

A COMPUTER PROGRAM FOR THE DETERMINATION OF KINETIC PARAMETERS FROM SIGMOIDAL STEADY-STATE KINETICS

H.-J. WIEKER, K.-J. JOHANNES and B. HESS

*Max-Planck-Institut für Ernährungsphysiologie,
46 Dortmund, Germany*

Received 23 April 1970

1. Introduction

Current computer programs for the determination of kinetic parameters [1] are either limited only to the treatment of hyperbolic v vs. s curves* [2, 3] or based on specified models [4, 5] which are not generally applicable to non-hyperbolic saturation curves.

In many cases, sigmoidal v vs. s curves are analysed on the basis of models developed for the treatment of allosteric enzymes [6, 7]. However, it should not be overlooked that also models of non-allosteric enzymes could yield sigmoidal saturation curves [5, 8, 9]. Generally, it should be noted that, if the parameter-finding process is model-oriented, the data-processing procedure and the interpretation of the results obtained are mixed, which might lead to erroneous results. In principle, therefore, it is useful to separate data-processing and interpretation.

Separation of the two operations is achieved, if the data-processing is based on a general function $v = f(s)$ with only the minimum number of parameters necessary for the description of the experimental data. We here describe a computer program, which exclusively serves the parameter fitting of sig-

moidal and hyperbolic v vs. s characteristics excluding any parameter interpretation based on models.

2. Equations, iteration procedure and computer program

Sigmoidal v vs. s curves can be described, analogously to the classical Hill equation [10], with the parameters V_{max} , $K_{0.5}$ and n_H :

$$v = \frac{V_{max}}{1 + \left(\frac{K_{0.5}}{s}\right)^{n_H}} \quad (1)$$

With the following transformations, the fit-problem can be reduced to a linear least-squares fit:

$$\log \frac{v}{V_{max} - v} = n_H \cdot \log s - n_H \cdot \log K_{0.5} \quad (2)$$

$$\frac{s^{n_H}}{v} = \frac{1}{V_{max}} \cdot s^{n_H} + \frac{K_{0.5}^{n_H}}{V_{max}} \quad (3)$$

V_{max} being known, n_H and $K_{0.5}$ can be computed according to equation (2); and n_H being known, V_{max} and $K_{0.5}$ are obtained from equation (3). However, only two parameters can be determined at a time, since one parameter is always part of the variables. All three parameters can be obtained by iterative applications of equations (2) and (3). The procedure is schematically represented in fig. 1.

* The following symbols are used:

v = initial velocity,
 s = substrate concentration,
 V_{max} = maximal velocity,
 $K_{0.5}$ = substrate concentration giving $v = V_{max}/2$,
 n_H = "interaction coefficient",
 α = limitation factor.

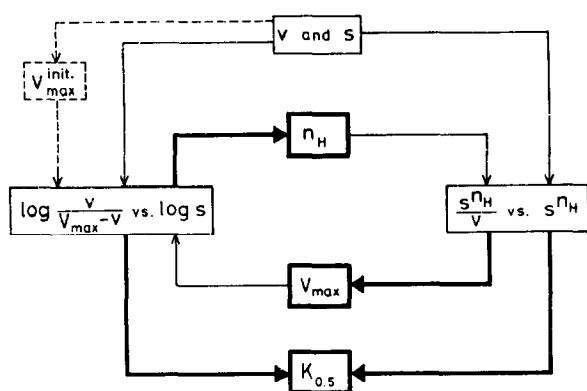


Fig. 1. Scheme of the iteration procedure.

From the experimental data v and s , an initial value of V_{max} (V_{max}^{init}) is produced and a least-squares fit of $\log(v/(V_{max}-v))$ vs. $\log s$ according to equation (2) is performed. The resulting value of n_H is used in equation (3), which on least-squares-fitting of s^{n_H}/v vs. s^{n_H} delivers a better value of V_{max} , which again is incorporated into equation (2). This procedure is repeated, until the changes of n_H and V_{max} in two consecutive loops pass a limit which is fixed in the program. $K_{0.5}$ is not used for iteration. For this estimation of the parameters, all the experimental data were used.

In order to analyse for asymmetry of the v vs. s curves as well as for linearity in equation (2), a second iteration procedure follows (see general remarks). Now, only the experimental data are used, which fulfil the condition $(1-\alpha) \cdot V_{max} < v < \alpha \cdot V_{max}$. The limitation factor α can arbitrarily be chosen.

The results of both iteration procedures are collected in form of tables (see figs. 2 and 3) and graphically (fig. 4). For further evaluation of the results of fitting, two additional linear transformations from equation (1) are used: $1/v$ vs. $1/s^{n_H}$ and v vs. v/s^{n_H} (fig. 2).

The least-squares procedure implies that the experimental error of the data is exclusively connected with the determination of v and that s is accurately determined (see [1]). However, the error prolongation resulting from s^{n_H} is considered. In contrast to the program of Cleland [4] no weighting factor for v is used, however, an indirect weighting is

implied by the quotient s^{n_H}/v in equation (3) [1, 11].

On routine application we have found that the iteration procedure is convergent with respect to n_H and V_{max} . Nevertheless, errors implied in the data set ($n_H < 0.5$; $V_{max} < v$) will result in specified comment statements, in order to avoid erroneous computations. The program written in FORTRAN IV is developed for an IBM 360/44 H-level computer and occupies 32 K bytes of system 360 memory. It is available at cost from the authors.

3. General remarks

Any interpretation of the results obtained necessitates a critical inspection of the computed and experimental data, looking for the following discrepancies in the curves of numerical values.

- 1) A large deviation of each individual experimental point from the respective computed point, as numerically given in the output (fig. 3, "error"),
- 2) a serial error along the curve with equal sign,
- 3) a larger error accumulated in the derived parameters,
- 4) discrepancies of V_{max} resulting from the three linear transformations (fig. 2),
- 5) discrepancies between the first and second (limited) iteration procedures.

Any error may result from the experimental data or from the inapplicability of the Hill equation. Whereas a discussion of errors in the experimental data set is out of the scope of this paper, the applicability of the equation system must always be questioned. As mentioned above, equation (1) displays a sigmoidal v vs. s curvature with the minimum number of three parameters. For this reason, equation (1) cannot be used for systems which can only be described with at least four parameters, such as e.g. substrate inhibition mechanism, systems which operate with a mixture of positive and negative cooperativity [12] or systems which exhibit asymmetry.

Equation (1) implies simplifications which are not fulfilled in all enzymic reactions, especially the assumption that the "interaction coefficient" n_H is

```

                                PDKP 1000

                                DATA CALCULATED ACCORDING TO
                                V=VMAX*(S**NH)/((S**NH)+(K(0.5)**NH))
                                VMAX=200, K(0.5)=1.905, NH=2.15
                                DEVIATIONS OF +-5% ADDED TO V

1. ITERATION
=====
NO LIMIT

*****
*
*          NH = 2.155          VMAX = 199.687          K(0.5) = 1.9049
*
*****

RESULTS OF THE LINEAR TRANSFORMATIONS

LOG V/VMAX-V VS. LOG S
NH = 2.155 +- 0.691 %          K(0.5) = 1.902 +- 0.950 %          SIGMA = 0.03784

(S**NH)/V VS. (S**NH)
VMAX = 199.687 +- 0.228 %          K(0.5) = 1.905 +- 1.123 %          SIGMA = 0.00178

1/V VS. 1/(S**NH)
VMAX = 181.273 +- 9.690 %          K(0.5) = 1.808 +- 4.548 %          SIGMA = 0.00289

V VS. V/(S**NH)
VMAX = 199.090 +- 0.892 %          K(0.5) = 1.897 +- 0.991 %          SIGMA = 5.52309

NUMBER OF ITERATIONS 31          INITIAL VMAX = 212.454

```

Fig. 2a

```

                                PDKP 1000

                                DATA CALCULATED ACCORDING TO
                                V=VMAX*(S**NH)/((S**NH)+(K(0.5)**NH))
                                VMAX=200, K(0.5)=1.905, NH=2.15
                                DEVIATIONS OF +-5% ADDED TO V

2. ITERATION
*****
LIMIT      0.100*VMAX  -  0.900*VMAX

*****
*
*          NH =  2.141          VMAX = 199.918          K(0.5) =  1.9086          *
*
*****

RESULTS OF THE LINEAR TRANSFORMATIONS

LOG V/VMAX-V  VS.  LOG S
NH      =    2.141 +-  1.406 %          K(0.5) =  1.907 +-  1.481 %          SIGMA =    0.02969

(S**NH)/V  VS.  (S**NH)
VMAX = 199.918 +-  0.812 %          K(0.5) =  1.909 +-  1.760 %          SIGMA =    0.00138

1/V  VS.  1/(S**NH)
VMAX = 197.854 +-  3.987 %          K(0.5) =  1.893 +-  2.465 %          SIGMA =    0.00065

V  VS.  V/(S**NH)
VMAX = 199.016 +-  1.679 %          K(0.5) =  1.899 +-  2.016 %          SIGMA =    6.32407

NUMBER OF ITERATIONS    59          INITIAL VMAX = 199.687

```

Fig. 2b

Fig. 2. Computer output: results of the linear transformations, Sigma = standard deviation = $\sqrt{\sum(x_i - \bar{x})^2 / n - 2}$: a) First iteration, b) Second iteration.

PDKP 1000

DATA CALCULATED ACCORDING TO

$$V = V_{\text{MAX}} * (S^{*NH}) / ((S^{*NH}) + (K(0.5) * NH))$$

VMAX=200, K(0.5)=1.905, NH=2.15

DEVIATIONS OF +-5% ADDED TO V

1. ITERATION

=====

NO LIMIT

```

*****
*                                     *
*      NH = 2.155      VMAX = 199.687      K(0.5) = 1.9049      *
*                                     *
*****

```

| S | V | V(CALC) | ERROR | |
|---------|-----------|-----------|------------|---|
| 0.10000 | 0.35335 | 0.34744 | 1.70194 % | + |
| 0.20000 | 1.52761 | 1.53862 | -0.71560 % | + |
| 0.40000 | 6.87252 | 6.67697 | 2.92873 % | + |
| 0.50000 | 10.66490 | 10.58248 | 0.77887 % | + |
| 0.70000 | 19.76716 | 20.68761 | -4.44927 % | + |
| 0.80000 | 28.13981 | 26.66595 | 5.52714 % | + |
| 0.90000 | 31.57753 | 33.09545 | -4.58650 % | + |
| 1.00000 | 40.00000 | 39.84947 | 0.37774 % | + |
| 1.10000 | 46.02166 | 46.80733 | -1.67852 % | + |
| 1.20000 | 56.71321 | 53.85885 | 5.29970 % | + |
| 1.40000 | 64.61767 | 67.87107 | -4.79350 % | + |
| 1.50000 | 74.82569 | 74.68340 | 0.19052 % | + |
| 1.60000 | 83.05723 | 81.29217 | 2.17125 % | + |
| 1.70000 | 89.54476 | 87.65869 | 2.15161 % | + |
| 1.80000 | 92.00235 | 93.75567 | -1.87009 % | + |
| 2.00000 | 105.19386 | 105.08058 | 0.10780 % | + |
| 2.20000 | 121.08892 | 115.21788 | 5.09559 % | + |
| 2.40000 | 121.81598 | 124.20262 | -1.92157 % | + |
| 2.60000 | 134.85774 | 132.11667 | 2.07473 % | + |
| 2.80000 | 132.20238 | 139.06370 | -4.93394 % | + |
| 3.00000 | 145.25167 | 145.15301 | 0.06797 % | + |
| 3.30000 | 156.07275 | 152.90793 | 2.06975 % | + |
| 3.60000 | 156.21403 | 159.28930 | -1.93062 % | + |
| 4.00000 | 166.24229 | 166.11668 | 0.07561 % | + |
| 5.00000 | 177.66999 | 177.51097 | 0.08958 % | + |
| 5.50000 | 185.05156 | 181.24895 | 2.09800 % | + |
| 6.00000 | 184.34455 | 184.15680 | 0.10195 % | + |
| 6.50000 | 182.92140 | 186.45469 | -1.89499 % | + |
| 7.00000 | 188.50716 | 188.29673 | 0.11175 % | + |
| 8.00000 | 191.24980 | 191.02180 | 0.11936 % | + |
| 9.00000 | 193.14020 | 192.89853 | 0.12528 % | + |

Fig. 3a

```

                                PDKP 1000

                                DATA CALCULATED ACCORDING TO

                                V=VMAX*(S**NH)/((S**NH)+(K(0.5)**NH))

                                VMAX=200, K(0.5)=1.905, NH=2.15

                                DEVIATIONS OF +-5% ADDED TO V

2. ITERATION
=====
LIMIT      0.100*VMAX  -  0.900*VMAX

*****
*
*          NH =  2.141          VMAX =  199.918          K(0.5) =  1.9086
*
*****

      S          V          V(CALC)          ERROR
0.10000      0.35335      0.36165      -2.29578 %  -
0.20000      1.52761      1.58515      -3.63020 %  -
0.40000      6.87252      6.80668      0.96731 %  -
0.50000     10.66490     10.75079     -0.79888 %  -
0.70000     19.76716     20.90769     -5.45508 %  -
0.80000     28.13981     26.89557      4.62621 %  +
0.90000     31.57753     33.32329     -5.23885 %  +
1.00000     40.00000     40.06506     -0.16239 %  +
1.10000     46.02166     47.00167     -2.08506 %  +
1.20000     56.71321     54.02477      4.97630 %  +
1.40000     64.61767     67.96632     -4.92692 %  +
1.50000     74.82569     74.73999      0.11467 %  +
1.60000     83.05723     81.30991      2.14897 %  +
1.70000     89.54476     87.63855      2.17508 %  +
1.80000     92.00235     93.69955     -1.81132 %  +
2.00000    105.19386    104.96050      0.22233 %  +
2.20000    121.08892    115.04644      5.25221 %  +
2.40000    121.81598    123.99272     -1.75554 %  +
2.60000    134.85774    131.88019      2.25777 %  +
2.80000    132.20238    138.81082     -4.76075 %  +
3.00000    145.25167    144.89209      0.24817 %  +
3.30000    156.07275    152.64688      2.24431 %  +
3.60000    156.21403    159.03810     -1.77572 %  +
4.00000    166.24229    165.88785      0.21366 %  +
5.00000    177.66999    177.35378      0.17829 %  +
5.50000    185.05156    181.12711      2.16669 %  -
6.00000    184.34455    184.06757      0.15048 %  -
6.50000    182.92140    186.39489     -1.86351 %  -
7.00000    188.50716    188.26322      0.12958 %  -
8.00000    191.24980    191.03232      0.11384 %  -
9.00000    193.14020    192.94370      0.10184 %  -

```

Fig. 3b

Fig. 3. Computer output of tabulated input data (s, v), calculated values of v (v calc) and the relative error between v and v (calc), for a) first and b) second iteration. + indicates the values used in the iteration procedures.

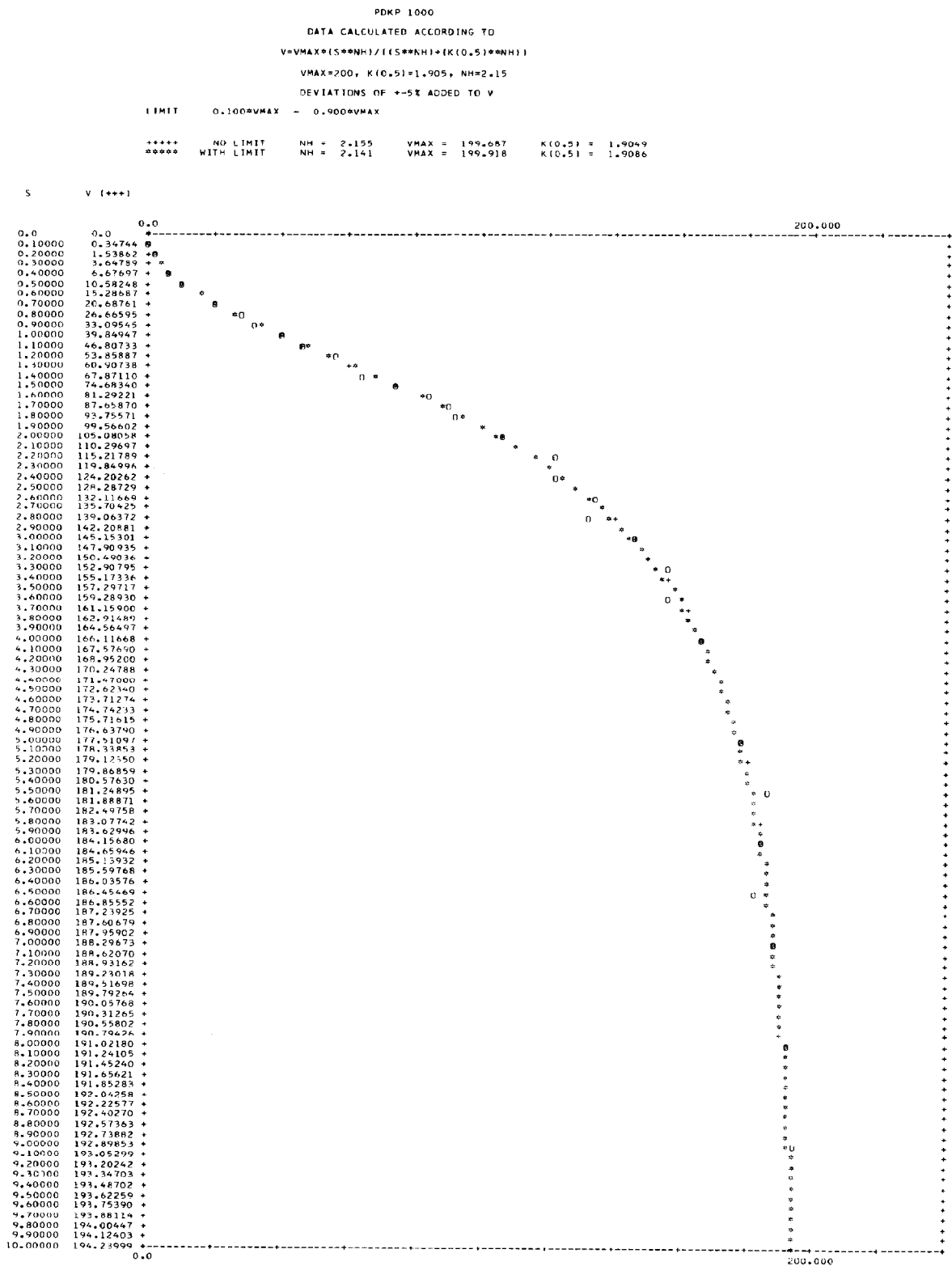


Fig. 4. Graphical output of ν vs. s . Experimental values: \circ , fitted values: + first iteration, * second iteration.

independent of s . If n_H is a function of s , the Hill plot (equation 2) cannot be linear over the total range of $\log s$ [8, 13]. However, as pointed out by Frieden [14], Hill plots usually are linear in the optimum experimental range (0.1 to 0.9 V_{max}), which indeed is approached in the second iteration procedure used here. It should be pointed out that for theoretical reasons [7, 13] this range of the Hill plot served for the interpretation of the "interaction coefficient" n_H . Thus, on practical and theoretical grounds, the introduction of the limitation factor α is fully justified and serves finally as the test of the linearity of the Hill plot. For greater refinement [7], the magnitude of the factor α can be varied arbitrarily.

The three parameters obtained with the procedure described here immediately lead to the process of parameter interpretation by one of the known models in enzyme kinetics, as exemplified in the case of the steady-state kinetics of yeast pyruvate kinase [15]. Furthermore, the program can be used for parameter evaluation in any type of binding studies.

References

- [1] W.R.Gardiner and J.H.Ottaway, FEBS Letters 2, Suppl. (1969) S34.
- [2] K.R.Hanson, R.Ling and E.Havir, Biochem. Biophys. Res. Commun. 29 (1967) 194.
- [3] W.W.Cleland, Nature 198 (1963) 463.
- [4] W.W.Cleland, Adv. Enzym. 29 (1967) 1.
- [5] J.Kowalik and J.F.Morrison, Mathem. Biosciences 2 (1968) 57.
- [6] J.Monod, J.Wyman and J.-P.Changeux, J. Mol. Biol. 12 (1965) 88.
- [7] D.E.Koshland Jr., G.Némethy and D.Filmer, Biochemistry 5 (1966) 365.
- [8] C.C.Griffin and L.Brand, Arch. Biochem. Biophys. 126 (1968) 856.
- [9] W.Ferdinand, Biochem. J. 98 (1966) 278.
- [10] A.V.Hill, Biochem. J. 7 (1913) 471.
- [11] G.N.Wilkinson, Biochem. J. 80 (1961) 324.
- [12] A.Levitzki and D.E.Koshland Jr., Proc. Natl. Acad. Sci. U.S. 62 (1969) 1121.
- [13] M.M.Rubin and J.-P.Changeux, J. Mol. Biol. 21 (1966) 265.
- [14] C.Frieden, J. Biol. Chem. 242 (1967) 4045.
- [15] H.-J.Wieker, K.-J.Johannes and B.Hess, Abstract No. 398, VI. Meeting of FEBS, Madrid, 1969.