Study of Clusters and Hypernuclei production within PHSD+FRIGA model at the NICA energies

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I - Introduction

Experimental data shows that at 3 A.GeV even in central collisions almost 20% of the baryons are bound in the clusters [1]. If we do not describe the dynamical formation of fragments we cannot properly describe the nucleon observables like v1, v2, transverse momentum spectra, and we cannot explore the new physics opportunities like hyper-nucleus formation, 1-st order phase transition, fragment formation at midrapidity (RHIC, LHC). Modelling the clusters formation is a complicated problem, and therefore many present microscopic approaches fail to describe fragments at NICA/FAIR (and higher) energies.

II – PHQMD+FRIGA model

In our studies we are using the *"Fragment Recognition In General Application"* (FRIGA)[2], an improved version of SACA, within the *Parton-Hadron Quantum Molecular Dynamics* (PHQMD) model, which is is a non equilibrium microscopic transport model that describes n-body dynamics based on QMD [3] propagation with collision integrals from **PHSD** [4].

The FRIGA algorithm consists of the following steps:

1) the algorithm takes the positions and momenta of all nucleons at time t to determine clusters within a phase space coalescence approach using the Minimum Spanning Tree technique (MST).

2) the MST clusters and individual particles are recombined in all possible ways into fragments or left as single nucleons, such as to choose that configuration which has the highest binding energy. This procedure is repeated many times (within a Metropolis procedure) and it automatically leads to the most bound configuration.





III – Model predictions



actively under development and soon we expect new results.

VII – References

W. Reisdorf et al., Nucl. Phys. A 846 (2010) 366-427 [1] A. Le Fèvre et al., J. Phys.: Conf. Ser. 668 (2016) 012021. [2] J. Aichelin and H. Stöcker, Phys. Lett. B 176 (1988) 14 [3] [4] E.L. Bratkovskaya and W. Cassing, Nucl.Phys. A 856 (2011) 162-182. [5] Courtesy of the ALADIN Collaboration (new S254 data).