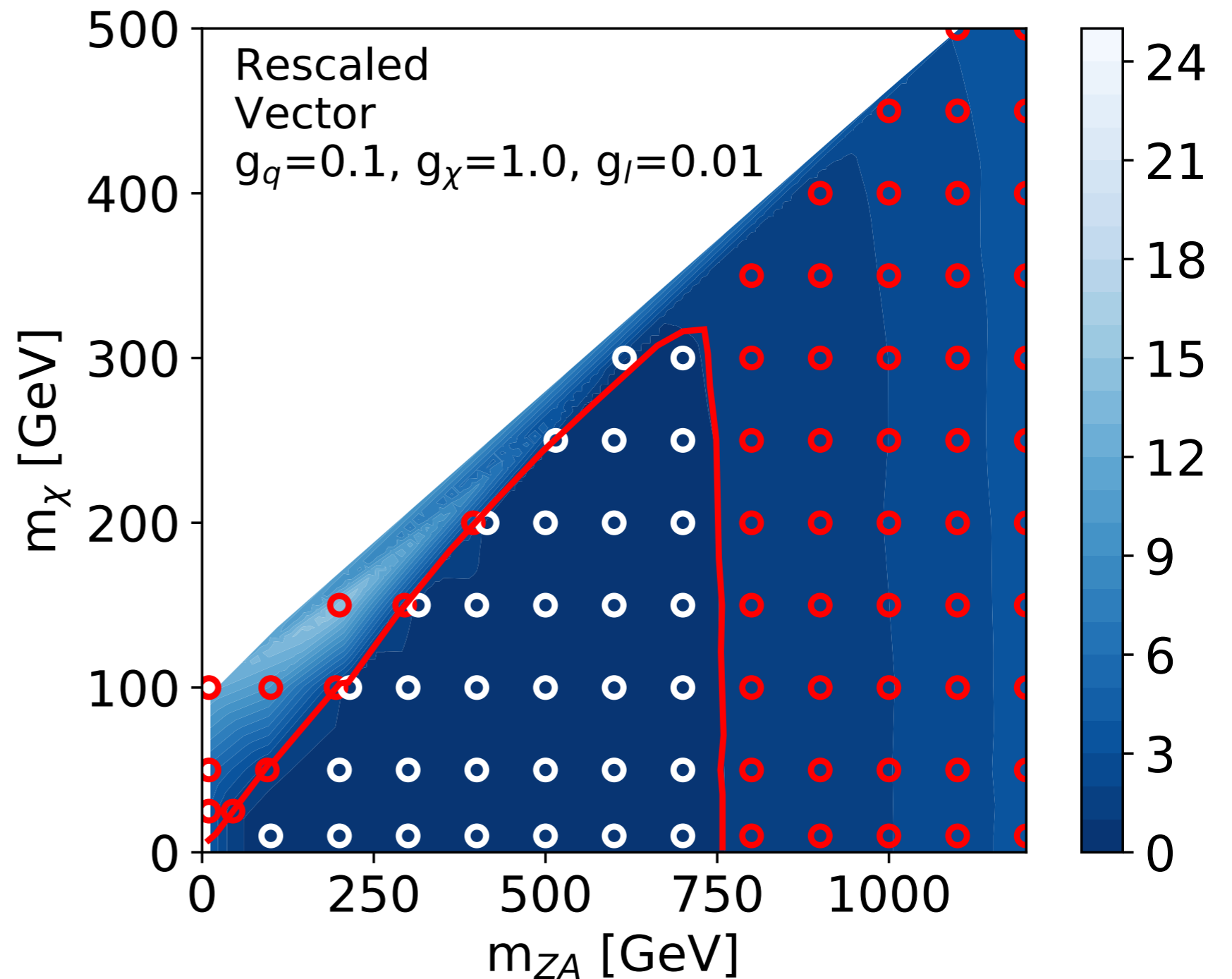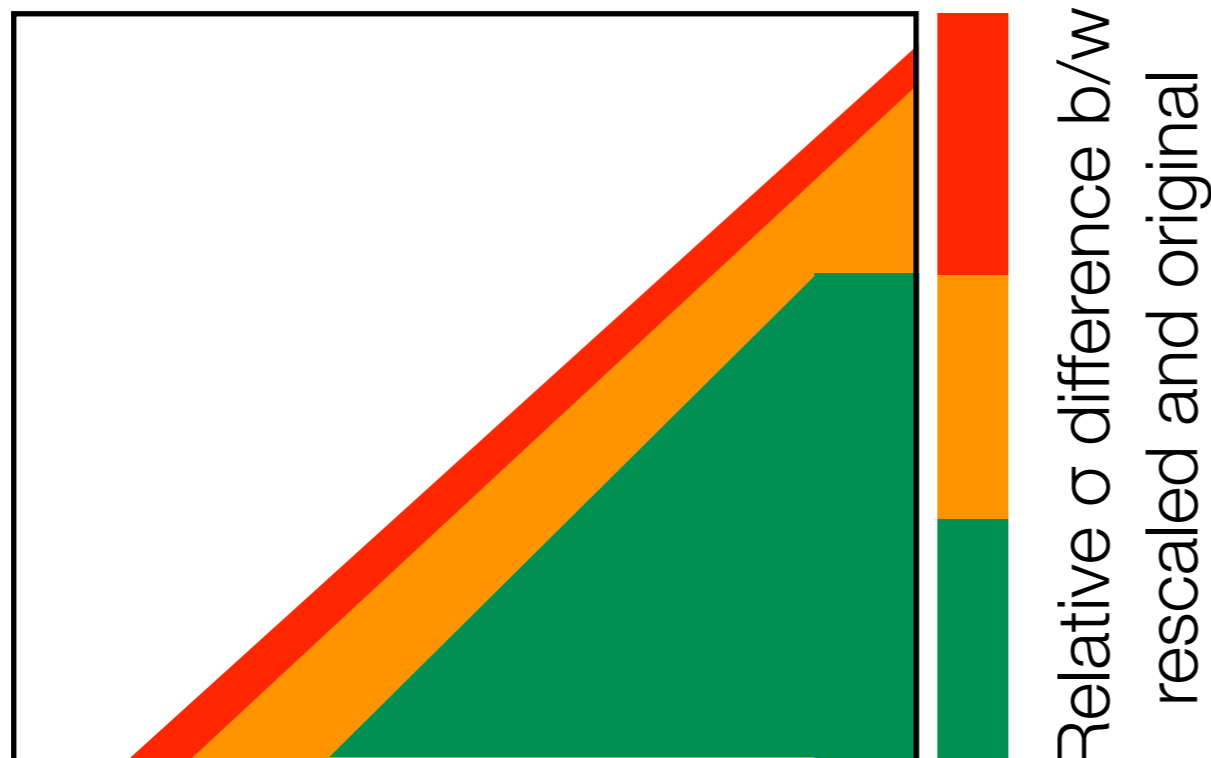# What I've got in the draft so far

- 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)

Rescaled Vector
g_q=0.1, g_χ=1.0, g_l=0.01

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

Relative σ difference b/w rescaled and original

- 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?