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THE $\chi^{2}$ CRITERION IN THE PHASE SHIFT
ANALYSIS FOR CHOOSING STATISTICALLY PERMISSIBLE

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In a phase shift analysls of experimental data in accordance with the general method of a regressive analysis phase shift evaluations are found from the condition of minimization of the further square functionai:

$$
\begin{equation*}
M\left(y_{1} ; \delta_{1}\right)=\sum_{1=1}^{\mathbf{N}} \frac{\left[y_{1}-\hat{\eta}_{1}\left(\delta_{1}\right)\right]^{2}}{\sigma_{1}^{2}}, \tag{1}
\end{equation*}
$$

where $y_{1}$ is experimentally obtained values,
$\sigma_{i}$ is root-mean-square errors of measurements of the given values, and
$\hat{\eta}_{1}\left(\delta_{1}\right)$ is theoretical values of the same quantities which depend upon the selection of the estimate of the phase shifts $\delta_{1}, \delta_{2}, \ldots, \delta_{m}$.

If experimental data $y_{i}$ is subordinated to the normal law of distribution, the minimum values of the functional $M$ should be respectively described by the $x^{2}$-distribution. Consequently, the mean value and the dispersion of this quantity should be, respectively, $E M=m-n$ and $D M=2(n-m)$.

For a particular minimum value of $M_{0}$ corresponding to the glven selection of experimental data we can determine basing on the $x^{2}$-distribution the degree of statistical confidence $P\left(M \geq M_{0}\right)$ which defines the rate of acquiring the minimum values of $M_{j}>M_{0} \quad$ In a series of mulitple repeated measurements of the whole assembly of experimental data $y_{11}$.

The acquisition of the minimum value $M_{0}$ corresponding to a low degree of confidence can show the imperfection of the theoretical curve taken for estmating the true regression line, e.g., due to a small number of the parameters varled. The reason of a high value of $\mathrm{m}_{0}$ can be the presence of systemaHic errors not taken into account in the measurements of experimental values.

It might seem that the same rellability criterion is acceptable also for clear-
ing out the possibillty of negdecting some of the solutions in such a case when the functional manifests some minima in various argument regions. Just due to the seeming valldity the $x^{2}$-criterion has been for many years erraneously taken for clearing out the permissibilty of ambiguous solutions obtained. (See, e.g., refs. $/ 1-6 /$ ). The erraneous recomendation on the use of the $x^{2}$-criterion in considering the problems of neglecting ambiguous solutions can be found in monographs dedicated specially to the regressive analysis (see, e.g., ref. 7/).

In fact the $x^{2}$-distribution dominates only over the values of absolute minima obtalned in the analysis of independent series of repeated experiments. Therefore, the application of the $x^{2}$-criterion is justiflable only for determining the statistical reliability of a solution corresponding to the deepest minimum of the functional $m$ or the compatiblity of various evaluations obtained in separate series of experiments.

The $x^{2}$-distribution has no bearing on the value of the relative minima of the functional M .

We have no right to consider a large value of the obtained relative minimum of the functional $M$ to be due to the realization (in a given series of measu rements) of statistically permissible, large in sums deviations of experimental values from the theoretical curve. Just for this series of experimental values we have the indication to the realization of smaller summed deviations from the curve obtained under the same theoretical assumptions but corresponding to the absolute minimum of the functional M. A selection among the general assembly has been performed, dice have been cast, so to say, the game has been made and, therefore, one cannot refer to the statistical possibility of appearing large in sums errors in repeated selections, in general, if for a particular selection the realization of considerably smaller summed errors by the deepest absolute minimum of the functional $M$ is a fact.

The statistical reliability of solutions corresponding to the relative minimum of the functional $M$, can be easily determined by the maximum likellhood method. Say, apart from the solution $\hat{\eta}$ (0) corresponding to the absolute mimimum $M_{0}$, there is another solution $\hat{\eta}$ (1) corresponding to the relative minimum $M_{1}>M_{0} \quad$. The probabllity of obtaining the given selection (i.e., the preset experimental values $y_{1}, y_{2}, \ldots, y_{n}$ in the intervals $d_{1}, d y_{2}$ $\ldots$.. $d_{n}$, respectively) from the general assembly presented by some true values $\eta_{1}, \eta_{2}, \ldots \quad \eta_{n}$ is expressed by the likelihood function

$$
\begin{equation*}
L\left(y_{i} ; \eta_{i}\right)=\frac{1}{(2 \pi)^{n / 2} \prod_{i=1}^{\pi} \sigma_{i}} \exp -y_{i=1}^{n} \frac{\left(y_{i}-\eta_{i}\right)^{2}}{\sigma_{i}^{2}} \tag{2}
\end{equation*}
$$

## multiplied by the corresponding intervals $\mathrm{dy}_{1}, \mathrm{dy}_{2} \ldots \mathrm{dy}_{\mathrm{n}}$

If for true values we have hypothetically two series of values $\eta_{i}(-1)$ and $\eta_{i}(-I I)$, then having calculated the probabluties of obtaining the glven selection from different general assemblies

$$
\begin{align*}
& W_{1}=L\left[y_{i} ; \eta_{1}(l)\right] \prod_{i=1}^{n} d y_{i}  \tag{3}\\
& W_{n I}=L\left[y_{i} ; \eta_{1}(I)\right] \prod_{i=1}^{n} d y_{i}
\end{align*}
$$

we can define the statistical reliability of accepting each of the hypotheses.
Accepting one of the hypotheses, e.g., the first one to be true, we can determine the probability that the accepted mypothesis is correct

$$
\begin{equation*}
P_{I}=\frac{\dot{W}_{I}}{W_{I}+W_{I I}}=\frac{L\left[y_{1} ; \eta_{1}(I)\right]}{\left.L\left(y_{i} ; \eta_{i}(I)\right]+L_{y_{1}} ; \eta_{1}(I I)\right]}=\frac{1}{1+\exp -y_{2}\left(M_{I I}-M_{I}\right)} \tag{4}
\end{equation*}
$$

The probabillty of rejecting the correct hypothesis will be then

$$
a_{I}=1-P_{I}=P_{I I}=\frac{W_{I I}}{W_{I}+W_{I I}} .
$$

If hypotheses 1 and II are different, noncompatible with each other estimates of the general essembly determined on the basis of the given selection $y_{1}, y_{2}, \ldots y_{n}$, the corresponding likellhood functions are of another form:

$$
\begin{align*}
& L\left[y_{i} ; \hat{\eta}_{i}(I)\right]=\frac{1}{(2 \pi)^{\frac{n-m}{2}}\left(\sum_{i=1}^{n} \frac{1}{\sigma_{1}^{2}}\right)^{m / 2} \prod_{i=1}^{n} \sigma_{i}} \exp -\frac{M_{I}}{2}  \tag{5}\\
& L\left[\bar{y}_{1} ; \hat{\eta}_{1}(I I)\right]=\frac{1}{(2 \pi)^{\frac{n-m}{2}}\left(\sum_{i=1}^{n} \frac{1}{\sigma_{1}^{2}}\right)^{m / 2} \prod_{i=1}^{n} \sigma_{i}} \exp -\frac{M_{I_{1}}^{2}}{l}
\end{align*}
$$

It can be easily seen that this account of the dependence of the enalysed mypotheses from the experimental data selection itself does not change the expressions for the probabilities $P_{I}$ and $P_{1 I}$. Thus, the problem of comparing the
rellabillties of two hypotheses on the general assembly on the basis of selection data does not differ from that of comparing the rellablilties of two noncompatible estimates of the same general assembly.

The above probabilities $P_{1}$ and $P_{11}$ are related to a statistical sub-as sembly consisting of the selections made according to a certain characteristic from the general assembly of possible selections. This sub-assembly consists of selections within the intervals $d_{1}, d y_{2}, \ldots d_{m}$ colnciding with the glven selection or with that symmetrical to the given one with respect to both the hypotheses.

We permit ourselves to explain the reason of introducing such statistical sub-assembly by very simple examples which in their essence do not differ from more complicated ones, As justlfication of the talk started on a statistical assembly let us note that a mess in essence in simple problems on statistics is most of ten due to the usage of the concept "probability" without rendering concrete a statistical assembly to which it refers. First of all, we should clear out in what sense, for instance, a small probability $W_{I I}$ of obtaining a given selection from
general assembly II compared to the probability $\boldsymbol{W}_{1}$ of obtalining the same selection from, general assembly I allows one to speak about the unreliability of hypothesls II.

We permit ourselves to dwell upon this problem in more detail apologizing to the reader beforehand for the simpllcity of examples taken for illustration since a mess is committed as a result of introducing reciprocal probabilities.

Indeed, one can speak about a small probabllity of fulfilling hypothesis II and, respectively, about a great probability of fulfilling hypothesis If one knows deliberatly that only one of them is carried out in nature.

All the above problems including the unlawfulness of using the : $x^{2}$-criterion in the phase shift analysis for choosing permissible ambiguous soiutions, ale lllustrated in full measure by various versions of a more vivid example on the determination of the target position by the results of shooting. Suppose, for instan$c e$, that we are to determine the motion line $\eta$. ( $t$ ) of a point target, if the coordinates of hitting $y_{1}\left(t_{1}\right), y_{2}\left(t_{2}\right) \cdots y_{n}\left(t_{n}\right)$ are known for $n$ shots at a mov ing target made with the root-mean-square errors $\sigma_{i}$. Thls problem coincides in full measure with the discussed statistical problems on the phase shift analysis of experimental data on the nuclear particle interactions This example allows one to more vividly determine simple statistical questions not related to those of special complex problems on nuclear interaction.

We shall not violate the truth if for simplicity of narration in the above example of shooting we introduce some additional simplifications. Assume that an
immovable target is fired at and the root-mean-square error $\sigma$ is constant. Then instead of comparing the rellabilities of two motion lines of the target we have, respectively, the comparison of the rellabillites of two gtven positions $\eta_{1}$ and $\eta_{\text {II }}$ of a point target by the results of one given serles from equally accurate shots.

It is clear that we can easily calculate the probabilities $W_{1}$ and $W_{11}$ of obtaining just this series of results corresponding to the target positions at the points $\eta_{1}$ and $\eta_{\mathrm{II}}$. These probabilities in statistical assemblies of repeated serles on $\mathbf{n}$-shots performed separately on the first and second targets define the corresponding rates of appearing serles of results coinciding with the given one ( $y_{1}, y_{2} \cdots y_{n}$ ) within the selected limits $d y_{1}, d_{2}, \ldots d y_{n}$. It should be noted that here we do not mean the general coincidence of results of hitting the target with the given hittings but the coincidence of the results in strict consequence when the result of the first shot coincided just with the first result of the given series, etc.

The probabilities we are interested in
are easily treated in the case when the analyzed series of results is known to be a selection from the assembly in which series of results of shooting at both the targets are presented in equal measure. Indeed, in this case the value $P_{1}$ Is simply a probability that the given series of results was obtained in shooting at the first target. From $N$ cases of acquiring selections coinclding with the given one, $N P_{1}$ cases on average arose in shooting at the first target and $N_{11}$ cases in shooting at the second target. But despite its clarity; this application of the probabilities $P_{1}$ and $P_{11}$ is rather limited as it requires the a priori knowledge of probabilities of carrying out each hypothesis in a general statistical body.

Thus, the above $N P_{1}$ and $N P_{I I}$ rates of obtaining results coinciding with the given series are carried out in testing only in the case when a rifle-man
equally often shoots on average at both the targets (the case of equality of a priory probabilities). The equallty of a priorl probabllities ls not important since in the case of their difference one can always make a proper recalculation of the $P_{1}$ and $P_{I I}$ values. But one must know the values of a priori probablifties for the above application of $\because P_{I}$ and $P_{I I}$. When there is no informa tion on a priori probabilities this particular use of the $P_{I}$ and $P_{\text {II }}$ probabi-
lities is impossible in the framework of a specially made mbed statistical body. However, the possibilities of statistical estimate on the belonging of the given selection to one of the general assemblies is not changed. The authors of monograph /8/ stating that it is impossible to solve the problem on the statistical eveuation of hypotheses without the assumption on the a priori probablulty are absolutely not right.

In the case of interest when it is known that in nature only one hypothesis is brought about (a rifle-man shoots all the time only at one of the targets) one cannot in general complle a mixed statistical body of selections from various general assemblies. The problem of optimal statistical estimate whether a certain sèlection belongs to a definite assembly or not has, of course, a strict solution In this case. The degree of distingulshing between the two hypotheses depends not only uodn $\eta_{1}-\eta_{I I}$ (the distance between the targets) and the value $\sigma$ (the inaccuracy of shooting), but also upon an accidental result of the analyzed selection. Thus, in the case of hitting exactly the middle between the two targets ( $W_{I}=W_{I I}$ ) a shot does not carry ary additional information for distinquishing between the two hypotheses since with a $50 \%$ probability all the more it ls possible to simply guess the belonging of a selection to one of the general assemblies. When hitting moves off the central point the degree of possible distinguishing between the two hypotheses is increased. It is of interest that due to the square structure of the value. II when the point of hitting la moved off from the positions of both the targets, the degree of possible distinguishing between the two hypotheses is increased. Thus, exact hitting the centre of one of the targets does not provide a maximum rellability for selecting this target.

All the selctions from the general assembly in the example of choosing one of the hypotheses should be classified, separated in groups on the degree of possible distingulshing between the two hypotheses given by them. In this case one and the same group, one and the same statistical sub-assembly will include cases referring symmetrically to both the hypotheses. For example, one statistica sub-assembly will contain the cases of exact hitting any of the targets.

These sub-assemblies can be made up according to the given characteristic both when only one target is shot at and when two targets are shot at in turn without any assumptions on a priorl probabilities. The probabillties $P_{1}$ and $P_{11}$ obtained in accordance with relations (6) refer to such sub-assembles. Some value $\xi$. will be accidental in this problem on the statistical deter mination of a correct solution by one selection, it is equal to either 1 or 0 , if the chosen hypothesis turned out to be correct or false, respectively. The analo-
gous statement of the question is possible in the case of the infinite number of hypotheses continuously turning one into another. Just in this case the artificiall ty and the lack of logic of introducing the so-called reciprocal probabilities become understandable ${ }^{/ 12 /}$.

A statistically grounded reliabillty of the second solution $P_{I I}=\frac{1}{1+\exp \frac{M_{I I}-M_{I}}{2}}$
out to be considerably smaller than the reliability erraneously turns out to be considerably smaller than the reliability erraneously
determined by the $x^{2}$-distribution in the cases when the value $2(a-m)$ essentially exceeds 1. Thus, solutions found in the 310 MeV phase-shift analysis of pp-scattering, $1,2 /$ have in fact the following relabilities:

| $P_{1}(M=17.9)=0.80$ | $P_{g}(M-34.2)=0.002$ |
| :--- | :--- |
| $P_{11}(M=21.7)=0.12$ | $P_{6}(M-34.6)=0.002$ |
| $P_{8}(M=23.8)=0.045$ | $P_{7}(M-31.3)=2.10^{-5}$ |
| $P_{4}(M=24.5)=0.03$ | $P_{B}(M=52.3)=3.10^{-7}$. |

At the same time according to the $x^{2}$-criterion the first four solutions entered into the $90 \%$ reliability region; the 5 -th and the 6 - th solutions entered Into the $1 \%$ rellability region.

The incorrectness of a widely spread appllcation of the $x^{2}$-criterion for determining the reliability of ambiguous solutions of the phase shift analysis was cleared out by the author in discussing the paper by F.Lehar and V.V.Fyodorov 9/ in which planning of experiments for the determination of a single solution is calculated. I take an opportunity to gratify N.P. Klepikov and V.V. Fyodorov for this discussion.

It is worth noting that the incorrect logic of the error found coincides with the earlier mentioned fallubility of using the $x^{2}$-criterion $/ 10 /$ for deflning the permissible phase shift reglons in the analysis by the method of LM. Gelfand $11 /$.

One has to regret and wonder that this circumstance has been left without any attention and the criticism of paper $/ 11$ / has not been automatically propageted to the method accepted by everybody which determines the reliability of the ambiguous solutions of the phase shift analysis.

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