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Subject: Re: Moving to Genfit2

Posted by [Elisabetta Prencipe \(2\)](#) on Sat, 14 Feb 2015 16:49:07 GMT

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Hello Alicia,

I attach here some files, as you asked: config.sh is from my genfit\_branch/build; the CMakeList.txt is from my genfit\_branch. CMakeList.txt\_genfitAlicia is that from the package /genfit/ inside the branch. You will see, it is different from the CMakeList.txt of the standard pandaroot release. In the new branch, 'genfit' is the name of the folder; but genfit2 is the package there in. I just preserved the same name: the content is different from the other pandaroot release/trunk revisions. The file CMakeList.txt\_branchAlicia comes from the branch.

I think that something went wrong when you uploaded

a) apr13 external package in you home

or

b) the intallation of the new branch somehow did not succeeded.

My easy suggestion is to delete them, and start all process from scratch. In other words:

1) cd /your\_home/<...>/

2) svn co <https://subversion.gsi.de/fairroot/fairsoft/release/apr13>

and you get installed again the apr13 external packages;

3) follow instructions as they are addressed to you, once you do 2)

4) install the genfit2 branch as follow:

```
svn co https://subversion.gsi.de/fairroot/pandaroot/development/genfit2 <my_branch_name>
```

5) follow the instructions how to install pandaroot on your pc, as indicated here:

<https://panda-wiki.gsi.de/foswiki/bin/view/Computing/PandaRootInstall>

6) make sure to export your SIMPATH env. variable as follow:

```
export SIMPATH=/you_home/your_pandaroot_repository_on_your_pc/apr13.build/
```

7) configure your environment with:

```
your_new_branch_build> ./config.sh
```

and you should be ready to run your macros.

Please, let me know if you still get problems.

ciao, Elisabetta

### File Attachments

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- 1) [config.sh](#), downloaded 542 times
  - 2) [CMakeLists.txt](#), downloaded 572 times
  - 3) [CMakeLists.txt\\_genfitAlicia](#), downloaded 578 times
  - 4) [CMakeLists.txt\\_branchAlicia](#), downloaded 585 times
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