JAM3D

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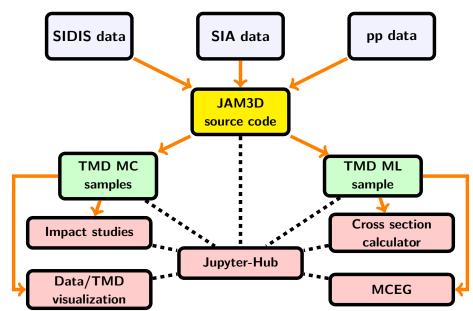




Project Scope

- TMD fits
 - + single fits → Maximun Likelihood (ML)
 - + MC fits → Nested Sampling (NS)
- Impact studies for future measurements
 - + Reweighting MC samples with new pseudo data sets
- Data visualization
 - + Dedicated data vs theory visualization
 - + TMD plotter
- Cross section simulation
 - + Evaluation of cross sections
 - + Single particle MCEG

Road Map



TMD fitter

Features

- + Fast evaluation of residuals \rightarrow powered by dedicated parallelization scripts (based on ZMQ)
- + Parallelization can take advantage of cluster environments:
 - JLab HPCs
 - Amazon web services (AWS): EC2, ECS via docker images
- + Modularized framework. Easy to incorporate new observables

Methodologies

- + Maximun likelhood analysis (ML)
 - Terminal based ("input.py") + jupyter-notebooks
 - Also via jupyter-notebook (useful for jupyterhub enviroments)
- + MC analysis based on Nested Sampling (NS)
 - Terminal based ("input.py")
 - o Ideal to run on a cluster enviroment

Impact studies for future measurements

Methodology

- + Simulation of new observables based on existing MC samples
- + Bayesian reweighting → fastest implementation

MC samples repository

- + Dedicated TMD MC parameter samples generated from existing data
- + Access to the input files for the MC samples in case new samples are need within a different setup (change in parametrization, TMD theory, etc.)
- + Dedicated jupyer-notebooks templates for simulation and impact studies

Jupyter-hub frontend

- + No local software installation
- + Dedicated jupyer-hub server packed with JAM3D + MC samples repositories
- + Users needs to upload jupyter-notebooks from from repository

Data visualization

Gallery of TMD studies

- + Dedicated repo for jupyter-notebooks to display:
 - o data vs theory
 - o 2D, 3D TMDs plots (nucleon imaging)

Jupyter-hub frontend

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Cross section simulation

Features

- + Dedicated jupyter-notebooks for cross section evaluation
- + Dedicated jupyter-notebooks for single particle event generator

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