independent polynomials of degrees $0, 1, \dots, N-1$. Thus

$$(x - \alpha_n) p_{n-2} \equiv \sum_{k=0}^{n-1} \mu_k p_k(x) \quad (44)$$

and (by hypothesis) $p_n \perp$ every term of the rhs of Eq. 44, hence it follows (by mathematical induction) that

$$p_{n+1} \perp \{p_{n-2}, p_{n-3}, \dots\}.$$

Let us illustrate the process for Legendre polynomials, defined by weight $w(x) = 1$, interval $[-1,1]$:

$$p_0 = \left(\frac{1}{2}\right)^{1/2},$$

$$p_1 = \left(\frac{3}{2}\right)^{1/2} x,$$

$$p_2 = \left(\frac{5}{2}\right)^{1/2} \left(\frac{3}{2}x^2 - \frac{1}{2}\right),$$

.....

These are in fact the first three (normalized) Legendre polynomials, as any standard reference will confirm.

Now we can discuss Gram polynomials. While orthogonal polynomials are usually defined with respect to an **integral** as in Eq. 40, we might also define orthogonality in terms of a **sum**, as in Eq. 39a. That is, suppose we define the polynomials such that

$$\sum_{k=0}^{M-1} p_n(x_k) p_m(x_k) \frac{1}{\sigma_k^2} \equiv \delta_{nm} = \begin{cases} 1, & m=n \\ 0, & m \neq n \end{cases} \quad (45)$$

Then we can construct the Gram polynomials, calculating the coefficients by the algebraic steps of the Gram-Schmidt process, except now we evaluate sums rather than integrals. Since $p_n(x)$ satisfies 45 by construction, the coefficients γ_n in our fitting polynomial are simply

$$\gamma_n = \sum_{k=0}^{M-1} p_n(x_k) f_k \frac{1}{\sigma_k^2}; \quad (46)$$

they can be evaluated without solving any coupled linear equations, ill-conditioned or otherwise. Roundoff error thus becomes irrelevant.

The algorithm for fitting data with Gram polynomials may be expressed in flow-diagram form:

```

Read in points  $f_k, x_k$  and  $w_k \equiv 1/\sigma_k^2$ .

DO n = 1 to N-1      (outer loop)
  Construct  $a_n, c_n$  :

    DO k = 0 to M-1 (inner loop)
       $p_{n+1}(x_k) = (x_k - a_n)p_n(x_k) - c_n p_{n-1}(x_k)$ 

       $sum = sum + (p_{n+1}(x_k))^2 w_k$ 

       $c_{n+1} = c_{n+1} + f_k p_{n+1}(x_k) w_k$ 
    LOOP              (end inner loop)
     $c_{n+1} = c_{n+1} / sum$ 

  )

  DO k = 0 to M-1    (normalize)
     $p_{n+1}(x_k) = p_{n+1}(x_k) / \sqrt{sum}$ 
  LOOP

LOOP

```

Fig. 8-5 Construction of Gram polynomials

The required storage is 5 vectors of length M to hold $x_k, p_n(x_k), p_{n-1}(x_k), f_k$ and $w_k \equiv 1/\sigma_k^2$. We also need to store the coefficients a_n, c_n and the normalizations b_n —that is, 3 vectors of length $N \ll M$ — in case they should be needed to interpolate. The time involved is approximately $7M$ multiplications and additions for each n , giving $7NM$. Since N can be no greater than $M-1$ (M data determine at most a polynomial of degree $M-1$), the maximum possible running time is $7M^2$, which is much less than the time to solve M linear equations.

In practice, we would never wish to fit a polynomial of order comparable to the number of data, since this would include the noise as well as the significant information.

We therefore calculate a statistic called $\chi^2/(\text{degree of freedom})^{24}$. With M data points and an N 'th order polynomial, there are $M-N-1$ degrees of freedom. That is, we evaluate Eq. 38 for fixed N , and divide by $M-N-1$. We then increase N by 1 and do it again. The value of N to stop at is the one where

$$\sigma_{M,N}^2 = \frac{\chi_{M,N}^2}{M-N-1}$$

stops decreasing (with N) and begins to increase.

The best thing about the $\chi_{M,N}^2$ statistic is we can increase N without having to do any extra work:

$$\begin{aligned} \chi_{M,N}^2 &= \sum_{k=0}^{M-1} \left(f_k - \sum_n \gamma_n p_n(x_k) \right)^2 w_k \\ &\equiv \sum_{k=0}^{M-1} (f_k)^2 - \sum_{n=0}^N (\gamma_n)^2 \end{aligned} \quad (47)$$

The first term after \equiv in Eq. 47 is independent of N , and the second term is computed as we go. Thus we could turn the outer loop (over N) into a **BEGIN ... WHILE ... REPEAT** loop, in which N is incremented as long as $\sigma_{M,N}^2$ is larger than $\sigma_{M,N+1}^2$. (Incidentally, Eq. 47 guarantees that as we increase N the fitted curve deviates less and less, on the average, from the measured points. When $N = M-1$, in fact, the curve goes through the points. But as explained above, this is a meaningless fit, since all data contain measurement errors. A fitted curve that passes closer than σ_k to more than about $1/3$ of the points is suspect.)

The code for Gram polynomials is relatively easy to write using the techniques developed in Ch. 5. The program is displayed in full in Appendix 8.4.

24. That is, "chi-squared per degree of freedom".

§§3 Simplex algorithm

Sometimes we must fit data by a function that depends on parameters in a nonlinear manner. An example is

$$f_k = \frac{F}{1 + e^{\alpha(x_k - X)}} \quad (48)$$

Although the dependence on the parameter F is linear, that on the parameters α and X is decidedly nonlinear.

One way to handle a problem like fitting Eq. 48 might be to transform the data, to make the dependence on the parameters linear. In some cases this is possible, but in 48 no transformation will render linear the dependence on all three parameters at once.

Thus we are frequently confronted with having to minimize numerically a complicated function of several parameters. Let us denote these by $\theta_0, \theta_1, \dots, \theta_{N-1}$, and denote their possible range of variation by \mathbf{R} . Then we want to find those values of $\{\theta\} \subset \mathbf{R}$ that minimize a positive function:

$$\chi^2(\theta_0, \dots, \theta_{N-1}) = \min_{\{\theta\} \subset \mathbf{R}} \chi^2(\theta_0, \dots, \theta_{N-1}) \quad (49)$$

One way to accomplish the minimization is *via* calculus, using a method known as **steepest descents**. The idea is to differentiate the function χ^2 with respect to each θ_k , and to set the resulting N equations equal to zero, solving for the N θ 's. This is generally a pretty tall order, hence various approximate, iterative techniques have been developed. The simplest just steps along in θ -space, along the direction of the local downhill gradient $-\nabla\chi^2$, until a minimum is found. Then a new gradient is computed, and a new minimum sought²⁵.

Aside from the labor of computing $-\nabla\chi^2$, steepest descents has two main drawbacks: first, it only guarantees to find a minimum, not necessarily the minimum — if a function has several local

25. This is not by itself very useful. Useful modifications can be found in Press, *et al.*, *Numerical Recipes*, *ibid.*, p. 301ff.

minima, steepest descents will not necessarily find the smallest. Worse, consider a function that has a minimum in the form of a steep-sided gulley that winds slowly downhill to a declivity – somewhat like a meandering river's channel. Steepest descents will then spend all its time bouncing up and down the banks of the gulley, rather than proceeding along its bottom, since the steepest gradient is always nearly perpendicular to the line of the channel.

Sometimes the function χ^2 is so complex that its gradient is too expensive to compute. Can we find a minimum *without* evaluating partial derivatives? A standard way to do this is called the **simplex method**. The idea is to construct a **simplex** – a set of $N + 1$ distinct and **non-degenerate** vertices in the N -dimensional θ -space (“non-degenerate” means the geometrical object, formed by connecting the $N + 1$ vertices with straight lines, has non-zero N -dimensional volume; for example, if $N = 2$, the simplex is a triangle.)

We evaluate the function to be minimized at each of the vertices, and sort the table of vertices by the size of χ^2 at each vertex, the best (smallest χ^2) on top, the worst at the bottom. The simplex algorithm then chooses a new point in θ -space by the a strategy, expressed as the flow diagram, Fig. 8-8 on page 203 below, that in action somewhat resembles the behavior of an amoeba seeking its food. The key word **MINIMIZE** that implements the complex decision tree in Fig. 8-8 (given here in pseudocode) is

```

: )MINIMIZE ( n.iter - - 87: rel.error - - )
  INITIALIZE
  BEGIN done? NOT N N.max < AND .
  WHILE
    REFLECT r >= best?
    IF r >= 2worst?
      IF r < worst? IF STORE.X THEN
        HALVE r < worst?
        IF STORE.X ELSE SHRINK THEN
      ELSE STORE.X THEN
    ELSE DOUBLE r >= best?
    IF STORE.XP ELSE STORE.X THEN
  THEN
  N 1+ IS N SORT
  REPEAT ;

```

used in the format

USE(f.name 20 % 1.E-4)MINIMIZE

Fleshing out the details is a — by now — familiar process, so we leave the program *per se* to Appendix 8.5. We also include there a FORTRAN subroutine for the simplex algorithm, taken from

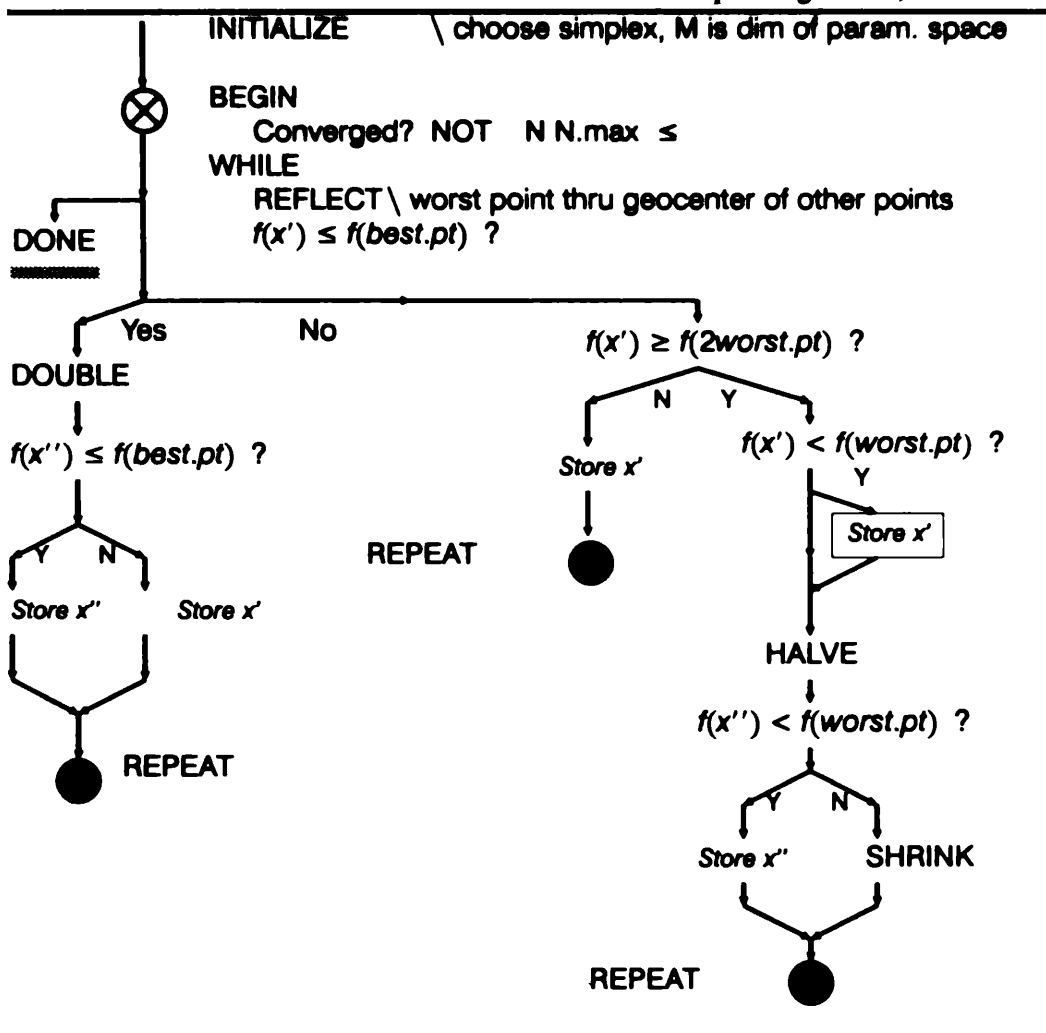


Fig. 8-8 Flow diagram of the simplex algorithm

*Numerical Recipes*²⁶, as an example of just how indecipherable traditional languages can be.

§3 Appendices

§§1 Gaussian quadrature

Gaussian quadrature formulae are based on the following idea: if we let the points ξ_n and weights w_n be $2N$ free parameters (n runs from 1 to N), what values of them most accurately represent an integral by the formula

$$I = \int_A^B dx \sigma(x) f(x) \approx \sum_{n=1}^N w_n f(\xi_n) \quad ? \quad (50)$$

In Eq. 50 $\sigma(x)$ is a (known) positive function and $f(x)$ is the function we want to integrate. This problem can actually be solved, and leads to tables of points ξ_n and weight coefficients w_n specific to a particular interval $[A,B]$ and weight function $\sigma(x)$. **Gauss-Legendre** integration pertains to $[-1, +1]$ and $\sigma(x) = 1$. (Note any interval can be transformed into $[-1, +1]$.)

The interval $[0, \infty)$ and $\sigma(x) = e^{-x}$ leads to **Gauss-Laguerre** formulae, whereas the interval $(-\infty, +\infty)$ and $\sigma(x) = e^{-x^2}$ leads to **Gauss-Hermite** formulae.

Finally, we note that the more common integration formulae such as Simpson's rule or the trapezoidal rule can be derived on the same basis as the Gauss methods, except that the points are specified in advance to be equally spaced and to include the end-points of the interval. Only the weights w_n can be determined as free fitting parameters that give the best approximation to the integral.

For given N Gaussian formulae can be more accurate than equally-spaced rules, as they have twice as many parameters to play with.

26. Press, et al., *Numerical Recipes*, *ibid.*, p. 289ff.

Some FORTH words for 5-point Gauss-Legendre integration:

% 0.906179845938664	FCONSTANT	x2	
% 0.538469310105683	FCONSTANT	x1	: }integral (87: A B -- l)
% 0.568688686868689	FCONSTANT	w0	scale (87: A B -- [A+B]/2 [B-A]/2)
% 0.478628670498366	FCONSTANT	w1	FOVER F(x) w0 F* F-ROT (87: -- l a b)
% 0.2386286865056189	FCONSTANT	w2	XDUP x1 rescale F(x) w1 F* F3R +
			XDUP x1 FNEGATE rescale
: scale (87: A B -- [A+B]/2 [B-A]/2)			F(x) w1 F* F3R +
FOVER F- F2/ FUNDER F+ ;			XDUP x2 rescale F(x) w2 F* F3R +
: rescale (87: a b x -- a+b*x) F* F+ ;			XDUP x2 FNEGATE rescale
: F3R+ (87: a b c x -- a+x b c) F3R F+ F-ROT ;			F(x) w2 F* F3R +

§§2 The trapezoidal rule

Recall (Ch. 8, §1.1) how we approximated the area under a curve by capturing it between rectangles consistently higher- and lower than the curve; and calculating the areas of the two sets of rectangles. In practice we use a better approximation: we average the rectangular upper and lower bounds. The errors tend to cancel, resulting in²⁷

$$\frac{w}{2} \sum_{n=0}^{(B-A)/w} (f(A+nw) + f(A+nw+w)) \quad (51)$$

$$= \int_A^B dx f(x) + \frac{1}{12} \left(\frac{B-A}{w} \right) w^3 \max_{A < x < B} |f''(x)|$$

Now the error is much smaller²⁸ — of order w^2 — so if we double the number of points, we decrease the error four-fold. Yet this so-called **trapezoidal rule**²⁹ requires no more effort (in terms of the number of function evaluations) than the rectangle rule.

27. See, e.g., Abramowitz and Stegun, *HMF*, p. 885.

28. $f''(x)$ is the second derivative of $f(x)$, i.e. the first derivative of $f'(x)$.

29. Called so because the curve $f(x)$ is approximated by straight line segments between successive points $x, x+w$. Thus we evaluate the areas of trapezoids rather than rectangles.

§§3 Richardson extrapolation

In the program)**INTEGRAL** in Ch.8 §1§§5.3 the word **INTERPOLATE** performs Richardson extrapolation for the trapezoidal rule. The idea is this: If we use a given rule, accurate to order w^n , to calculate the integral on an interval $[a,b]$, then presumably the error of using the formula on each half of the interval and adding the results, will be smaller by 2^{-n} . For the trapezoidal rule, $n=2$, hence we expect the error from summing two half-intervals to be $4\times$ smaller than that from the whole interval.

Thus, we can write $(I_0 = \int_a^b, \quad I'_0 = \int_a^{(a+b)/2}, \quad I_1 = \int_{(a+b)/2}^b)$

$$I_0 = I_{\text{exact}} + R \quad (52a)$$

$$I'_0 + I_1 = I_{\text{exact}} + \frac{R}{4} \quad (52b)$$

Equation 52b is only an approximation because the R that appears in it is not exactly the same as R in Eq. 52a. We will pretend the two R 's are equal, however, and eliminate R from the two equations (8.52a,b) ending with an expression for I_{exact} :

$$I_{\text{exact}} \approx \frac{4}{3} (I'_0 + I_1 - I_0) \quad (53)$$

Equation 53 is exactly what appears in **INTERPOLATE**.

§4 Linear least-squares: Gram polynomials

Here is a FORTH program for implementing the algorithm derived in §2§2 above.

ASK GRAM1

CODEed words to simplify fstack management

```
CODE G(N+1) 4 FMUL FCHS. 2 FLD. 6 FSUB.
2 FMUL 1 FADDP.
DX DS MOV. DS POP. R64 DS: [BX] FST.
DS DX MOV. BX POP. END-CODE
seg off -- 87: s a b w x g[n] g[n-1] -- s a b w x g[n] g[n+1]
```

```
CODE B(N+1) 2 FXCH. 2 FMUL 3 FMUL
1 FXCH. 1 FMUL DX DS MOV. DS POP.
R64 DS: [BX] FADD. DS: [BX] FSTP.
DS DX MOV. BX POP. END-CODE
seg off --
87: s a b w x g[n] g[n+1] -- s a b w g[n+1] w x g[n+1]
```

```
CODE A(N+1) 1 FMUL DX DS MOV. DS POP.
R64 DS: [BX] FADD.
DS: [BX] FSTP. DS DX MOV. BX POP. END-CODE
seg off -- 87: s a b w g[n+1] w x g[n+1] -- s a b w g[n+1]
```

```
CODE C(N+1) 1 FXCH. 2 FMUL. 2 FMUL
2 FXCH. 1 FMULP.
DX DS MOV. DS POP.
R64 DS: [BX] FADD. DS: [BX] FSTP.
DS DX MOV. BX POP. 3 FADDP. END-CODE
seg off -- 87: s a b w g[n+1] f -- s = s + w g[n+1]**2 a b)
```

-----Gram polynomial coding

REAL SCALAR DELTA

VAR Nmax

usage: A{ B{ C{ G{ (Nmax) FIT

FIRSTAB's F=0 a{0} G! F=0 b{0} G! ;

```
INIT.DELTA (87:--g{{11}}) F=0 F=0
MODO w{1} G@ y{1} G@ F**2 FOVER F*
(87:ss'--ss'w w**2)
FROT F+ FROT F+ FSWAP
(87:--s=s+w s'=s'+w**2)
LOOP DELTA G! FSQRT 1/F ;
```

```
: FIRST.G's (87: g{{11}}--g{{11}})
MODO F=0 g{{01}} G! FDUP g{{11}} G! LOOP ;
```

```
: SECONDAB's (87: g{{11}}--g{{11}})
F=0 b{10} G! FDUP F**2
F=0 MODO w{10} G@ x{10} G@ F* F+ LOOP
F* a{10} G! ;
```

```
: FIRST.&SECOND.C's (87: g{{11}}-- )
F=0 c{00} G! F=0
MODO w{10} G@ y{10} G@ F* F+ LOOP
F* c{10} G! ;
```

```
: INITIALIZE FINIT IS Nmax IS g{{ IS c{ IS b{ IS a{
FIRSTAB's INIT.DELTA FIRST.G's SECONDAB's
FIRST.&SECOND.C's ;
```

0 VAR N 0 VAR N+1

```
: inc.N N+1 DUP IS N 1+ IS N+1 ;
```

```
: DISPOSE R DDUP R@ G@ R ;
```

```
: inc.OFF #BYTES DROP + ;
```

```
: ZERO.L DDUP F=0 R64!L ;
```

: START.Next.G

```
(--[c(n+1)][a(n+1)][b(n+1)] 87:--s=0 a{n} b{n})
```

```
FINIT F=0 c{N+10} DROP ZERO.L
```

```
a{N0} DISPOSE inc.OFF ZERO.L
```

```
b{N0} DISPOSE inc.OFF ZERO.L ;
```

```
: SET.FSTACK w{r0} G@ x{r0} G@ g{{Nr}} G@
g{{N1-r}} G@ ;
```

```
: )@*! (adr n--87: x--x) FDUP 0) DISPOSE F* G! ;
```

: NORMALIZE (87: sum--)

```
1/F a{N+1}@*! FSQRT
```

```
b{N+1}@*! c{N+1}@*!
```

```
MODO FDUP g{{N+1}} DISPOSE F* G! LOOP
FDROP ;
```

```
CODE 6DUP OPT* 6 PICK 6 PICK 6 PICK
```

```
6 PICK 6 PICK 6 PICK* END-CODE
```

```
CODE 6DROP OPT* DDROP DDROP DDROP* END-CODE
```

```

\ GRAM POLYNOMIAL LEAST-QUARES (CONTD)

Next.G START.Next.G
M 0 DO @DUP SET.FSTACK
  g{{ N+1 }} DROP G(N+1) B(N+1) A(N+1)
  y{10} G@ C(N+1)
  LOOP @DROP FDROP FDROP NORMALIZE ;

: New.DELTA (:: - old.delta new.delta)
  DELTA G@ F DUP c{ N 0 } G@ F**2 F- F DUP
  DELTA G! ;

: NOT.ENUF.G's? New.DELTA M N+1- S-F
  FDUP F=1 F-
  ( CR .FS ." NEXT ITERATION?" ?YN 0= IF ABORT THEN )
  (:: - d' m-n-1 m-n-2)
  FROT F- F-ROT F/ (:: - d'/(m-n-2) d/(m-n-1))
  FOVER F0 IF F ELSE FDROP FDROP 0 THEN ;

: }FIT (X{ Y{ S{ Nmax A{ B{ C{ G{{ - )
  INITIALIZE 1 IS N 2 IS N+1
  BEGIN NOT.ENUF.G's? N Nmax AND
  WHILE Next.G Inc.N
  REPEAT FINIT ;

\----- end of code

: RECONSTRUCT M 0 DO CR x{10} G@ F. y{10} G@ F.
  F=0 N+11+ 1 DO
    c{10} G@ g{{1J}} G@ F* F+
  LOOP F.
  LOOP ;

```

§§5 Non-linear least squares: simplex method

A FORTRAN program for the simplex method is given below on page 209. The FORTH version, as discussed in §§§3 given on pages 210 and 211.

```

SUBROUTINE AMOEBA(P,Y,MP,NP,NDIM,FTOL,FUNK,ITER)
PARAMETER (NMAX = 50,ALPHA = 1.0,
; BETA = 0.5,GAMMA = 2.0,ITMAX = 500)
DIMENSION P(NP,NP),Y(NP),PR(NMAX),PRR(NMAX),PBAR(NMAX)
MPTS = NDIM + 1
ITER = 0
ILO = 1
IF(Y(1).GT.Y(2))THEN
IH = 1
INH = 2
ELSE
IH = 2
INH = 1
ENDIF
DO 11 I = 1,MPTS
IF(Y(I).LT.Y(ILO)) ILO = I
IF(Y(I).GT.Y(IH))THEN
INH = IH
IH = I
ELSE IF(Y(I).GT.Y(INH))THEN
IF(I.NE.IH) INH = I
ENDIF
1 CONTINUE
RTOL = 2.*ABS(Y(IH)-Y(ILO))/(ABS(Y(IH)) + ABS(Y(ILO)))
IF(RTOL.LT.FTOL)RETURN
IF(ITER.EQ.ITMAX) PAUSE 'Amoeba exceeding maximum iterations.'
ITER = ITER + 1
DO 12 J = 1,NDIM
PBAR(J) = 0.
2 CONTINUE
DO 14 I = 1,MPTS
IF(I.NE.IH)THEN
DO 13 J = 1,NDIM
PBAR(J) = PBAR(J) + P(I,J)
3 CONTINUE
ENDIF
4 CONTINUE
DO 15 J = 1,NDIM
PBAR(J) = PBAR(J)/NDIM
PR(J) = (1. + ALPHA)*PBAR(J)-ALPHA*P(IH,J)
5 CONTINUE
YPR = FUNK(PR)
IF(YPR.LE.Y(ILO))THEN
DO 16 J = 1,NDIM
PRR(J) = GAMMA*PR(J) + (1.-GAMMA)*PBAR(J)
6 CONTINUE

```

```

YPRR = FUNK(PRR)
IF (YPRRLT.Y(ILO)) THEN
  DO 17 J = 1,NDIM
    P(IH,J) = PRR(J)
17  CONTINUE
    Y(IH) = YPRR
  ELSE
    DO 18 J = 1,NDIM
      P(IH,J) = PR(J)
18  CONTINUE
      Y(IH) = YPR
    ENDF
  ELSE IF (YPR.GE.Y(IH)) THEN
    IF (YPR.LT.Y(IH)) THEN
      DO 19 J = 1,NDIM
        P(IH,J) = PR(J)
19  CONTINUE
        Y(IH) = YPR
      ENDF
      DO 21 J = 1,NDIM
        PPR(J) = BETA*P(IH,J) + (1.-BETA)*PBAR(J)
21  CONTINUE
        YPRR = FUNK(PPR)
      IF (YPRRLT.Y(IH)) THEN
        DO 22 J = 1,NDIM
          P(IH,J) = PRR(J)
22  CONTINUE
          Y(IH) = YPRR
        ELSE
          DO 24 I = 1,MPTS
            IF (LINELO) THEN
              DO 23 J = 1,NDIM
                PR(J) = 0.5*(P(I,J) + P(ILO,J))
                P(I,J) = PR(J)
23  CONTINUE
                Y(I) = FUNK(PR)
              ENDF
            24  CONTINUE
          ENDF
        ELSE
          DO 25 J = 1,NDIM
            P(IH,J) = PR(J)
25  CONTINUE
            Y(IH) = YPR
          ENDF
        GO TO 1
      END
    END
  END

```

```
\ MINIMIZATION BY THE SIMPLEX METHOD
\ VERSION OF 20:51:19 4/30/1991
```

```
TASK AMOEBA
```

```
\----- FUNCTION NOTATION
```

```
VARIABLE <F>
:USE( [COMPILE] ' CFA <F> !;
:F(X) EXECUTE@;
BEHEAD' <F>
```

```
\----- END FUNCTION NOTATION
```

```
\----- DATA STRUCTURES
```

```
3 VAR Ndim
0 VAR N
0 VAR N.max
```

```
CREATE SIMPLEX({ Ndim 4 (bytes) * Ndim 1+ (# points)
* ALLOT
```

```
CREATE F{ Ndim 1+ 4 (bytes) * ALLOT \ residuals
CREATE index Ndim 1+ 2* ALLOT \ array for scrambled indices
```

```
:>index (I-I') 2*index + @;
```

```
DVARIABLE Residual
```

```
DVARIABLE Residual'
```

```
DVARIABLE Epsilon
```

```
CREATE X{ Ndim 4 (bytes) * ALLOT \ trial point
```

```
CREATE XP{ Ndim 4 (bytes) * ALLOT \ 2nd trial point
```

```
CREATE Y{ Ndim 4 (bytes) * ALLOT \ geocenter
```

```
:} (adr n - adr + 4n) 4* + ; \ part of array notation
:} (adr m n - adr + [m*Ndim + n]*4) SWAP Ndim * + ;
```

```
\----- END DATA STRUCTURES
```

```
\----- ACTION WORDS
```

```
: RESIDUALS
```

```
Ndim 1+ 0 DO SIMPLEX({10}) F(X) F(1) R32!
LOOP;
```

```
:<index> Ndim 1+ 0 DO I index I 2* + I LOOP; \ fill index
```

```
: ORDER <index>
```

```
Ndim 1+ 0 DO F(I >index) R32@
```

```
I 1+ BEGIN Ndim 1+ OVER
```

```
WHILE F( OVER >index) R32@ F OVER F OVER F >
```

```
IF FSWAP
```

```
I >index OVER >index
```

```
index I 2* + I index 3 PICK 2* + I
```

```
THEN FDROP 1+
```

```
REPEAT FDROP DROP
```

```
LOOP;
```

```
CENTER (-) FINIT Ndim S-F (87: - Ndim)
```

```
Ndim 0 DO F=0 \ loop over components
```

```
Ndim 0 DO \ average over vectors
```

```
SIMPLEX({ I index J }) R32@ F+
```

```
LOOP F OVER F/
```

```
Y(1) R32! \ put away
```

```
LOOP FDROP;
```

```
\ note: Worst.Point is SIMPLEX({ Ndim index J })
```

```
\ -- excluded!
```

```
: DONE! CR ." We're finished.";
```

```
: TOO.MANY CR ." Too many iterations.";
```

```
: V.MOVE (src.adr dest.adr -) \ move vector
```

```
Ndim 0 DO OVER I} OVER I} 2 MOVE
```

```
LOOP DDROP;
```

```
: STORE (adr1 adr2 --) 0}
```

```
SIMPLEX({ Ndim index 0 }) V.MOVE
```

```
F{ Ndim index } 2 MOVE;
```

```
: STOREX Residual X{ STORE;
```

```
: STOREXP Residual' XP{ STORE;
```

```
: New.F X{ F(X) Residual R32!;
```

```
: EXTRUDE (87: scale.factor --)
```

```
\ extend pseudopod
```

```
Ndim 0 DO DUP I} R32@ (87: -- s.f x)
```

```
Y(1) R32@ FUNDER F- (87: -- s.f y x-y)
```

```
FROT FUNDER F* (87: -- y s.f [x-y]*s.f)
```

```
FROT F+ (87: -- s.f y + [x-y]*s.f)
```

```
X(1) R32!
```

```
LOOP DROP FDROP New.F;
```

```
: F=1/2 F=1 FNEGATE F=1 FSCALE FPLUCK;
```

```
: F=2 F=1 FDUP FSCALE FPLUCK;
```

```
\----- DEBUGGING CODE
```

```
0 VAR DBG
```

```
: DEBUG-ON -1 IS DBG;
```

```
: DEBUG-OFF 0 IS DBG;
```

```
: V 3 SPACES Ndim 0 DO DUP I} R32@ F.
```

```
LOOP DROP;
```

```
: M (--)
```

```
Ndim 1+ 0 DO DBG CR
```

```
IF I. 2 SPACES THEN
```

```
SIMPLEX({ I index 0 }) .V
```

```
DBG IF F{ I index } R32@ F. THEN
```

```
LOOP CR
```

```
DBG IF CR X{ .V Residual R32@ F. THEN;
```

```
: F Ndim 1+ 0 DO CR F{ I index } R32@ F.
```

```
LOOP;
```

```
\----- END DEBUGGING CODE
```

```
: REFLECT CENTER
```

```
F=1 FNEGATE \ scale.factor = -1
```

```
SIMPLEX({ Ndim index 0 }) \ worst pt.
```

```
EXTRUDE \ calculate x, f(x)
```

```
DBG IF CR ." REFLECTING" THEN .M ;
```

AMOEBA, CONT'D

DOUBLE Residual Residual' 2 MOVE

```

\ save residual
X( XP( V.MOVE \ save point
  FINIT F=2 FNEGATE (?) \ scale factor = -2
  SIMPLEX({ Ndim Index 0 }) \ worst pt.
  EXTRUDE \ calculate x, f(x)
  DBG IF CR." DOUBLING" THEN .M ;

```

```

HALVE FINIT F=1/2 \ scale factor = 0.5
SIMPLEX({ Ndim Index 0 }) \ worst pt.
EXTRUDE \ calculate x, f(x)
DBG IF CR." HALVING" THEN .M ;

```

```

SHRINK SIMPLEX({ 0 > Index 0 }) \ best pt.

```

```

Y( V.MOVE \ save it
  Ndim 1+ 1 DO \ by vector
    Ndim 0 DO \ by component
      SIMPLEX({ J > Index 1 }) DUP
      R32@ Y{1} R32@ FINDER F-
      F=1/2 F* F+ R32!
    LOOP
  LOOP RESIDUALS ORDER
  DBG IF CR." SHRINKING" THEN .M ;

```

```

----- END ACTION WORDS
----- TEST WORDS

```

```

(test) FINIT Residual R32@
F( SWAP >Index } R32@ F> NOT ;
>=best? 0 (test) ;

```

```

<worst? Ndim (test) NOT ;
>=2worst? Ndim 1- (test) ;

```

```

done? F( Ndim Index } R32@
F{ 0 Index } R32@
FOVERFOVER F- F2*
F-ROT F+ F/ FABS Epsilon R32@ F> ;
----- END TEST WORDS

```

```

: MINIMIZE ( n.iter -- 87: error -- )
  IS N.max Epsilon R32! 0 IS N \ initialize
  RESIDUALS \ compute residuals
  ORDER \ locate best, 2worst, worst
  BEGIN \ start iteration
    done? NOT N N.max < AND
  WHILE
    REFLECT r> = best?
    IF r> = 2worst?
      IF r<worst? IF STOREX THEN
        HALVE r<worst?
        IF STOREX ELSE SHRINK THEN
      ELSE STOREX THEN
    ELSE DOUBLE
      r> = best?
      IF STOREXP ELSE STOREX THEN
    THEN
    N 1+ IS N ORDER
  REPEAT DONE! ;

```

----- EXAMPLE FUNCTIONS

```

: F1 ( adr - 87: F ) DUP 0 } R32@ FDUP F**2
  DUP 1 } R32@ F**2 F2* F+
  2 } R32@ F**2 F2* F2* F+
  36 S-F F- F**2 F2/ FSWAP 6 S-F F* F- ;
\ f1 = .5*(x*x + 2*y*y + 4*z*z - 36) ^ 2 - 6*x

```

```

: F2 ( adr - 87: F ) DUP DUP 0 }
  R32@ FDUP F**2 1 } R32@ F**2
  F2* F+ 36 S-F F- F**2 F2/ FSWAP 6 S-F F* F-
  DUP 1 } R32@ 0 } R32@ F/ FATAN F2* FCOS
  F**2 F* ;
\ f2 = [(x^2 2*y^2 - 36)^2 - 6*x] * cos(2*atan(y/x)) ^ 2
\ Usage: USE(MYFUNC 10 1.E-3) MINIMIZE
FLOATS

```

```

5. SIMPLEX({ 0 0 }) R32!      5. SIMPLEX({ 1 0 }) R32!
-3. SIMPLEX({ 0 1 }) R32!     3. SIMPLEX({ 1 1 }) R32!
7. SIMPLEX({ 0 2 }) R32!     -1.5 SIMPLEX({ 1 2 }) R32!

```

```

-10. SIMPLEX({ 2 0 }) R32!    5. SIMPLEX({ 3 0 }) R32!
1. SIMPLEX({ 2 1 }) R32!      3. SIMPLEX({ 3 1 }) R32!
3. SIMPLEX({ 2 2 }) R32!      3. SIMPLEX({ 3 2 }) R32!

```