Investigate number of extra tracks for $\gamma \textbf{\textit{p}} \to \eta \pi^- \Delta^{++}, \ \eta \to \gamma \gamma$

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Motivation

- Recent discussions on how many extra tracks are needed to be kept
- Pros: smaller analysis trees saves disk space and processing time
- Cons: May lose some information
- Procedure: run ReactionFilter with 5 different flags:
 - 1. B4_M17 ("standard")
 - 2. B4_M17_S0_T0 (no extra showers or tracks)
 - 3. B4_M17_S1_T0 (1 extra shower, no extra tracks)
 - 4. B4_M17_S0_T1 (no extra showers, 1 extra track)
 - 5. B4_M17_S1_T1 (1extra shower, 1 extra track)
- Try to get a handle on how much information we lose by cutting on tracks

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Distributions after Reaction Filter and cuts (unused E<100 MeV)



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Distributions

Size (GB)	%	DSelector Time (4 threads)	N $\eta\pi$	%
16	100	42m	33561	100
0.6	3.5	2m	24051	72
2.0	12.5	бт	28314	84.4
0.8	4.7	2m	26947	80.3
2.7	16.9	7m	31694	94.5
	Size (GB) 16 0.6 2.0 0.8 2.7	Size (GB) % 16 100 0.6 3.5 2.0 12.5 0.8 4.7 2.7 16.9	Size (GB) % DSelector Time (4 threads) 16 100 42m 0.6 3.5 2m 2.0 12.5 6m 0.8 4.7 2m 2.7 16.9 7m	$\begin{array}{llllllllllllllllllllllllllllllllllll$

- Save 83% of disk space yet only lose 5% of $\eta\pi$ events with keeping only 1 extra shower and track
 - For keeping analysis trees locally (eg IU cluster, own space on farm), this option could be ideal
 - Does not create a bias in mass and angular distributions
 - 2018-01 trees go from 5TB too 1TB (for S=1, T=1)
 - Would go to 196 GB for S=0, T=0 (my be good option for first looks)

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