

# Extending SIMP Search to L1L2

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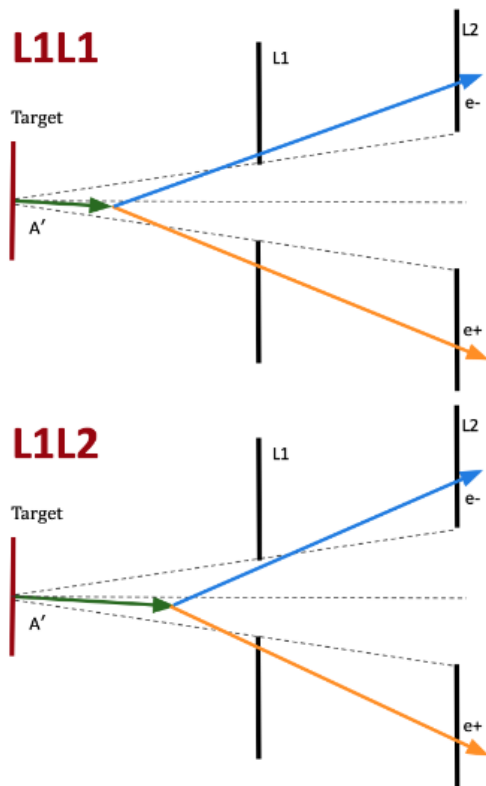


Fig 30 from PRD

## Samples

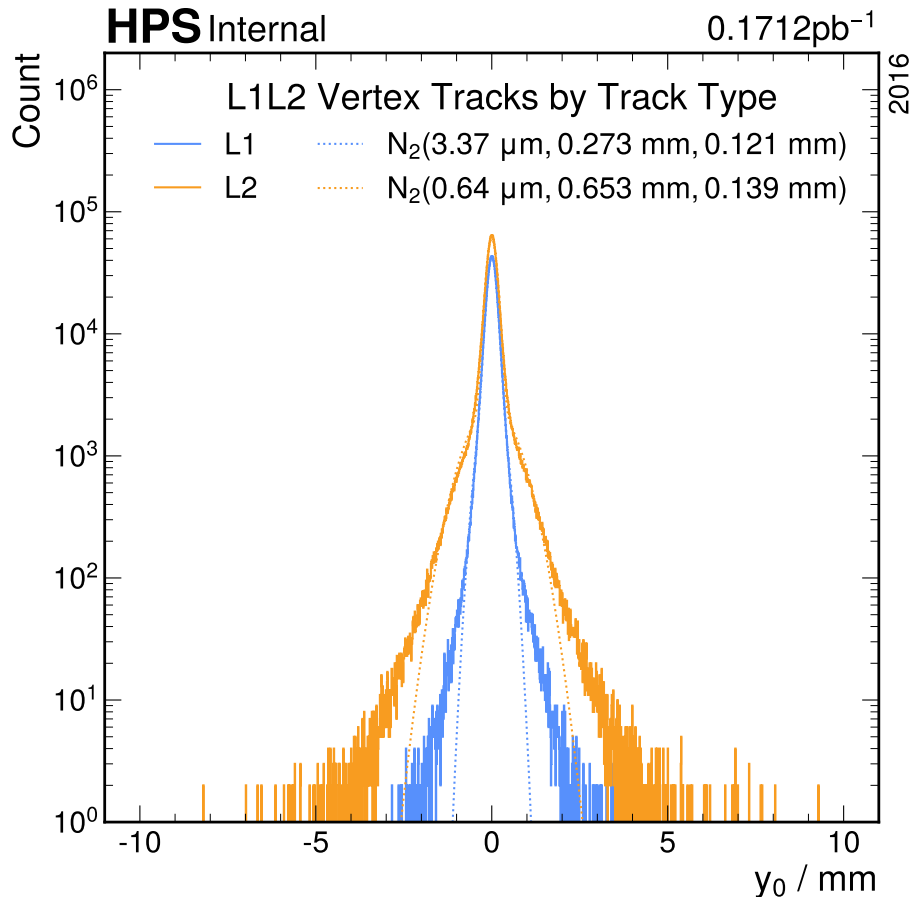
Same data and simulation samples as Alic's SIMP (L1L1) search.

## Selections

Rely on Alic's thorough study and validation, copy pre-selection and start with same final selection variables.

## Search and Exclusion

- Search for excess in  $m_{\text{reco}}$  vs  $\min(|y_{0,e^-}|, |y_{0,e^+}|)$  space
- Exclude by using OIM on the  $z$  distribution after final selections



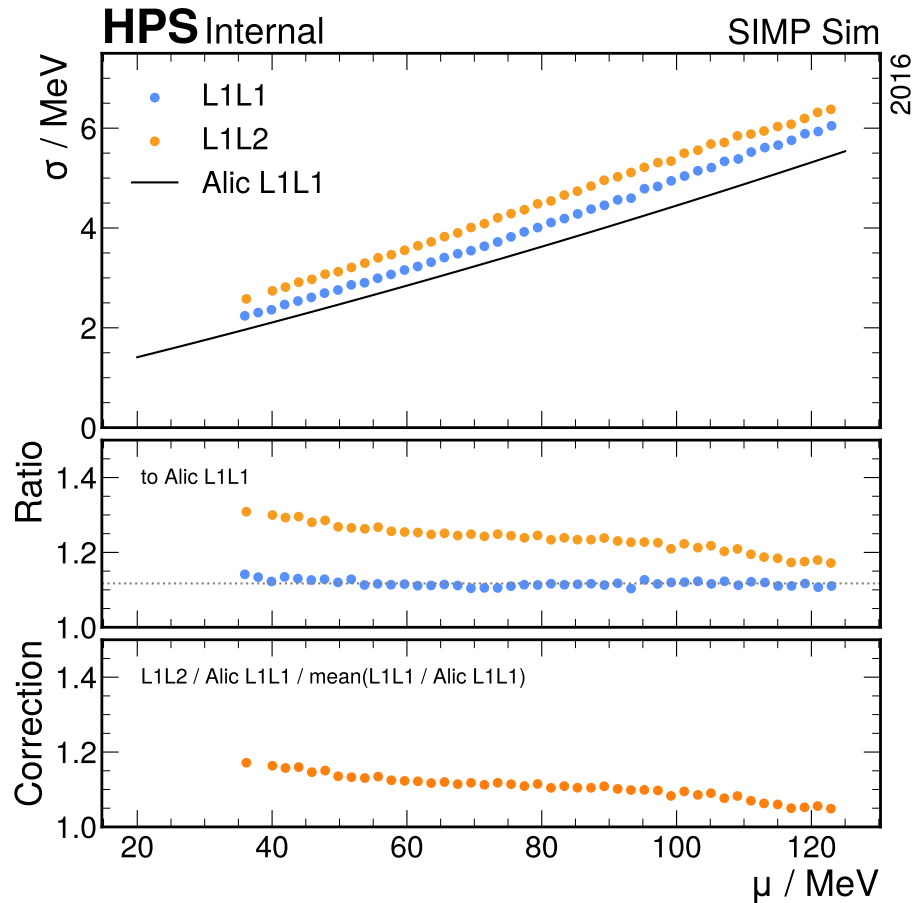
## Selection

Pre-selected vertices and required to be L1L2 (i.e. tracks for L1L1 or L2L2 vertices are not included in this plot).

$N_2(\mu, \sigma_1, \sigma_2)$  is the sum of two normal distributions that share the same mean  $\mu$ .

- L2 tracks slightly broader than L1 tracks (expected)
- Both centered on 0 within  $5 \mu\text{m}$
- Width of core  $y_0$  distribution is only  $\sim 18 \mu\text{m}$  larger for L2 tracks compared to L1 tracks (while the tail ends up being  $\sim 3\times$  wider)

# Mass Resolution Comparison



## Selection

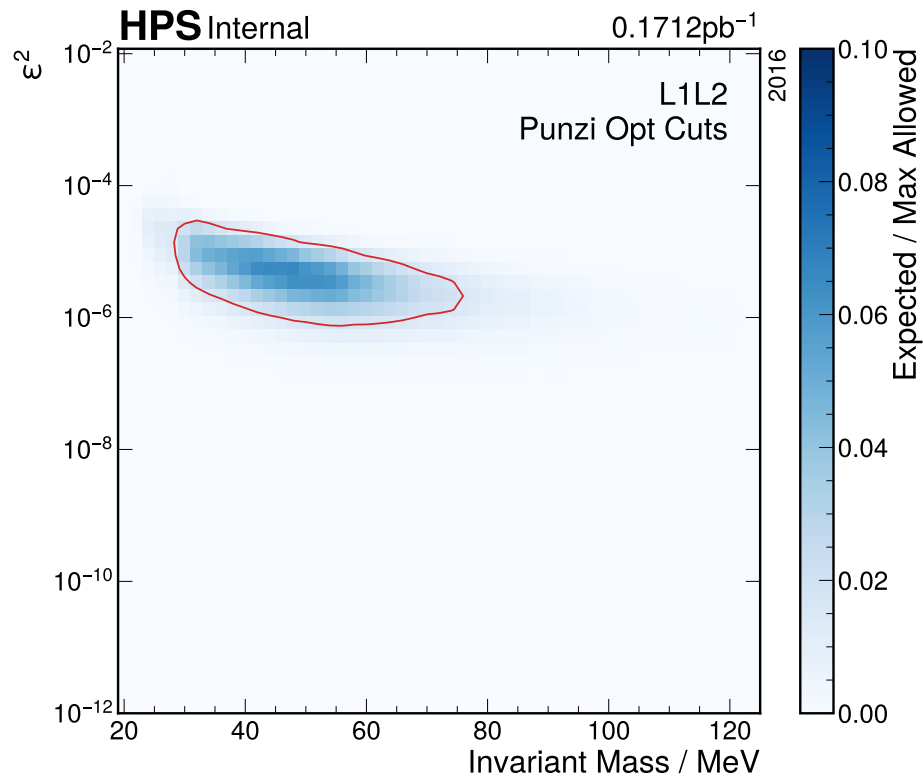
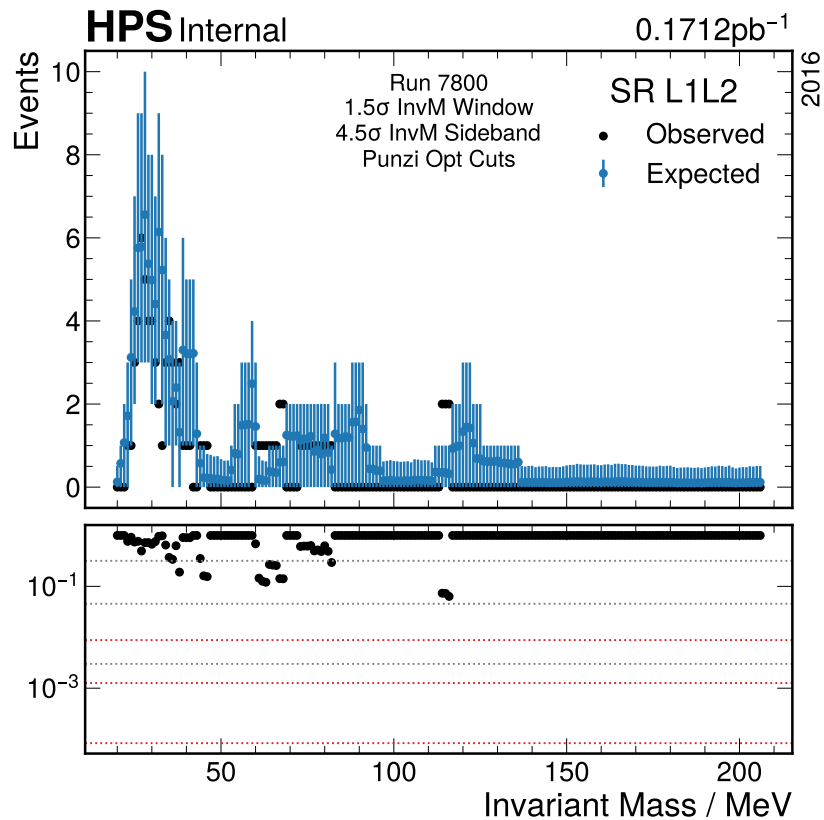
Pre-selected vertices – no truth matching done (hence the separation between L1L1 here and Alic’s L1L1 in black)

- Seeing L1L2 widening in mass resolution by  $< 20\%$  across mass range
- Separation shrinks as mass is increased

## Not Used

The following reach comparisons use “Alic L1L1” in order to maintain comparability with prior estimates. Updating to an L1L2 mass resolution would not be difficult but is left for the future when the cuts and optimization strategy have been more established.

# Reach of Punzi-Optimized Cuts



Replicating optimization procedure as developed by Alic – a one-tailed binomial p-value test (with the p-value converted to a significance).

$$Z_{Bi} = \sqrt{2}\text{erf}^{-1}(1 - 2p_{Bi})$$

where

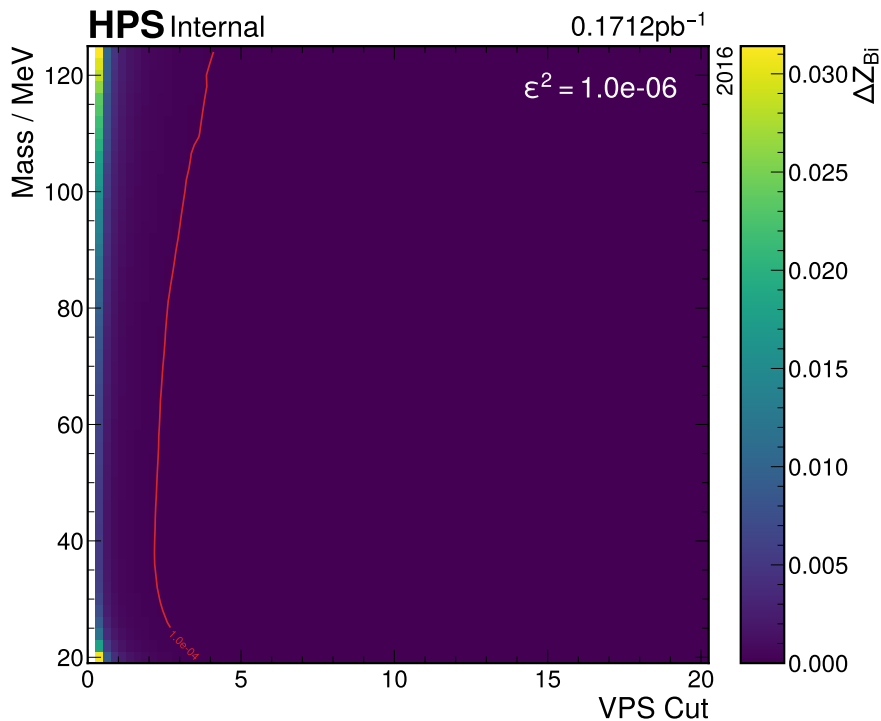
$$\begin{aligned} p_{Bi} &= \sum_{j=n_{on}}^{n_{tot}} P(j|n_{tot}; 1/(1 + \tau)) \approx B(1/(1 + \tau), n_{on}, 1 + n_{off})/B(n_{on}, 1 + n_{off}) \\ &= B_{reg}(1/(1 + \tau), n_{on}, 1 + n_{off}) \end{aligned}$$

where  $B(a, b)$  is the complete beta function,  $B(x, a, b)$  is the incomplete beta function, and  $B_{reg}(x, a, b)$  is the regularized incomplete beta function. Used  $\tau = 1$ ,  $n_{on} = S + B$ , and  $n_{off} = B$  for this evaluation, so in summary the FoM is

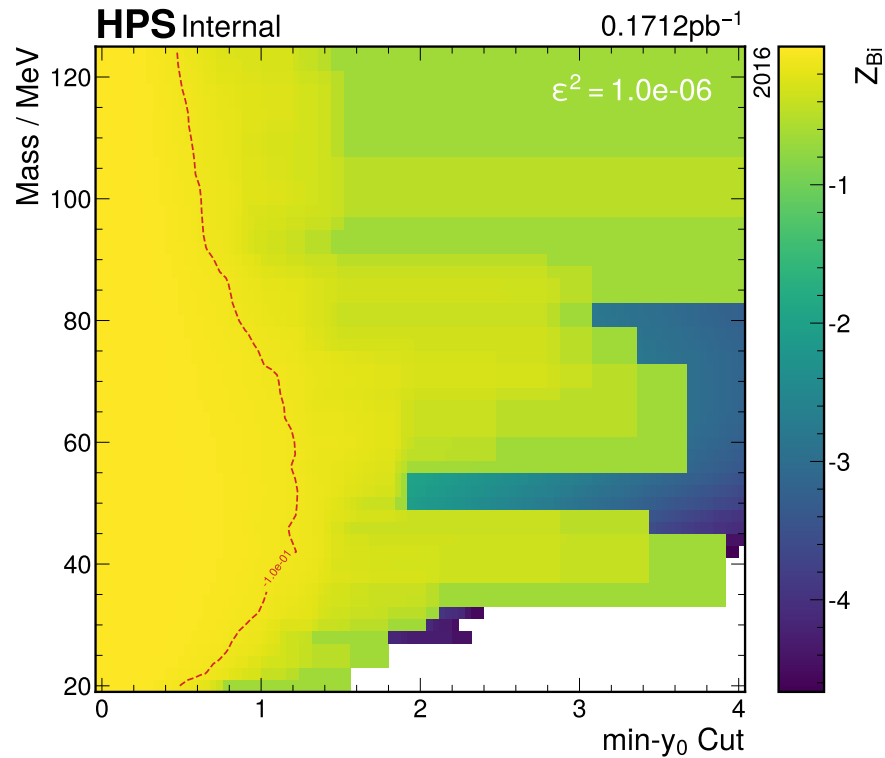
$$Z_{Bi} = \sqrt{2}\text{erf}^{-1}(1 - 2B_{reg}(0.5, S + B, 1 + B))$$

# Optimize Cuts Independently

Choosing  $\epsilon^2 = 10^{-6}$

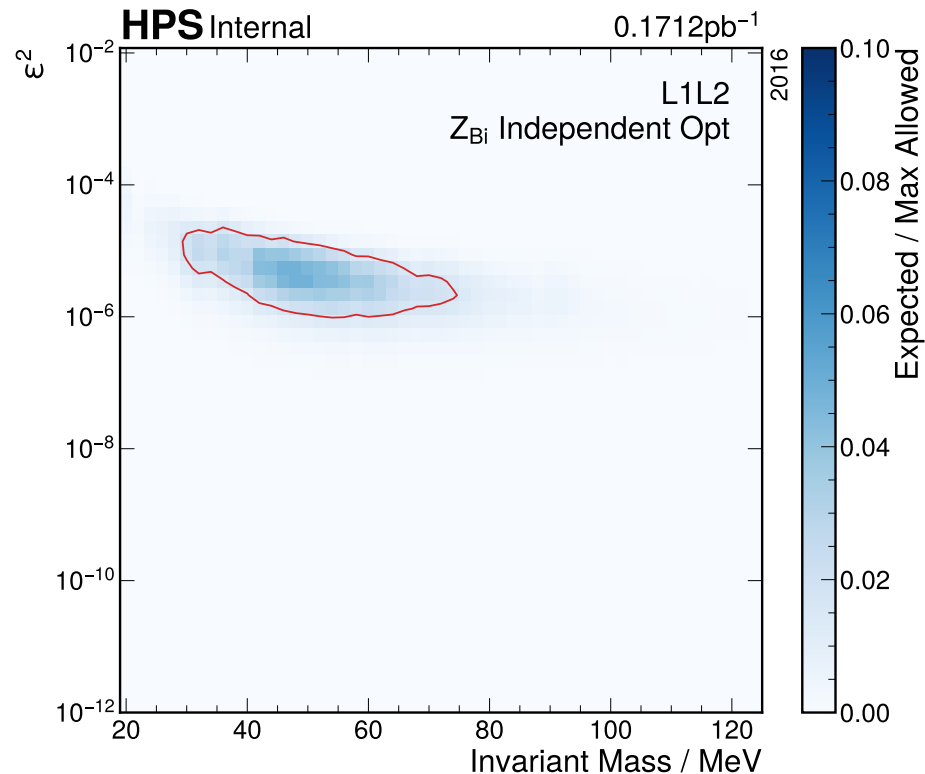
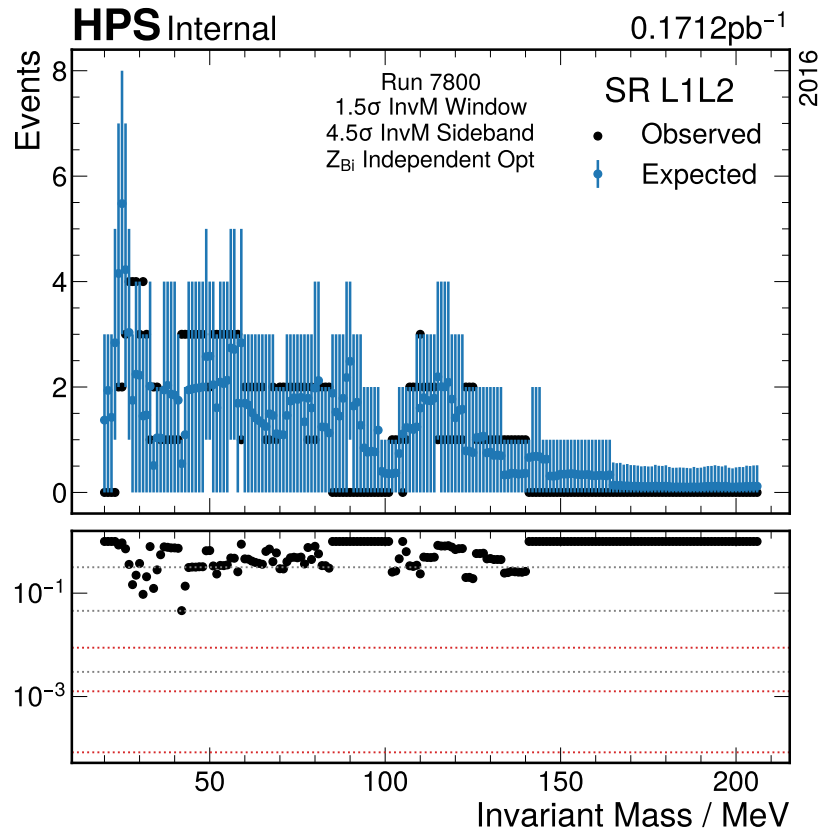


Choose cuts (red line) where  $\Delta Z_{Bi} < 10^{-4}$



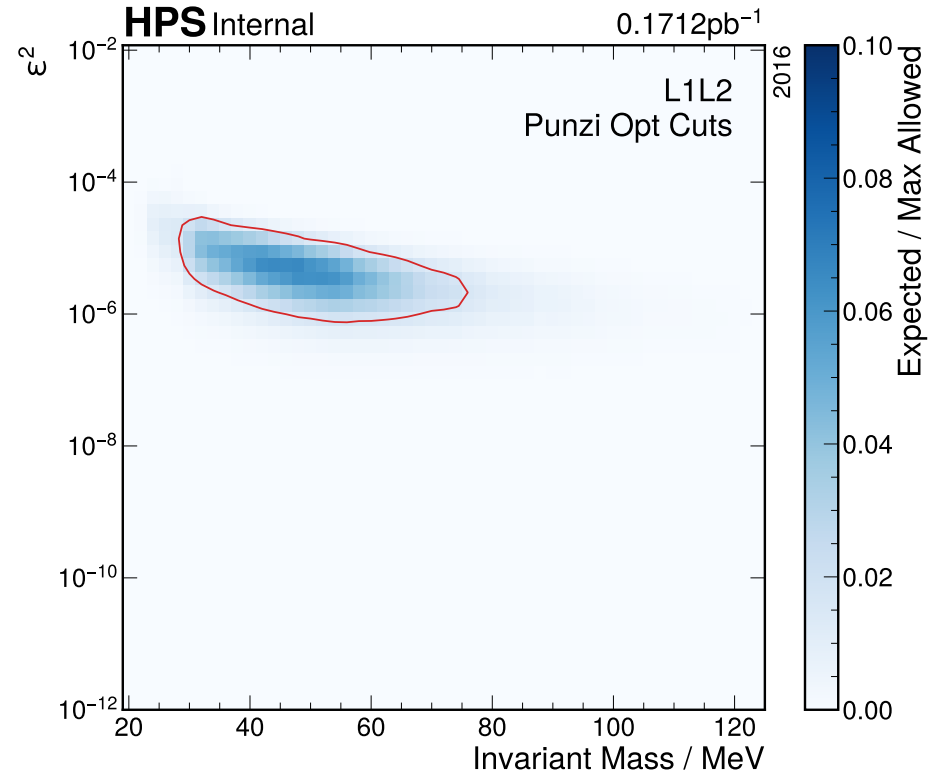
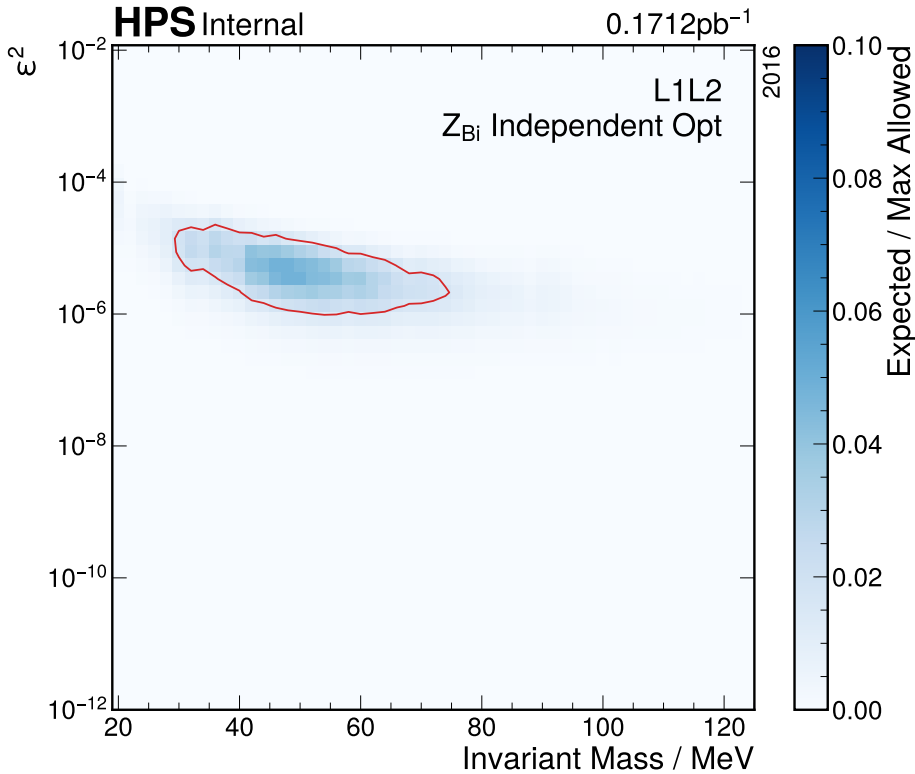
Choose cuts (red line) where  $Z_{Bi} > -0.1$

# Reach of $Z_{Bi}$ -Optimized Cuts





# Comparison of Reaches



## Optimization Strategy

- Staged approach – optimize  $\min-y_0$  *after applying* VPS cut
- Could weight Punzi FoM by decay weighting along  $z$  (pretty close algorithmically to  $Z_{Bi}$ ) or optimize Punzi within each  $z$  bin (cuts are functions of  $z$ )

## Exclusion Estimate

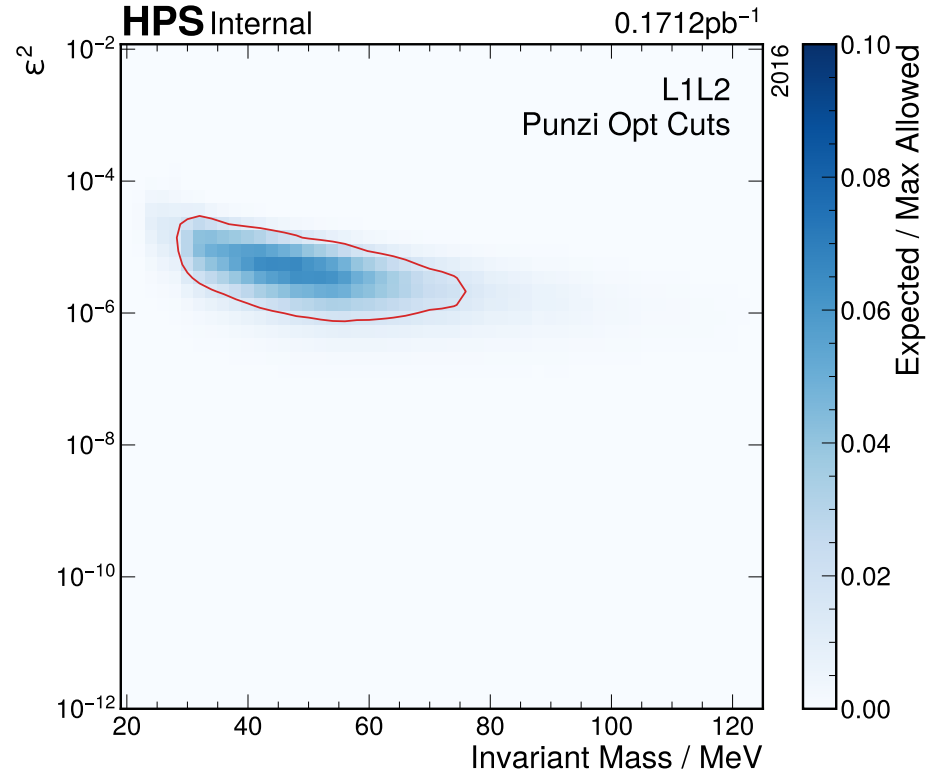
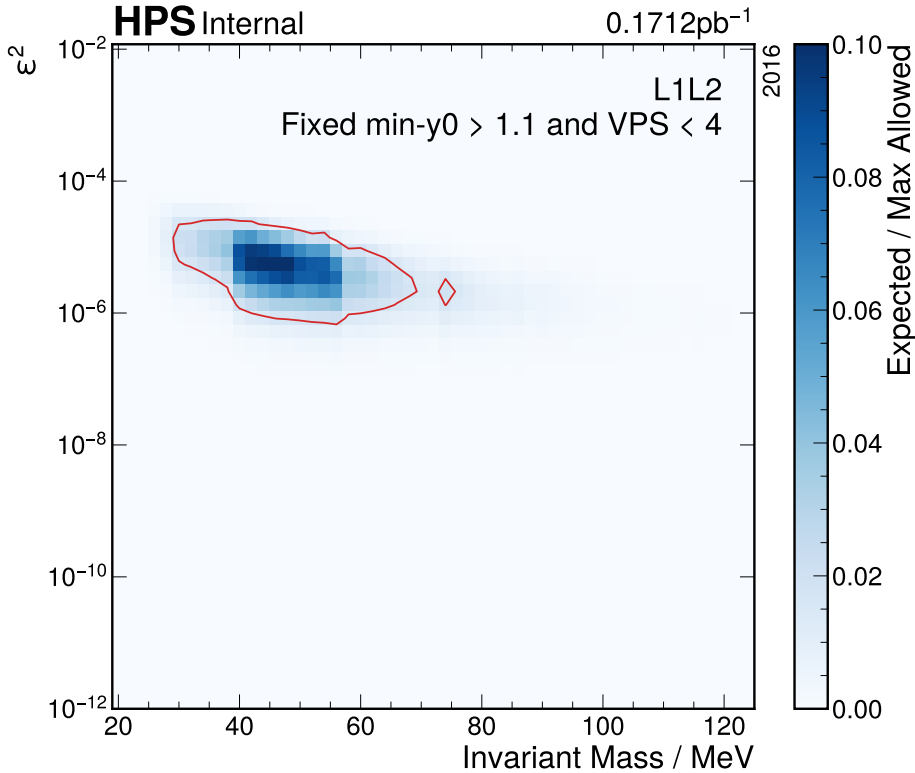
- Alic has been seeing hints that drawing the exclusion estimate at 0.1 in Expected / Allowed for 10% is overly optimistic

## Additional Material

- Various distributions (mass vs  $z$ ,  $z$  vs  $\min-y_0$ ) as selections are made
- Statistical combination of L1L1 and L1L2 exclusion estimates

# Questions

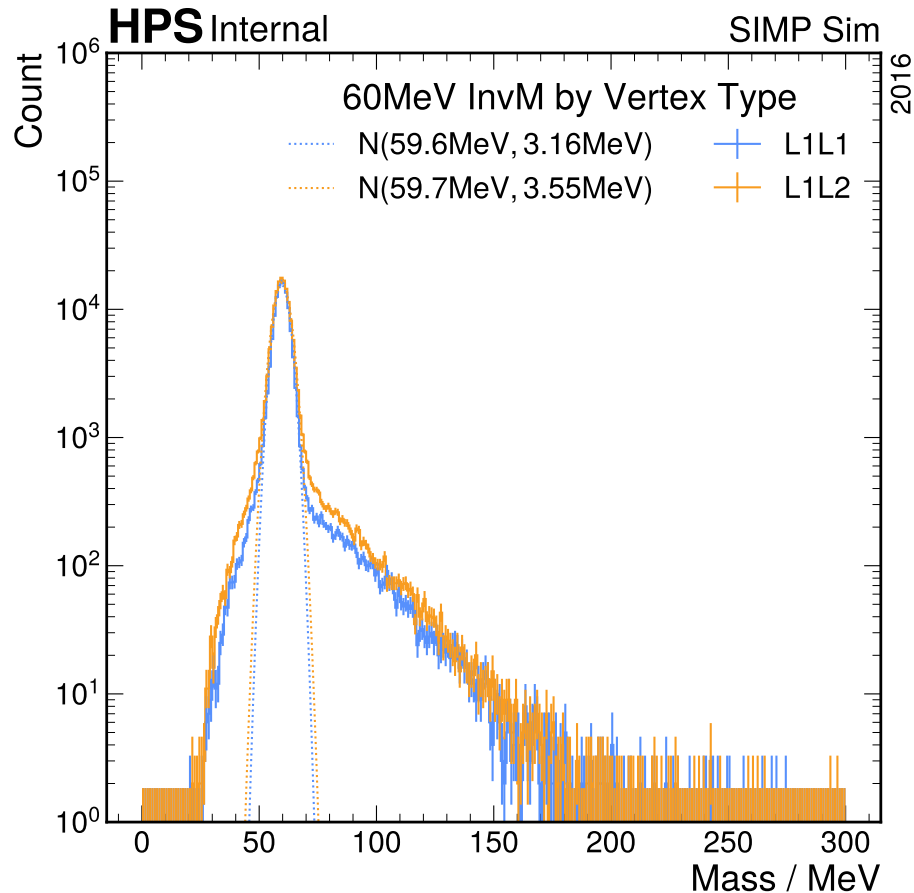
# Comparison of Reaches



To help contextualize reach estimates, it is helpful to compare the luminosity of these two reconstruction categories.

Category	$N_{7800}$	Ratio to L1L1
L1L1	2216982	1.0
L1L2	1445740	0.65

Table: Luminosity comparison between the two reconstruction categories being studied.  $N_{7800}$  is the number of pre-selected vertices that correspond to the given reconstruction category. No other selections (for example, on  $P_{\text{sum}}$ ) were made.

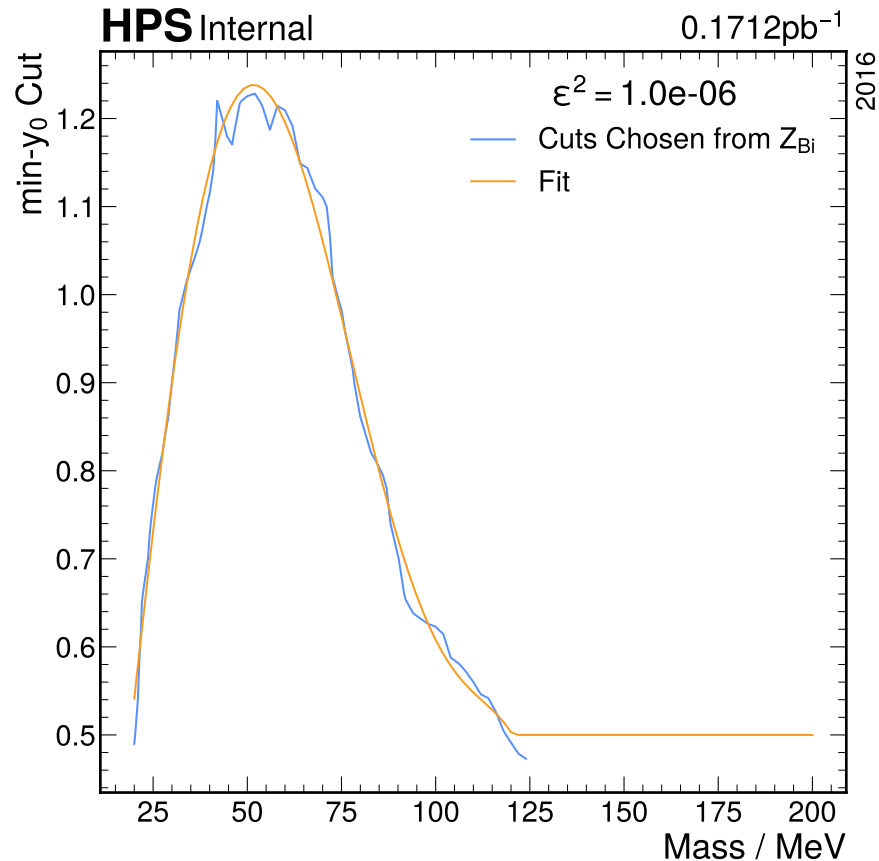
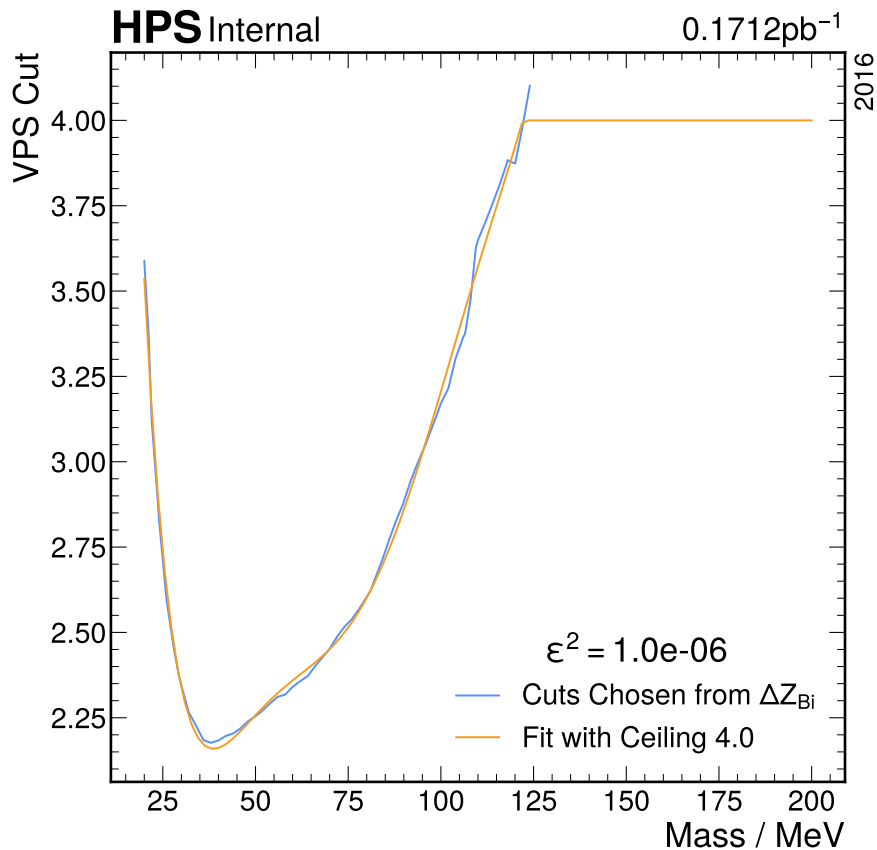


## Selection

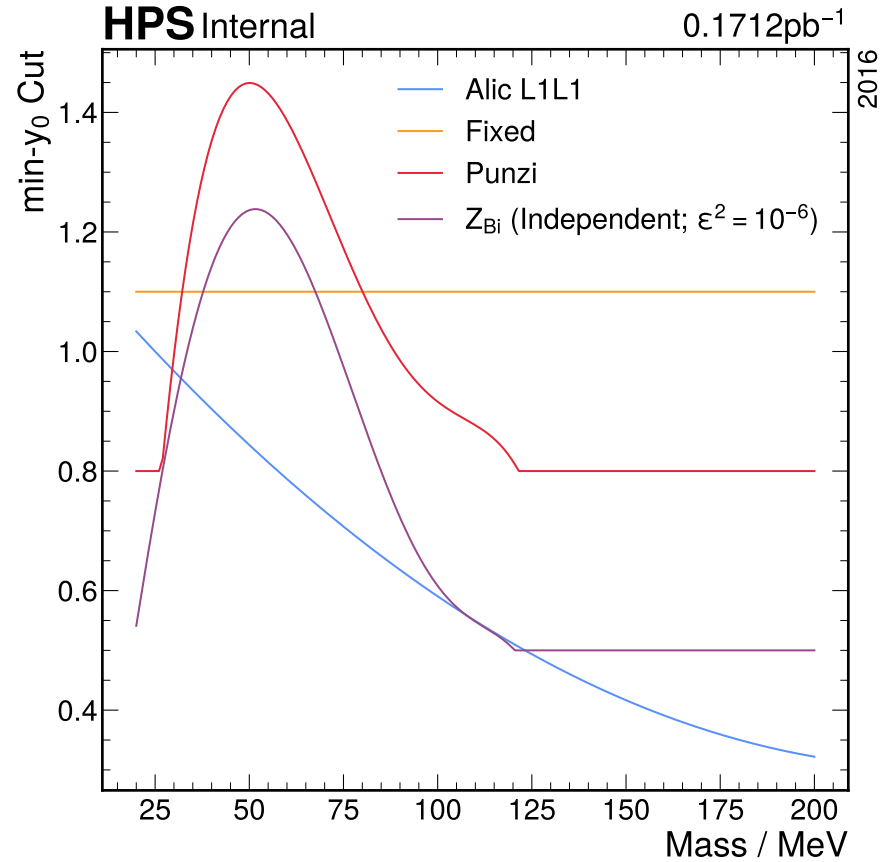
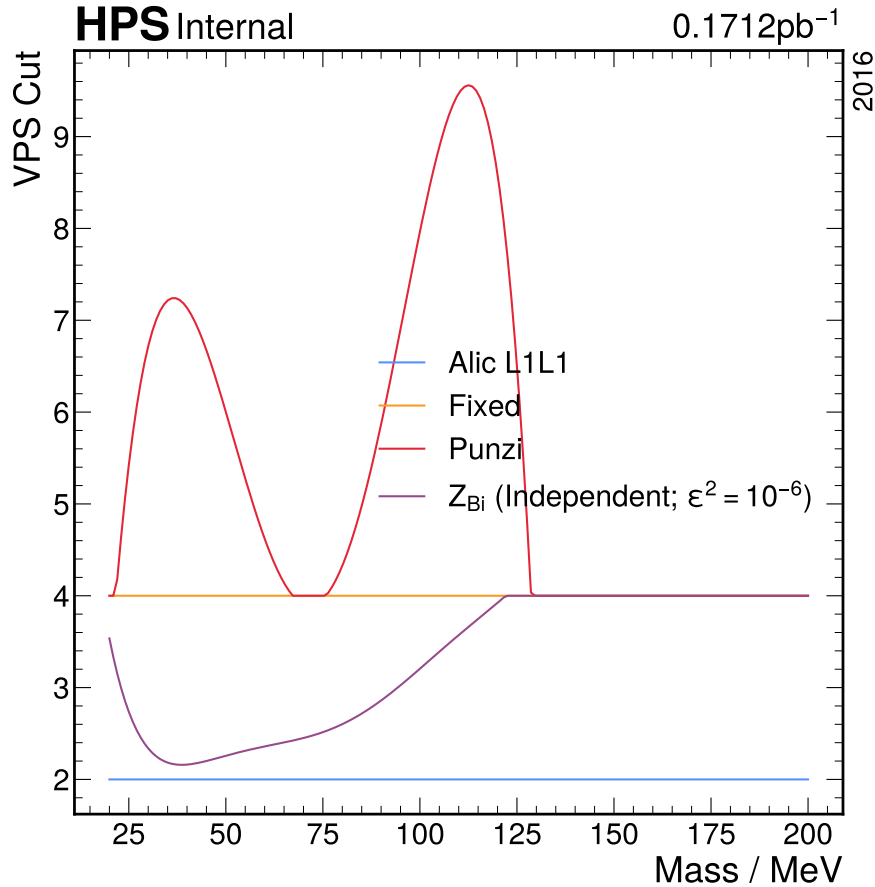
Pre-selected vertices – no truth matching done (hence the separation between L1L1 here and Alic’s L1L1 in black)

- Select core of distribution by calculating mean of histogram and dropping bins further than  $3\sigma$  away from mean
- Fit this core with a normal distribution to obtain  $\mu$  and  $\sigma$

# $Z_{Bi}$ Cut Values and Fits

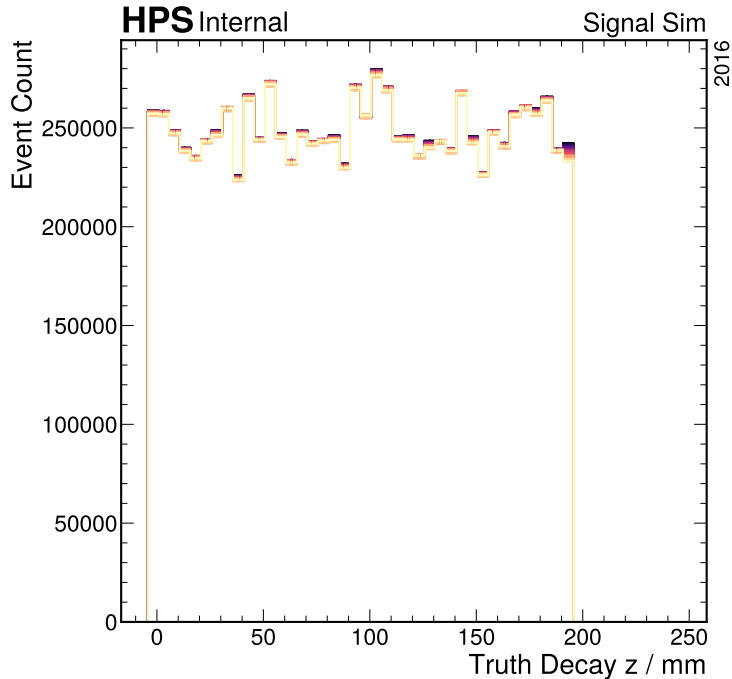


# Cut Comparisons



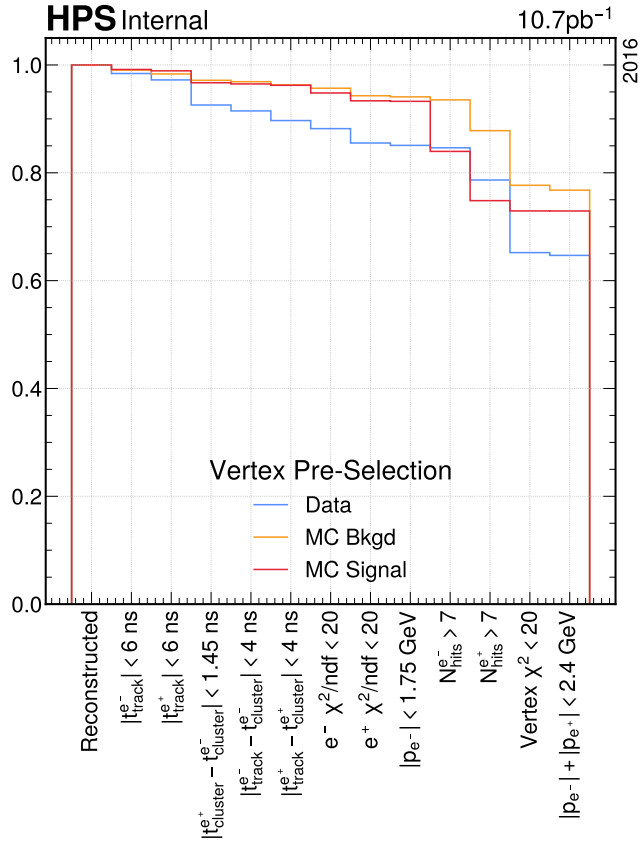


# Displacement of SIMP Signal Events

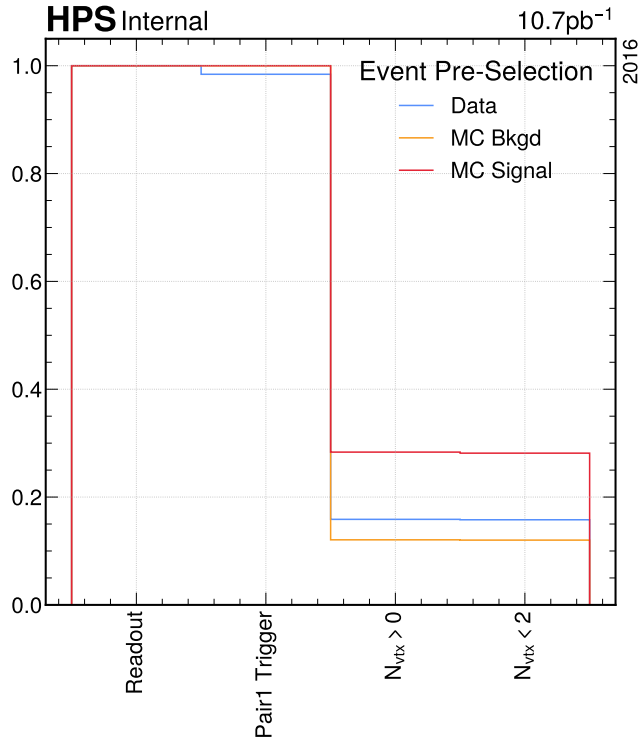


One of the first things I did to make sure samples didn't need to be created.

- Different colors correspond to different mass points
- Truth-level decay vertices sampled out until  $\sim 200$  mm
- Close to the same z position as L1, so I think these samples can be faithfully used to study the L1L2 selection
- Identical distribution across colors makes me think that the random seed determining the decay length was not changed, but I think that is okay since we normalize by this distribution during exclusion estimates anyways

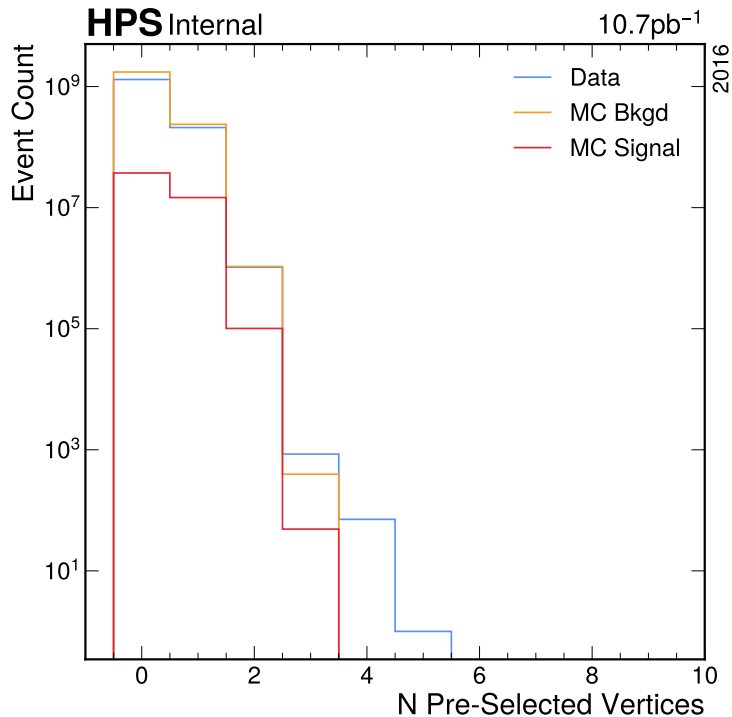


- Same Pre-Selection on vertices as developed and validated by Alic
- Seeing same efficiencies as documented within Alic's SIMP (L1L1) note

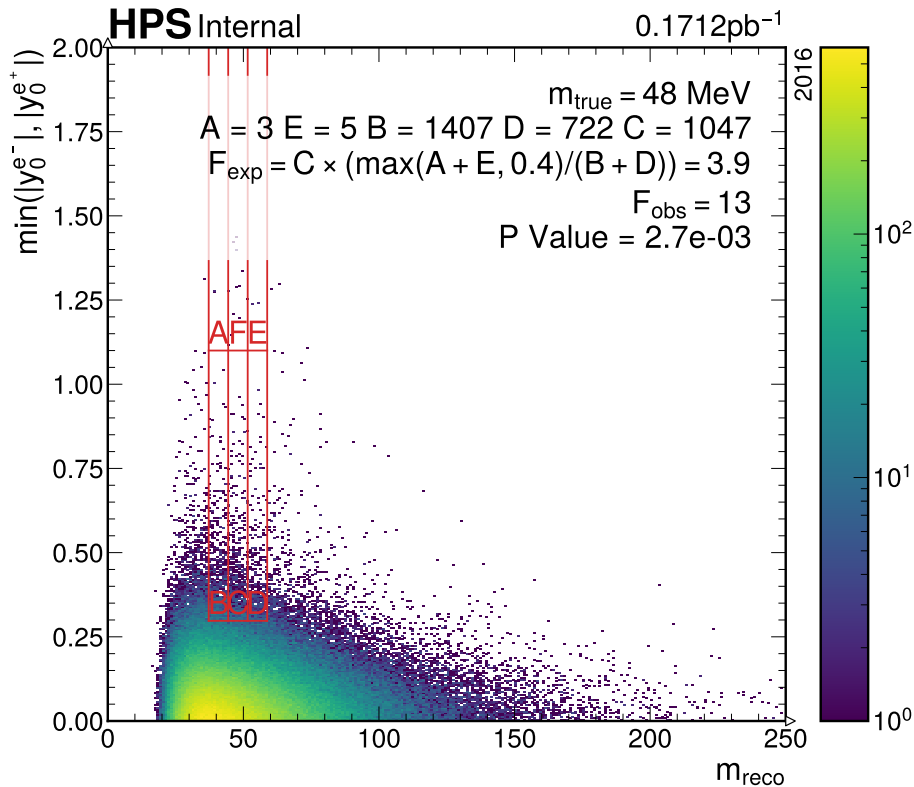


- Similar to first stage of Alic's event selection, although dropping reconstruction category requirement
- Largest effect is requiring at least one pre-selected vertex

# Number of Pre-Selected Vertices per Event



Event Pre-Selection basically amounts to choosing the events falling into the  $N = 1$  bin.  
Data is the only sample which has the additional requirement of the Pair1Trigger which has a small effect.



1. Fill histogram with data
2. Set mass edges at  $1.5\sigma$  and  $4.5\sigma$  (values optimized by Alic)
3. Set upper min- $y_0$  edge at cut value
4. Lower other min- $y_0$  edge (a.k.a.  $y_0$  “floor”) from the cut value until there are at least 1k events in region C
5. Calculate expected number of events in F and compare to observed number of events
6. Estimate p-value by throwing toy experiments in A+E (Poisson), B+D and C (Normal) and re-calculating F from these toys

## Basis

Proposed in [▶ PHYSTAT2003](#) by Giovanni Punzi where a FoM is designed to be maximized while improving *both* search and exclusion potential.

$$f_{\text{punzi}} = \frac{E}{\frac{a}{2} + \sqrt{B}}$$

where  $E$  is the signal efficiency,  $B$  is the background yield, and  $a$  is the desired confidence level of search or exclusion (in number of  $\sigma$ , currently using 3).

Two main benefits (from my perspective)

- Does not diverge as  $B \rightarrow 0$
- Does not require knowledge of absolute rate of signal

First idea is to simply define a new FoM that includes the decay weighting function.

$$f_{\text{DW}} = \frac{1}{\frac{a}{2} + \sqrt{B(t)}} \int_{z_{\text{target}}}^{\infty} D(z) E(z, t) dz$$

where

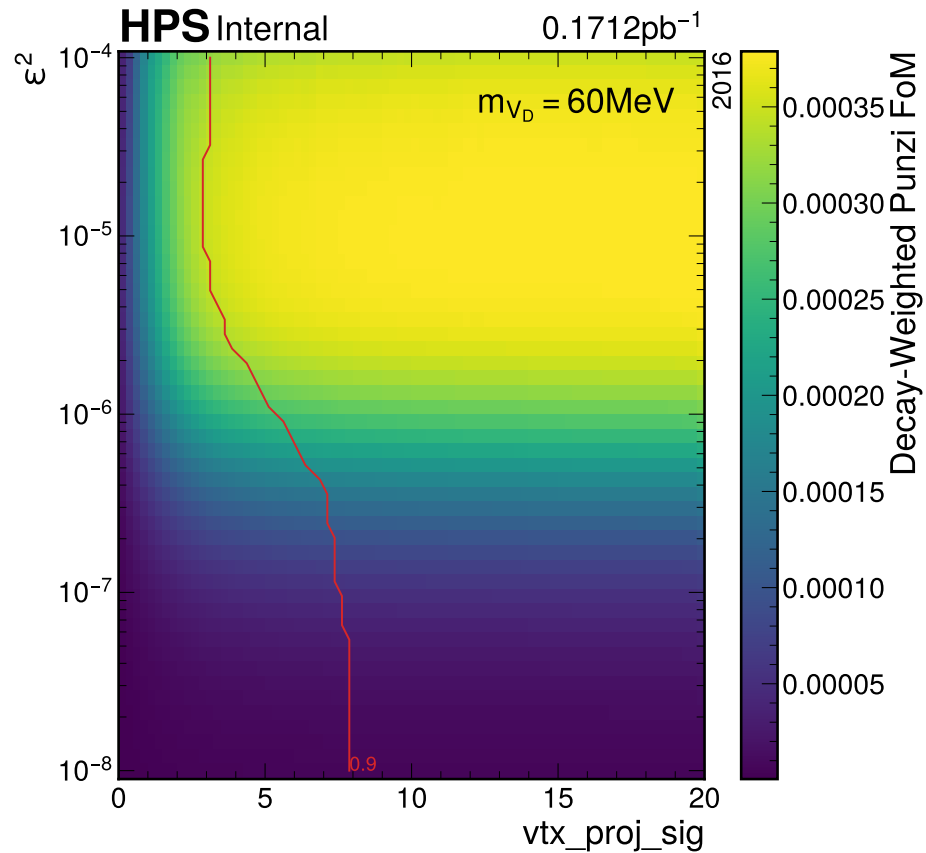
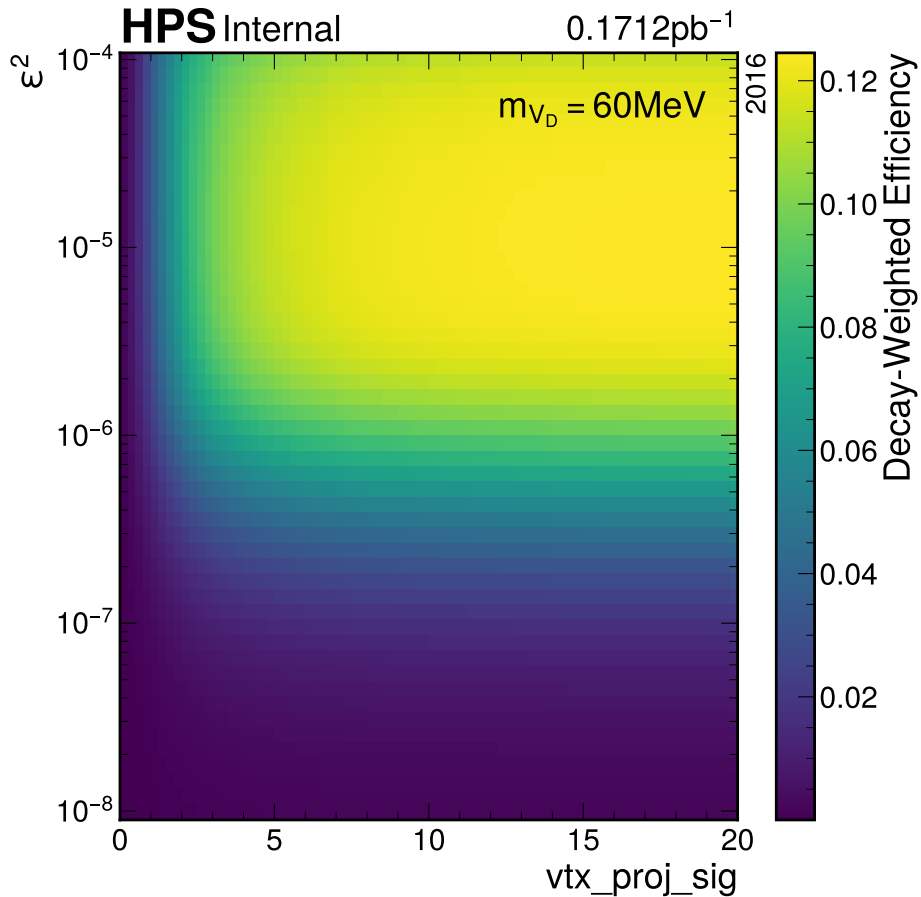
$$D(z) = \sum_{V \in \{\rho_D, \phi_D\}} \text{BR}(A' \rightarrow V \pi_D) \frac{\exp((z_{\text{target}} - z)/(\gamma c \tau_V))}{\gamma c \tau_V}$$

This becomes equivalent to  $f_{\text{punzi}}$  in the  $\epsilon \rightarrow 0$  limit where  $D(z)$  becomes flat and the events are equally weighted along  $z$ . Calling this “Decay-Weighted Punzi FoM” and the integral “Decay-Weighted Efficiency”.

## Issues

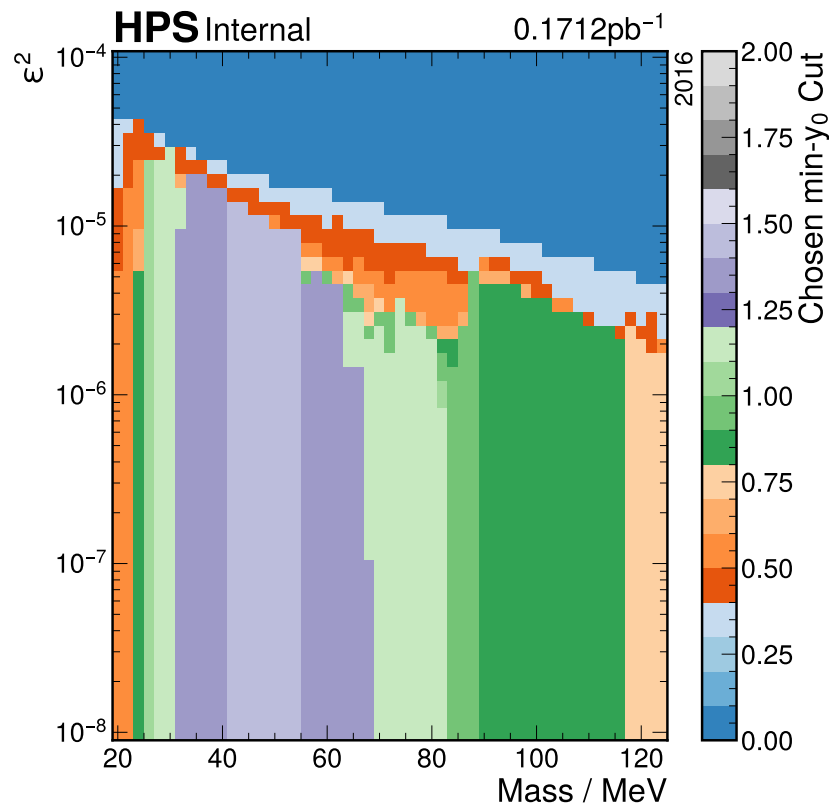
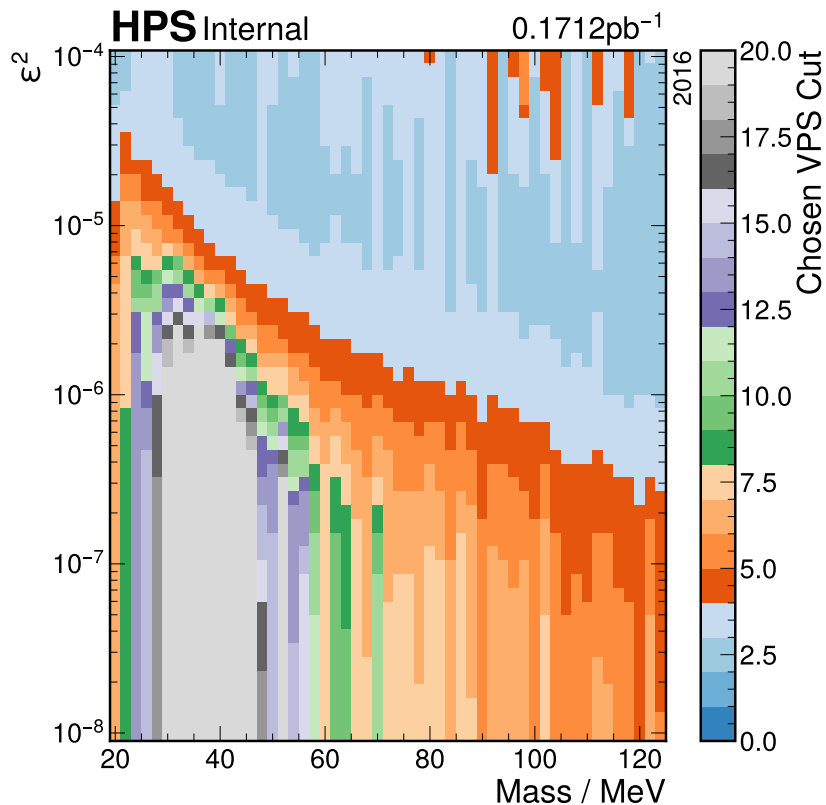
- May have to choose an  $\epsilon^2$  value to optimize for
- Pure maximum is often attained by removing the cut  $\rightarrow$  look for where the FoM “flattens” out (i.e. tightening the cut does not improve the FoM much anymore)  $\rightarrow$  choose cut that is the tightest cut getting to 90% of the maximum

# Example Decay-Weighted Punzi Calculation





# Cut Choices by $\epsilon^2$ and $m_{V_D}$



Chose VPS  $< 4$  and then applied it to optimize min- $y_0$ .