

The fitting method is widely used in the computational practice for approximate solution of the ill-posed problems.⁸ This method can be efficiently used for solution of the system of equations of laser photoacoustic gas analysis in the case that only the approximate solution, or the right-hand side $\Delta\mathbf{Y}$ of Eq. (3), rather than the exact one, is known and (or) the matrix of the absorption coefficients is inaccurate.

For such problems, the concept of quasi-solution is introduced and the fitting method allows one to find some approximation of a quasi-solution. The fitting method in this case consists in the following. The direct problem is solved for \mathbf{C} elements in some bounded region D satisfying the physical sense (the operator $\Delta K \mathbf{C}$ is calculated) and the element of $\tilde{\mathbf{C}} \in D$ minimizing the discrepancy between $\Delta K \tilde{\mathbf{C}}$ and $\Delta\mathbf{Y}$ is called the quasi-solution of Eq. (3), that is, the quasi-solution is determined by the following equation:

$$\rho(\Delta K \tilde{\mathbf{C}}, \Delta\mathbf{Y}) = \inf_{\mathbf{C} \in M} \rho(\Delta K \mathbf{C}, \Delta\mathbf{Y}) \quad (4)$$

under the condition that $0 \leq \tilde{c}_i \leq 1$; $\sum_{i=1}^N \tilde{c}_i \leq 1$.

The evolution-genetic method (EGM)^{15,16} is a promising computational fitting method that allows determination of a quasi-solution to Eq. (3). Its advantage is the possibility of efficiently solving complex optimization problems. Although the discrepancy functional has one extreme (minimum), it acquires the ravine shape (strongly elongated along one direction) in the case of ill-posed problems, and application of traditional minimum search algorithms becomes inefficient.¹⁷

In doing this work, we have realized a modification of the algorithm of evolution-genetic strategy described in Ref. 6. Let us introduce the needed terms: chromosome is some realization of the vector of unknown concentrations, the conditions of the domain of definition and the physical sense of the sought solution are imposed on a chromosome; population is a set of chromosomes.

Let the discrepancy functional

$$F(\tilde{\mathbf{C}}) = \rho(\Delta K \tilde{\mathbf{C}}, \Delta\mathbf{Y})$$

be a target (optimized) function calculated for every chromosome.

The algorithm of search for the quasi-solution involves the following stages:

1. At the initial stage, the starting parent population of P chromosomes is formed in a random manner, that is, P versions of the vectors of initial gas concentrations are generated: $\mathbf{C}_0^p (c_{01}^p, c_{02}^p, \dots, c_{0i}^p, \dots, c_{0N}^p)$, $p = 1 \dots P$. The values of Δy_i are calculated from the initial concentrations c_{0i}^p by the system of equations (2). The starting population is sorted by the target function $F(\mathbf{C}_0^p)$ in the decreasing order (the smallest superscript p corresponds to the smallest value of the target function).

2. The next stage involves reproduction. At the stage of reproduction, L new vectors are generated instead of the P initial concentration vectors: every new concentration vector \mathbf{C}_{1r}^l ($l = 1 \dots L$) is formed of two randomly selected old vectors \mathbf{C}_0^{p1} and \mathbf{C}_0^{p2} ($p1$ and $p2$ are values from the range $1 \dots P$), and the vectors with the smaller value of the target function $F(\mathbf{C}_0^p)$ are taken with higher probability. The components of the vectors are formed in the following way:

$$c_{1ri}^l = \begin{cases} c_{0i}^{p1}, & \text{if } s = 0, \\ c_{0i}^{p2}, & \text{if } s = 1, \end{cases}$$

where c_{1ri}^l are the concentration values after the first reproduction; s is a random value equal to 0 or 1.

3. Then the vectors obtained as a result of reproduction mutate. At this stage, the concentration vectors \mathbf{C}_{1r}^l are changed. The number of versions of the concentration vectors keeps equal to L , while the concentrations change by the following law:

$$c_{1mi}^l = \begin{cases} c_{1ri}^l, & \text{if } s_1 = 0, \\ c_{1ri}^l + w, & \text{if } s_1 = 1, \end{cases}$$

where c_{1mi}^l are the concentration values after the first mutation; s_1 is a random value equal to 0 or 1; w is a random parameter distributed normally with the mathematical expectation equal to 0 and the variance σ^2 . The values of Δy_i and new values of the target function $F(\mathbf{C}_{1m}^l)$ are calculated from the obtained concentrations c_{1i}^m by the system of equations (2).

4. Of all L versions of the concentrations \mathbf{C}_{1m}^l , P the best ones (from the viewpoint of smallness of the target function) are selected and, at the next stage, they are used as the initial ones for mutation and reproduction.

5. The reproduction and mutation cycle are reiterated a fixed number of times (generations) G . During a cycle, the variance σ^2 of a random parameter w varies in the process of evolution depending on the frequency of decrease of the target function calculated for several iterations.

6. As the cycle is terminated, only one, the best (from the viewpoint of the smallest value of the target function), vector of concentrations \mathbf{C}_{Gm}^l is taken as a quasi-solution.

For comparison of EGM operation with the Tikhonov regularization method and the standard method of solution of a system of linear algebraic equations, we conducted mathematical simulation and processing of experimental LPGA data. As a method for solution of a system of linear algebraic equations, we used the Gauss method with column-by-column selection of the main element¹⁷ (hereinafter in this paper, it is referred to as the method of direct solution). When constructing the regularized solution, the discrepancy method was used for determination of the regularization parameter.¹³

Mathematical simulation was conducted in a closed cycle: the composition of a gas mixture was specified, the vector of reduced signals of the system (1) was calculated for this mixture composition, the components of the vector were disturbed by additive noise distributed by the normal law with the given relative standard deviation. Then the concentrations of the mixture components were reconstructed from these model data. The measured data were obtained with LPGA based on a tunable CO₂ low-pressure laser and a nonresonance measurement cell described in Ref. 10.

Figure 1 depicts the results of algorithm operation with a few number of mixture components $N = 3$ [method of direct solution (1), search for quasi-solution over 1000 generations with the use of EGM (2), and the regularization method (3)].

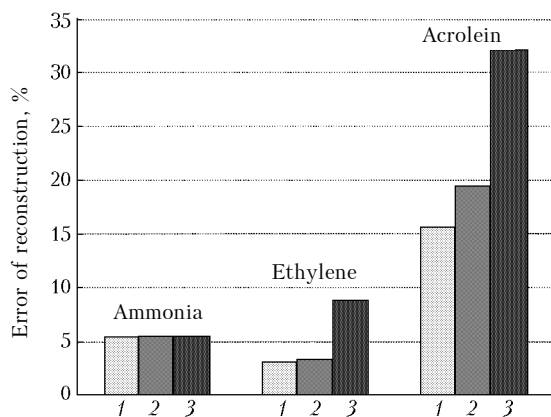


Fig. 1. Errors of reconstruction of gas concentration in a three-component mixture.

The following mixture was taken for analysis: ammonia–ethylene–acrolein ($10^{-3} : 5 \cdot 10^{-3} : 9 \cdot 10^{-4}$). The pairs of spectral measurement channels were 10R24–10R18, 10P34–10P30, and 10P6–10P4. The “measurement” data are the results of numerical simulation. The condition number for the matrix of difference of absorption coefficients was 5.3. The relative standard noise in all measurement channels was 10%. For reconstruction we used the data averaged over a series of 10 measurements. Simulation was repeated 10 times, and the mean error for 10 series of experiments was calculated.

It is seen from Fig. 1 that at a small number of mixture components, application of the processing algorithms based on the regularization method is inefficient (this was also noticed in Ref. 13). The mean error of reconstruction with the use of the search for quasi-solutions is roughly at the same level with the mean error of the method of direct solution, and the error of reconstruction by the regularization method is much larger.

Figures 2 and 3 illustrate how the error of reconstruction of gas mixture components depends on the composition of the analyzed mixture (with unchanged number of mixture components). Figures 2 and 3 depict the results of mathematical simulation of reconstruction

of gas concentrations in two six-component mixtures [method of direct solution (1), search for quasi-solution over 1000 generations with the use of EGM (2), search for quasi-solution over 200 generations with the use of EGM (3), and the regularization method (4)]. The relative standard noise was 5%, and no averaging was conducted. The reconstruction error calculated from a series of ten single measurements is presented.

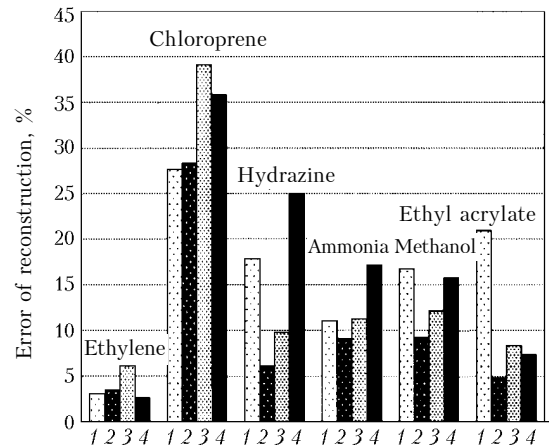


Fig. 2. Errors of reconstruction in a six-component mixture ethylene – chloroprene – hydrazine – ammonia – methanol – ethyl acrylate.

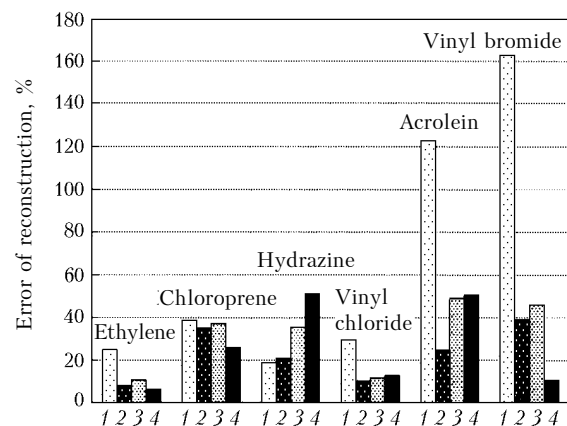


Fig. 3. Errors of reconstruction in a six-component mixture ethylene – chloroprene – hydrazine – vinyl chloride – acrolein – vinyl bromide.

Figure 2 shows the results of reconstruction for the mixture ethylene–chloroprene–hydrazine–ammonia–methanol–ethyl acrylate ($5 \cdot 10^{-4} : 9 \cdot 10^{-5} : 4 \cdot 10^{-4} : 10^{-4} : 2 \cdot 10^{-4} : 3 \cdot 10^{-4}$). The pairs of the measurement channels were: 10R6–10R12, 10P14–10P10, 10R18–10R22, 9P14–9P22, 10P22–10P18, and 9R14–9R22.

Figure 3 shows the results of reconstruction for the mixture ethylene–chloroprene–hydrazine–vinyl chloride–acrolein–vinyl bromide ($10^{-4} : 9 \cdot 10^{-5} : 2 \cdot 10^{-4} : 5 \cdot 10^{-4} : 4 \cdot 10^{-4} : 3 \cdot 10^{-4}$). The pairs of measurement channels were: 10P14–10P10, 10P22–10P18, 10R18–10R22, 10P32–10P28, 9P14–9P12, and 9R20–9R12.

For the mixture shown in Fig. 2, the errors of reconstruction are not very large and application of specialized reconstruction algorithms does not provide a

significant advantage as compared with the method of direct solution. However, even here the search for quasi-solutions over 1000 generations with the use of EGM is preferable as compared with the regularization method and the method of direct solution.

For the mixture shown in Fig. 3, two components (acrolein and vinyl bromide) cannot be reconstructed by the method of direct solution (errors are higher than 100%) and here the mean error of reconstruction for most gases decreases markedly with the use of the regularization method and the search for quasi-solutions (the errors for these two methods are on the same order). The increase in the error of reconstruction for EGM with the decrease of the number of generations from 1000 to 200 (the computational time in this case decreased five times) is not very significant due to irregular convergence rate of this algorithm. With the further decrease of the number of generations, the reconstruction error increased drastically and it is not shown in Figs. 2 and 3 because of too large difference in the values.

It should be noted that the condition number of the matrix of difference of the absorption coefficients for the gas mixture in Fig. 2 equals 17.8, and for Fig. 3 it is 1176. The condition number characterizes the maximum possible coefficient of transfer of the error in the right-hand side or the error of the matrix in the left-hand side to the error of solution of the system of linear equations.¹⁸ Thus (as is seen from Figs. 2 and 3), the condition number can serve a qualitative criterion of the need to apply specialized algorithms for solution of equations of laser gas analysis.

Figure 4 depicts the results of processing the LPGA experimental data. In this case, we analyzed a six-component mixture of ethylene – carbon dioxide – ammonia – methanol – ethanol – isopropanol ($1.6 \times 10^{-4} : 9.6 \times 10^{-2} : 7.3 \cdot 10^{-3} : 8.5 \cdot 10^{-3} : 6.6 \cdot 10^{-3} : 9.2 \cdot 10^{-3}$). (The results on reconstruction of carbon dioxide are not shown, since we failed to reconstruct it in all cases.) The pairs of the spectral measurement channels were 10P14–10P20, 9R18–9R10, 9R16–9R12, 9P34–9P32, 9P40–9P42, and 10R12–10R30. The relative standard noise value in the spectral measurement channels at single measurements was from 1 to 9%. Reconstruction was conducted from averaged (over ten measurements) data. The condition number of the matrix of difference of the absorption coefficients was $4.7 \cdot 10^4$. For simulation of the effect of errors in absorption coefficients, the system matrix was distorted (imperfect values of the absorption coefficients for isopropanol and carbon dioxide at 10R30 wavelength were input). The input distortions of the matrix of absorption coefficients were small:

$$\frac{\|\Delta K_{ERR}\|_M}{\|\Delta K\|_M} = 0.34\%,$$

where $\|\bullet\|_M$ is the matrix norm; ΔK is the matrix of difference of absorption coefficients; ΔK_{ERR} is the matrix of errors in the absorption coefficients.

The condition number of the distorted matrix was $2.4 \cdot 10^5$.

Figure 4 shows the errors of reconstruction performed with the use of the distorted [method of direct solution (1), search for quasi-solution over 1000 generations with the use of EGM (2), and the regularization method (3)] and actual matrix of absorption coefficients [method of direct solution (4), search for quasi-solution over 1000 generations with the use of EGM (5), and the regularization method (6)].

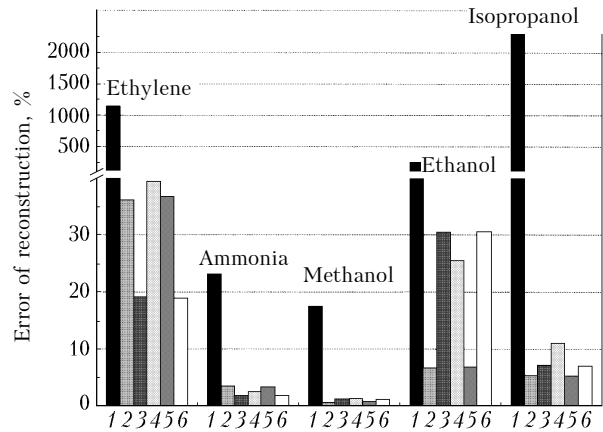


Fig. 4. Errors of gas reconstruction in a six-component mixture ethylene – carbon dioxide – ammonia – methanol – ethanol – isopropanol.

It can be seen from Fig. 4 that for the distorted matrix the method of direct solution gives very large errors (hundreds and thousands percent) for most gases. (The reconstructed concentrations not only differ from the actual ones by several orders of magnitude, but they may even have physically meaningless values.) At the same time, these errors are not too large for the actual matrix of absorption coefficients (although they can achieve 40%).

It is interesting to note that the level of reconstruction errors for the method of search for quasi-solutions and the regularization method is almost independent of errors in the matrix of gas absorption coefficients – both of these methods allow reconstruction of the concentrations of component gases with acceptable accuracy (errors of both methods are on the same order of magnitude).

Thus, the results of mathematical simulation and processing of measurements show that the method of search for quasi-solutions (using the evolution-genetic fitting algorithm) allows reconstruction of gas concentrations in laser photoacoustic gas analysis for both few-component and many-component gas mixtures. This method does not demand extra information about the level of noise in measurement channels.

A disadvantage of the method of search of quasi-solutions is relatively long (as compared with the regularization method) computational time. However, this disadvantage becomes insignificant when using up-to-date computers. Besides, with the use of EGM, it

becomes possible to regulate, to a certain extent, the computational time by varying the number of generations. The EGM speed can be increased due to paralleling of this algorithm and use of multiprocessor systems.^{15,16}

In conclusion, we would like to note one more significant advantage of the method of search for quasi-solutions – it allows solution of not only the system of linear algebraic equations (1), but also more general (without the condition of the small spectral width of a laser line as compared with the width of a gas absorption line) system of integral equations of the laser gas analysis.

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