

Isoscaling in central Sn + Sn collisions at 270 MeV/nucleon[†]

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Experimental study on the early stage of heavy-ion collision is challenging as excited fragments produced from the collision decay into lighter particles before they are detected. The nuclear isoscaling phenomenon is a useful tool as the ratio of yields from two different reactions is weakly affected by the fragment de-excitation process.¹⁾

Rare isotope Tin beams ¹³²Sn and ¹⁰⁸Sn were produced from RIBF and impinged onto the isotopically enriched Tin targets. Hydrogen and helium isotopes were detected in S π RIT time projection chamber²⁾ placed inside the SAMURAI dipole magnet.³⁾ Particles were identified from the magnetic rigidity and mean energy loss.⁴⁾ Most central events with impact parameter $b < 1.5$ fm and mid rapidity range $y_0 = 0-0.4$ are chosen for this analysis.

The yield ratios between two systems $R_{21} = Y(^{132}\text{Sn} + ^{124}\text{Sn})/Y(^{108}\text{Sn} + ^{112}\text{Sn})$ as a function of p_T/A are shown in Fig. 1. Given that the collision systems are thermally equilibrated, R_{21} follow the isoscaling law $R_{21} = C \exp(\alpha N + \beta Z)$ where α and β are the fit parameters. Empirically, α and β have similar values with opposite signs, therefore, particles with the same $(N-Z)$ value show similar R_{21} values. Figure 1 show isoscaling effect for $p_T < 280$ MeV/c (left side of vertical dashed line). In this region, the isoscaling fit gives $\alpha = 0.29$ and

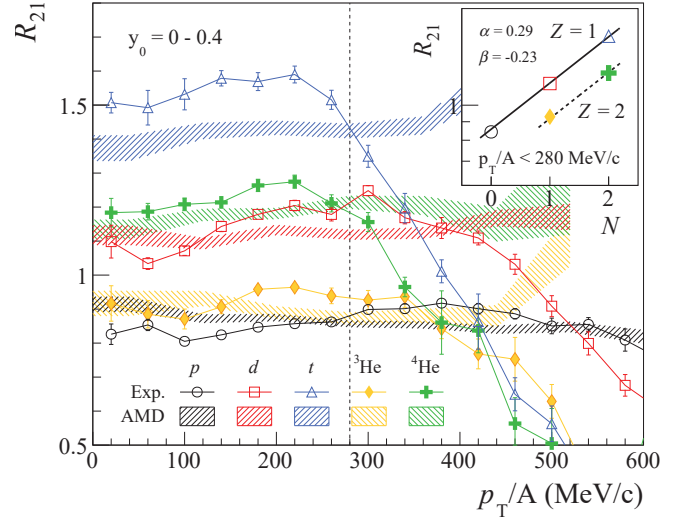


Fig. 1. The yield ratio R_{21} between two systems ¹³²Sn + ¹²⁴Sn and ¹⁰⁸Sn + ¹¹²Sn. Experimental data (markers) are compared with AMD (shaded area). Inner panel: Isoscaling fit for $p_T/A < 280$ MeV/c.

$\beta = -0.23$ (inner panel of Fig. 1). On the other hand, the triton and helium R_{21} values breakdown above this limit, and the isoscaling phenomenon vanishes. This suggests that high p_T/A particles come from the non-equilibrated environment.

The antisymmetrized molecular dynamics (AMD) model^{5,6)} is employed with Skyrme SLy4 effective interaction and symmetry energy slope parameter $L = 46$ MeV. The AMD result qualitatively explains isoscaling for $p_T/A < 280$ even though AMD is a dynamical model. However, AMD underestimate triton R_{21} and do not reproduce breakdown of R_{21} .

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[†] Condensed from the article in Eur. Phys. J. A **58**, 201 (2022)

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