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Results on open charm from NA50

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Abstract

The NA50 and NA38 experiments have studied muon pair production in p–A and nucleus–nucleus collisions at the CERN SPS. Results on dimuons in the

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invariant mass region between the ϕ and the J/ψ masses (IMR) are presented. The standard sources of muon pairs in the IMR are the Drell–Yan process and the semileptonic decay of charmed mesons. A new four-dimensional unfolding method is applied to the data. The p–A results are well described in terms of these known sources, which, in contrast, fail to explain nucleus–nucleus data: an extra yield of muon pairs is observed. Although this excess is compatible with an enhanced production of charmed hadrons, other possible sources cannot be excluded. In particular, a comparison between data and predictions from a theoretical model which calculates the production of thermal dimuons is presented in this paper.

1. Introduction

The invariant mass distribution of dimuons $(M_{\mu\mu})$ produced in hadronic interactions at SPS energies is characterized by a continuous spectrum and by peaks from two-body decays of vector mesons ($\rho, \omega, \phi, J/\psi$ and ψ'). The study of the non-resonant part of the spectrum leads to several experimental challenges: the cross section at high mass is very low and muon pairs are produced by the overlap of various sources. Moreover, the major source of muon pairs at low masses ($M_{\mu\mu}$ < 2 GeV c^{-2}) is the combinatorial background from charged pion and kaon decays. In this paper an analysis of dimuons in the mass interval between the ϕ and the J/ψ masses is presented. The study is performed on data collected by the NA50 and NA38 collaborations at the CERN SPS, namely p-A collisions at 450 GeV, S-U (NA38) at 200 A GeV and Pb–Pb at 158 A GeV. The relevance of this intermediate-mass region (IMR) arises from the fact that thermal radiation from early and hot stages of the collision could give a detectable signal in the form of muon pairs of relatively low mass. In the IMR the contribution of low-mass resonances is negligible and the other known sources are hard processes for which an evaluation is possible. An excess of dimuons in the IMR for nucleus-nucleus collisions with respect to a linear extrapolation of the standard sources in proton-induced collisions has already been demonstrated by a previous analysis of these data [1]. Moreover, the kinematical distributions of the observed excess turned out to be compatible with those expected from semileptonic decays of charmed mesons. An excess in the IMR was also found in S-W collisions by the Helios-3 experiment [2].

The analysis presented here differs from the previous one in the unfolding method. The unfolding, i.e. the procedure used to go from measured quantities to physical ones, was done separately for each kinematical variable ($M_{\mu\mu}$, $p_T Y_{CM}$ and²⁰ cos Θ_{CS}). In the present analysis a four-dimensional approach [4] allowing for correlations among the kinematical variables is adopted. In the following, we will focus on the peculiarities of this analysis. The description of the apparatus and the data reduction together with the discussion of the charge correlation effects on the normalization of the combinatorial background can be found in [1].

2. The analysis

Measured distributions are affected by detector effects such as acceptance and resolution. In more quantitative terms, let us define x as the set of kinematical variables used to describe a

 $^{^{20} \}Theta_{CS}$ is the polar angle of one of the muons in a particular rest frame of the muon pair, known as the Collins–Soper reference frame [3].

dimuon and $\Phi(x)$ as the physical distribution that has to be determined experimentally. The measurement is a finite sampling of the distribution *D* defined as

$$D(x') = \int S(x'|x)A(x)\Phi(x) \,\mathrm{d}x \tag{1}$$

where A(x) is the acceptance and S(x'|x) is the smearing function. S(x'|x) dx' is the probability that a dimuon characterized by x is mapped by the apparatus in an elementary cell of size dx' centred at x'. The unfolding procedure consists in the resolution of the integral equation (1). To solve equation (1), both the smearing and the acceptance must be known. The solution of equation (1) is based on a method proposed by Richardson [5] and, independently, by Lucy [6]. Equation (1) can be rewritten as

$$D(x') = \int S(x'|x)A(x)\Phi(x) \, dx = \int S(x'|x)\Psi(x) \, dx$$
(2)

with $\Psi(x) = A(x)\Phi(x)$. The *inverse* conditional probability density Q(x|x') dx can be related to S(x'|x) through the Bayes theorem:

$$Q(x|x') = \Psi(x) \frac{S(x'|x)}{D(x')} \implies \Psi(x) = \int D(x')Q(x|x') \,\mathrm{d}x'. \tag{3}$$

The determination of the physical distribution Ψ can be achieved iteratively. At the *r* th iteration, the physical distribution is approximated by $\Psi_r(x)$. The next step is computed as follows:

$$\Psi_{r+1}(x) = \int D_0(x) Q_r(x|x') \, \mathrm{d}x' = \int D_0(x) \frac{\Psi_r(x) S(x'|x)}{D_r(x')} \, \mathrm{d}x$$

where

$$D_r(x') = \int \Psi_r(x) S(x'|x) \,\mathrm{d}x$$

and $D_0(x') \equiv D(x')$ is the measured distribution. To speed up the procedure of convergence, the starting value $\Psi_0(x)$ is chosen to be the measured distribution. This procedure leaves the normalization of the distributions unchanged at each iteration.

The number of iterations needed to find the physical distribution has to be determined with care, as described in [4], since noise in the distributions appears when the number of steps is too large. For mass distributions, the width of the J/ψ meson decreases with the number of steps in the procedure and flattens when the number of iterations is ~ 50. In order to avoid amplifications of the fluctuations, this number of steps has not been exceeded. The smearing function and the acceptance are computed by means of a detailed Monte Carlo simulation as a function of the four kinematical variables. The computation is done on a four-dimensional grid, hence the integrals in the iterative procedure are replaced by discrete sums.

The computation of the errors by means of the iterative method is practically impossible given the high number of elements of the matrices involved ($\sim 10^{13}$). Therefore, the errors of the data, corrected for the acceptance, are used. This assumption has been checked with a complete one-dimensional error matrix calculation for the mass spectrum.

The combinatorial background from π and K meson decays is more than 85% of the total $\mu^+\mu^-$ yield. The number of background muon pairs is calculated starting from the number of like-sign pairs as

$$N_{bkg}^{+-} = 2R\sqrt{N^{++}N^{--}} \tag{4}$$

where *R* expresses the effects of charge correlations and is equal to 1 in the limit of high multiplicities. Its proper value has been determined as described in [1]. In a four-dimensional (4D) analysis, however, detected like-sign pairs are not enough, as many cells of the 4D grid are empty. As a solution, equation (4) is still used to normalize the integrated $\mu^+\mu^-$ background, but the 4D grid is filled by means of the so-called fake opposite sign method (FOS). An exhaustive combination among muons coming from different like-sign pairs is used to increase the population of the grid. The deconvolution is applied to data after the background subtraction.

The kinematical region used for this analysis has been chosen in order to have an acceptance which is relatively high for each cell of the grid ($\ge 5\%$):

$$M_{\mu\mu} \ge 1.6 \text{ GeV } c^{-2}$$
 $0.2 \le Y_{CM} \le 0.8$ $|\cos \Theta_{CS}| \le 0.3.$

For p–A data $-0.2 \leq Y_{CM} \leq 0.4$. The standard sources contributing to the dimuon production in the IMR are the Drell–Yan process (*DY*) and the semileptonic decays of pairs of charmed hadrons ($D\bar{D}$). In the selected kinematical region, the contribution of the tails of the low-mass resonances is negligible. The mass spectra have been fitted in the range $1.6 \leq M_{\mu\mu} \leq 8$ GeV c^{-2} , assuming a Gaussian shape for the J/ψ and ψ' mesons and taking the shapes of the mass distributions for *DY* and $D\bar{D}$ processes from the Pythia 6.1 generator [7], with the MRS A set of parton distribution functions [8], $m_c = 1.5$ GeV c^{-2} , $\sigma_{k_T}^{DY} = 0.8$ GeV c^{-1} and $\sigma_{k_T}^{D\bar{D}} = 1.0$ GeV c^{-2} . The normalizations of the various contributions have been determined from the fit.

3. Results

The four p–A samples (where A denotes Al, Cu, Ag and W) were fitted together, imposing the same ratio $D\bar{D}/DY$, since, as hard processes, both DY and open charm are expected to scale linearly with the target mass. The ratio between the semileptonic decays of charmed particles and the DY process in the IMR ($1.6 \le M_{\mu\mu} \le 2.5 \text{ GeV } c^{-2}$) has been found to be (statistical errors only):

$$\left(\frac{DD}{DY}\right)_{\rm p-A, \, 450} = 4.13 \pm 0.15.$$

From the measured cross section for the DY process, the total open charm production cross section in p-p collisions can be estimated indirectly. Our result is compared with direct measurements in figure 1.

The S–U and Pb–Pb data samples have been subdivided, respectively, into five and seven centrality bins defined in terms of the measured electromagnetic transverse energy. In order to compare the experimental mass distributions with the predicted ones, it is necessary to determine an expected value for the open charm cross section in nucleus–nucleus collisions. To this purpose the p–A cross section was extrapolated to nucleus–nucleus collisions assuming a linear dependence on the colliding nuclei masses and using the \sqrt{s} dependence given by Pythia. This extrapolation, together with the measurement of the number of *DY* events from the high mass data (where *DY* is the only known source) allowed us to determine both the expected ratio $(D\bar{D}/DY)$ and the number of $D\bar{D}$ events for each centrality bin.

The total measured dimuon yield compared with the expected one in the IMR is shown in figure 2. There is clearly an excess of dimuons in the IMR that increases monotonically with

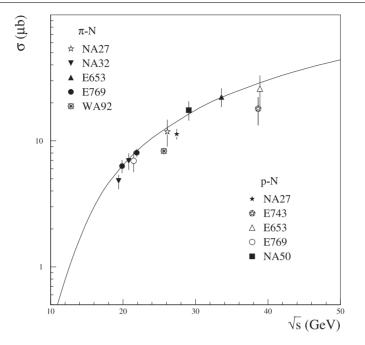


Figure 1. Compilation of charm hadro-production cross section measurements (forward production $x_F > 0$) [1]. The results from nuclear targets have been divided by the target mass number. The \sqrt{s} dependence of σ_{cc}^{pp} obtained with PYTHIA is also shown. The normalization of the curve is given by a fit to the experimental points.

the number of participants in the collision, reaching a value of the order of two for central Pb–Pb collisions. The observed excess cannot be ascribed to a bad evaluation of the combinatorial background, since the shape of the data cannot be reproduced by varying the normalization of the background as discussed in detail in [1].

Mass spectra can be fitted well if the normalization of the open charm contribution is taken as a free parameter of the fit. Moreover, also the p_T spectra can be well reproduced by a combination of *DY* and $D\bar{D}$ provided that the latter is left free to fit the data. The double ratio

$$E = \frac{D\bar{D}/DY\big|_{meas}}{D\bar{D}/DY\big|_{expected}}$$
(5)

can be taken as a measure of the excess. The charm enhancement needed to reproduce data is of the order of three for central Pb–Pb events.

The fact that data can be understood in terms of an extra production of open charm with respect to a linear extrapolation from the p–A collision does not imply necessarily that the excess is actually due to this source. Other possible phenomena that could lead to this *charm-like* excess in the IMR have been proposed. Spieles *et al* [9] evaluated the role of secondary *DY* processes in hadronic rescatterings by means of the UrQMD model and found a ~10% enhancement of the primordial *DY* for $M_{\mu\mu} \simeq 2$ GeV c^{-2} .

Lin and Wang [10] investigated the role of final-state rescattering of charmed mesons, which broadens their m_T spectra leading to an enriched $\mu^+\mu^-$ yield in the acceptance of the NA50 spectrometer. The degree of thermalization is parametrized in terms of the temperature T of the environment in which D mesons thermalize. This temperature is a local parameter

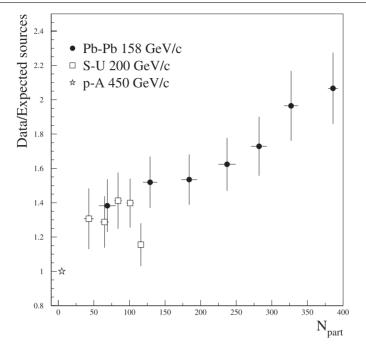


Figure 2. The ratio D/E (data/expected sources) computed in the mass interval $1.6 \le M_{\mu\mu} \le 2.5$ GeV c^{-2} is plotted as a function of the number of participant nucleons (N_{part}).

and should not be confused with the inverse slope parameter T_{eff} of m_T distributions. The open charm enhancement *E* is related by the model to the temperature T (E(T = 0) = 1) so it is possible to assign a temperature to each centrality bin and to calculate the $M_{\mu\mu}$ distribution for dimuons originating from $D\bar{D}$ decays, taking into account the rescattering effects. In figure 3(*a*) the IMR mass spectrum for central Pb–Pb collision is fitted with DY and $D\bar{D}$ contributions, with the latter taken as a free parameter, while in figure 3(*b*) the open charm contribution is evaluated with the rescattering model. The *D* meson rescattering is not sufficient to account for the observed excess.

Theoretical work on thermal dilepton radiation from an expanding fireball is in progress [11–13]. In figure 4 a comparison between central Pb–Pb data and the predictions of the thermal model from Rapp and Shuryak [13] is presented. The model is based on the assumption that the fireball formed in a Pb–Pb collision is in local thermal equilibrium. The dimuon yield is evaluated starting from the production rate that in the IMR is given by the perturbative quantum chromodynamics (QCD) result for the $q\bar{q}$ annihilation process for both the quark–gluon plasma (QGP) initial phase and the subsequent hadron gas phase. This rate has to be integrated over the spacetime evolution of the collision. Here a simplified thermal fireball model that includes the most relevant features of a complete hydrodynamical simulation is used. The parameters applied for the comparison are the following: initial temperature $T_i = 192$ MeV, fireball lifetime equal to 14 fm c^{-1} and critical temperature for the QGP $T_c = 175$ MeV. The impact parameter for the Pb–Pb collision is $b \leq 1$ fm, so the comparison is made with the most central bin of the data sample. The thermal contribution rapidly drops when $M_{\mu\mu}$ increases and it is completely negligible at the J/ψ mass. The agreement with data is rather good, even though the comparison is limited to a single centrality bin and the p_{T} analysis is still in progress.

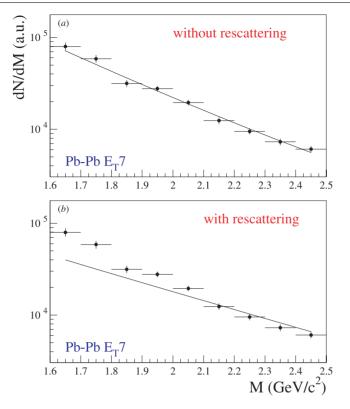


Figure 3. (*a*) The line represents a fit to the data with the $D\bar{D}$ contribution taken as a free parameter. The good quality of the fit is obtained with an open charm excess that is up to a factor of three for central Pb–Pb collisions. The excess is expressed by equation (5); (*b*) the line represents the dimuon yield due to DY and $D\bar{D}$, where the latter is computed with the rescattering model [10].

(This figure is in colour only in the electronic version, see www.iop.org)

4. Conclusions

The use of a new method for the data unfolding leads to results that are in quantitative agreement with those already published [1]. The advantages of the method consist in the fact that it accounts for correlations among kinematical variables and does not imply the assumption of specific shapes for their distributions. On the other hand, the method can be applied only to a relatively restricted kinematical region with high acceptance. Moreover, local small distortions of the distribution appear near the narrow J/ψ peak. This is a known drawback of the unfolding method (the so-called Gibbs ring effect): data points affected by this problem have been replaced by dots in figure 4.

The p–A dimuon yield in the IMR has been described as a combination of semileptonic decays of charmed hadrons and Drell–Yan. An indirect measurement for charm hadroproduction has been performed and the result is in good agreement with direct measurements. For nucleus–nucleus collisions, the dimuon yield is underestimated by a linear extrapolation of the processes that describe p–A data. The excess cannot be explained as a bad evaluation of the background and a local open charm enhancement due to the rescattering of charmed hadrons is insufficient to account for the measured yield. Mass and p_T distributions are consistent with

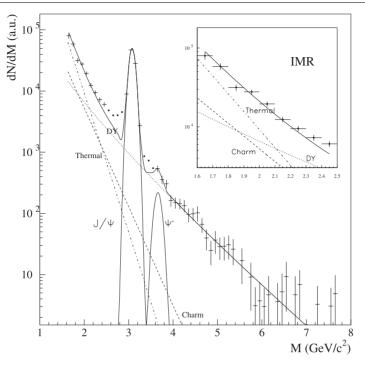


Figure 4. Central Pb–Pb mass spectrum with the estimated contributions. The full curve is the sum of all the contributions.

a combination of DY and open charm semileptonic decays, with a ratio $(D\overline{D})/(DY)$ up to three times higher than the expected value. The explicit introduction of thermal dimuons leads to results that are compatible with the data, even if a more systematic comparison is needed.

These results suggest that only a direct measurement of the open charm production in nucleus–nucleus collisions at SPS energies would allow one to draw final conclusions on the nature of the dimuon production in the intermediate-mass region.

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