LCFIPlus for 6-jets@500GeV



sw/ana meeting, 4th Apr. 2018

Concerns

LCFIPIus

Is LCFIPlus software (more focusing on vertexing for now) working well at least at the same level as in DBD era?

* Samples

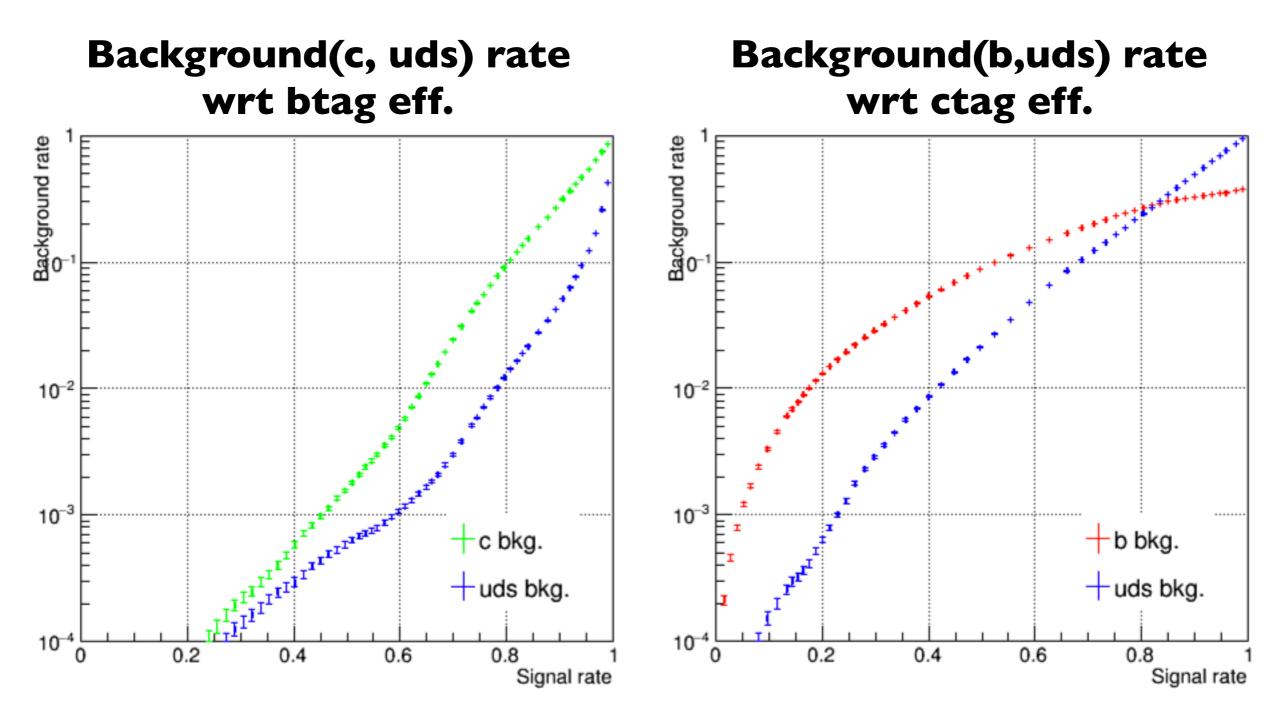
Is there any problem in sample production?

Detector models (Next step)

- Comparison between I5 and s5.
- ▶ I will not discuss in this presentation.

FlavourTagging

(Example plot just for explanation for later slides)



3

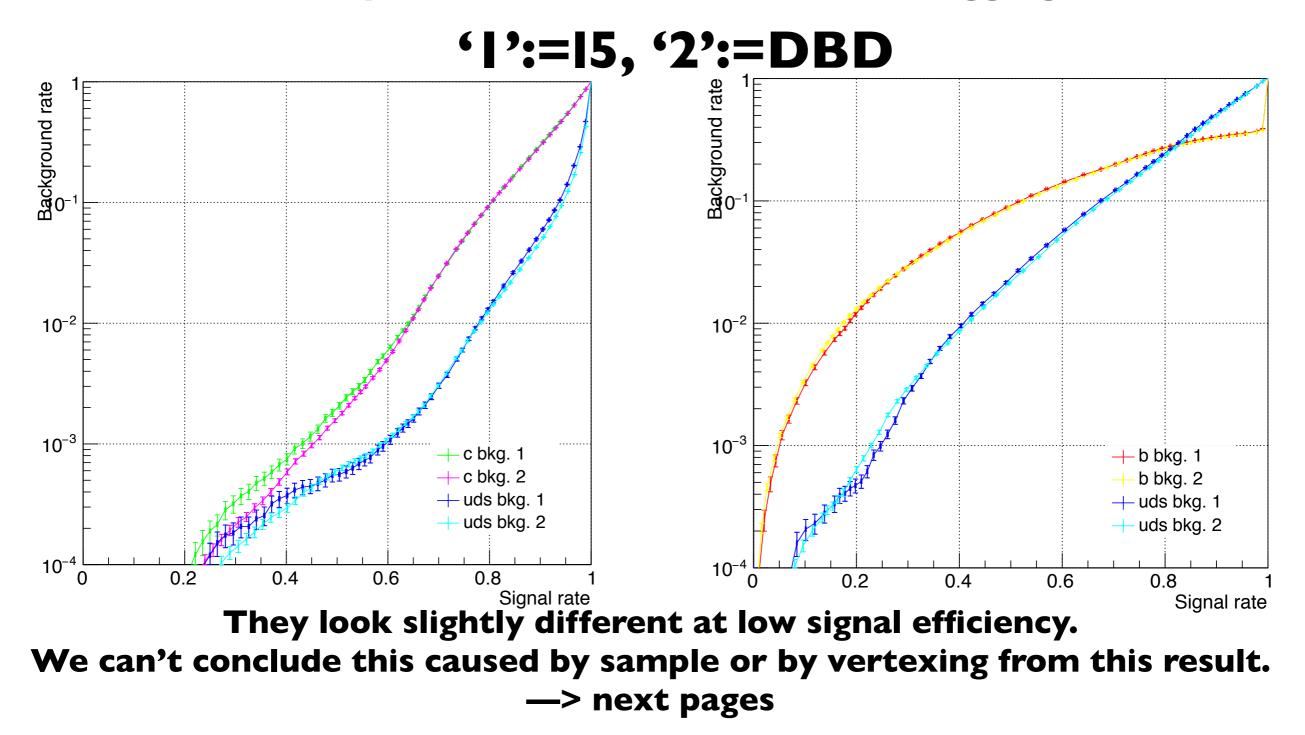
different sample, different version of vertexing

sample: |) Test production sample (last Dec.)2) DBD sample (2012?)

Both are $\sqrt{s=91.2GeV}$, bb, cc, qq samples

Software: 1) latest vertexing, 2) DBD vertexing,

Both are processed with latest flavour tagging.



vertexing comparison

sample : DBD sample, $\sqrt{s}=91.2$ GeV, bb, cc, qq samples for both.

Software: 1) latest vertexing, 2) DBD vertexing, Both are processed with latest flavour tagging.

> Unfortunately one of output files I got today was somehow broken. I couldn't investigate the cause in detail.

the error message :

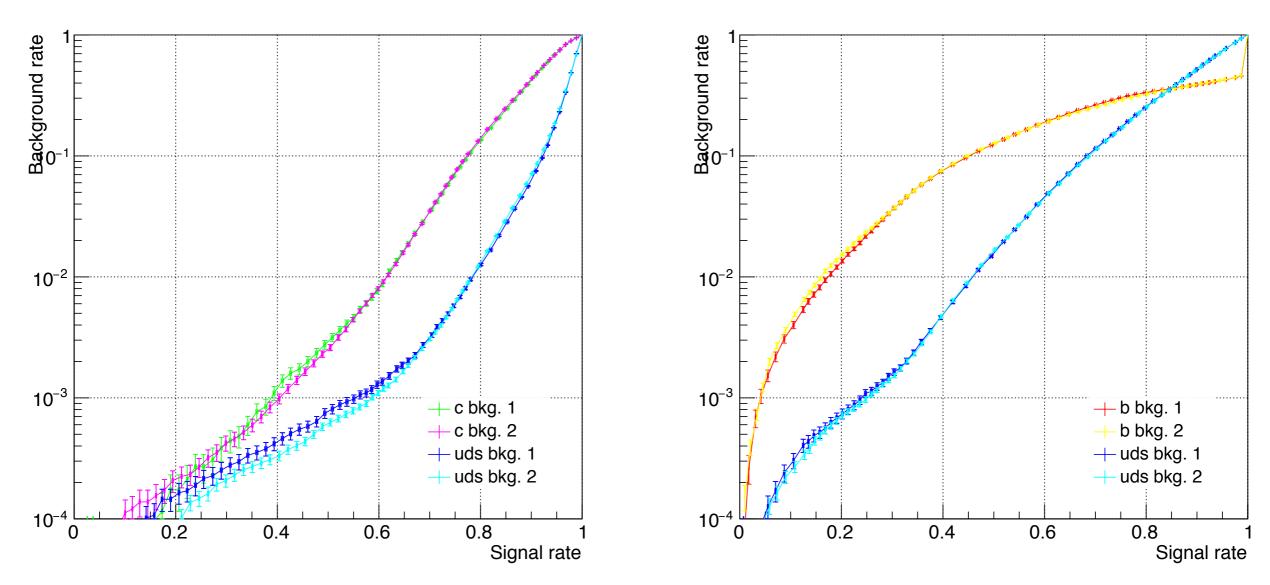
Error in <TFile::ReadBuffer>: error reading all requested bytes from file Training_DBD_newvtx/lcfiweights/ qq91_v03_p01_c1.root, got 260 of 300 Error in <TFile::Init>: Training_DBD_newvtx/lcfiweights/qq91_v03_p01_c1.root failed to read the file type data.

> (Please let me know if anyone knows how to fix this kind of problem.)

vertexing comparison

sample : DBD sample, √s=500GeV, 6b, 6c, 6q (6u+6d+6s) samples. <u>No beam bkg.</u>

Software: 1) latest vertexing, 2) DBD vertexing, Both are processed with latest flavour tagging.



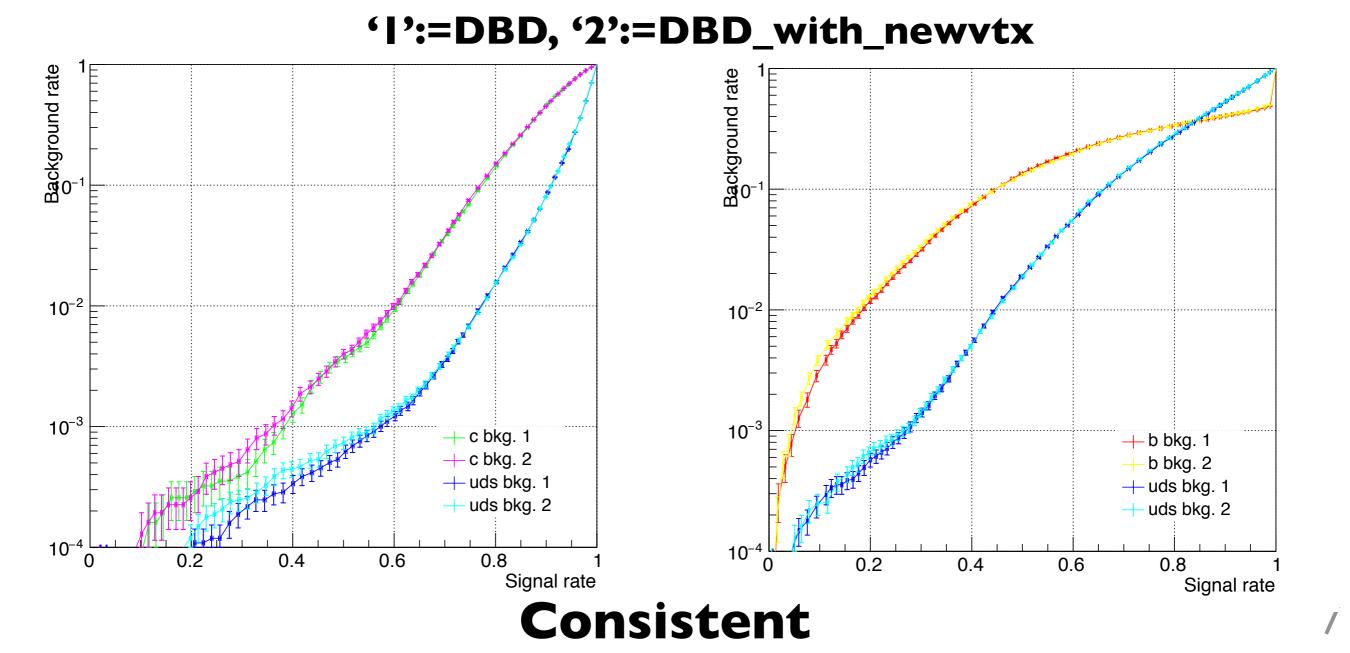
'I':=DBD, '2':=DBD_with_newvtx

Consistent

vertexing comparison

sample : DBD sample, √s=500GeV, 6b, 6c, 6q (6u+6d+6s) samples. Beam bkg. included

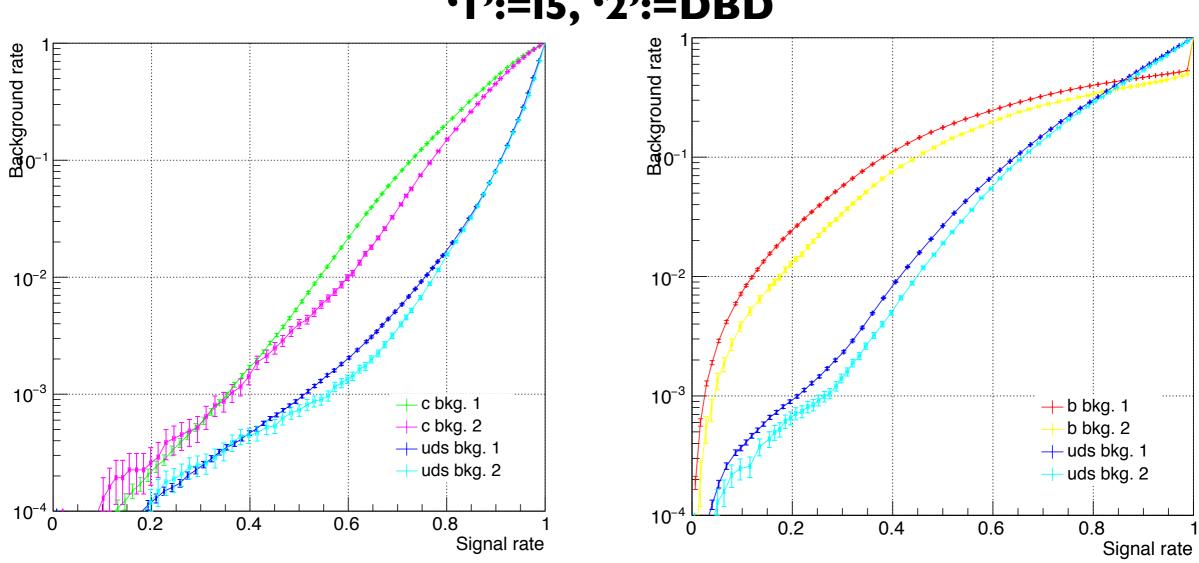
Software: 1) latest vertexing, 2) DBD vertexing, Both are processed with latest flavour tagging.



sample comparison

sample : Test production I5 sample (last Dec.) **2) DBD sample (2012?)**

Software: v01-19-05 (vertexing, jet clustering, flavour tagging)



'I':=I5, '2':=DBD

I'm confused with these new plots... Why larger differece than p4?? Any reasonable explanation? Need to understand.

My tentative conclusion

* According to the results I have,

- Looks no crucial problem on vertexing itself from p6 and p7.
- Need to understand the difference between DBD sample and the test samples from p8 to be more confident.

* Any suggestions are welcome!