fiTQun Seeding Values

- We want to study the impact of seeding values on fiTQun analyses like 1R-Fit etc.
- E.g. how does performance change if the seeding method changes?
- Idea: Extract seed and store it in a new Branch in the .root file. Later analyse fit results and how they compare to seed.
- Here: New Branch "fitqun1r_.fq1rseed" contains eight entries per event.

fitqunoutC.h

120	struct I	rrdnutt_common {
151	int	fqnse;
152	int	<pre>fqitwnd[MAXNPEAK];</pre>
153	int	<pre>fqipeak[MAXNPEAK];</pre>
154	int	<pre>fqnhitpmt[MAXNPEAK];</pre>
155	float	<pre>fqtotq[MAXNPEAK];</pre>
156	float	fq0rtotmu[MAXNPEAK];
157	float	<pre>fq0rnll[MAXNPEAK];</pre>
158	int	fqn50[MAXNPEAK];
159	float	fqq50[MAXNPEAK];
160	int	<pre>fq1rpcflg[MAXNPEAK][MAXPX];</pre>
161	float	<pre>fq1rmom[MAXNPEAK][MAXPX];</pre>
162	float	<pre>fq1rt0[MAXNPEAK][MAXPX];</pre>
163	float	<pre>fq1rtotmu[MAXNPEAK][MAXPX];</pre>
164	float	<pre>fq1rnll[MAXNPEAK][MAXPX];</pre>
165	float	<pre>fq1rpos[MAXNPEAK][MAXPX][3];</pre>
166	float	<pre>fq1rdir[MAXNPEAK][MAXPX][3];</pre>
167	float	<pre>fq1rdconv[MAXNPEAK][MAXPX];</pre>
168	float	<pre>fq1reloss[MAXNPEAK][MAXPX];</pre>
169	double	<pre>fq1rseed[MAXNPEAK][MAXPX][8];</pre>

FQROOTOut.cc

99	<pre>void fQR00TOut::InitTree(){</pre>
135	
136	<pre>trout->Branch("fqnse" , &fitqun1rfqnse, "fqnse/I");</pre>
137	<pre>trout->Branch("fqitwnd" , fitqunlrfqitwnd, "fqitwnd[fqnse]/I");</pre>
138	<pre>trout->Branch("fqipeak" , fitqunlrfqipeak, "fqipeak[fqnse]/I");</pre>
139	<pre>trout->Branch("fqnhitpmt" , fitqun1rfqnhitpmt, "fqnhitpmt[fqnse]/I");</pre>
140	<pre>trout->Branch("fqtotq" , fitqun1rfqtotq, "fqtotq[fqnse]/F");</pre>
141	<pre>trout->Branch("fq0rtotmu" , fitqun1rfq0rtotmu, "fq0rtotmu[fqnse]/F");</pre>
142	<pre>trout->Branch("fq0rnll" , fitqun1rfq0rnll, "fq0rnll[fqnse]/F");</pre>
143	<pre>trout->Branch("fqn50" , fitqun1rfqn50, "fqn50[fqnse]/I");</pre>
144	<pre>trout->Branch("fqq50" , fitqun1rfqq50, "fqq50[fqnse]/F");</pre>
145	<pre>trout->Branch("fq1rpcflg" , fitqun1rfq1rpcflg, Form("fq1rpcflg[fqnse][%d]/I",MAXPX));</pre>
146	<pre>trout->Branch("fq1rseed" , fitqun1rfq1rseed, Form("fq1rseed[fqnse][%d][8]/F",MAXPX));</pre>
147	<pre>trout->Branch("fglrmom" , fitgunlr .fglrmom, Form("fglrmom[fgnse][%d]/F",MAXPX));</pre>

fiTQun_shared.cc

```
:::SaveTestSnglTrkFit(double* fitparams, int istage, int itwnd, int iPID, double totmu, double nll, int PCflg)
        if (fitgun1rtest .fgtestn1r >= FQTESTMAX1R)
1009
1010
             ::cout << "In fiTQun shared::SaveTestSnglTrkFit: cannot save more than " << FQTESTMAX1R << " test 1 ring fits." << sid::endl;
                             ---> This test fit will not be saved (itwnd, iPID, totmu, nll) = (" << itwnd << "," << iPID << "," << totmu << "," << nll << ")." <<
1011
             ::cout << "
                                                                                                                                                                     ::endl;
1012
1013
        fitqun1rtest .fqtest1rstage[fitqun1rtest .fqtestn1r]
                                                                = istage;
1015
        fitqun1rtest .fqtest1rse [fitqun1rtest .fqtestn1r]
1016
                                                                = itwnd;
        fitgun1rtest .fqtest1rpid [fitgun1rtest .fqtestn1r]
1017
                                                                = iPID;
        for (int i=0; i<
                                     :::nSnglTrkParams; i++) {
1018
          fitgun1r .fg1rseed[0][iPID][i]=fitparams[i];
        // fitgun1rtest .fgtest1rpos [fitgun1rtest .fgtestn1r][0] = fitparams[0];
1021
        // fitgun1rtest .fqtest1rpos [fitqun1rtest .fqtestn1r][1] = fitparams[1];
1022
            fitqun1rtest .fqtest1rpos [fitqun1rtest .fqtestn1r][2] = fitparams[2];
```

"fitparams" corresponds to the values of "snglTrkParams" at the point in fiTQun.cc where fiTQun_shared::SaveTestSnglTrkFit gets called.

fiTQun_shared::SaveTestSnglTrkFit only saves "test" variables which don't get read into the .root file. Perhaps problematic to save the seed using this function?

fiTQun.cc

```
2500
             ::cout << "##########Initial parameters: ";</pre>
         for (int i=0; i<fi</pre>
                                        :::nSnglTrkParams; i++) {
2501
2502
               ::cout << X[i] << ", ";
2503
2504
             ::cout << "####################
                                                d::endl;
2505
         if (X0!=NULL) {
2506
           for (int isd=0; isd<fiTOun sha</pre>
                                              ::nSnglTrkParams; isd++) {
2507
             X0[isd]=X[isd];
2508
2509
2510
2511
2512
         lnlval=GetOneRngnglogL(iPID,X,PCflg);
         totmu=GetTotmu();
2513
         if (fgshared->GetFmskbkPMT()) fignrbkPMT=true;
2514
         fqshared->SaveTestSnglTrkFit(X,1,itwnd,iPID,totmu,lnlval,PCflg);
2515
         if (fgshared->GetDrawflg()) DrawPreddist(Form("%d %d %d 1R%s 0",ievent,itwnd,ipeak,
2516
                                                                                                               l::PIDnames[iPID].Data());
```

SaveTestSnglTrkFit gets called at the point where X[i] still corresponds to the seed, and afterwards assumes the values of the fit result.

The other relevant variables are saved using functions like *GetDefaultSnglTrkFit* or *SaveDefaultSnglTrkFit*, but they are called when X[i] is either already the previous fit result or the new fit result.

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- Recommendations/inputs?
- Right now used for testing purposes, but it would be useful for production. Do we want that?



Fit Result

Difference to Seed

