Probing formation time in p+p at LHC via HBT¹

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Motivation

- Hanbury-Brown Twiss (HBT) correlations provide a unique tool to extract the freeze-out configuration in high energetic collisions
- Bulk behaviour like radial flow can be tested by identical particle HBT correlations
 The correlations provide insight into the timescales of the string breakup

Correlation Function

We use the formula

$$C(\vec{q},\vec{K}) = 1 + \int d^4x \cos(q \cdot x) d(x,K)$$

Femptoscopic Correlations



Figure: Demonstrative picture [2] for HBT results in the Out-Side-Long System.

Region of Homogeneity



 Pairs analyzed in the Out-Side-Long system

 $C = 1 + \frac{\left|\int S(x, K)e^{i \cdot q \cdot x}d^{4}x\right|^{2}}{\left|\int S(x, K)d^{4}x\right|}$ $= 1 + \lambda \cdot \exp(-\sum_{i,j} R_{ij}^{2}q_{i}q_{j})$

to generate the 3D correlation function, where q and x are momentum and spacial distance of pions in a pair, \vec{x} and \vec{q} are the corresponding 3 vectors and d(x,K) is the separation distribution of the pairs.



Figure: Projections of an example correlation function for $dN_{ch}/d\eta = 23-29$ for $K_{\perp}=0.3-0.4$ GeV and the corresponding fit.

KT



Figure: Probed regions for different K_{\perp} intervals [3].

HBT correlation analysis is only sensitive to close momentum pairs
Radial flow leads to close momentum pairs being close in space (regions of homogeneity)

• Regions of homogeneity lead to pair momentum (K_{\perp}) dependence in the HBT radii

Model

We use the UrQMD = Ultra-relativistic quantum molecular dynamics model [4] to generate particle freezeout distributions. UrQMD features:

HBT for p+p at $\sqrt{s} = 7$ **TeV**



Figure: UrQMD results [1] are the lines with formation eigentimes of 0.3 fm/c (dashed), 0.8 fm/c (full - default) and 2 fm/c (dotted). The Symbols represent ALICE data [5] for different multiplicity classes.

Results & Outlook

Nonequilibrium transport model

- All hadrons and resonances up to 2.2 GeV included
- Full space-time dynamics of hadrons from beginning to end
 String excitation and fragmentation

String fragmentation

- Particle creation via string breakup takes time
- This time is often referred to as formation time
- So far, there is no measurement of the formation time
- The UrQMD model uses formation time as a parameter

- HBT measurements in p+p collisions allow to probe formation times
- UrQMD shows next to no multiplicity scaling in R_{out} and R_{long} and weak multiplicity scaling in R_{side}
- Constraints for a more refined treatment of formation times can be extracted from p+p HBT

References

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