Dear Klaus,

I think your argument that there is no perfect MC match is not right. For this purpose I have written the FairLinks which are working extremly well for comparison between reconstructed data and MC data. Maybe it is time to use them for the analysis as well.

For the rest of the discussion I see a discrepancy between what the designers of the code think their code is doing and the expectation of the users what the code should do according to the used names. I fear that the problem Simone and Martin had will happen to other users as well.

In addition I think there should be a method for an ideal list filling.

Cheers,

Tobias