

Coupling scan validation plans

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Introduction


- The coupling scan white paper must have two main purposes:
 1. explain rescaling methods, make code available
 2. demonstrate + quantify the performance of the methods
- This talk is about (2): What do we want to demonstrate?
- Few ideas immediately come to mind:
 - **Inclusive:**
 - In how far does the rescaling reproduce MG-level cross sections as a function of couplings? → next slides
 - **Differential:**
 - In how far are MET shapes in mono-X searches independent of couplings?
 - In how far are angular distributions for visible decay products of mediators independent of couplings?
 -

Step 1: Inclusive cross section validation

Results as .txt

At the lowest complexity level,
we need to show that we can
reproduce the inclusive XS

Study outline:

- Calculate coupling scan of XS in standalone MG 
- Compare to analytical method, e.g. BR-based rescaling

```
$ cat coupling_scan.txt

import DMSimp_s_spin1
generate p p > xd xd- j

output dmsimp_coupling_scan

launch

set param_card MXd 1.0
set param_card MY1 scan: [100, 250, 500, 1000, 1500, 2000]
set param_card gVh 0
set param_card gad11 0
set param_card gad22 0
set param_card gad33 0
set param_card gau11 0
set param_card gau22 0
set param_card gau33 0
set param_card gaxc 0
set param_card gaxd 0
set param_card gvd11 scan1: [0.005, 0.01, 0.025, 0.05, 0.1, 0.15, 0.2, 0.25, 0.3, 0.35, 0.4, 0.5, 0.75, 1.0]
set param_card gvd22 scan1: [0.005, 0.01, 0.025, 0.05, 0.1, 0.15, 0.2, 0.25, 0.3, 0.35, 0.4, 0.5, 0.75, 1.0]
set param_card gvd33 scan1: [0.005, 0.01, 0.025, 0.05, 0.1, 0.15, 0.2, 0.25, 0.3, 0.35, 0.4, 0.5, 0.75, 1.0]
set param_card gvu11 scan1: [0.005, 0.01, 0.025, 0.05, 0.1, 0.15, 0.2, 0.25, 0.3, 0.35, 0.4, 0.5, 0.75, 1.0]
set param_card gvu22 scan1: [0.005, 0.01, 0.025, 0.05, 0.1, 0.15, 0.2, 0.25, 0.3, 0.35, 0.4, 0.5, 0.75, 1.0]
set param_card gvu33 scan1: [0.005, 0.01, 0.025, 0.05, 0.1, 0.15, 0.2, 0.25, 0.3, 0.35, 0.4, 0.5, 0.75, 1.0]
set param_card gvxc 0
set param_card gvxd 1
set WY1 AUTO

set run_card ptj 100

$ ./bin/mg coupling_scan.txt
```

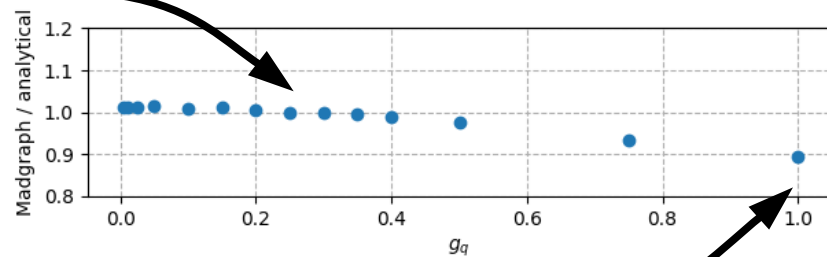
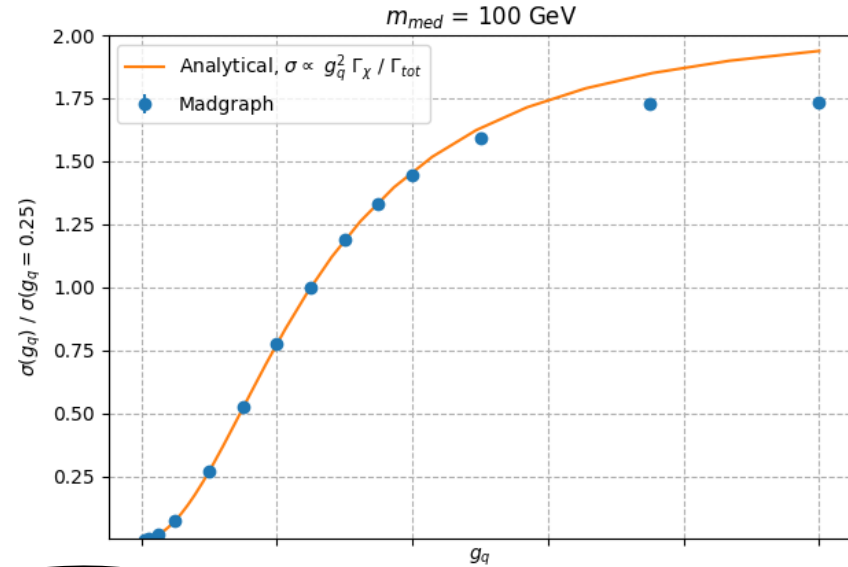
Comparison of analytical and simulated scaling (vector, $m_{dm} = 1$ GeV)

MG result (blue points) here compared to BR rescaling from mediator width dependence on g_q , g_{chi} (orange line)

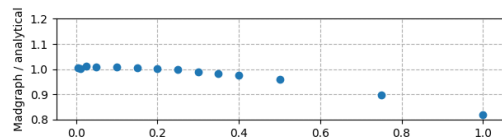
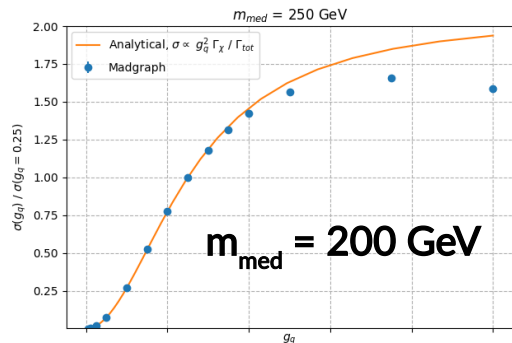
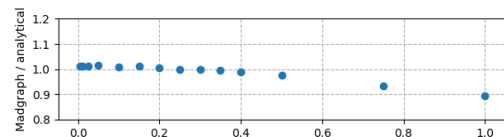
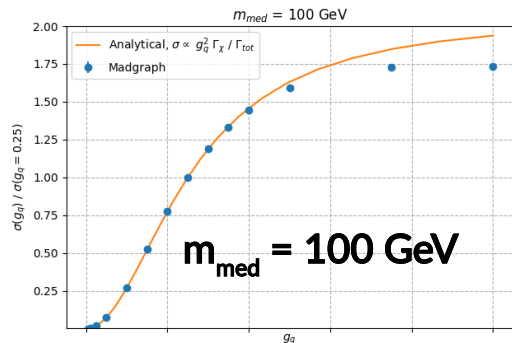
We only care about the ratios
→ y-axis is relative to $g_q=0.25$

Point here is vector med with
 $m_{med} = 100$ GeV, $m_{DM} = 1$ GeV

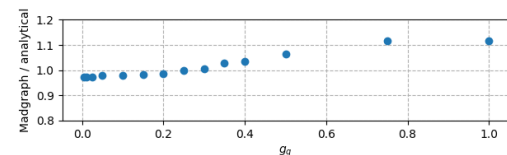
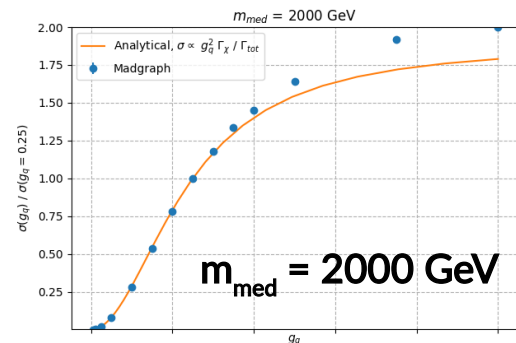
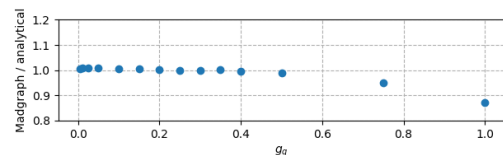
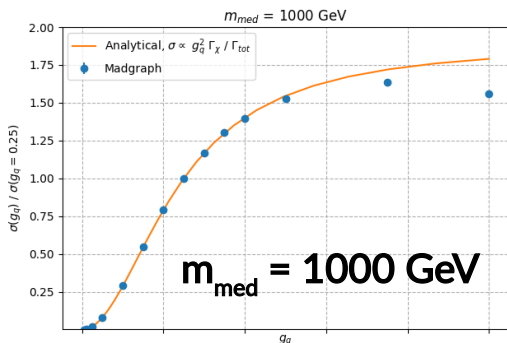
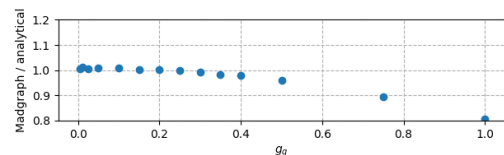
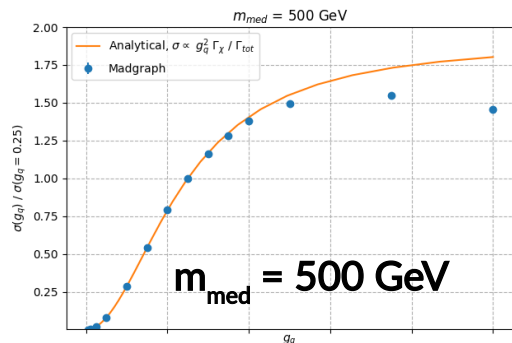
Quote agreement regions:
few % level for $g_q \in [0, 0.5]$,
10% level for g_q up to 1



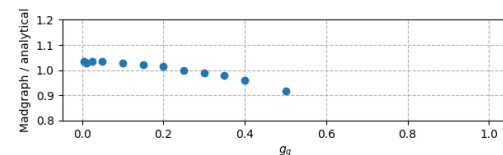
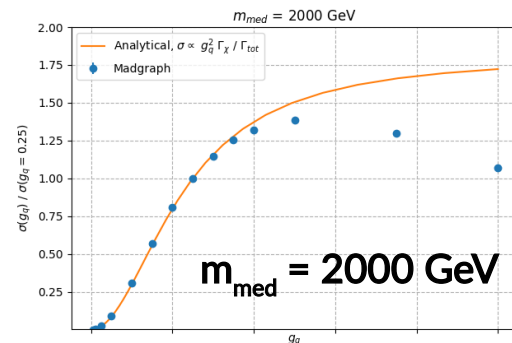
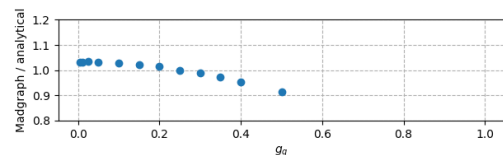
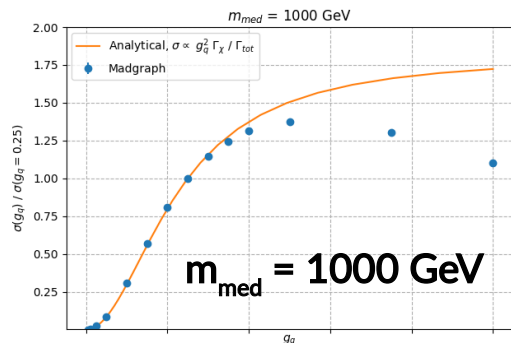
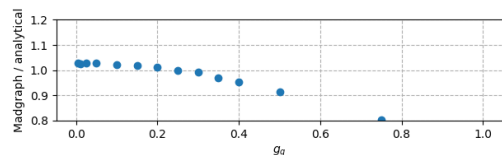
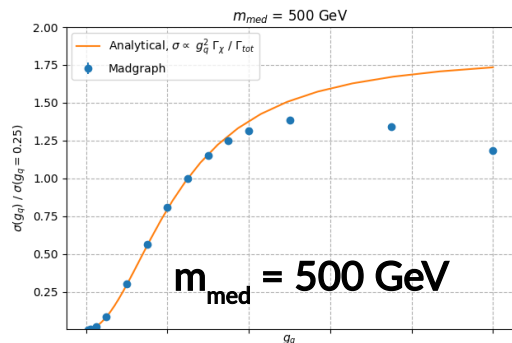
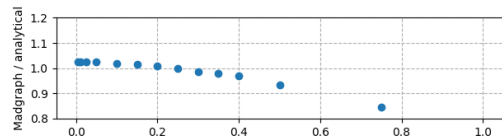
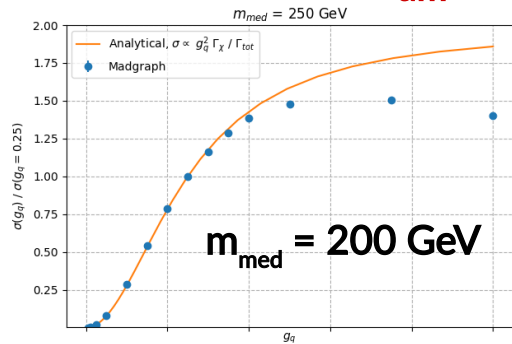
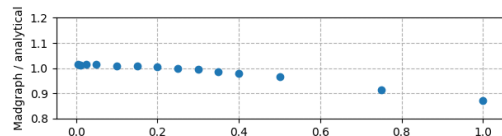
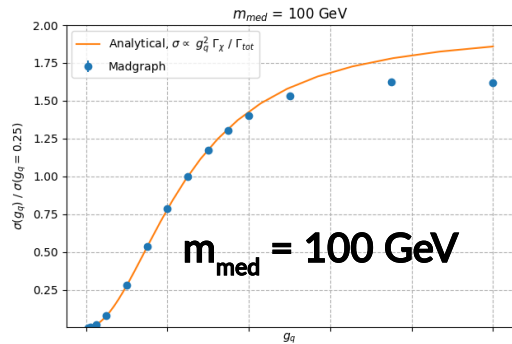
More points, vector $m_{DM} = 1$ GeV



Exact agreement varies with mediator mass, but overall very good until 0.5



Same as last slide, bt different DM mass: $m_{dm} = m_{med}/3$



- qualitative similar to $m_{dm} = 1$ GeV
- quantitatively the deviations are bigger
→ Should trust gq scaling until about 0.5

Extension?

- Plots based on simplest level of analytical rescaling
- **Ideal plot: Overlay different levels of rescaling on same plot.**
 - Demonstrates where exactly additional complexity is useful
 - Allows the user to gauge whether simple method might suffice within needed precision
- Presentation should explore major axes in which agreement develops: mmed, mdm/mmed

Step 2: Differential agreement

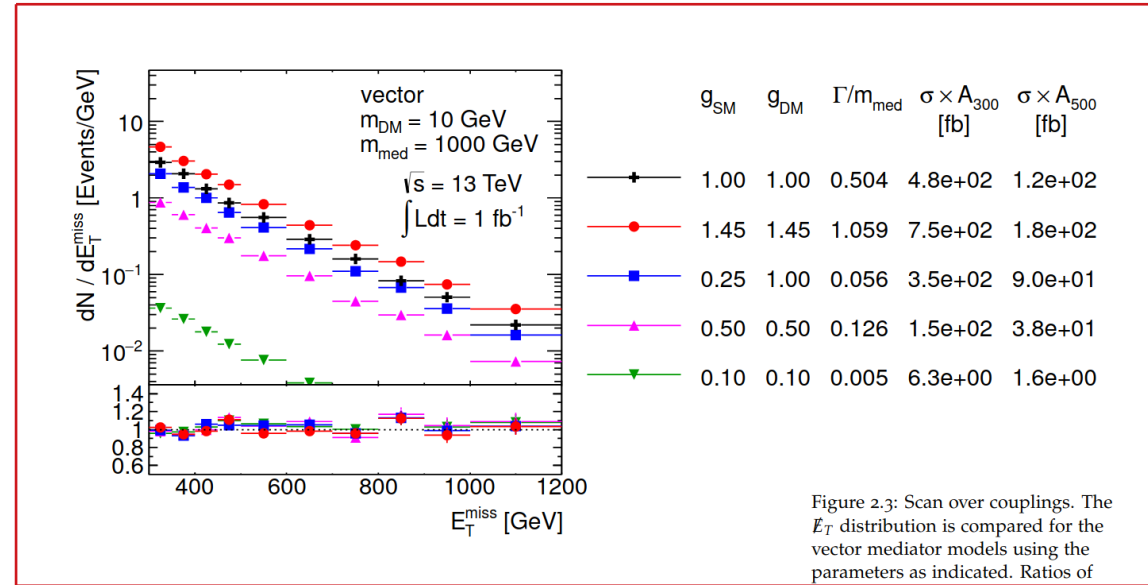
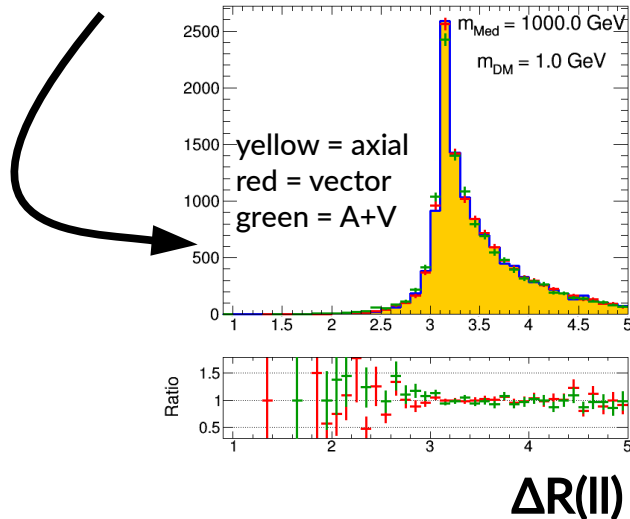
- Limit rescaling only works if change of differential distributions is understood
- For MET+X, this was previously explored in [2015 DMF report](#) (and probably elsewhere, too):

m_{DM} dependence not studied here AFAICT

→ Would be good addition

- For dilepton, differential dists of leptons must agree with DY as function of mass width

tested explicitly in CMS, easy to replicate standalone



Other aspects we should think about?

Summary

- **Validation is integral part of white paper: quantify validity in different parameter ranges**
- **Basis for inclusive validation ready**
 - extend with more complex rescaling methods
- **What do we want to do for differentials?**
 - MET shapes
 - Dilepton angular dists?
 - ?

Backup

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