

Hello Stefano,

I am checking the pandaroot trunk rev-25813. The goal is to perform the scan of Ds(2317) mass, using the full simulation.

Some months ago I had used the trunk rev-23845: results were presented at the collaboration meeting in March, this year. In those first tests, the Ds mass resolution obtained, using standard tutorial macros as in /tutorial/rho/*, was $\sim 16 \text{ MeV}/c^2$. Later on, I even performed comparison between fast- and full simulation tools, in the pandaroot rel-scrut14. Reconstruction efficiency, with $p(\gamma) > 50 \text{ MeV}/c$, was roughly 35% in both cases, if I remember correctly.

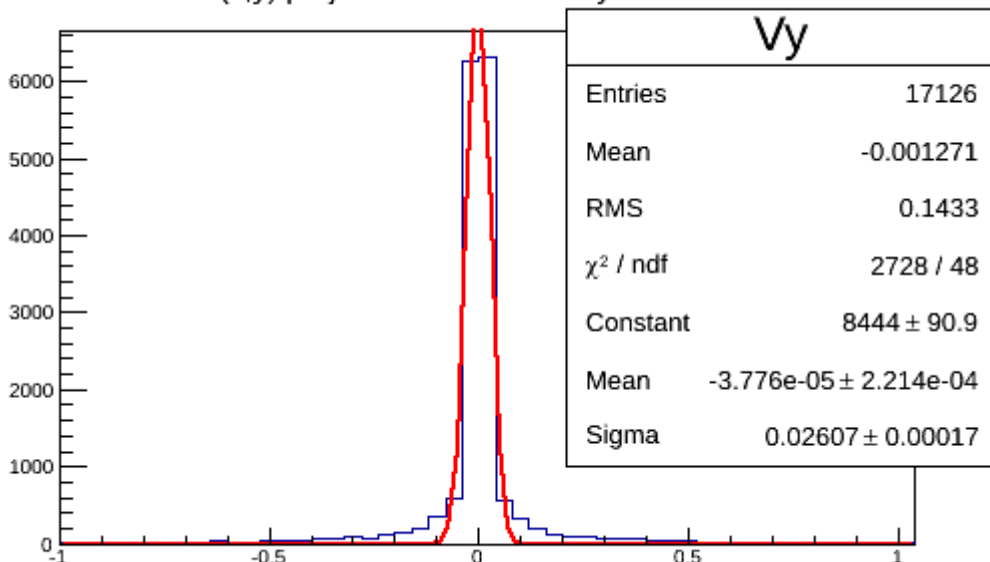
I've made use of the same analysis macro and I have repeated the tests in the new trunk rev-25813. Efficiency is slightly different: $\sim 30\%$, with same basic selection cuts; and the mass resolution becomes worse: $\sim 19 \text{ MeV}/c^2$. Also the vertex resolution is worse in this trunk. Is this something expected, due to the changes in the framework, or is it the signal of something that went wrong, in this new trunk revision? Except the changes in the GEM geometry, and the consequent update of the reco files, I am not warned of other additional changes.

Here attached are some plots. The basic test is performed running on 3000 generated events, EvtGen, $p = 8.8 \text{ GeV}/c$, for the process $p \bar{p} \rightarrow Ds^- Ds(2317)^+, Ds(2317)^+ \rightarrow Ds \pi^0, \pi^0 \rightarrow \gamma\gamma, Ds \rightarrow K^+ K^- \pi$.

cheers, Elisabetta

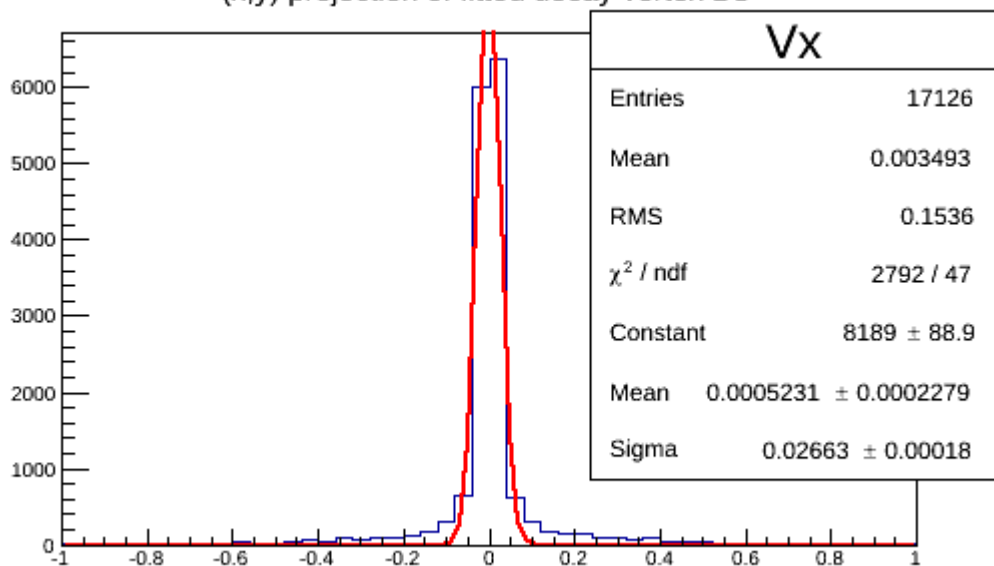
File Attachments

- 1) [Vy_checkRev25813.gif](#), downloaded 388 times
(x,y) projection of fitted decay vertex Ds-



- 2) [Vx_checkRev25813.gif](#), downloaded 383 times

(x,y) projection of fitted decay vertex Ds-



3) [Dsmissingmass_checkRev25813.gif](#), downloaded 394 times

