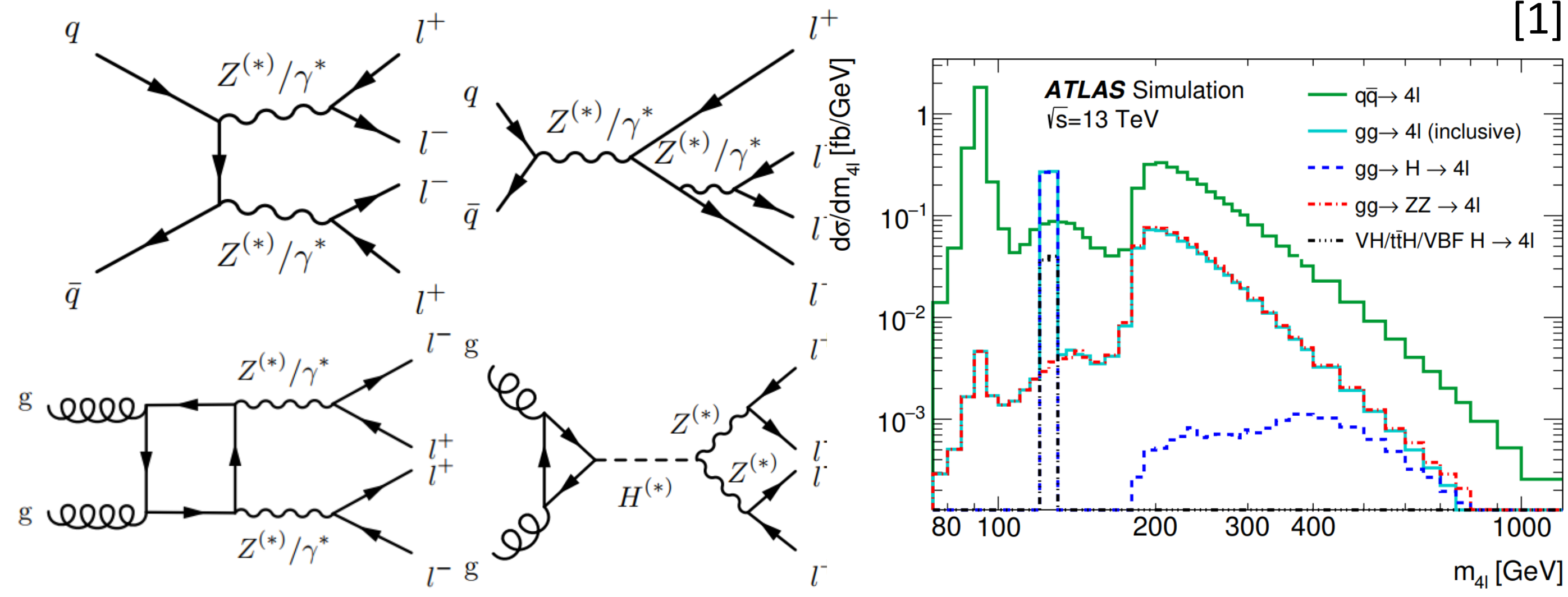


Motivation and Introduction

Motivation

- four-lepton (two same-flavor, opposite-sign lepton pairs, e or μ) final state is “clean” and has rich physics
- allow precision test on the SM (e.g. Higgs)
- sensitive to BSM



Fiducial definition

- no Z mass constraint in this analysis

Lepton selection	
Muon selection	Bare, $p_T > 5 \text{ GeV}$, $ \eta < 2.7$
Electron selection	Dressed, $p_T > 7 \text{ GeV}$, $ \eta < 2.47$
Event selection	
Four-lepton signature	At least 4 leptons, with 2 Same-Flavour, Opposite-Sign pairs
Lepton kinematics	$p_T > 20/10 \text{ GeV}$ for leading two leptons
Lepton separation	
J/ψ -Veto	$\Delta R_{ij} > 0.05$ for any leptons
Truth isolation	$m_{ij} > 5 \text{ GeV}$ for all SFOS pairs $p_{\text{cone}30}/p_T < 0.16$

Measured variables

- totally 10 variables measured in form of differential or double differential cross-sections
- invariant 4l mass (m_{4l}), invariant lepton-pair mass ($m_{12(34)}$), angular variable sensitive to polarization ($\cos\theta_{12(34)}^*$)^[2] etc.

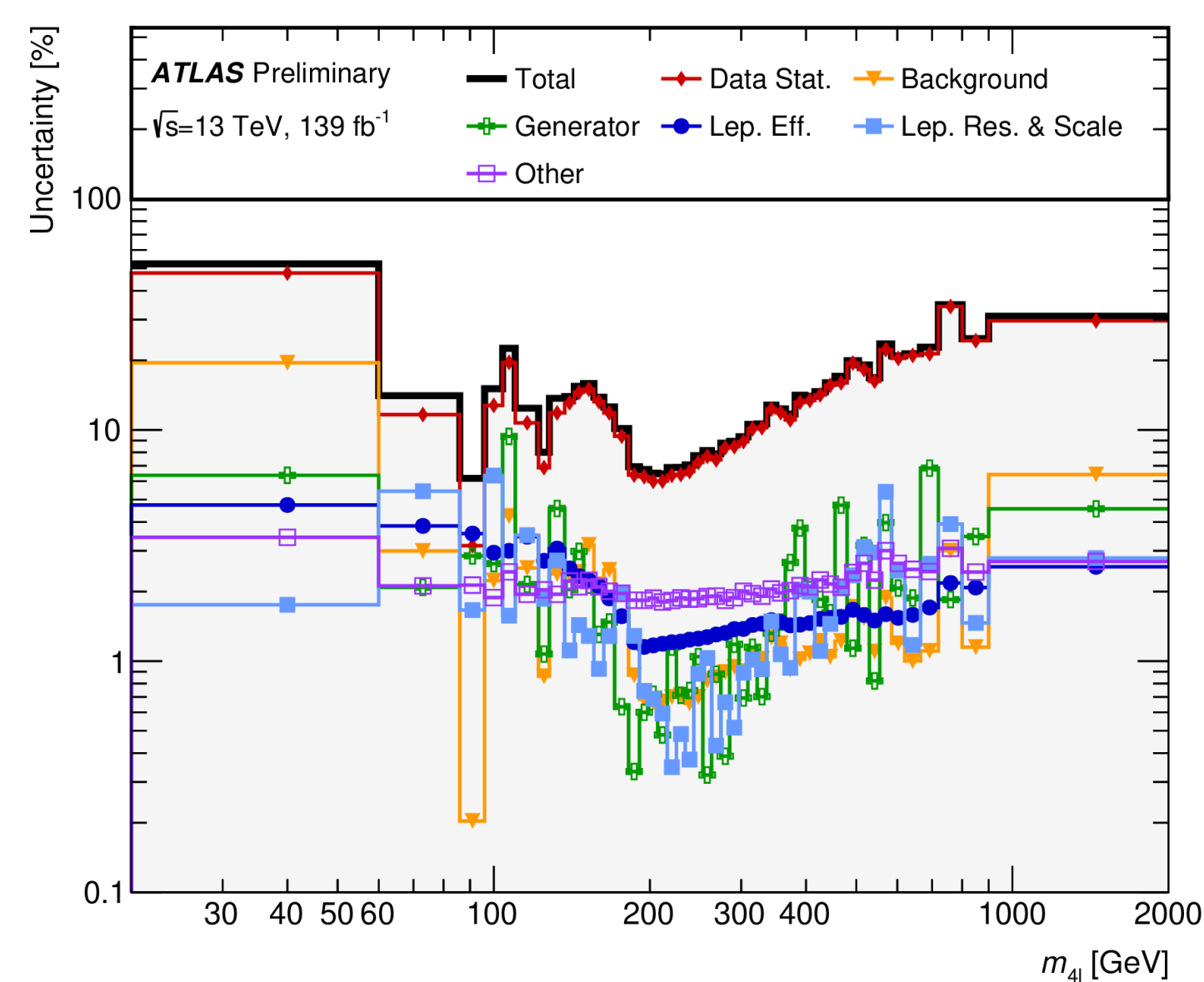
Techniques

- unfolding method used to obtain particle-level data distributions
- binning determination considering the unfolding purity^[3] as well as reconstruction-level statistics in each bin

Background and Uncertainties

Background

- Non-prompt background:
 - estimate in data
 - amounts to 4.4%, inclusive m_{4l}
 - largest background in off-shell ZZ region

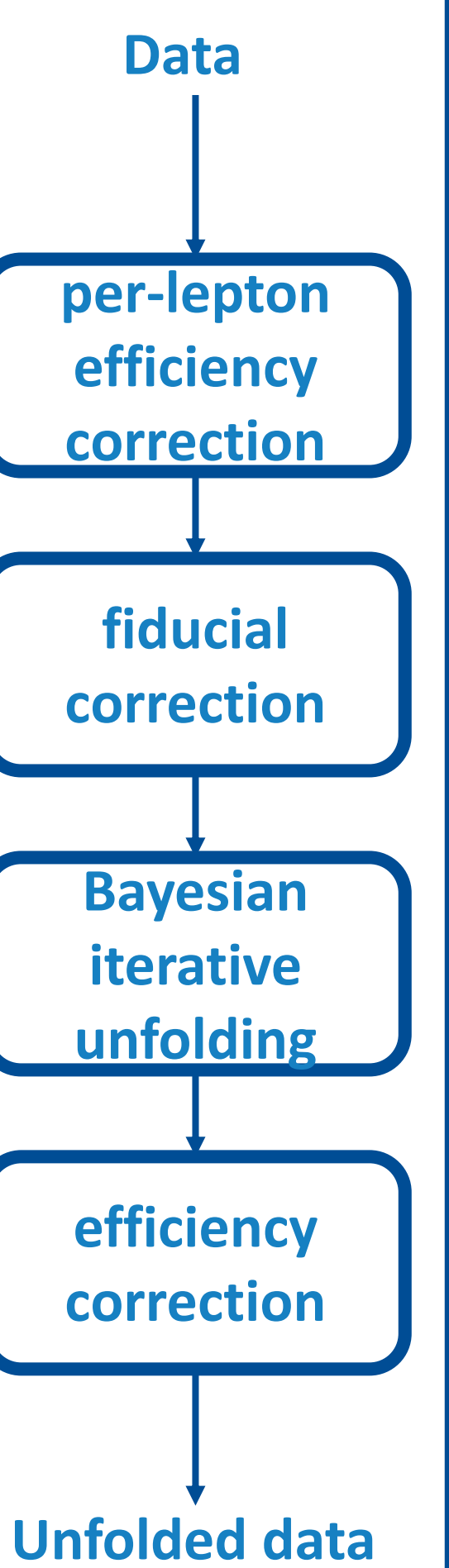
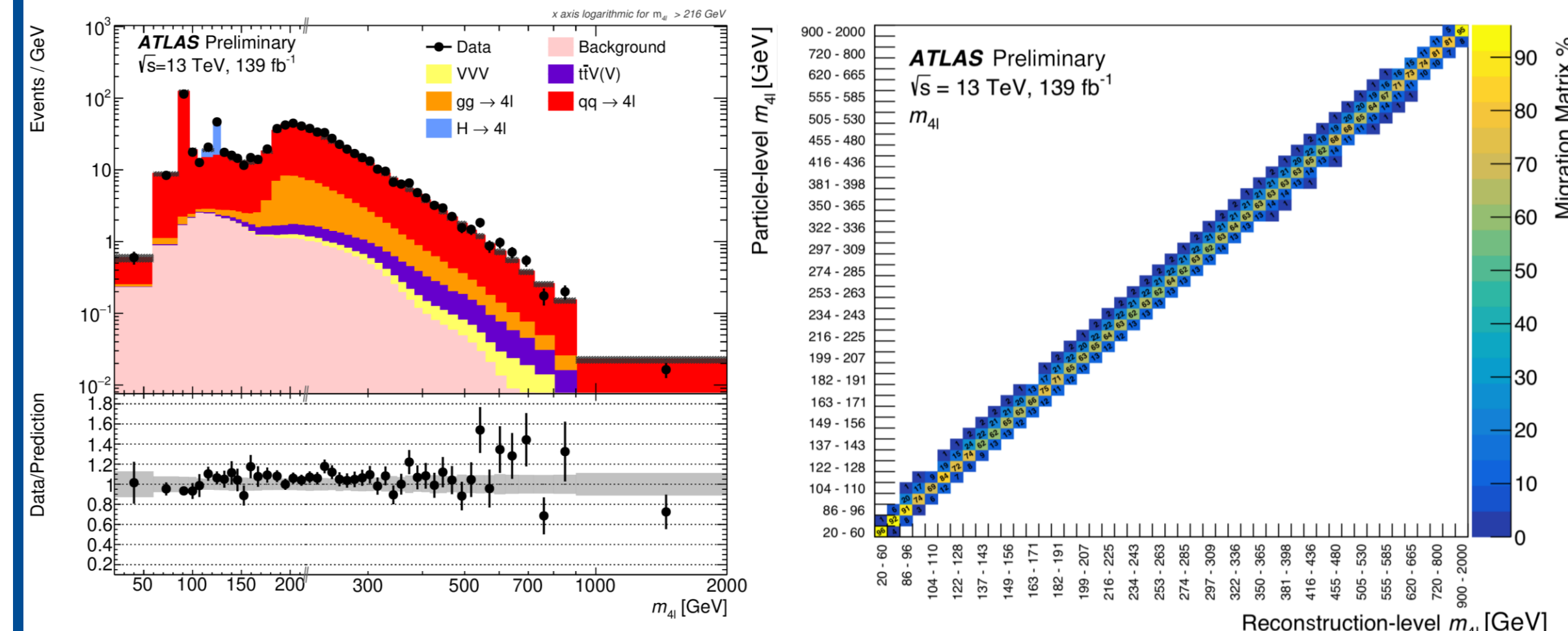


Uncertainties

- data statistical uncertainty dominant for most of the bins
- main systematic uncertainties:
 - 1.7% luminosity, inclusive m_{4l}
 - 1.3% generator, inclusive m_{4l}
 - largest systematic uncertainty for very low and high m_{4l}

Detector Correction

- detector effects: resolution of kinematic variables, inefficiency of reconstructing leptons and triggering of events
- corrections for inefficiencies and resolution effects at the fiducial phase space boundary
- Bayesian iterative unfolding to account for migrations within the fiducial phase space



Measurement Results

Cross-sections

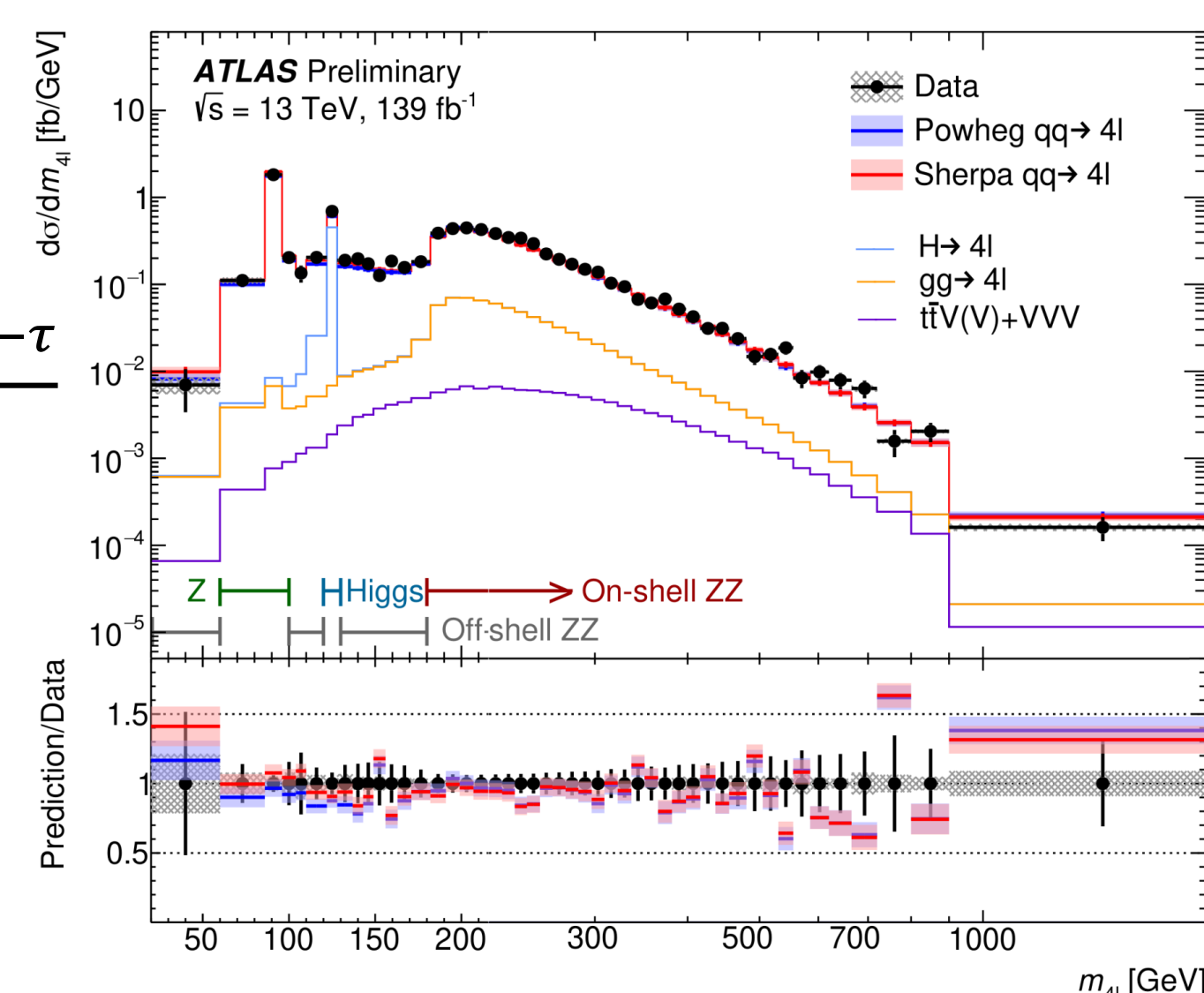
- fiducial cross-section inclusively and in slices of m_{4l}
- differential cross-section in m_{4l}
- the measurements extended the fiducial phase space and increased granularity compared to previous results^[1]

	Full	Region			
		$Z \rightarrow 4l$	$H \rightarrow 4l$	Off-shell ZZ	On-shell ZZ
Measured fiducial cross-section [fb]	88.9	22.1	4.76	12.4	49.3
	± 1.1 (stat.)	± 0.7 (stat.)	± 0.29 (stat.)	± 0.5 (stat.)	± 0.8 (stat.)
	± 2.3 (syst.)	± 1.1 (syst.)	± 0.18 (syst.)	± 0.6 (syst.)	± 0.8 (syst.)
	± 1.5 (lumi.)	± 0.4 (lumi.)	± 0.08 (lumi.)	± 0.2 (lumi.)	± 0.8 (lumi.)
	± 3.0 (total)	± 1.3 (total)	± 0.35 (total)	± 0.8 (total)	± 1.3 (total)
SHERPA	86 \pm 5	23.6 \pm 1.5	4.57 \pm 0.21	11.5 \pm 0.7	46.0 \pm 2.9
POWHEG + PYTHIA8	83 \pm 5	21.2 \pm 1.3	4.38 \pm 0.20	10.7 \pm 0.7	46.4 \pm 3.0

$Z \rightarrow 4l$ Branching ratio

$$\mathcal{B}_{Z \rightarrow 4l} = \frac{(\sigma_{\text{meas}} - \sigma_{\text{non-}q\bar{q} \rightarrow 4l}^{\text{pred}}) \times f_Z \times f_{\text{non-}\tau}}{\sigma_Z \times A_{\text{fid}}} = (4.41 \pm 0.30) \times 10^{-6}$$

- most precise result and compatible with previous study^[1]

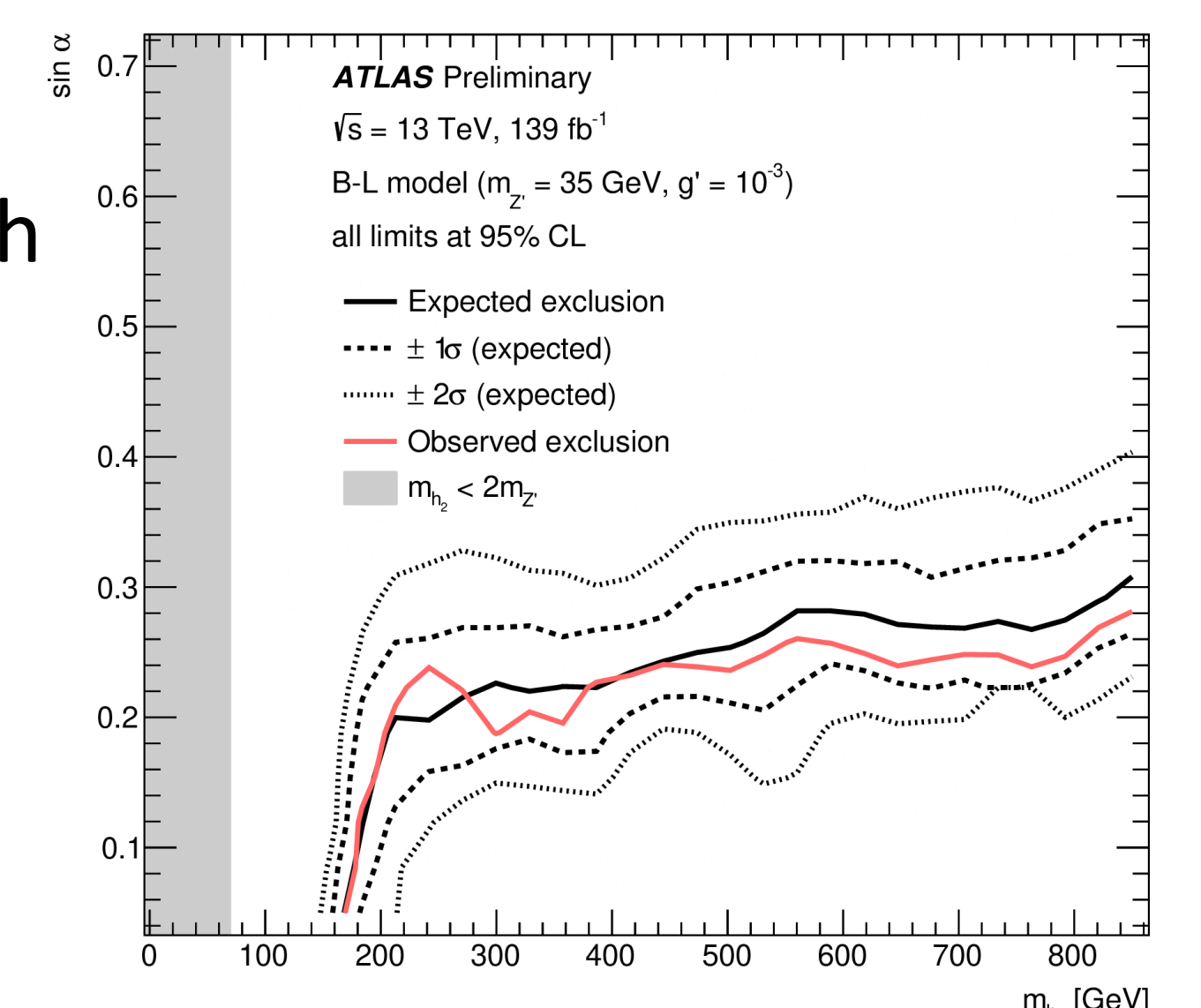


BSM Interpretation

- using measured differential cross-sections to constrain BSM model, with all information needed in the HEPData

B-L Model

- $U(1)_{B-L}$ gauge symmetry broken introduced to solve neutrino mass problem
- multi-Gaussian based statistical likelihood function
- exclusion on $m_{h_2} \sim \sin\alpha$ parameter-space @ 95% CL with fixed ($m_{Z'} = 35 \text{ GeV}$, $g' = 10^{-3}$)
- using the double differential distributions yields significantly stronger limits than the m_{4l} distribution alone



CONF note: ATLAS-CONF-2020-042

[1] previous m_{4l} analysis with ATLAS 2015-16 data: [https://doi.org/10.1007/JHEP04\(2019\)048](https://doi.org/10.1007/JHEP04(2019)048)

[2] the angle between the negative lepton in the di-lepton rest frame and the di-lepton pair system in the laboratory frame

[3] purity here defined as the fraction of events in a reconstruction-level bin that originate from the same bin at particle-level