



# pMSSM grand scan status

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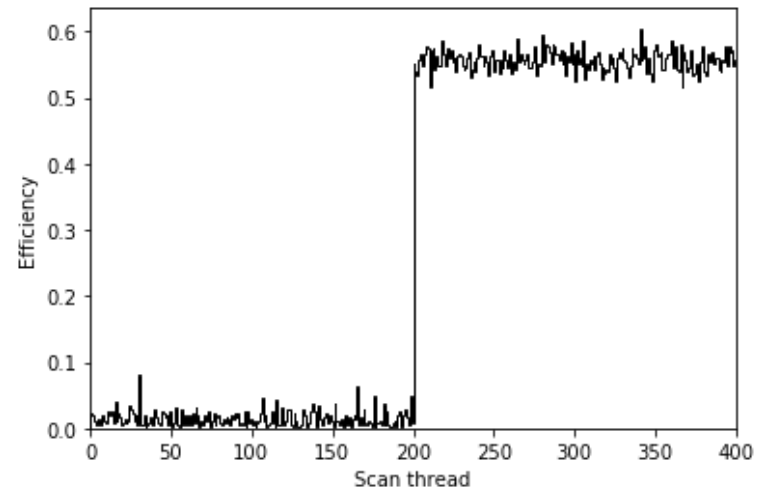
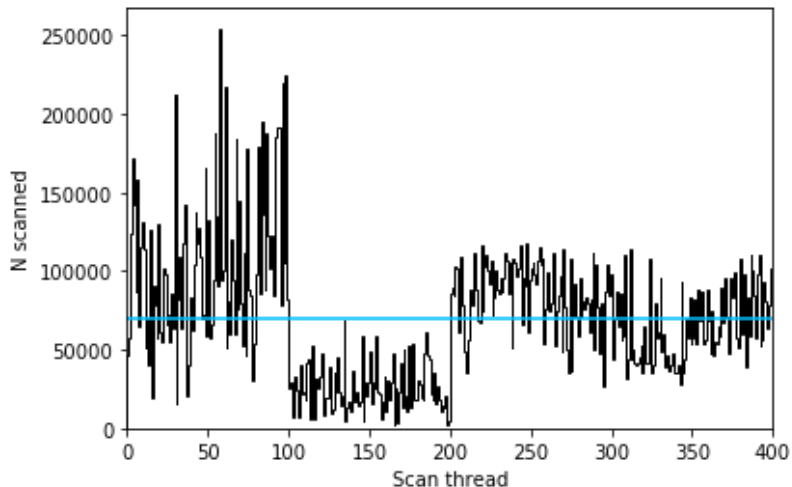
pMSSM Scan Meeting

# What's running: 400 parallel scans

- 200 threads with the nominal setup
  - Saw the first 100 converging to  $\Delta a_\mu \sim 250$  ([gif](#))
- 200 more threads with the same setup but without  $\Delta a_\mu$  in the MCMC likelihood, to populate  $\Delta a_\mu = 0$
- Test scans with different stepping functions
  - Amounts to only a few thousand points

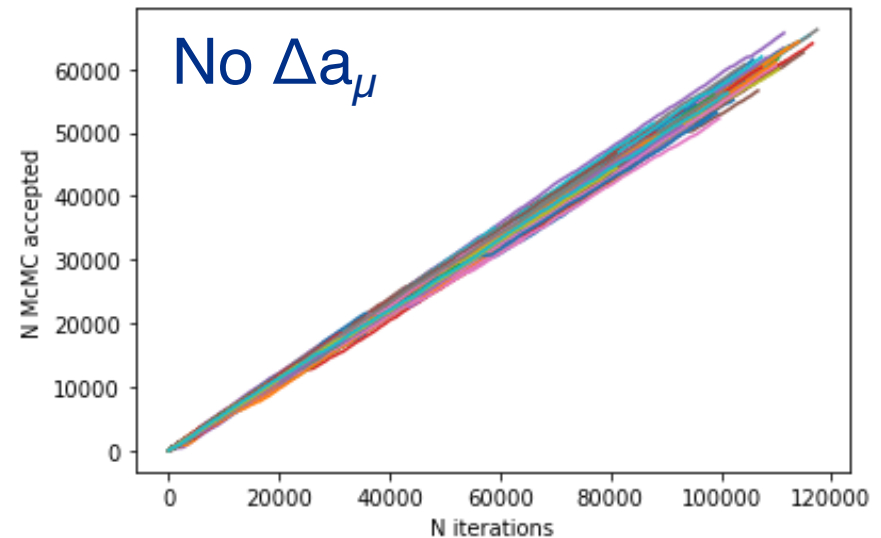
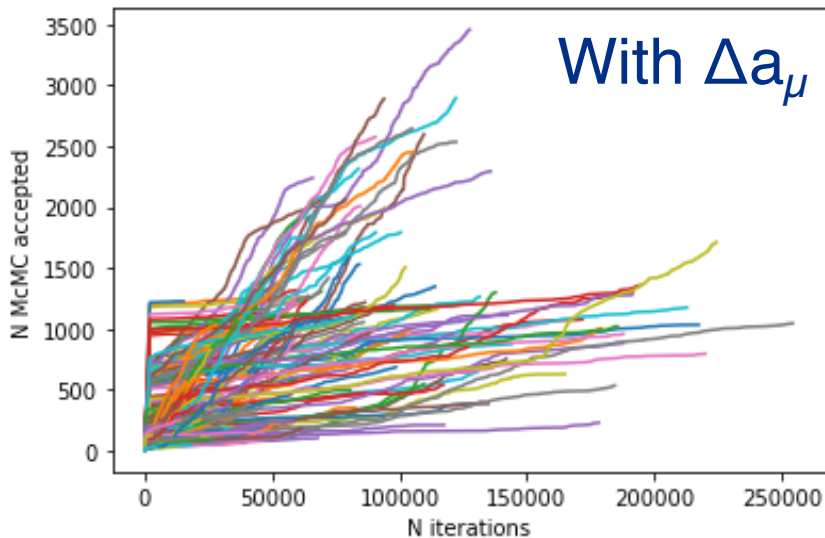
# Progress of the scan

- Threads 1-200: with  $\Delta a_\mu$ 
  - 1-100 started  $\sim$ Oct, 101-200 started in Jan
- Threads 201-400: no  $\Delta a_\mu$ 
  - Much higher efficiency

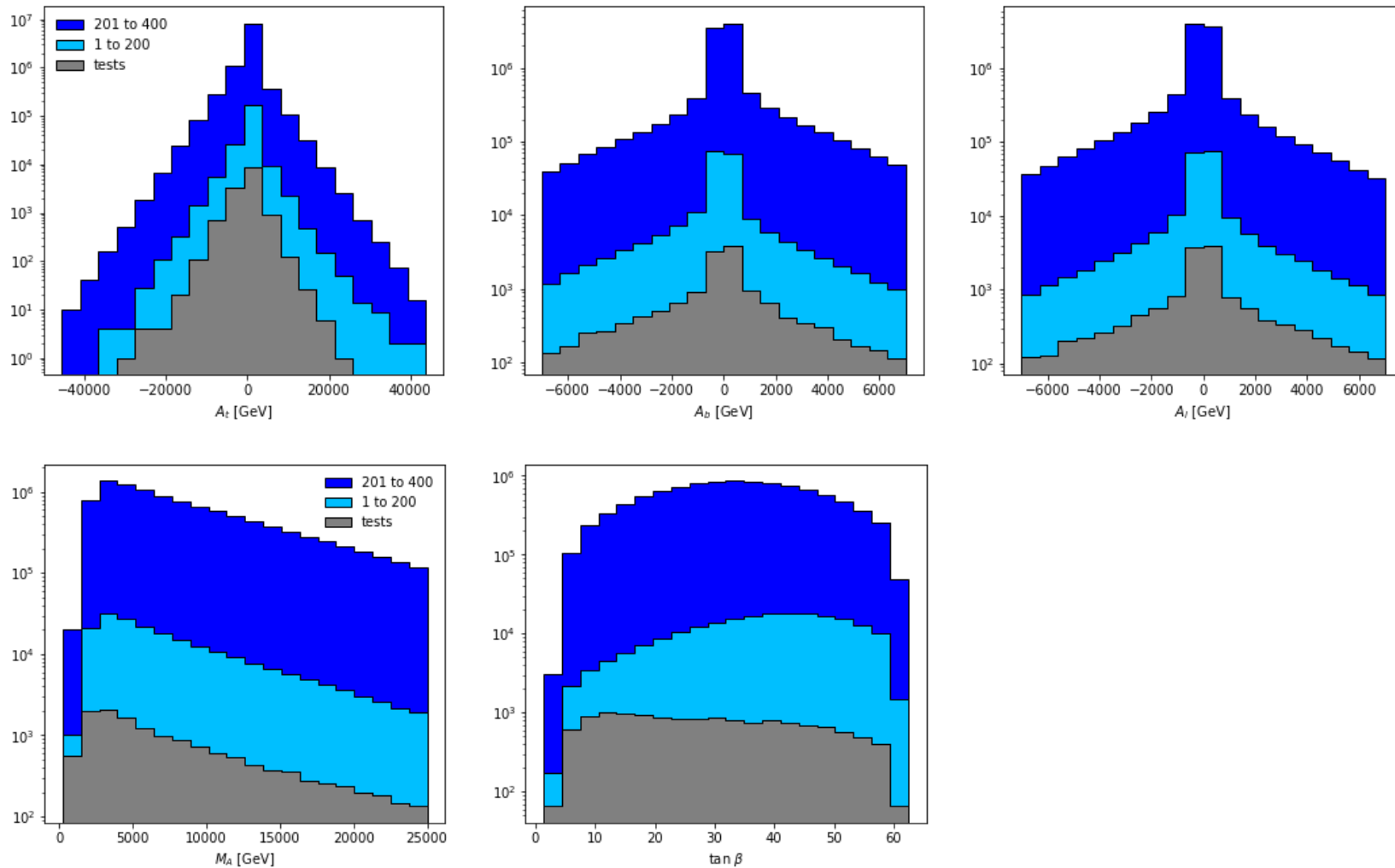


# Progress of the scan

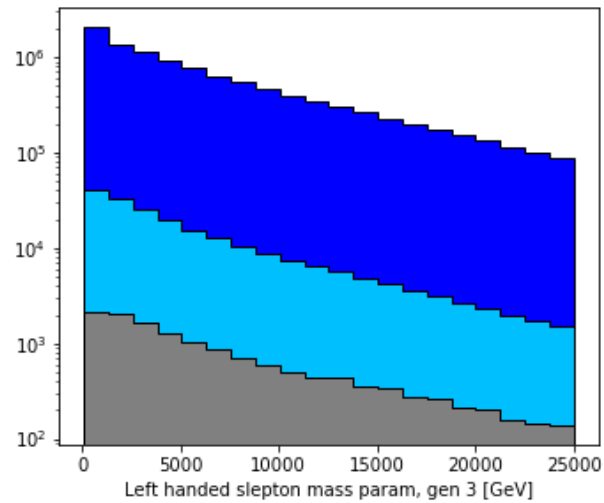
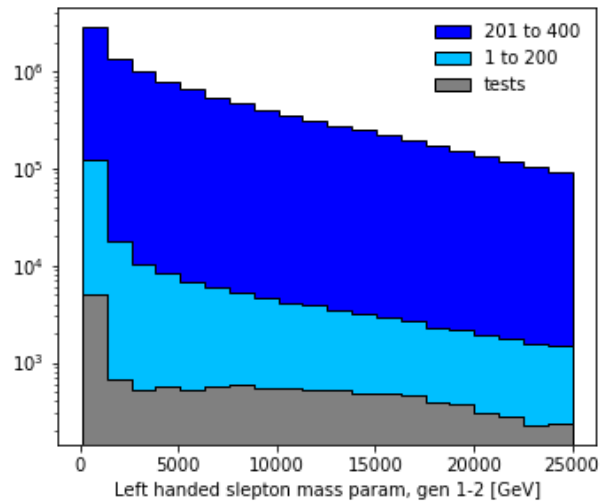
- So far, scanned 28,768,213 points (up from 5.6M on 01/12)
  - MCMC accepted 10,368,890 (up from 61k on 01/12)
  - At least 9,438 passing all cuts (HiggsBounds, Micromegas and SModelS). Likely many more
    - Have not run post-processing on points produced since mid-Jan



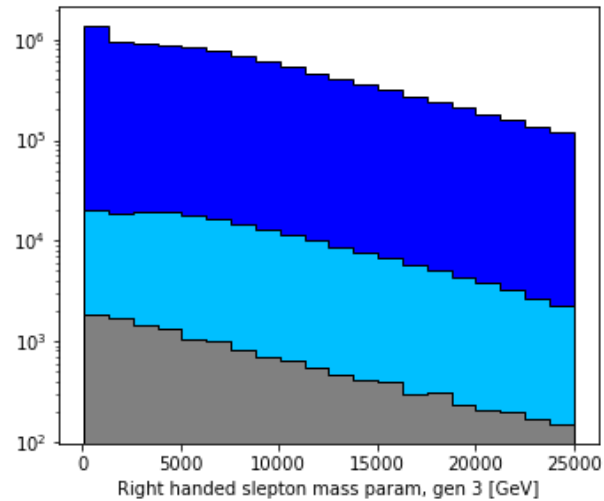
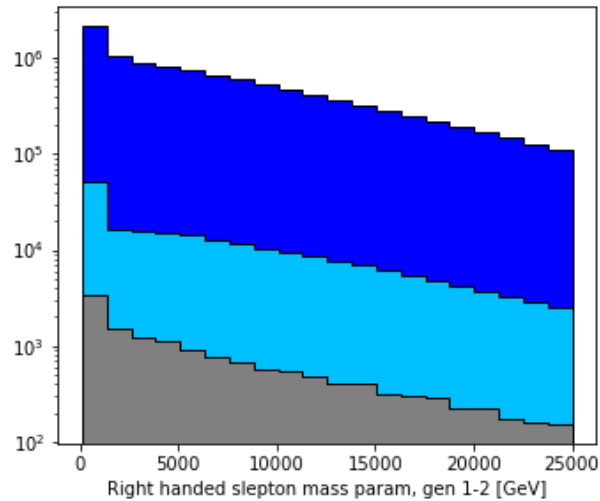
# Comparison of with / without $\Delta a_\mu$



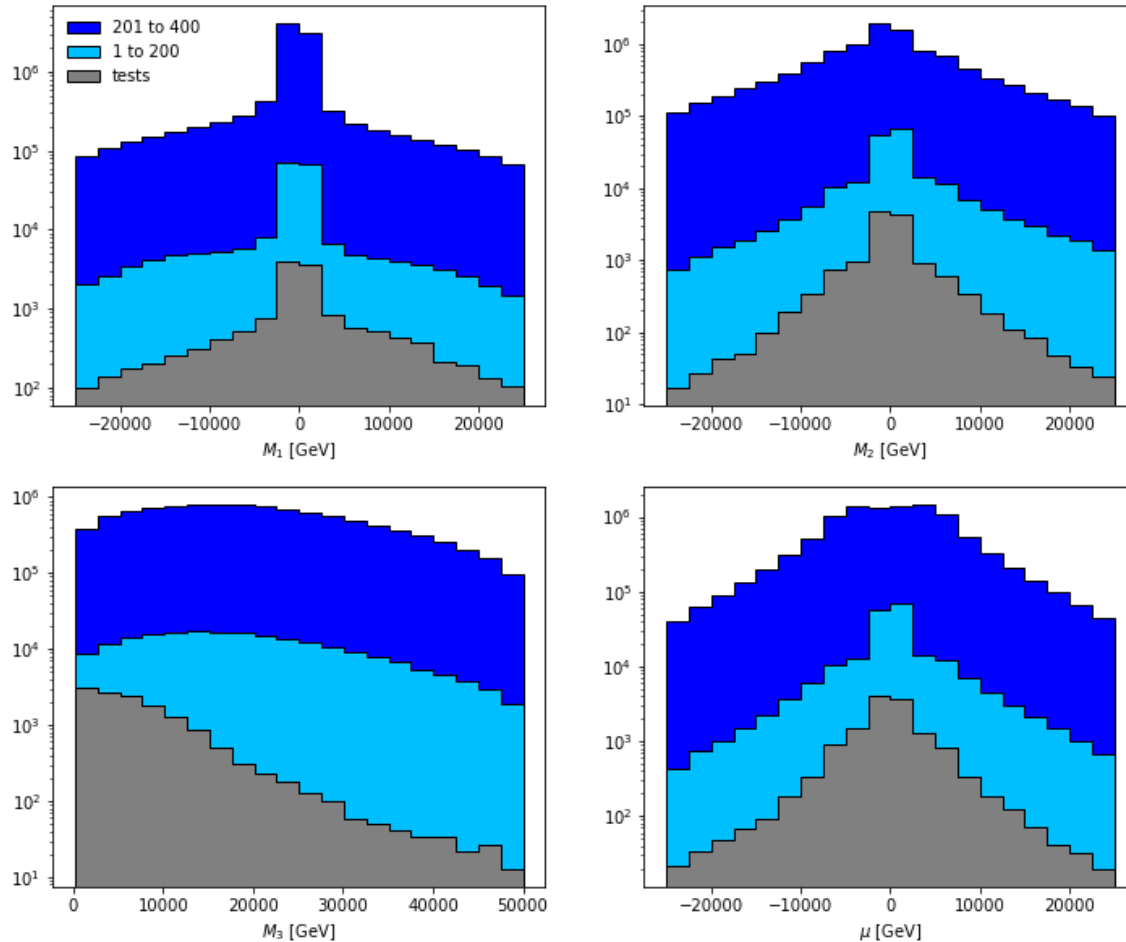
# Comparison of with / without $\Delta a_\mu$



More light  
smuons when  
 $\Delta a_\mu$  is included

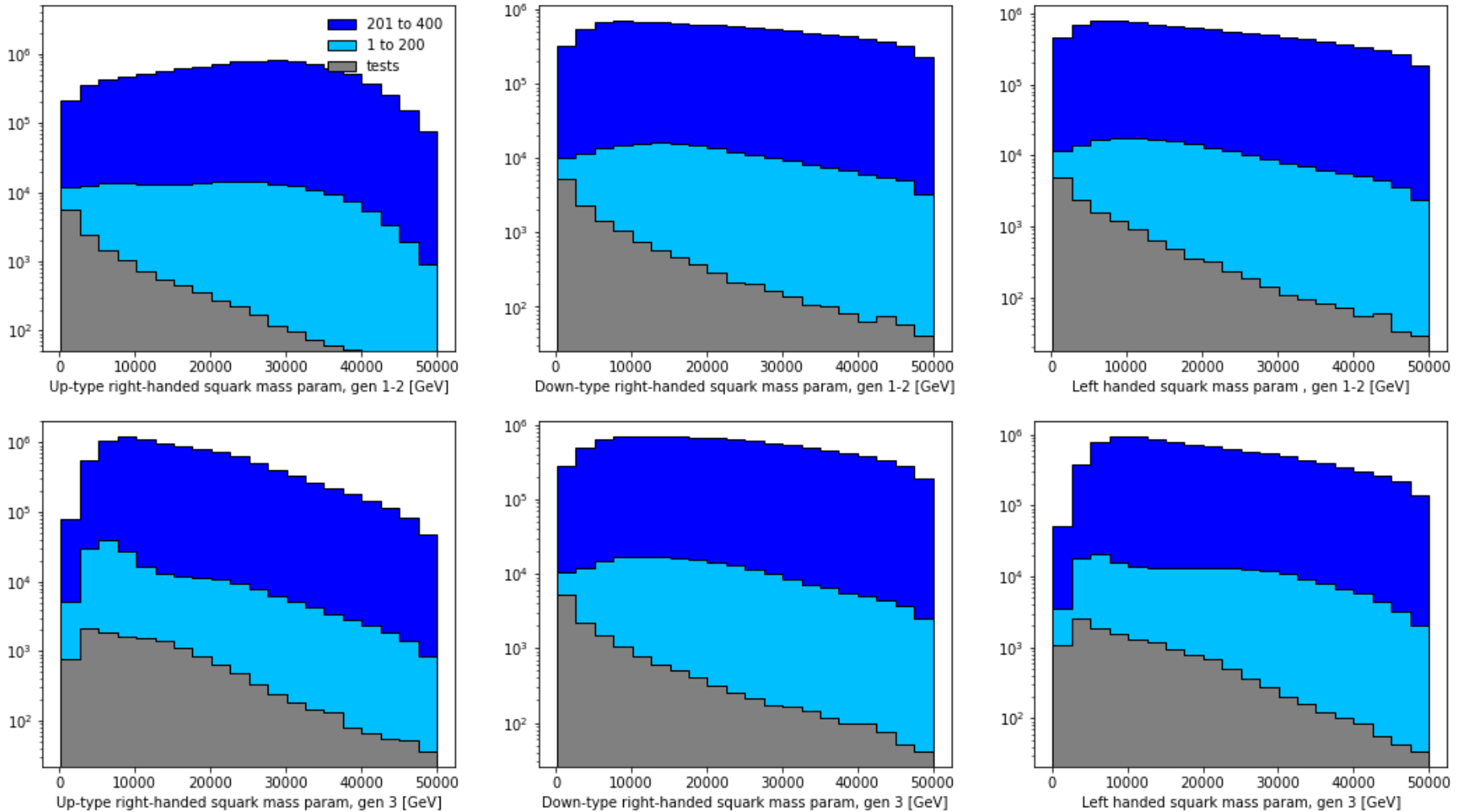


# Comparison of with / without $\Delta a_\mu$



Tests use log stepping in gluino mass param  $M_3$

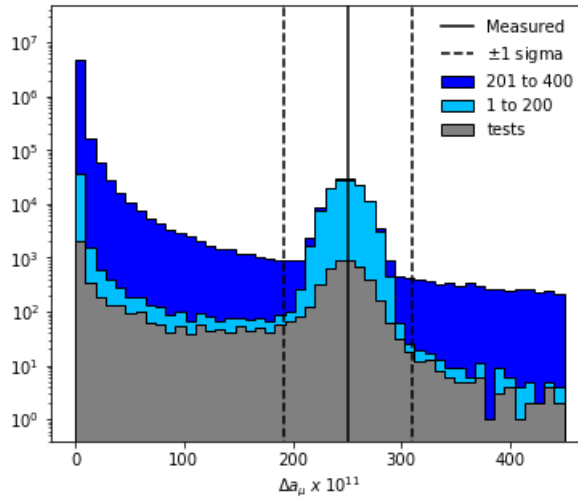
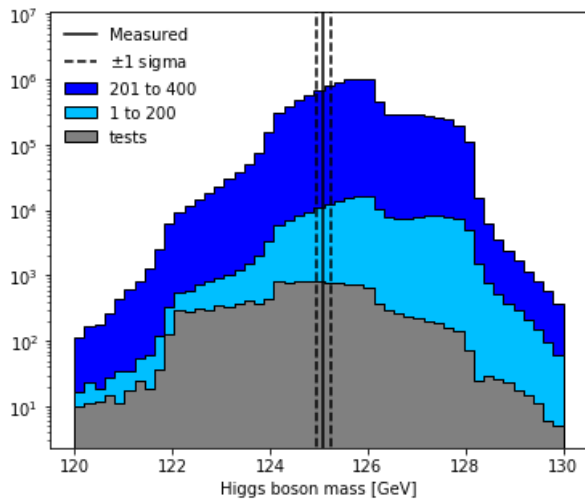
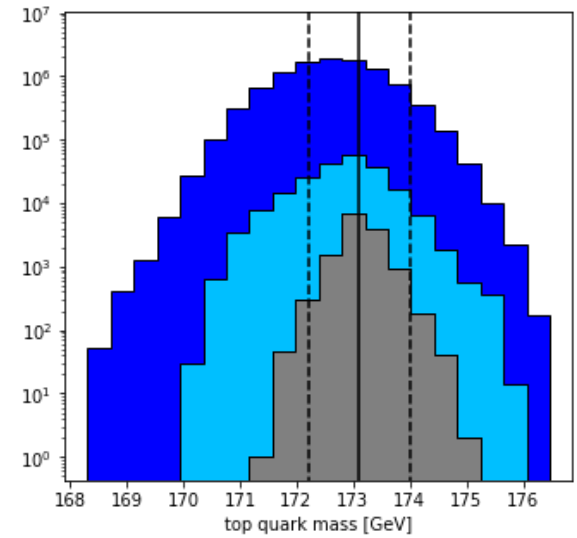
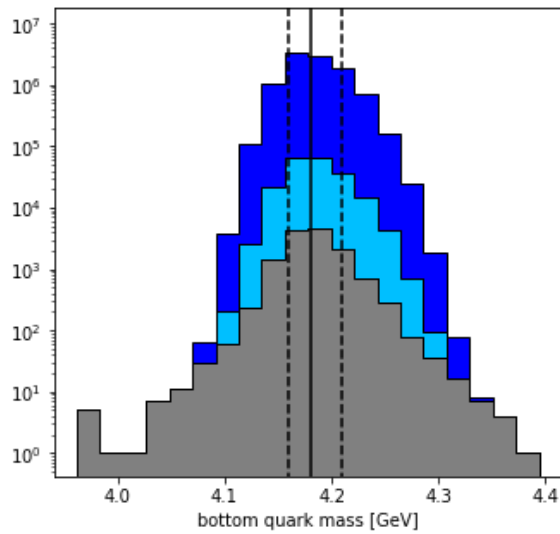
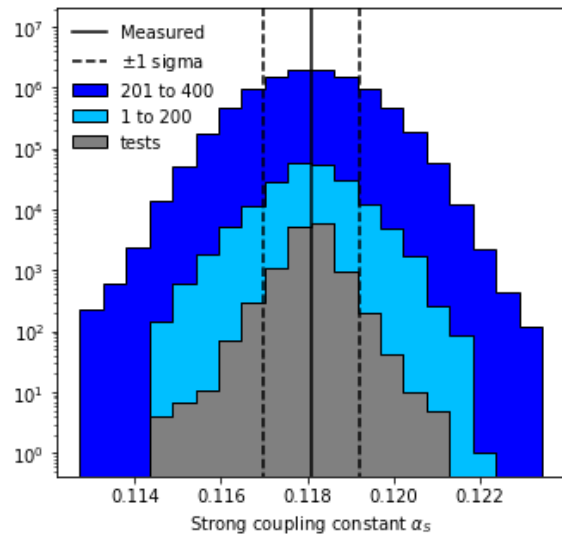
# Comparison of with / without $\Delta a_\mu$



Tests use stepping in squark mass params



# Comparison of with / without $\Delta a_\mu$



# Impact of future measurements

- For pMSSM paper: include the impact of projected future precision measurements on pMSSM space
  - Assume observable  $X$  falls into a narrow range projected at a future collider. How does this shape the allowed pMSSM space?
- Parameters to be considered (blue are missing)
  - Higgs mass
  - Higgs couplings:  $HHH$ ,  $HWW$ ,  $HZZ$ ,  $Htt$ ,  $Hbb$ ,  $H\tau\tau$ ,  $H\mu\mu$ ,  $Hcc$
  - $\Delta a_\mu$
  - DM quantities calculated by Micromegas
  - Flavor observables? (Belle II)