What I’ve got in the draft so far

• Very simple format for display of the method results:

• White points are those considered excluded by the rescaling technique

• Red line is boundary considered excluded by the analysis team (contour comes from ATLAS HEPData)
What we could add for more explicit validation

• 2D plot format probably worth hanging onto for its ability to highlight differences between regions (e.g. worst agreement closest to the diagonal, and so on)

• Could use this to validate 1) cross section, by comparison to generated values, and 2) limits, which is a convolution of cross section agreement and other factors like changing kinematics etc - only if not initially rescaled

• Drawbacks: does not show effects of scanning coupling values like validations Andreas suggests for resonances

• Challenges: is this info public? Do we need to regenerate signals ourselves? Do proper limits exist?