
Subject: Re: PndVtxPRG vs PndKinVtxFitter

Posted by [Ralf Kliemt](#) on Fri, 24 Oct 2014 14:31:38 GMT

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Hi Karin,

There is a potential bug (marked in red). Simply use bool checkb=vtxfitterlb.Fit(); as the fit statement in the beginning of the block.

Karin Schöning wrote on Fri, 24 October 2014 11:44 for (j=0;j<lamb.GetLength();++)

{

PndVtxPRG vtxfitterlb(lamb[j]);

```
vtxfitterlb.Fit();
double chi2_vtx = vtxfitterlb.GetChi2(); // access chi2 of fit
double prob_vtx = vtxfitterlb.GetProb(); // access probability of fit
h0b_chi2_vf->Fill(chi2_vtx);
hlamb_prob_vf->Fill(prob_vtx);
bool checkb=vtxfitterlb.Fit();
if(checkb)
    // when good enough, fill some histos
{
```

RhoCandidate *lambv = lamb[j]->GetFit(); // access the fitted cand

Do you apply other selections, such as a distance cut to the interaction point? The PndVtxPrg should be just a touch less accurate than the PndKinVtxFitter. I expect no different efficiency drops by fitting.

Cheers

Ralf
