

Results for profiled FC for $B^0 \rightarrow K^* \mu \mu$

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What's new since last week presentation at the BPAG

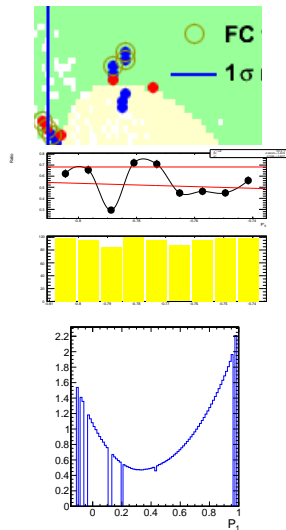
- We did some interaction with Phys Coords (“this is insane”)
- interaction with ARC: “ah, good, you did some progress, let us now when all is finished so we can look at it”
- interaction with Olaf: quite concerned with the 2d gauss fit
 - ▶ not because it can be tricky and maybe it is not the proper way to find the toy LogL minimum
 - ▶ but instead suggesting fancy way to fit a damn 2D gauss!
 - ▶ **You are trying to solve the WRONG problem man!!**
- **no comments whatsoever about the fact that this crazy method**
 - ▶ is insane
 - ▶ is not giving clear results
 - ▶ when it does, the 1σ is the same as what we called “custom minos”
- my personal feeling that no one is really taking care to look at the big picture anymore
 - ▶ We have this insane FC bul***it to perform and that's it
 - ▶ when we will finish we'll be not a inch closer to understanding the problem, but who cares?

- Run a non negligible number of new GenPoint
 - ▶ babysitting is now a bit easier but not painless
- Fixed (mostly) the problem of the non converging 2d gaussian fit
 - ▶ Alessio had put a requirements for $\sigma_{P_1} < 0.9$, $\sigma_{P'_5} < 0.9$, $\rho < 0.99$,
 - ▶ no idea why
 - ▶ removing that requirement allow to recover most of the 2d gaus fit, now only $\sim 5\%$ is failing (fit status 3)
- Added a linear fit to the ratio vs $P_1(P'_5)$
 - ▶ A possible way to determine the 1σ border
 - ▶ is not helping much if the ratio trend is not clear

Debug and GenPoint

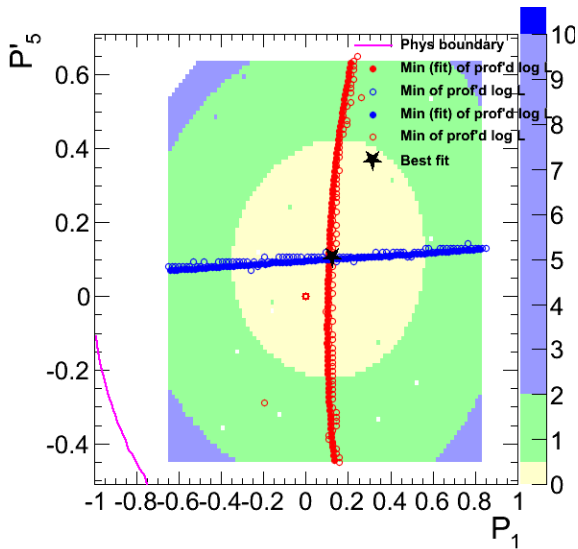
debug a number of problematic Gen Points

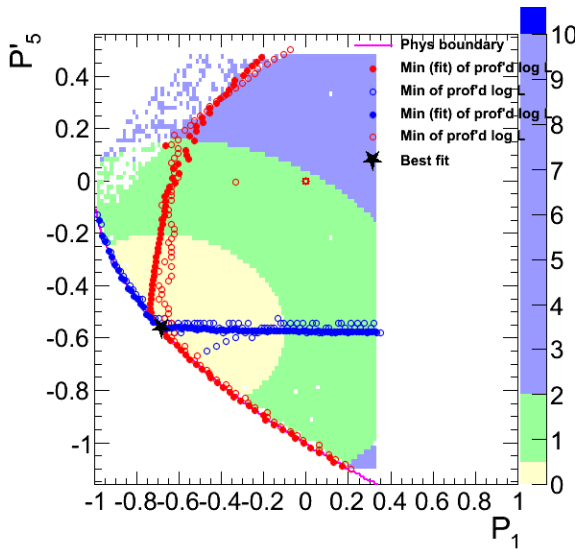
- eg Bin 2, region P'_5 up
- one point has ratio ~ 0.3 , between ones with ~ 0.7 : why?
- It is a “displaced point”, namely with a P_1 not aligned with the other
- actually it happens that that is the true minimum along the profile, but it does not correspond to the minimum of the parabola of the LogL.
- possibly a fluctuation or whatever.

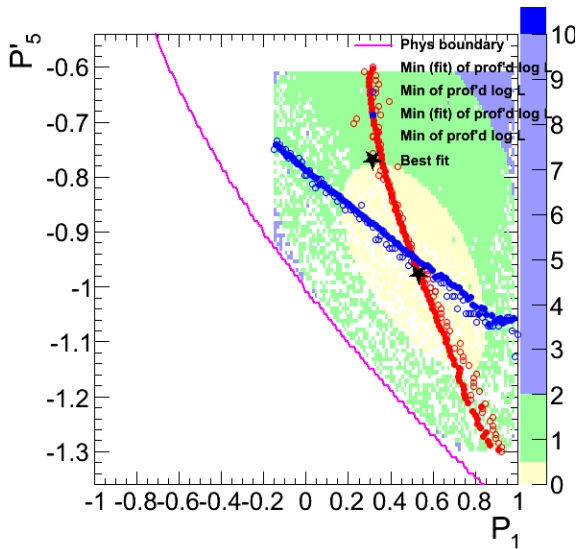


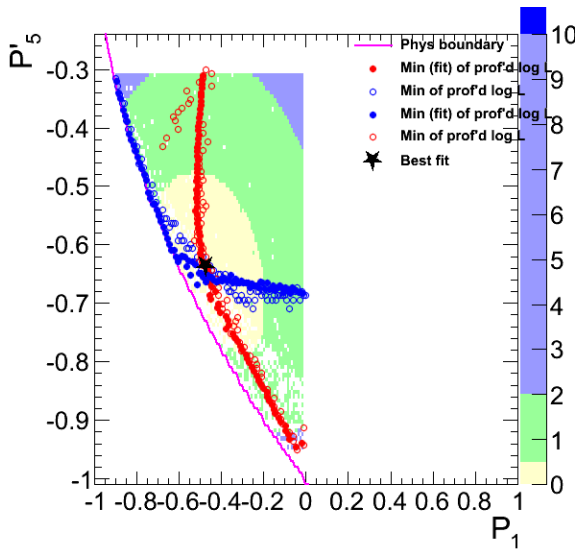
Are we using the correct Gen Point?

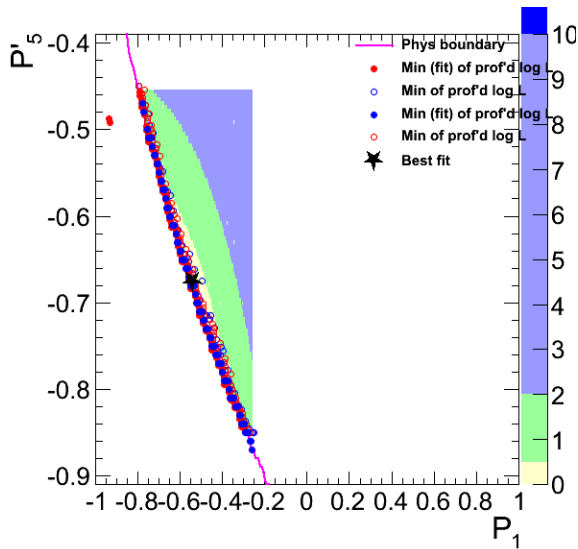
- So it seems that if we move just a bit away from the “ideal” line of profiled minimum the L is changing significantly
- and so the DLL computed for Data and Toys can be tricky
- it could also explain the non monotony of the ratio vs $P1/P5'$
- **solution**
 - ▶ Instead of considering the absolute minimum of a profile (that is the bin with the minimum LogL), fit a parabola to the LogL distribution and take the vertex
 - ▶ the vertex x is the coordinate of the Gen Point to be generated
 - ▶ the vertex y is the LogL to be considered for the $\Delta \log \mathcal{L}$ computation
 - ▶ Pros: these Gen Point are much more aligned than the one computed previously.
 - ▶ Cons: we (I) have to redo most of the work
 - ▶ Will try to do that for one region which is not problematic, and we will see

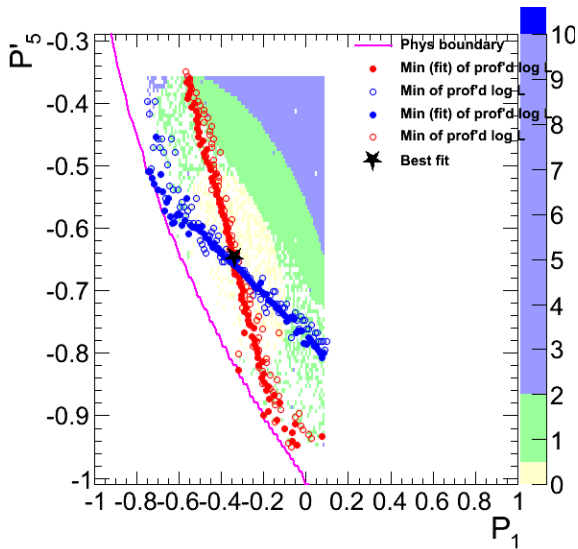


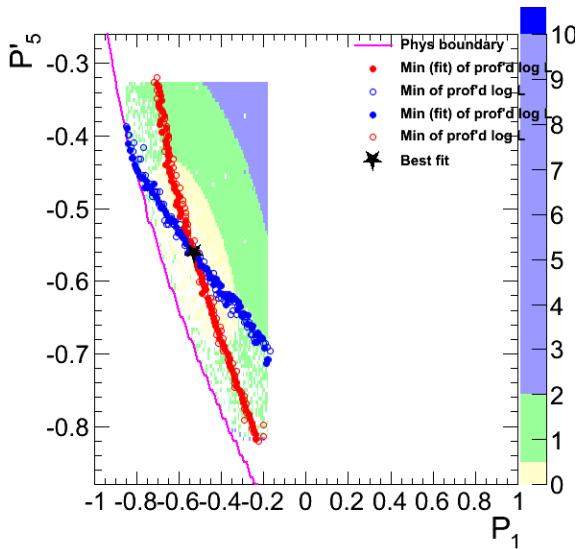






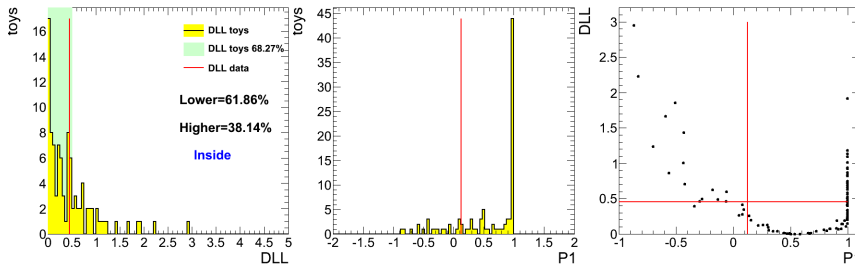


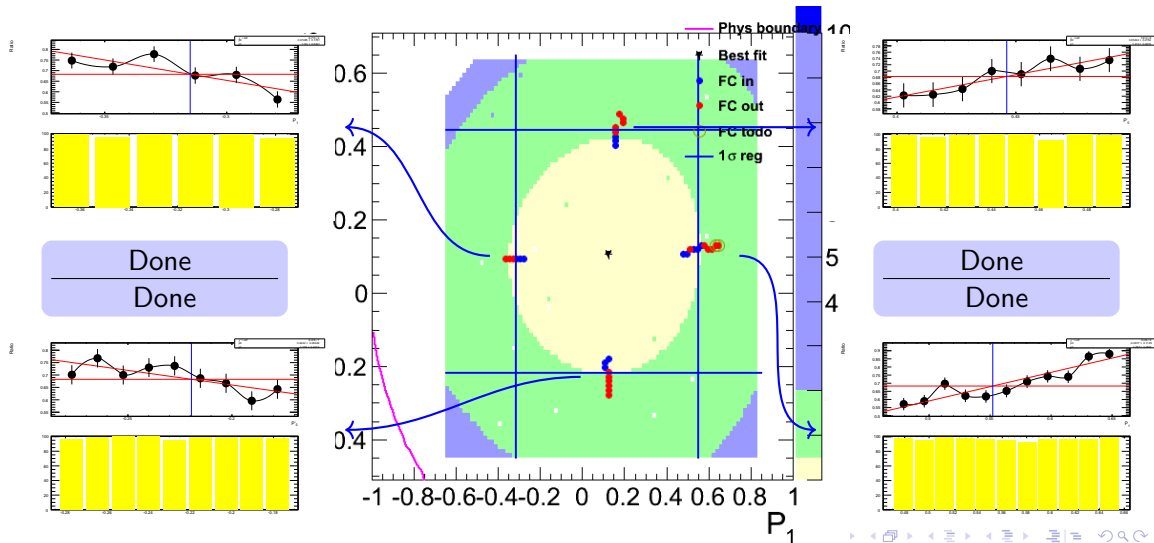


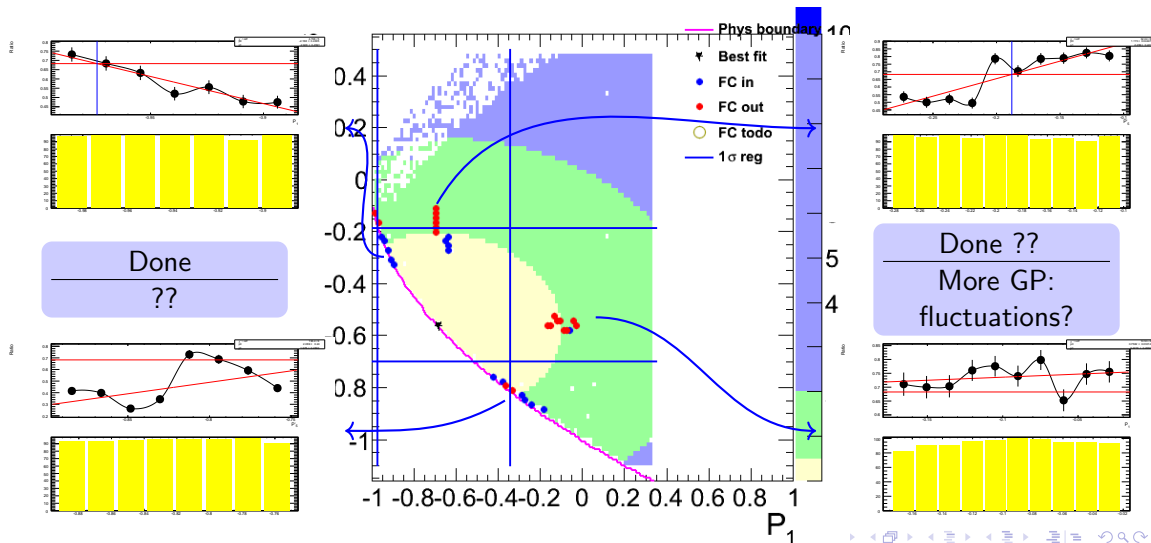


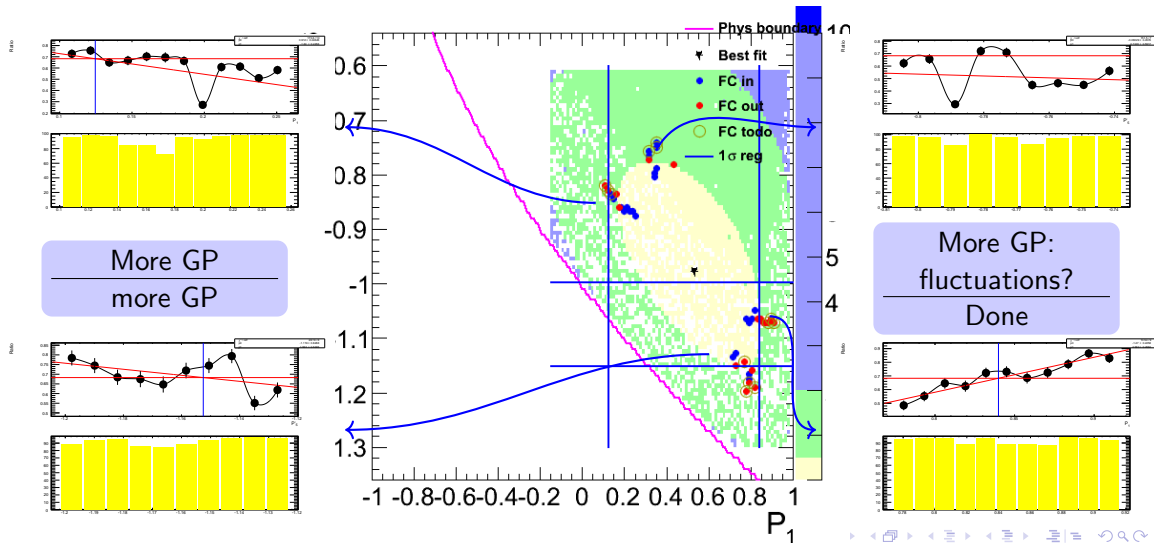
DLL vs P1 for one GenPoint

- Following Mauro's idea: look at P_1 distribution for toys, instead of DLL
- build a 68% region for P_1 and see if P_1 for data is inside that region or not
- problem with P_1 physical boundary
- DLL vs P_1 is parabolic (as expected?), so comparing DLL is the same as comparing P_1 ranges (but easier)

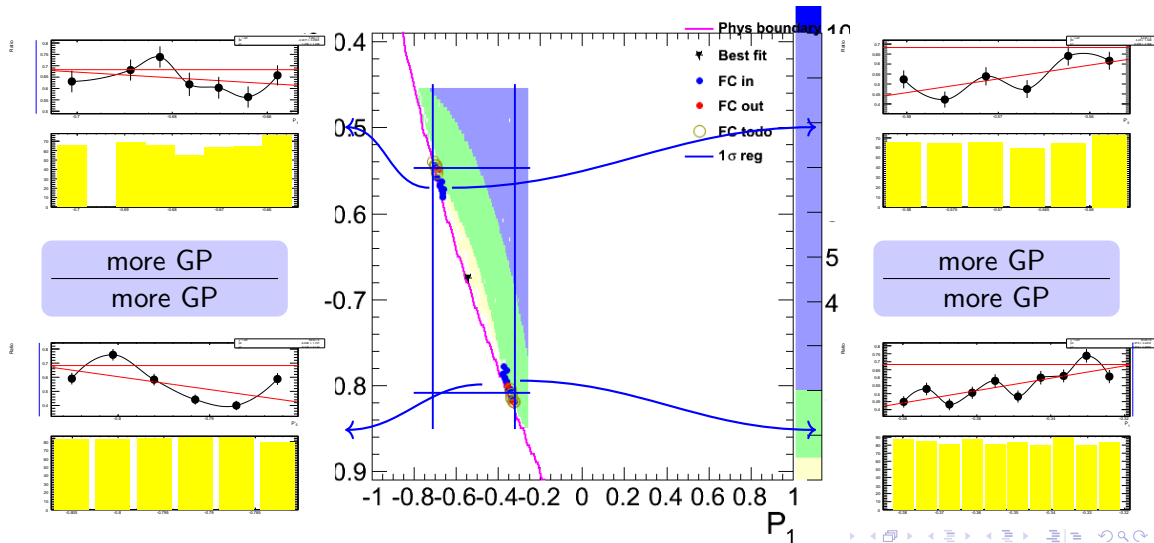


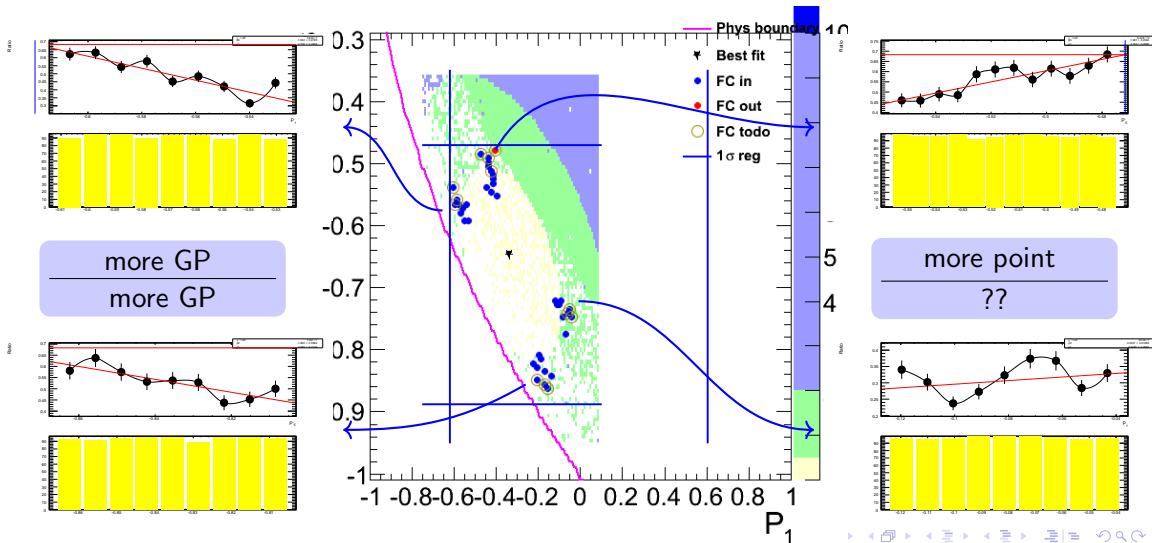


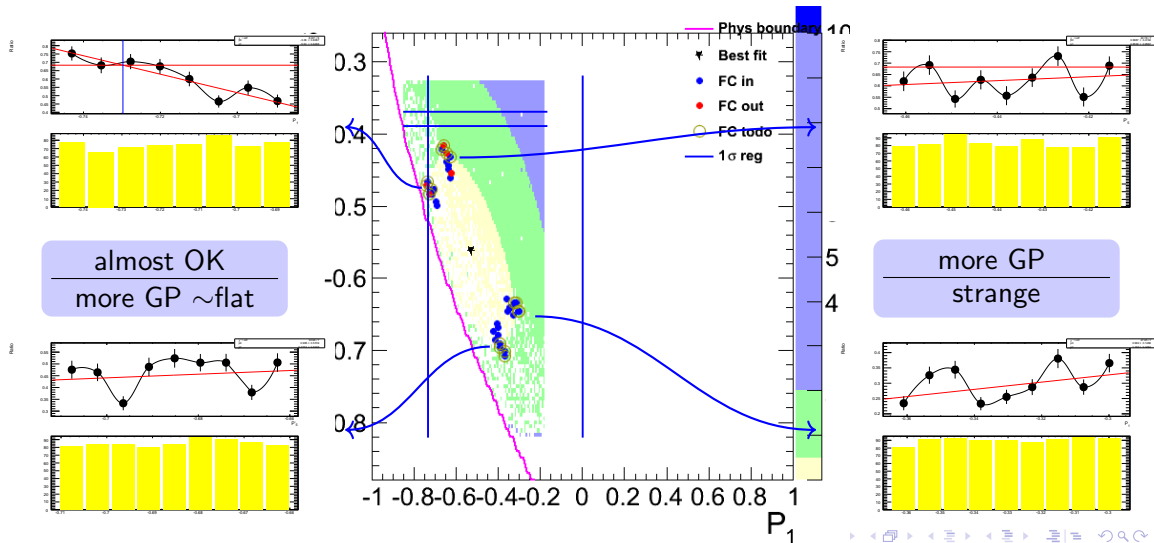














Additional stuff

Additional or backup slides



Bibliography I