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# Snowmass21 pMSSM scan: workflow overview and open items

Jennet Dickinson, Jim Hirschauer EF08 pMSSM Scan Meeting November 11, 2020

## **Goal for today**

- Agree on the outline of our pMSSM scan workflow and start to assign tasks to interested groups
- This talk
  - Summarizes the steps in our workflow
    - Inspired by last week's talks and discussion
  - Points out important open items
  - Makes a few concrete proposals as a starting point
    - Of course these are still up for discussion
- Later talks: introductions to some of the technical tools that fit in to this workflow

### Sampling pMSSM parameter space

- Scan the 19D pMSSM parameter space
  - Proposal: use a Markov Chain Monte Carlo
- The scan must span a *REALLY LARGE* range of pMSSM parameter space
  - Option 1: a single "grand scan" covering the OR of the accessible ranges of all future collider scenarios
    - Elegant, easily interpreted in a Bayesian framework, easy to compare the reach of different experiments
    - <sup>(2)</sup> The parameter space may be too large to perform oversampling with a meaningful resolution
  - Option 2: multiple scans targeting the accessible range of a smaller group of future collider scenarios
    - Can more easily get reasonable statistics in regions of interest
    - Solution Not straightforward to compare experiments across different scans

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## Sampling pMSSM parameter space (2)

- What is the best way to incorporate existing experimental results into the scan?
  - **Option 1**: directly into the likelihood
    - 😕 Potential for bias if experimental results change...
  - Option 2: by over-sampling in regions of interest
    - Scan could spend lots of time covering less interesting areas...
- In the end, some measurements will be included via Option 1, others via Option 2
  - Which ones in which way? We will need to discuss



### Signal generation and simulation

- Signal simulation with Delphes for all pMSSM points for all future colliders of interest
  - We can use Snowmass MC TF tools
- Additional steps could reduce the number of points to simulate. We should consider:
  - Rejecting points with small cross sections / low yield at fixed luminosity
  - Rejecting points based on a truth-based likelihood (with smearing), as ATLAS does



#### **Event counts**

- Obtain background yields (B) from Snowmass groups dedicated to specific future experiments
  - For missing searches, generate B ourselves using centrally produced SM MC samples
- Generate signal yields (S) from signal simulation
  - In complex cases, e.g. missing tracks analysis, ask other Snowmass groups for help
- Generation of S and B can be done with e.g. RECAST or MadAnalysis5
- Hypothesis testing using simplified likelihood based on S, B,  $\delta$ S, and  $\delta$  B



## **Today's meeting**

• We will get details of the technical implementation of some of the steps in this workflow:

<b>2:00 PM</b> → 2:10 PM	Introduction Speaker: Jennet Dickinson (Fermilab)	⊙10m 🖉 -
<b>2:10 PM</b> → 2:25 PM	Recent progress Speaker: Jeff Shahinian (University of Pennsylvania (US))	<b>⊙</b> 15m
<b>2:25 PM</b> → 2:40 PM	Recent progress Speaker: Malte Mrowietz (University of Hamburg)	©15m

