

Top Threshold Scan & Luminosity Spectrum at CLIC

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Motivation

- The top quark mass enters in precision SM calculations (Higgs mass, ...): Good to determine it with high precision
- Theoretically clean definitions of the mass ($1S$, $M\overline{S}$) are not what we typically measure in experiments (Invariant mass of decay products): Substantial theory uncertainty, on the level of several 100 MeV
 - With other systematics: LHC limit probably on the GeV level

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 - With other systematics: LHC limit probably on the GeV level
- Assume we want to do better than this (precision physics!):
 - Better theory - Promising first results, but this is highly non-trivial
 - Theoretically well understood measurement: Threshold scan with e^+e^-
ILC / Tesla studies show that substantially better than 100 MeV total uncertainty can be achieved - Does this also work for CLIC?

Threshold Scan - How it works

- The simplest version: Measure the cross section for top pair production at several energies, compare to NNLO QCD theory calculations
 - Need precise understanding of luminosity, and of the effects of the luminosity spectrum on the measured cross section
 - Need to be able to measure at several (O 5 - 10) energy points near threshold (within a 5 to 10 GeV energy window) with reasonable integrated luminosity per point (a few to a few 10 fb^{-1})

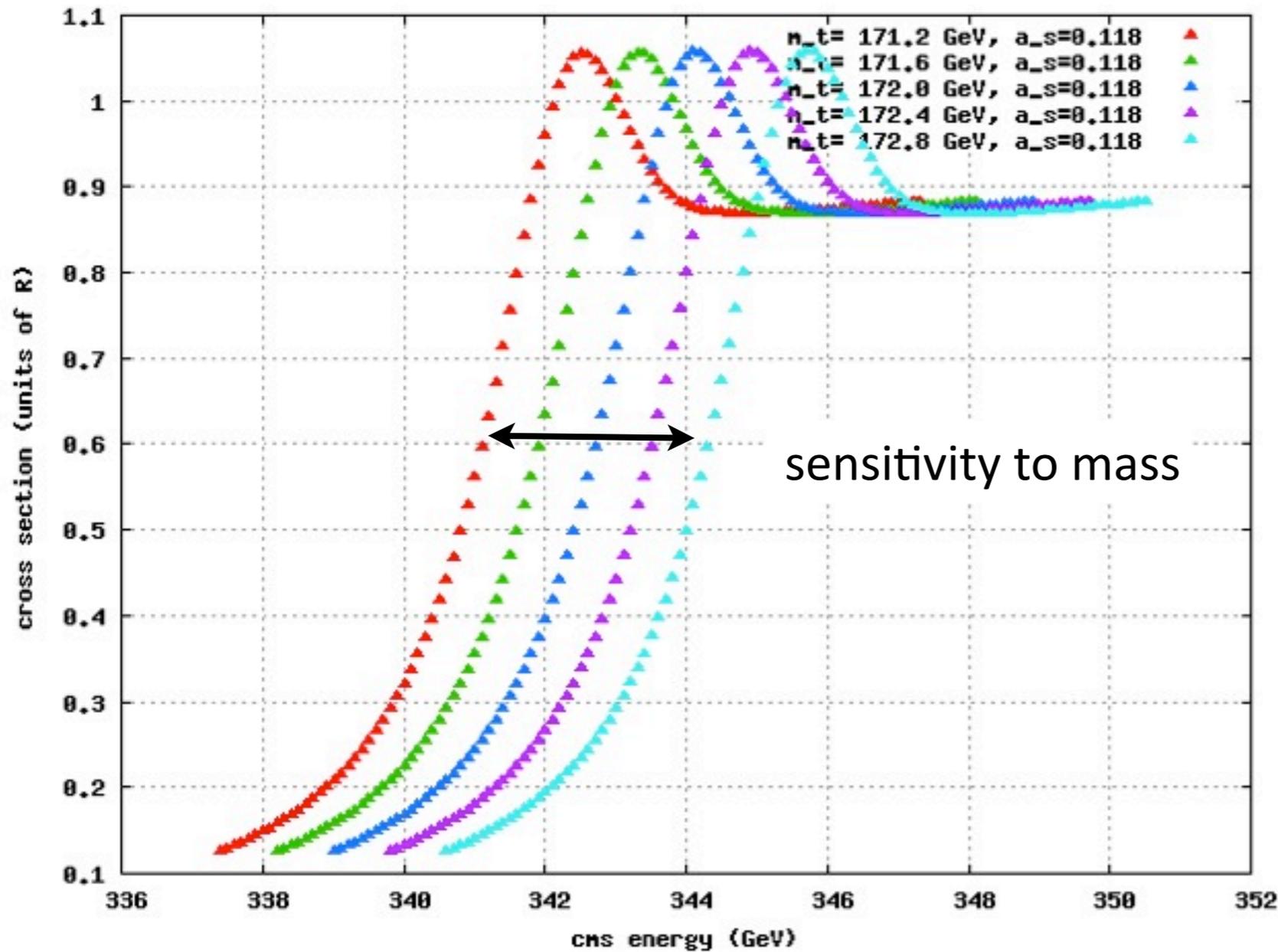
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Side remark:

- We do not need energy steps on the sub-100 MeV level
- Cross-sections are small: No point in measuring at more than ~ 10 energy points

Threshold Scan: How it works



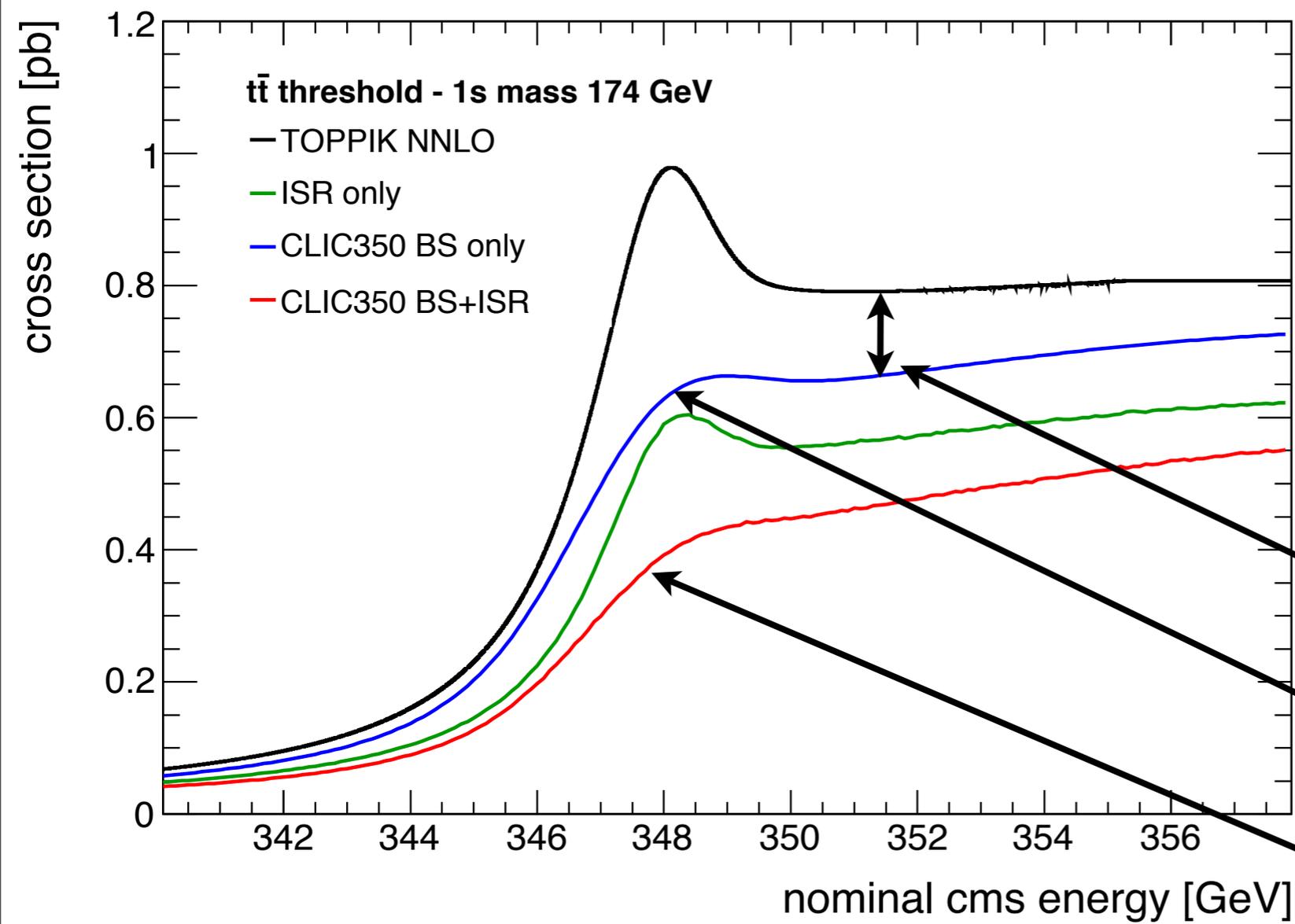
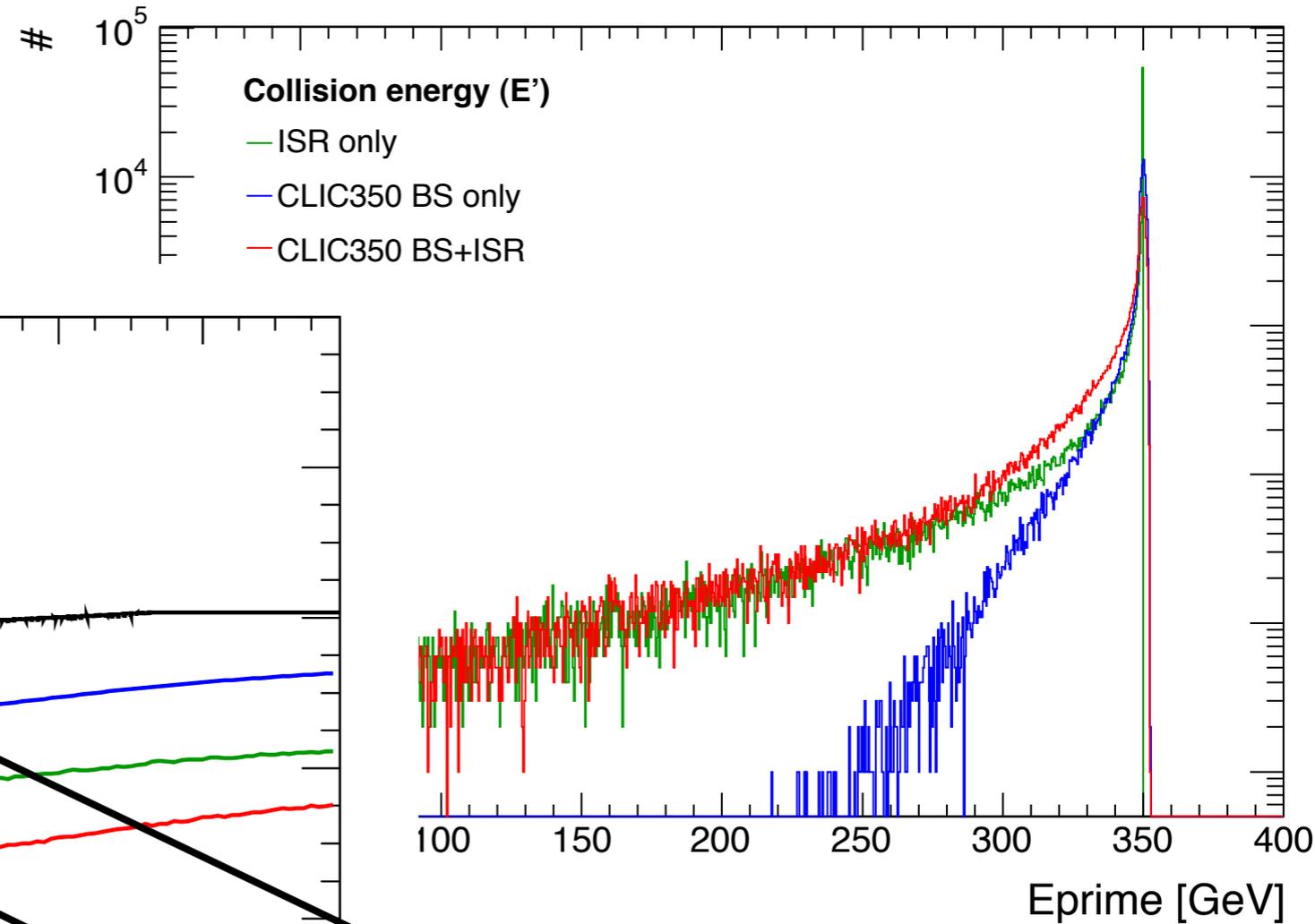
theory $t\bar{t}$ cross section
calculated with TOPPIK
(Hoang, Teubner)

- Access mass by determining position of edge - shifts linearly with top mass

Threshold and Luminosity Spectrum

- The threshold gets distorted by two effects:

- ISR
- Luminosity spectrum

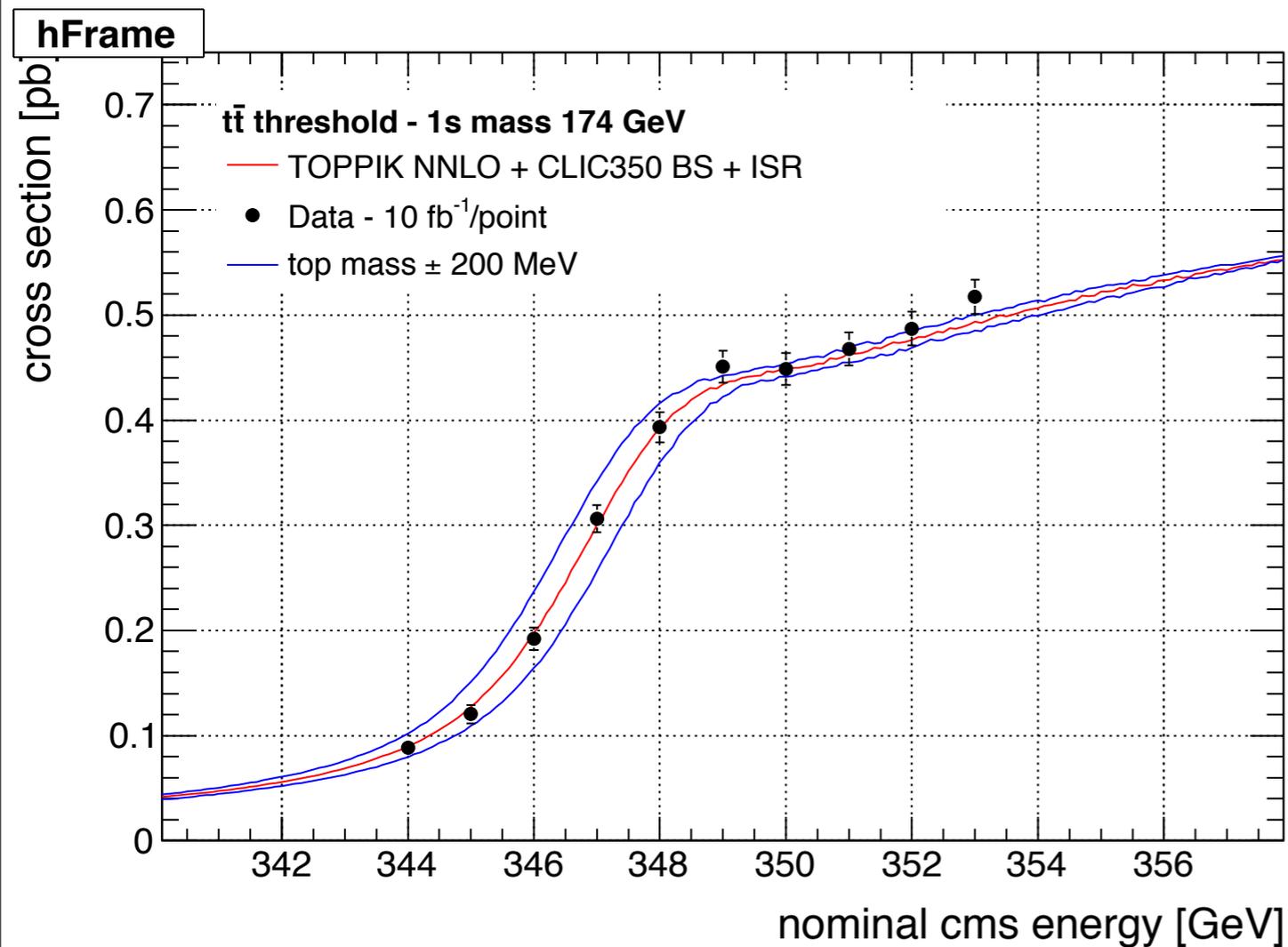


lowering of cross section
 smearing of peak,
 broadening of edge
 further smearing and
 reduction due to ISR

The Study of a Top Threshold Scan at CLIC

- The idea: Measure the top mass from the cross section at the threshold
 - other variables exist as well, and provide additional sensitivity (A_{FB} , top momentum distribution, ...) - not studied at present
- The problem: No full event generator for the top threshold publicly available
 - Generators like PYTHIA are leading order, plus hadronization: Incorrect threshold behavior (ignores binding effects, ...)
- The approach: Generator-level study, with fully simulated efficiencies
 - Obtain cross section (and with that signal yield) from TOPPIK calculations
 - Take selection efficiencies and background fractions from full simulations, including overlaid background, determines error bars for background-subtracted data points
 - At the moment: Simulations for top pairs in PYTHIA slightly above threshold still in progress - Using efficiencies and scaled background levels from 500 GeV study

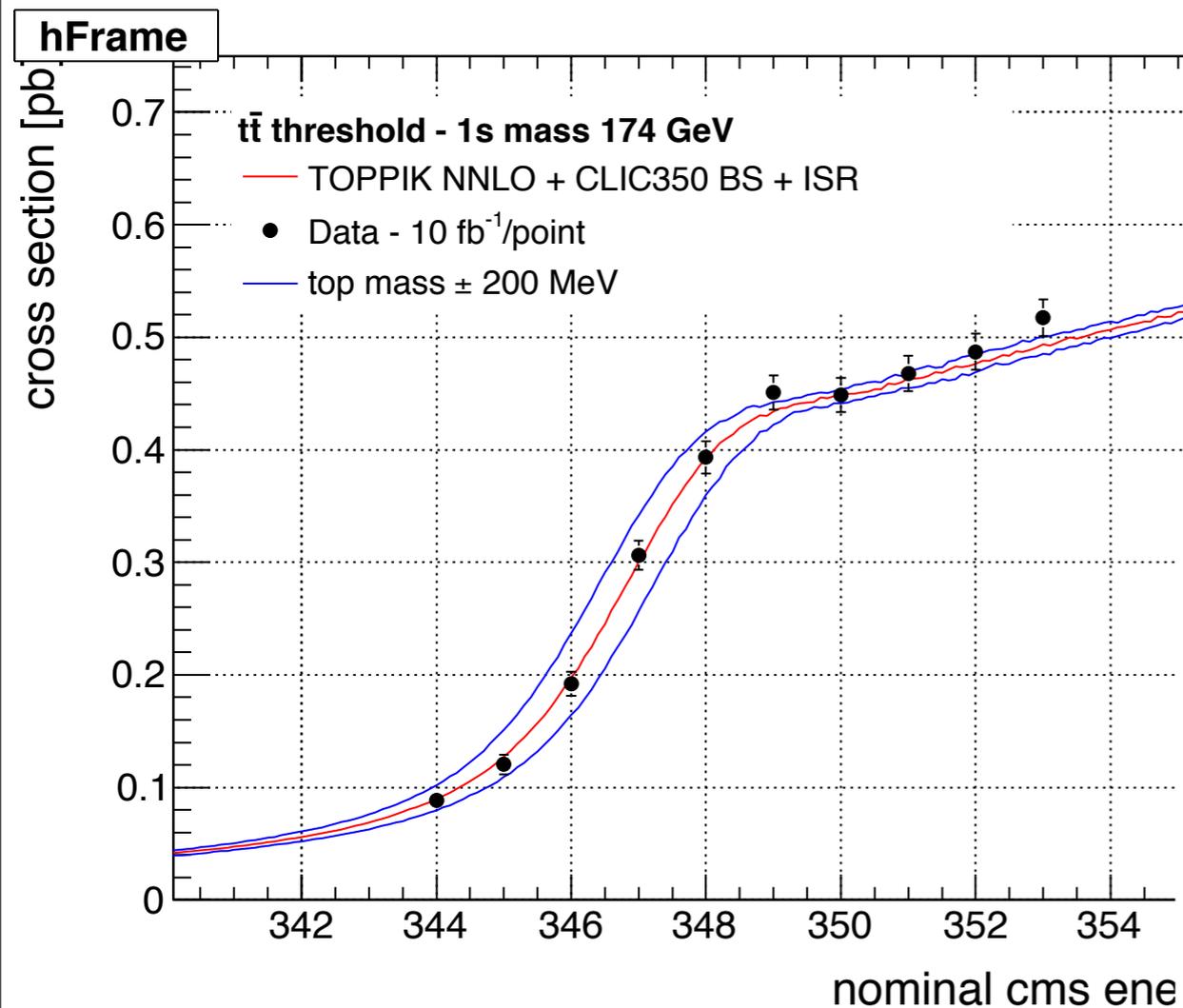
How Important is the Luminosity Spectrum?



- Preliminary results:
 - ~ 33 MeV statistical error on mass
 - ~ 20 MeV systematic uncertainty due to α_s (using PDG error)

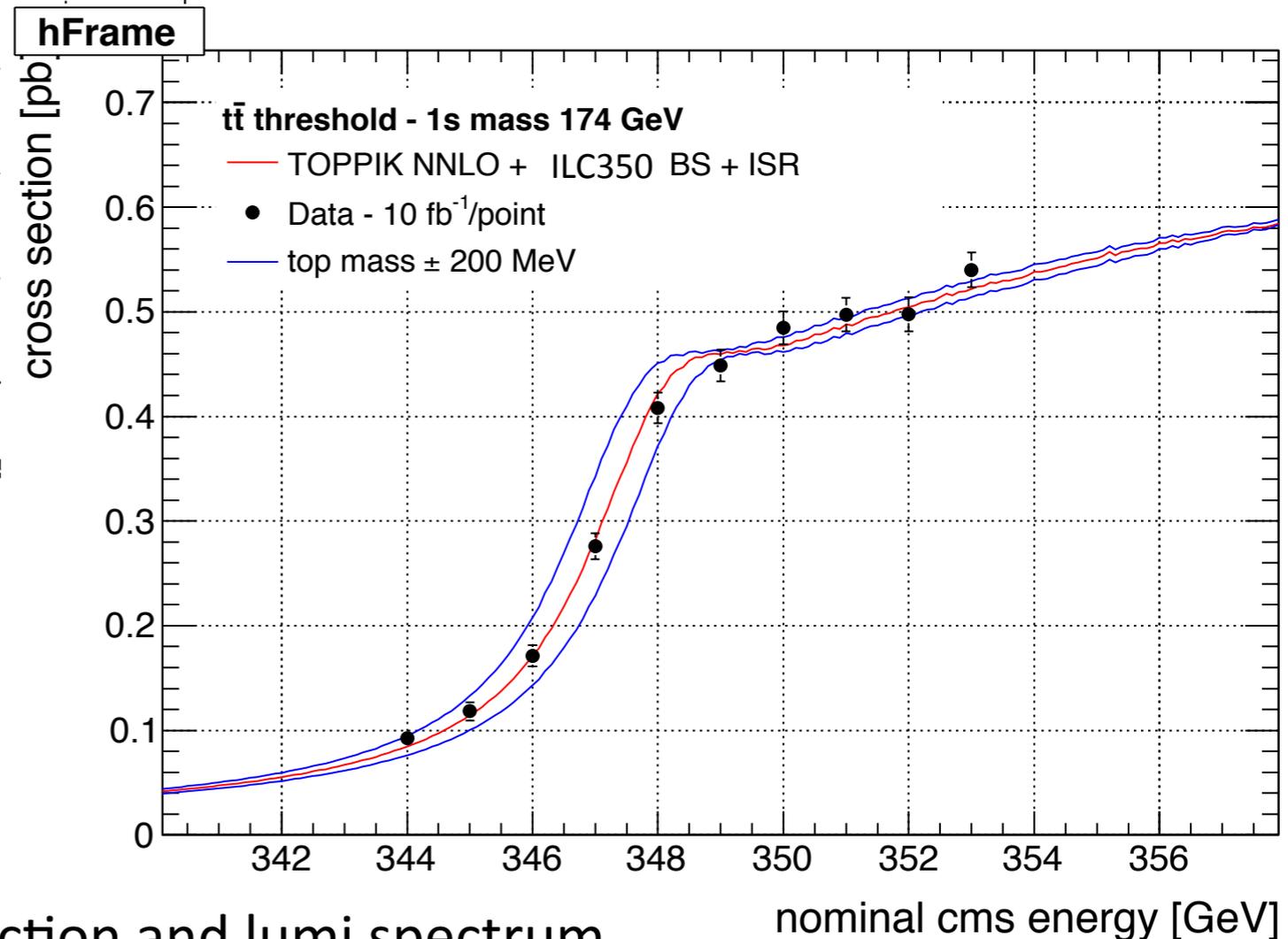
Fits assume perfect knowledge of cross section and lumi spectrum

How Important is the Luminosity Spectrum?



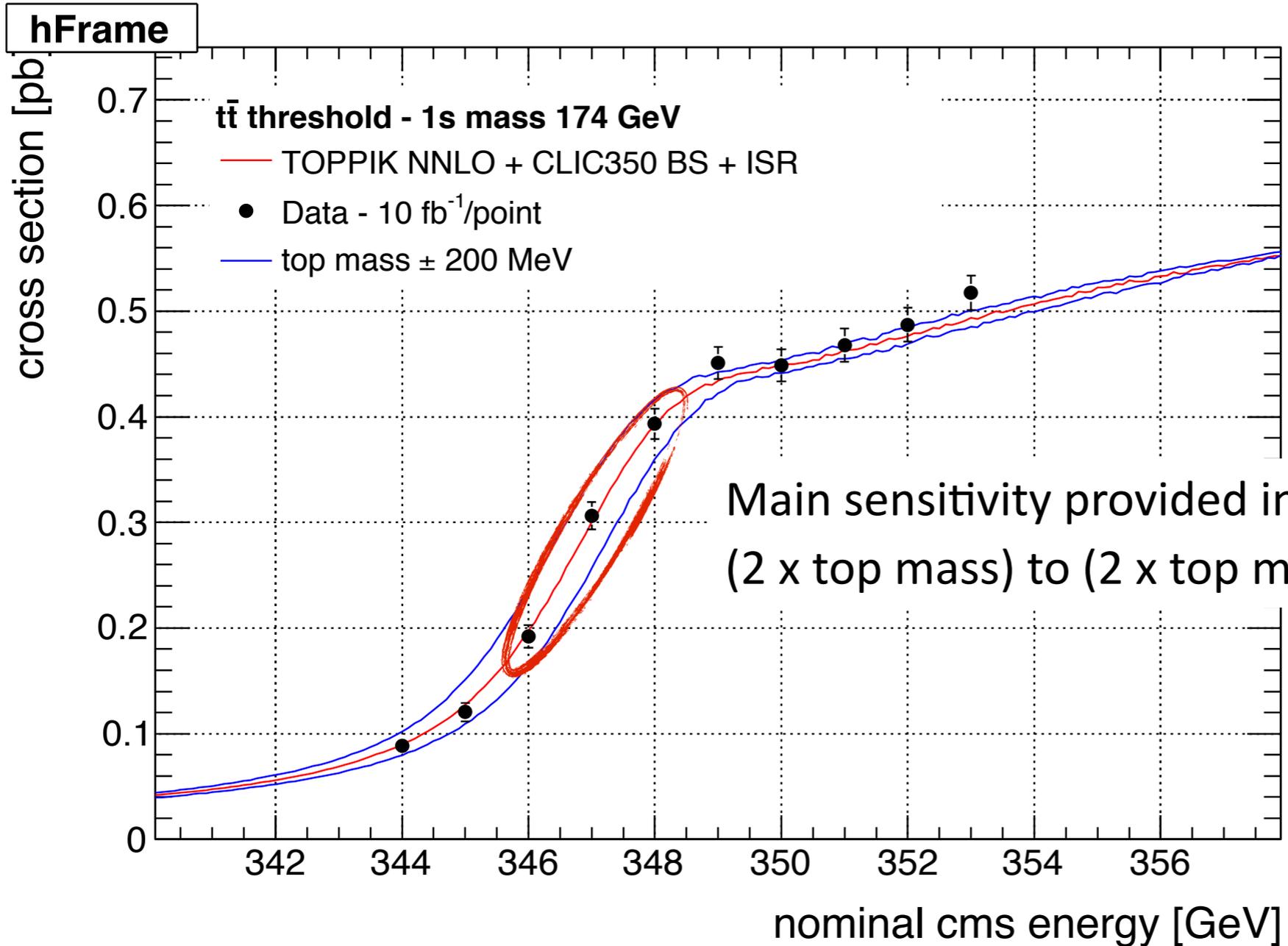
- ILC luminosity spectrum:
 - ~ 30 MeV statistical error on mass
 - ~ comparable uncertainty due to α_s

- Preliminary results:
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Fits assume perfect knowledge of cross section and lumi spectrum

Mass Sensitivity and Scan Points

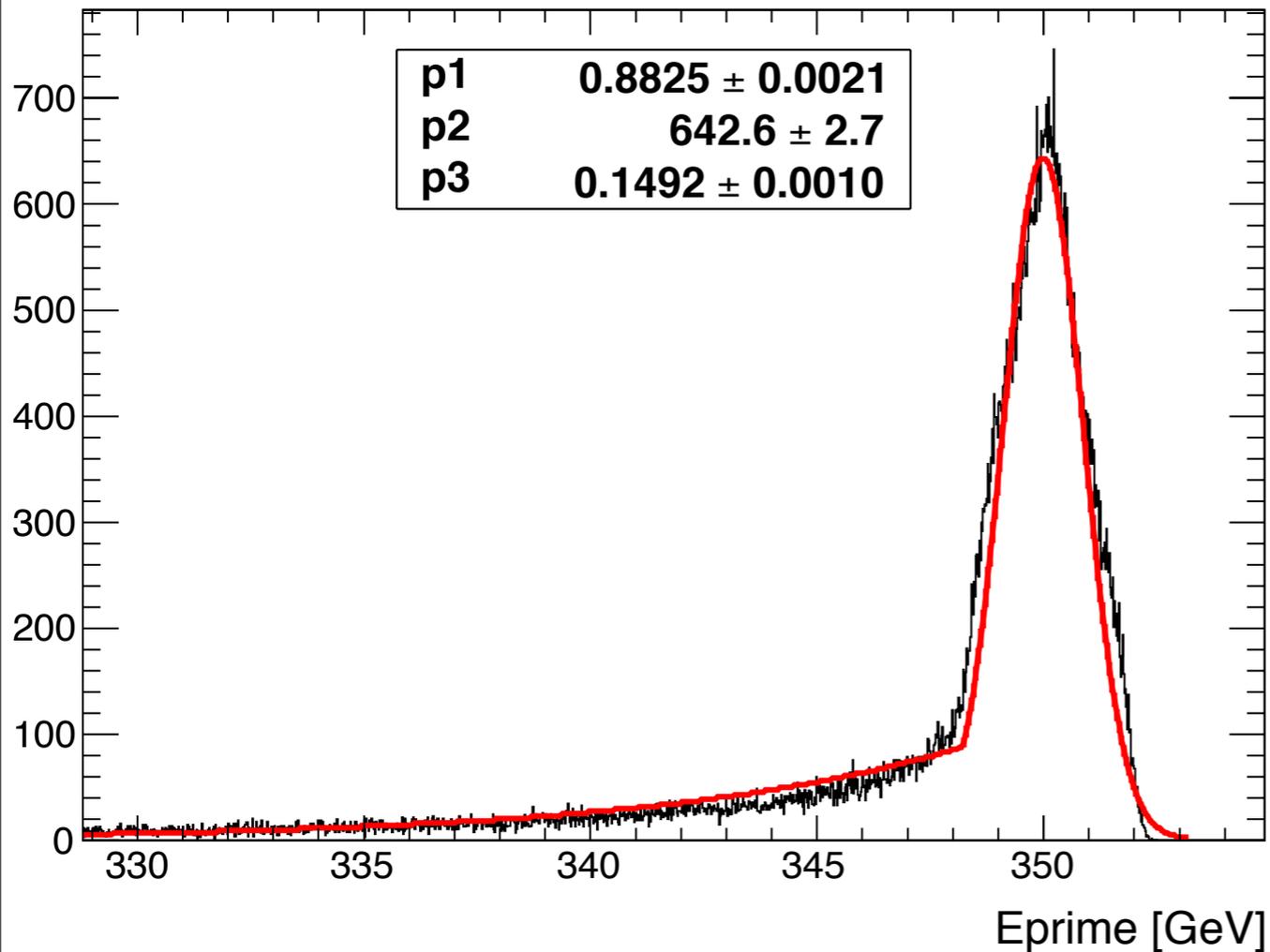


In this range:
0.2 pb change at CLIC
0.24 pb change at ILC
Not a dramatic difference!

- Optimization of scan points clearly possible
 - here: clearly too many points above threshold: potentially increases sensitivity to systematic effects from normalization uncertainties

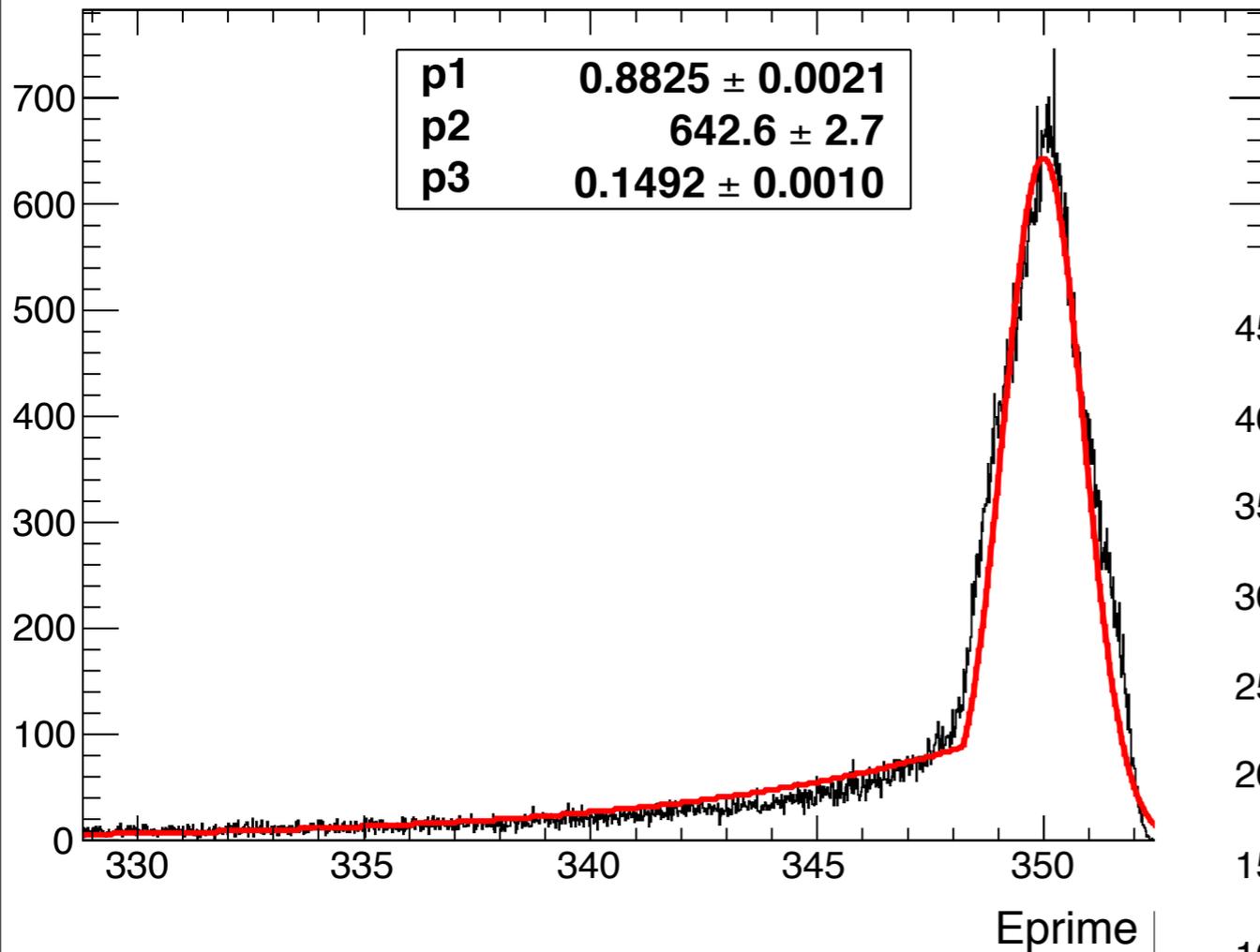
Crucial: Knowledge of the Luminosity Spectrum!

- A first simple test - Parametrize CLIC350 Lumi Spectrum
 - Gaussian peak, exponential tail (starting at -2σ for default spectrum)

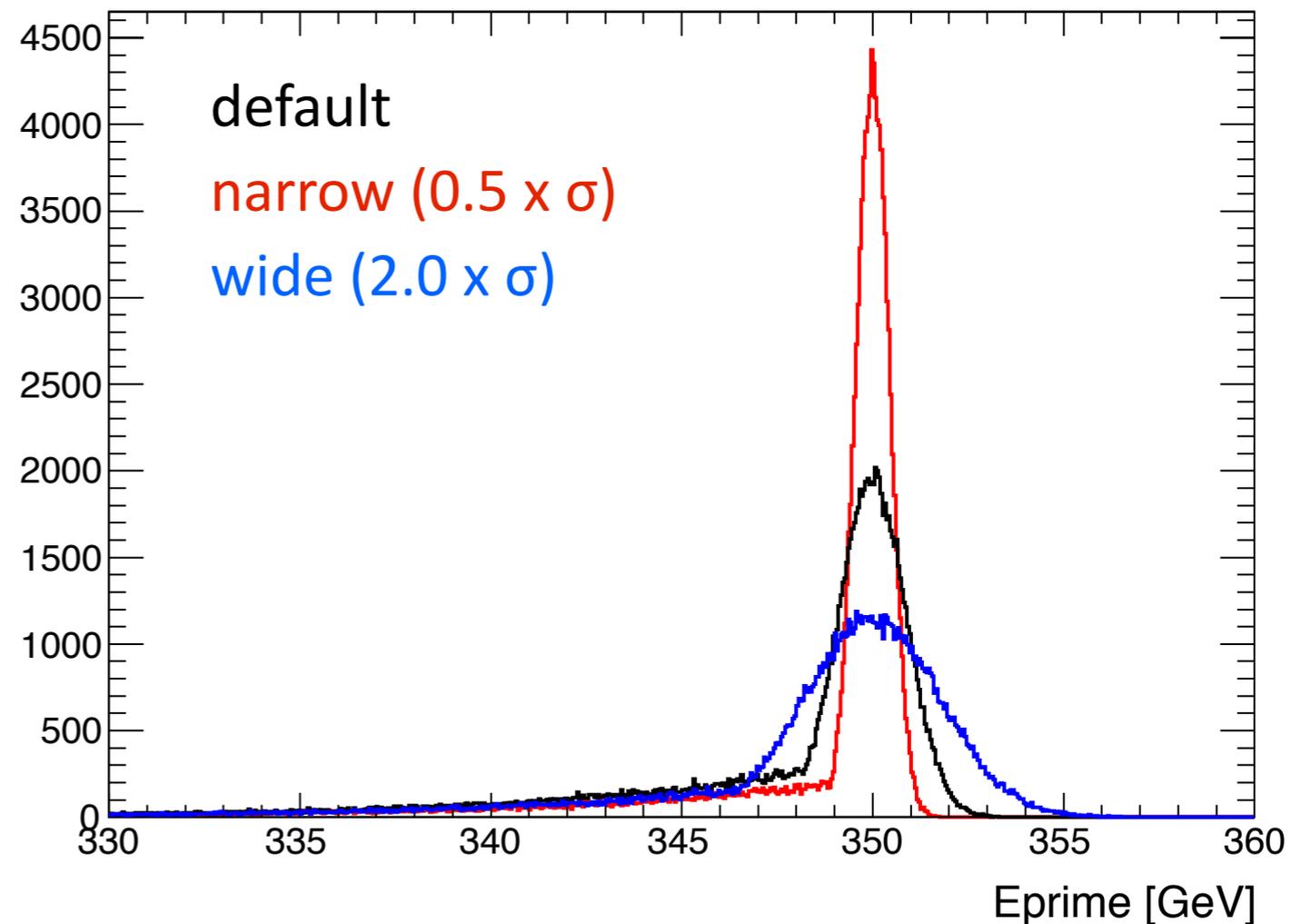


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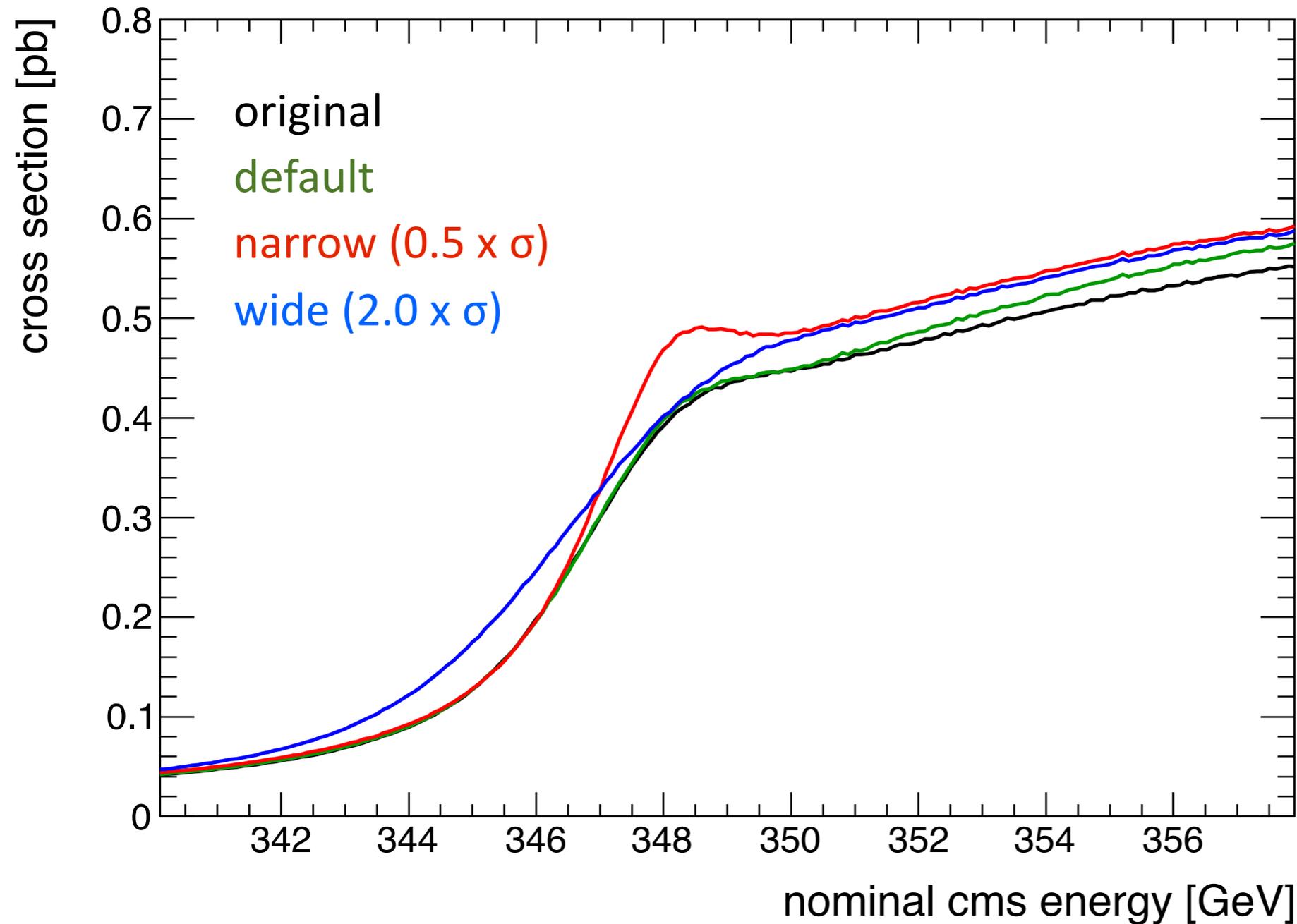
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Three parametrizations:



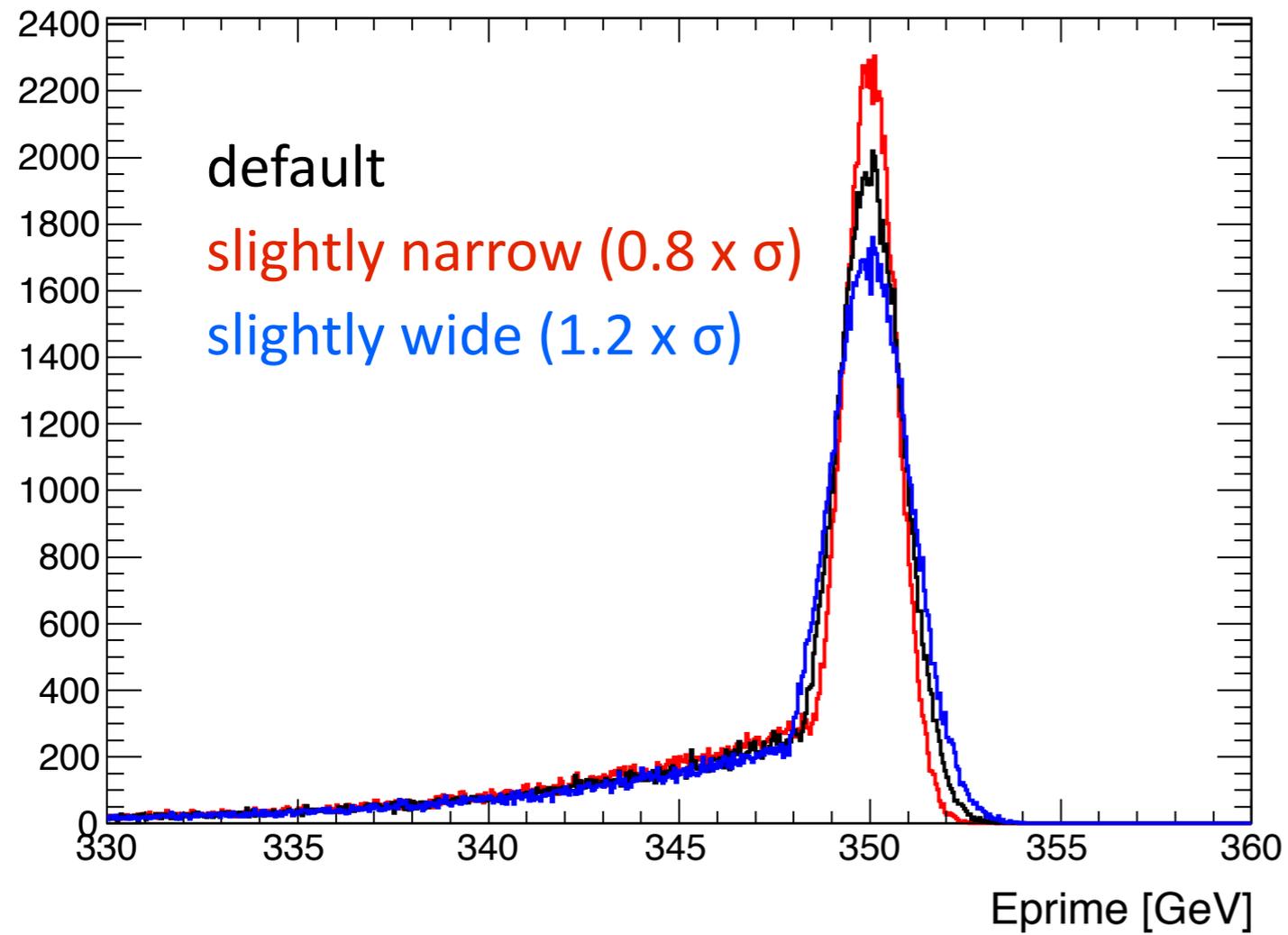
Effect on the Top Threshold



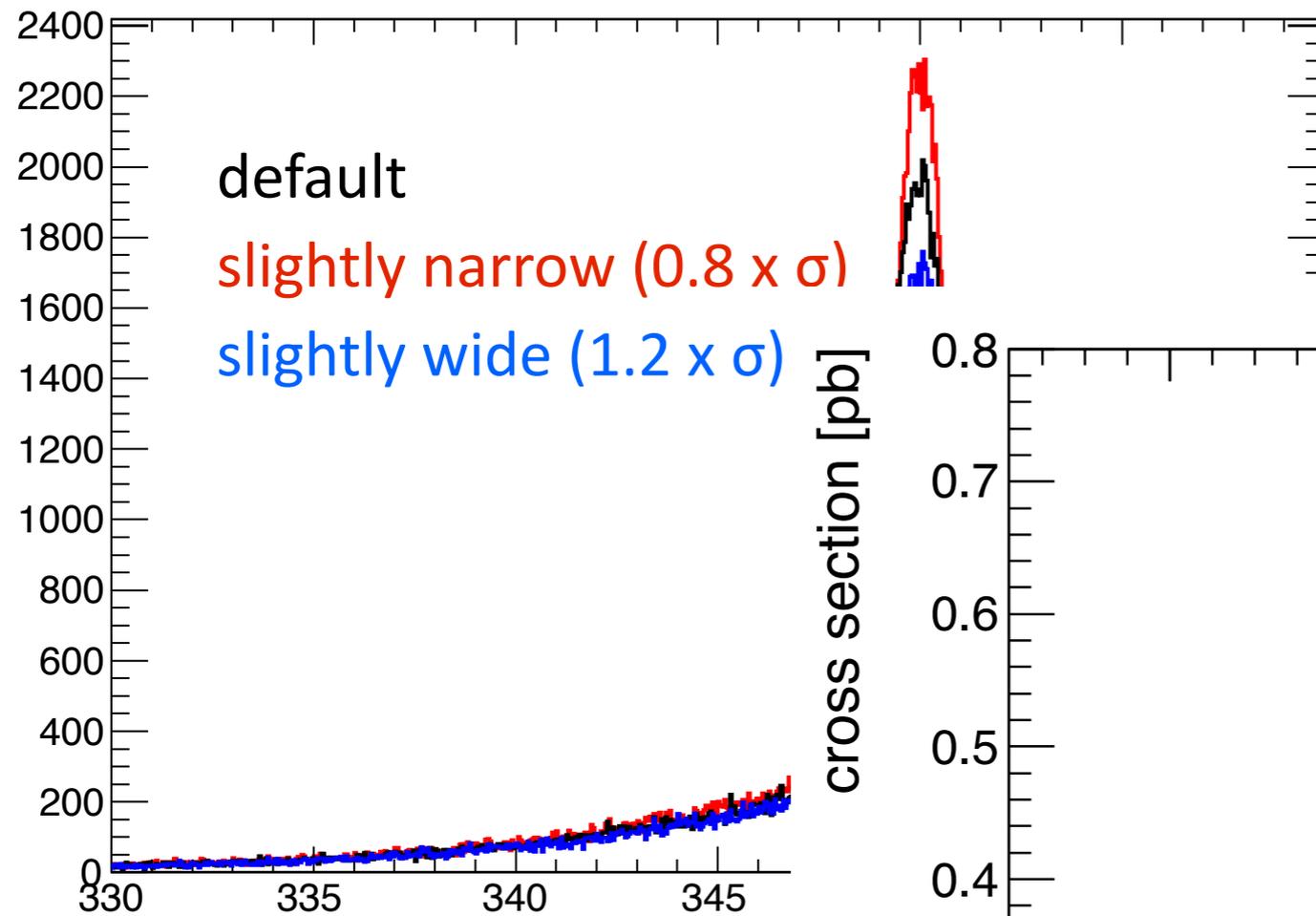
Quick study:
Fit with the **default**
spectrum, study data
distributions following
default, **narrow**, **wide**

- Impact of changes: Softer slope for wide spectrum, harder slope for narrow spectrum increased cross section above threshold in both cases

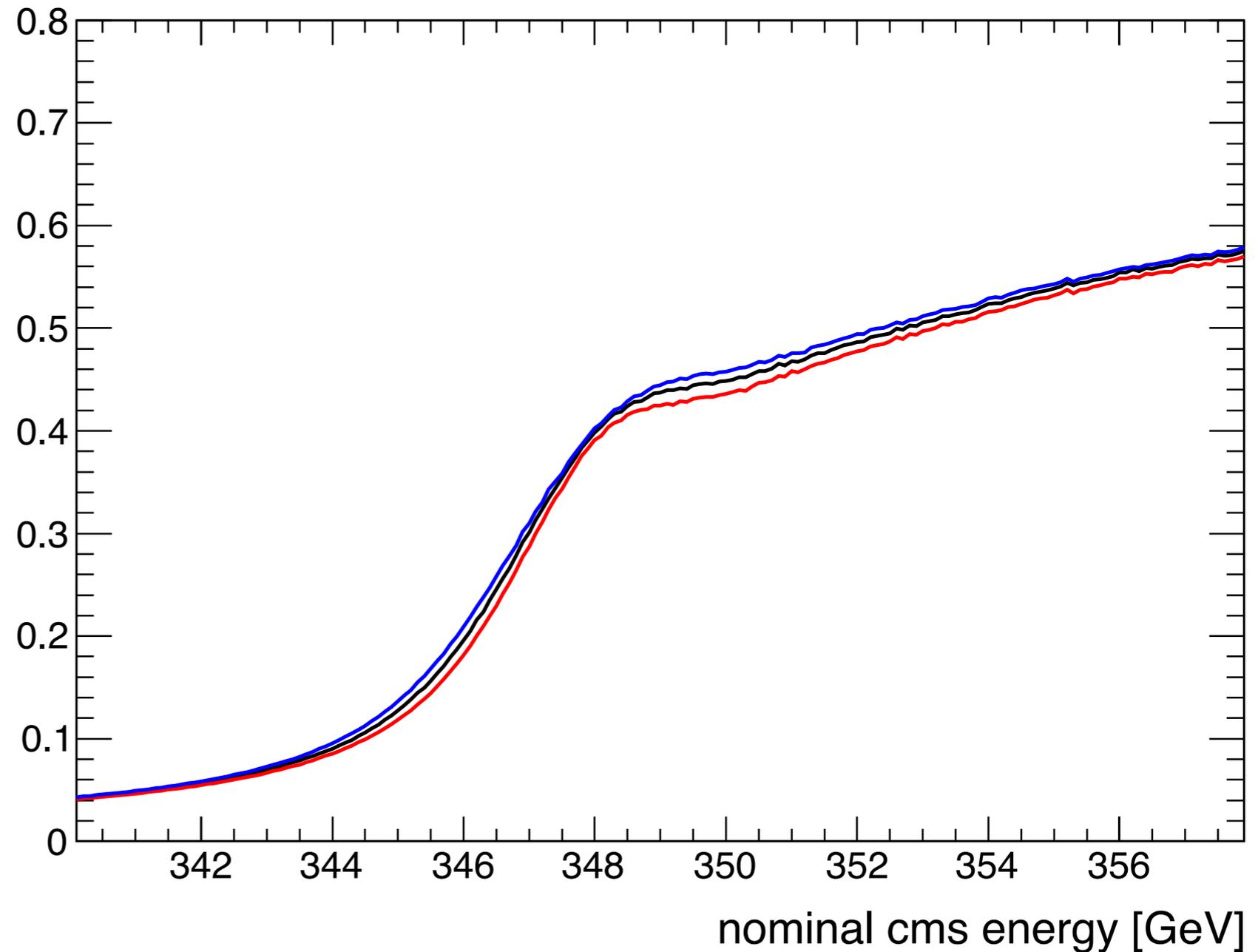
In addition: Less extreme Variations



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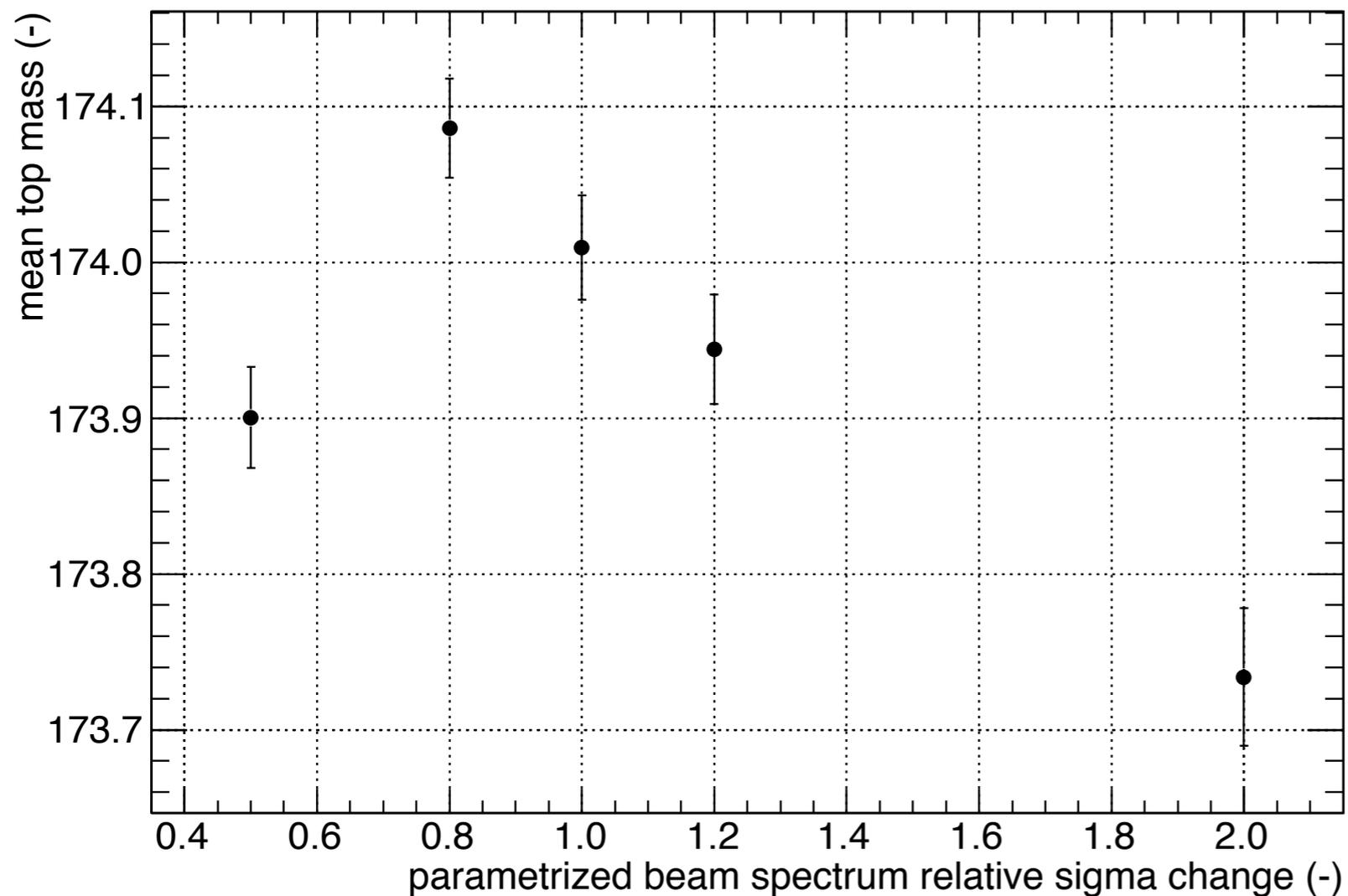


- Moderate impact on cross section: more intuitive ordering of curves



Impact of Changing Spectrum

- The study: Always perform fit with default distribution
 - Data points distributed according to various lumi spectra, always 10 points, 10 fb^{-1} each, in 1 GeV steps



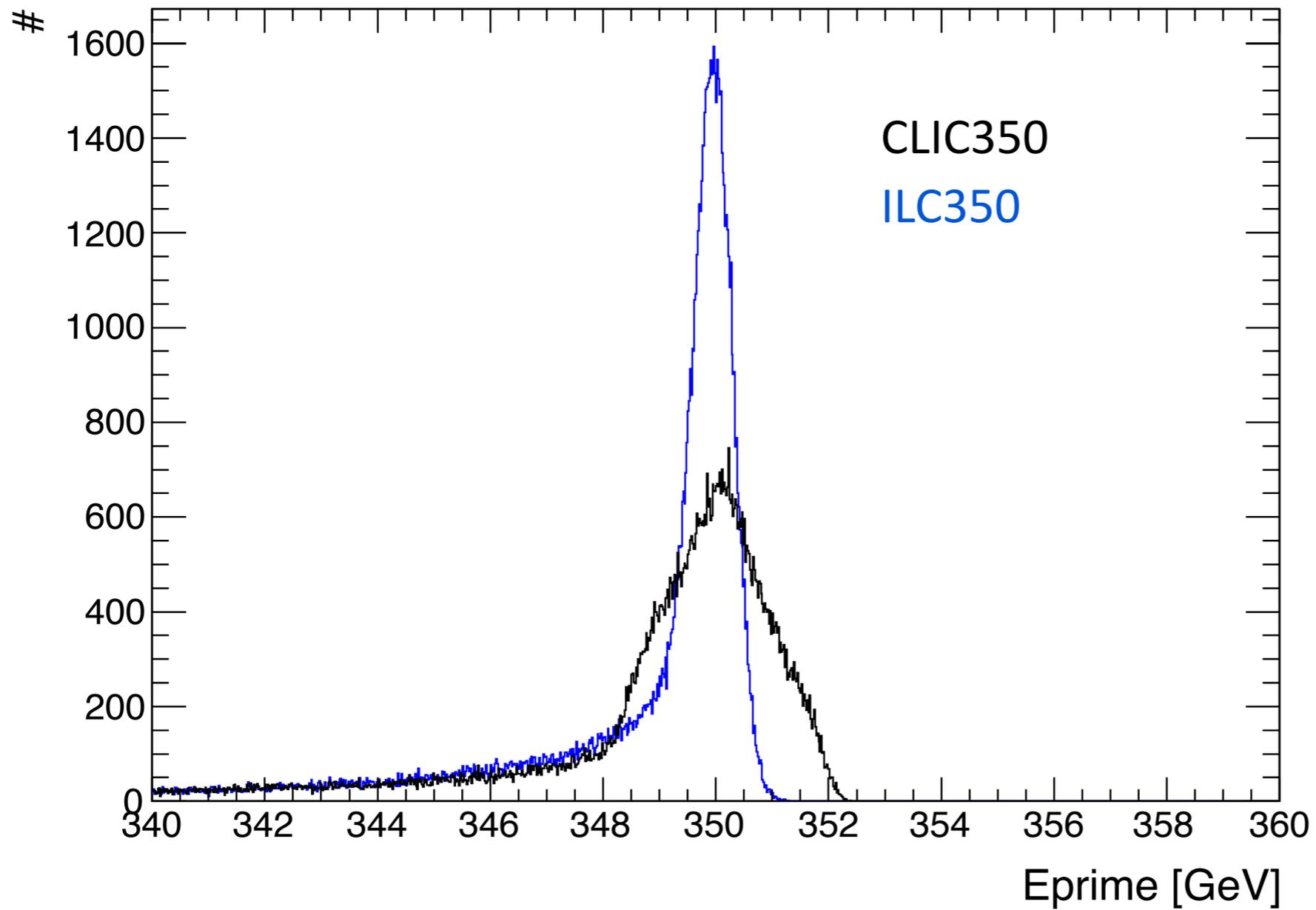
- Substantial bias of the top mass: Already for the smaller variations of the spectrum in excess of expected statistical uncertainty and theory systematics

Conclusions

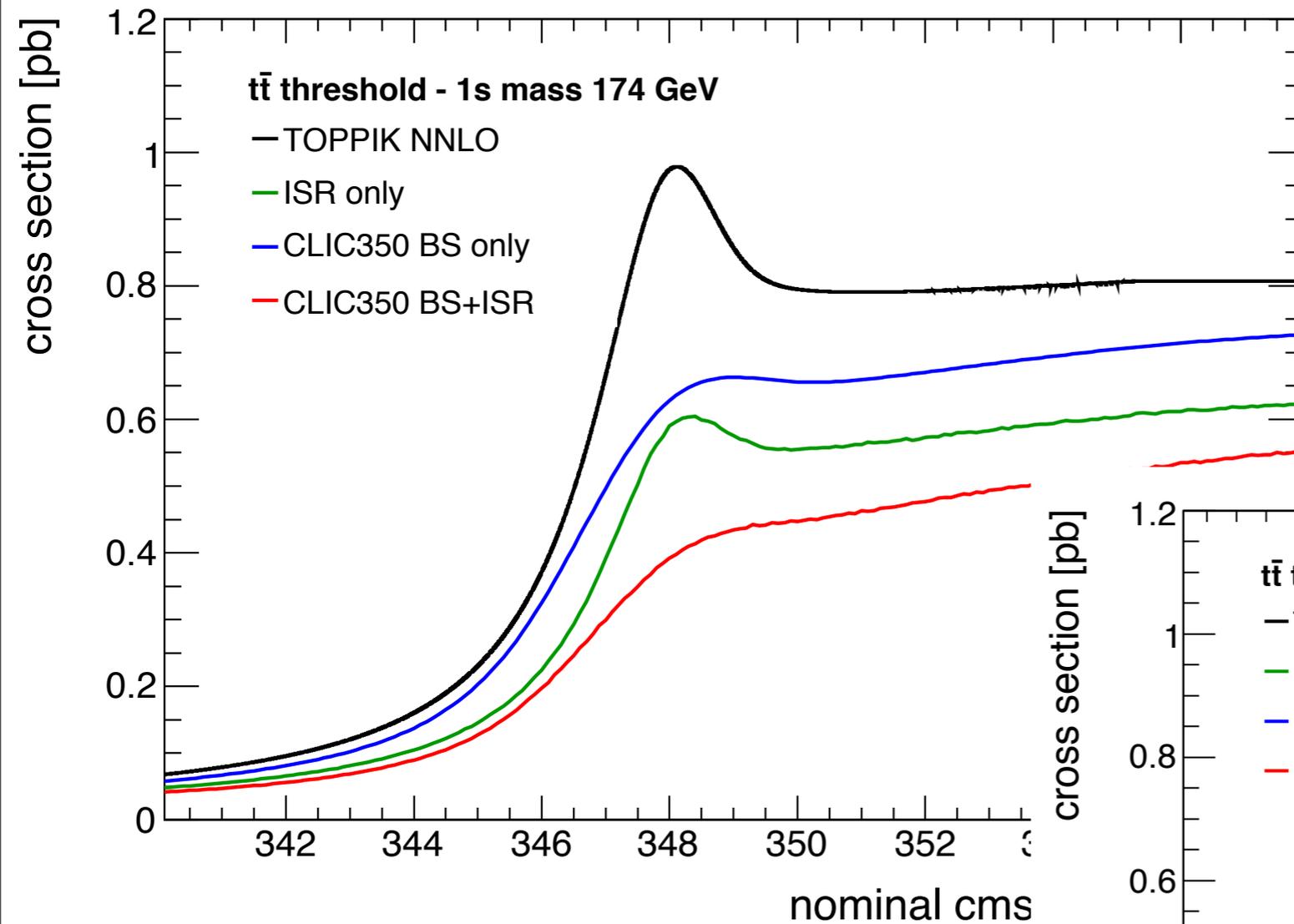
- The beam spectrum itself has a moderate effect on the mass resolution
- The knowledge of the beam spectrum is crucial: Strongly affect precision
 - Particularly important: The knowledge of the main peak of the luminosity distribution
- For a full study: Need to know the expected precision of the luminosity spectrum measurement - Ideally with two additional sets of lumi files (± 1 sigma)

Backup

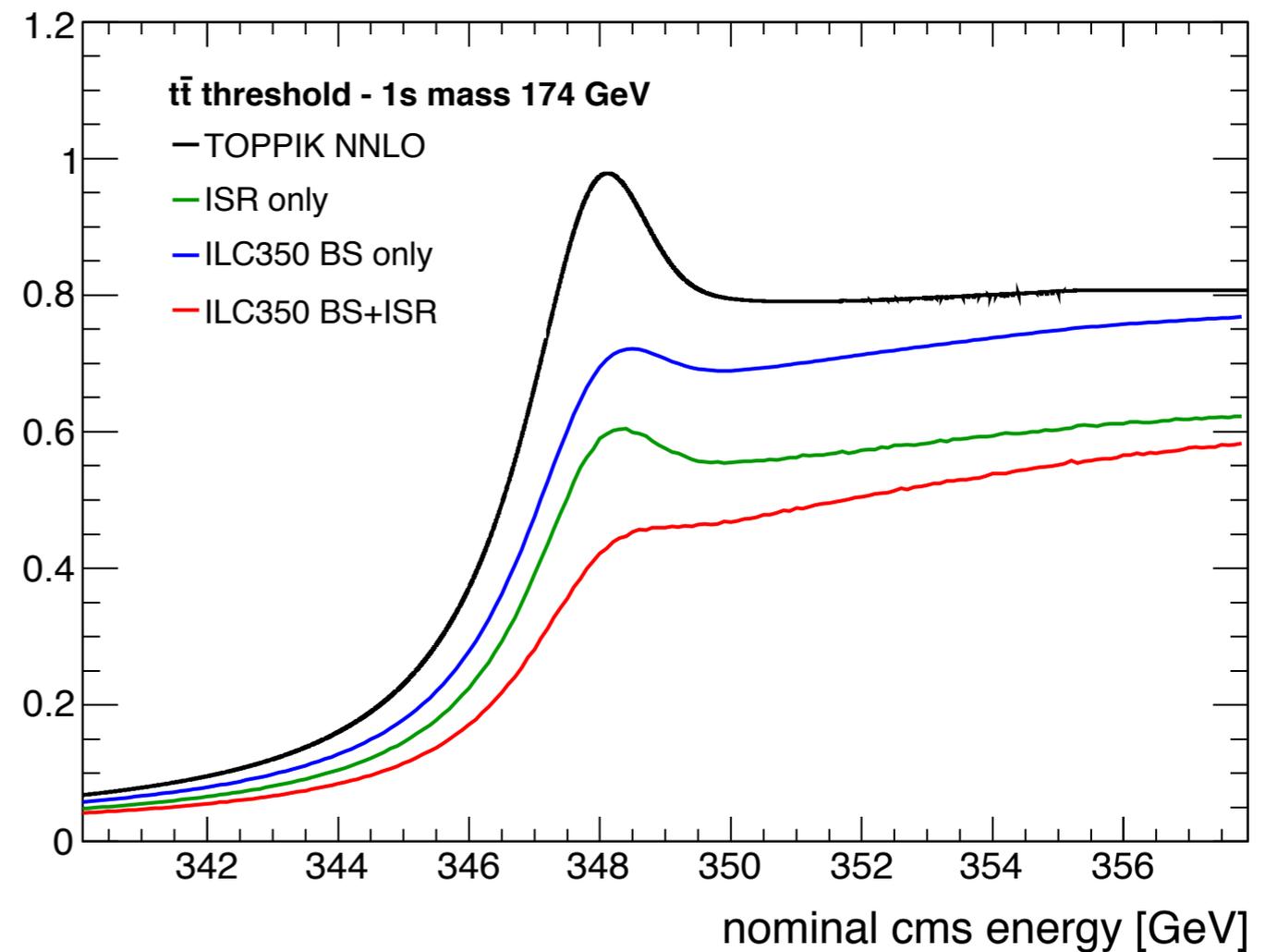
Beam Spectrum: CLIC vs ILC



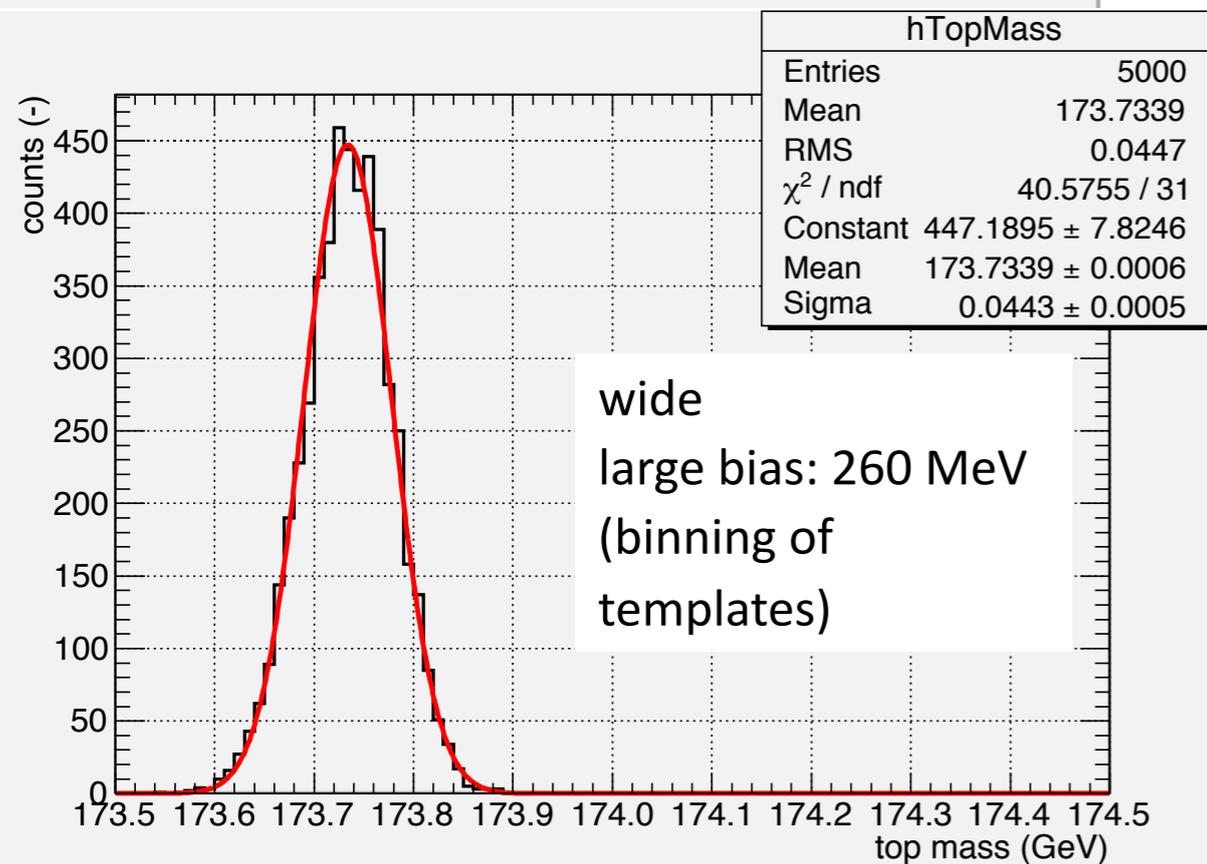
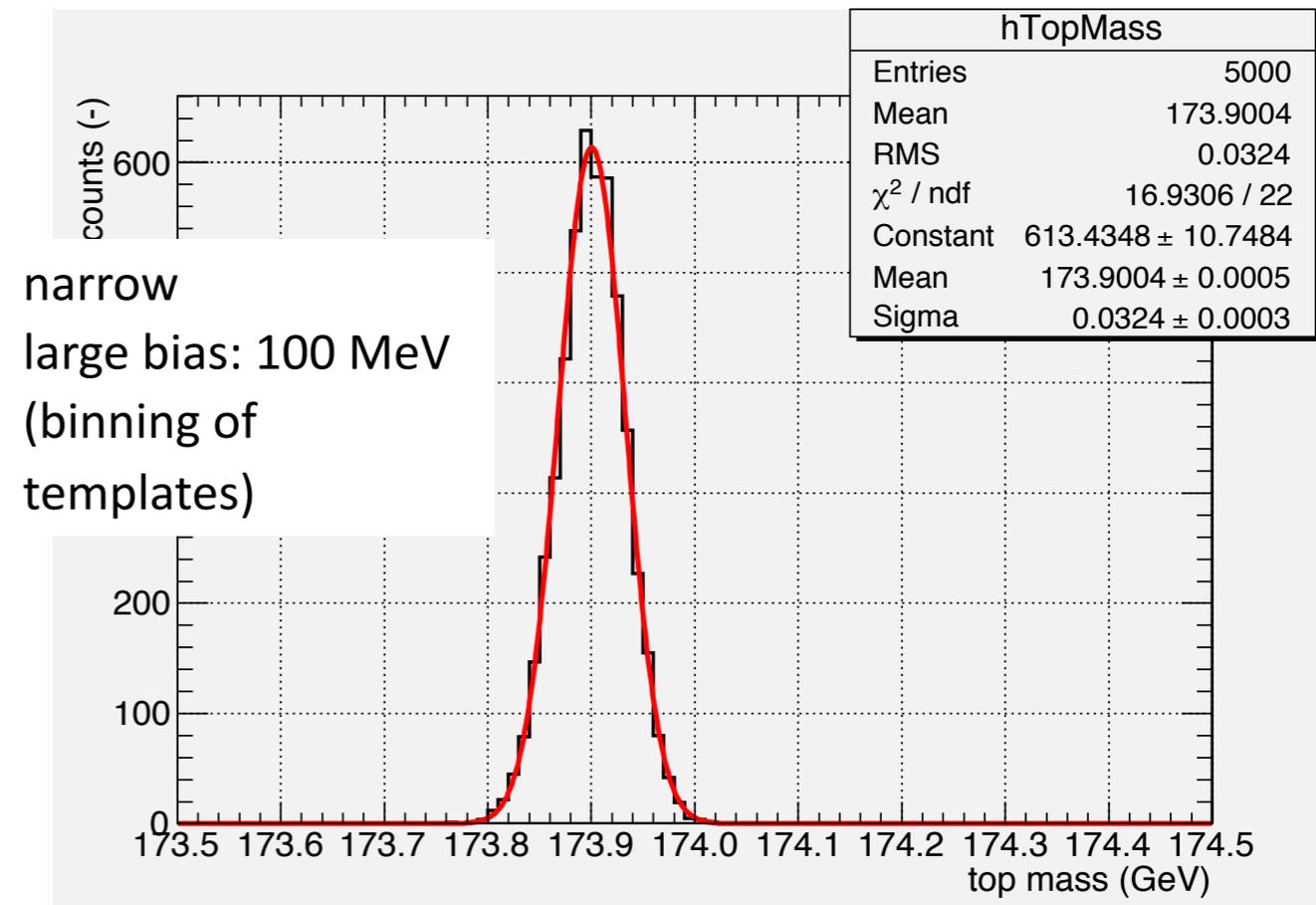
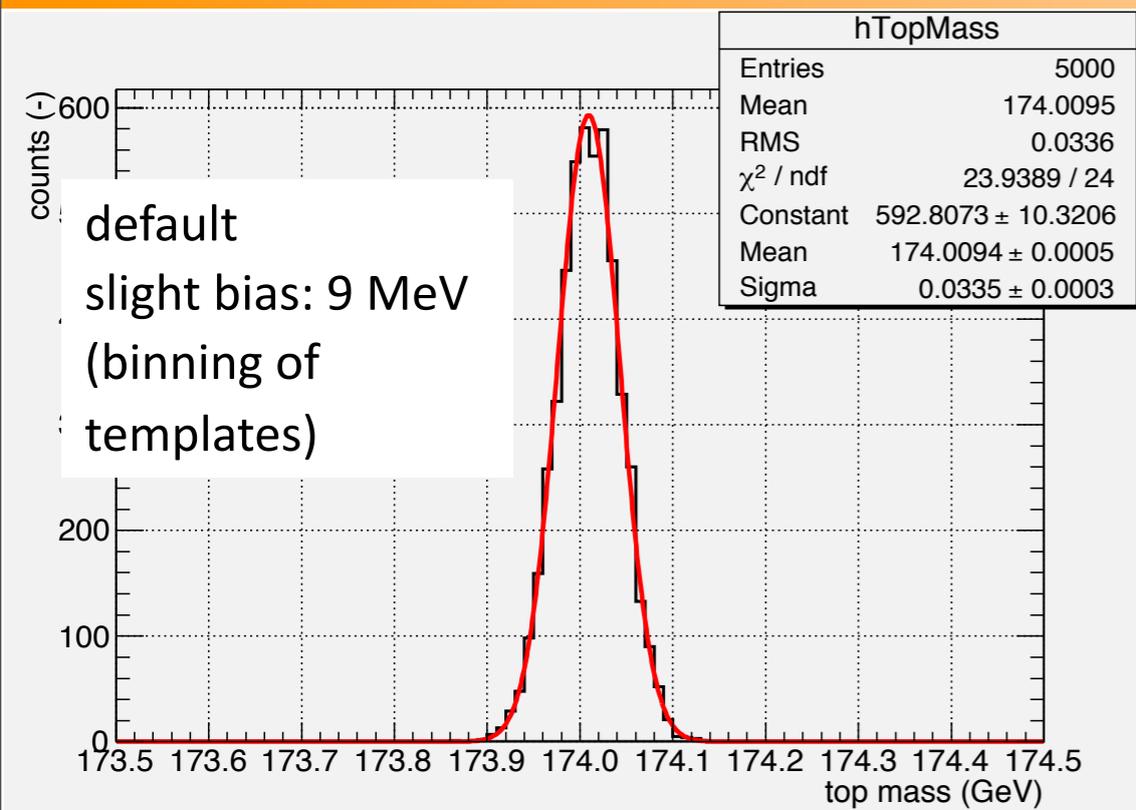
Top Threshold: CLIC vs ILC



Broader beam energy spectrum leads to softening of edge:



Template Fits - Results



- Substantial bias, far in excess of statistical uncertainty (can potentially be mitigated by different choice of scan points...)