

# Using Finesse to Model Mode-Mismatch in aLIGO

# The Setup

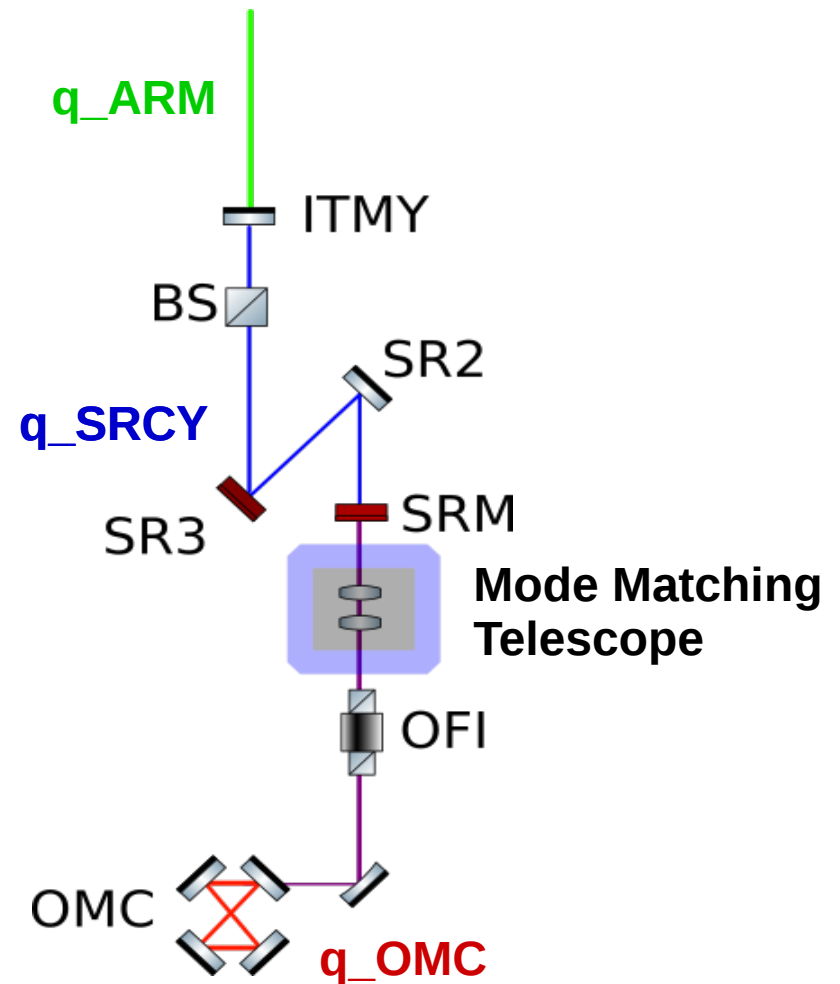
- Start with aLIGO standard .kat file
  - Includes all main optics
  - Added tunable squeezing (10db) + a filter cavity
  - Added ideal mode-matching optics between SRC and OFI

# The Big Questions

- We've seen SR3 and SRM RoC changes affect the mode-matching of the IFO:
  - How do those changes affect the  $h(t)$  w/ and w/o squeezing?
  - If we vary the mode of the SRC, but keep the Arms and OMC mode-matched, how does that effect  $h(t)$ ?
    - If the negative effect of changing the SRC mode is small, it could allow us to use SRM as a mode-matching actuator.

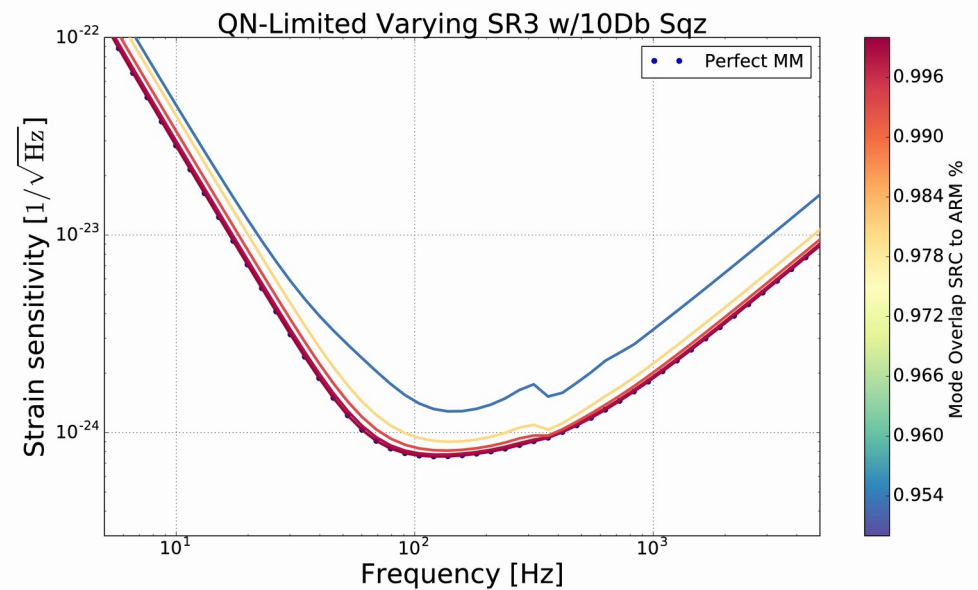
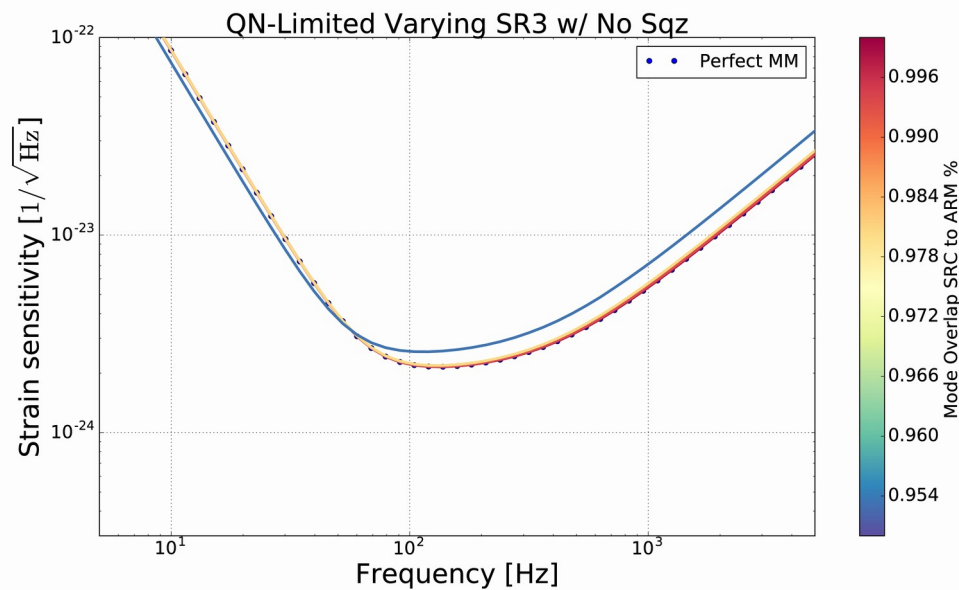
# The Modified Cavities

1. Vary the SRM **OR** SR3 RoC
2. Use the Mode Matching Telescope to keep the ARM and OMC overlap at 100% while changing the SRM RoC.
3. Find the ARM, SRC, and OMC modes at the BS to measure the overlaps between any two modes.
4. Output the QM-limited  $h(t)$  to see how badly an SRC mismatch hurts us.



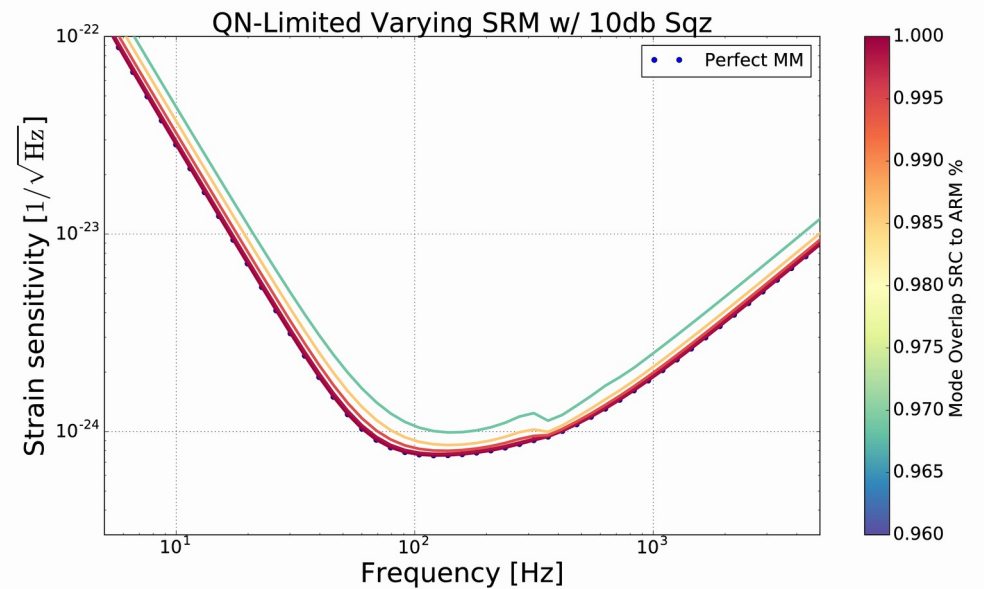
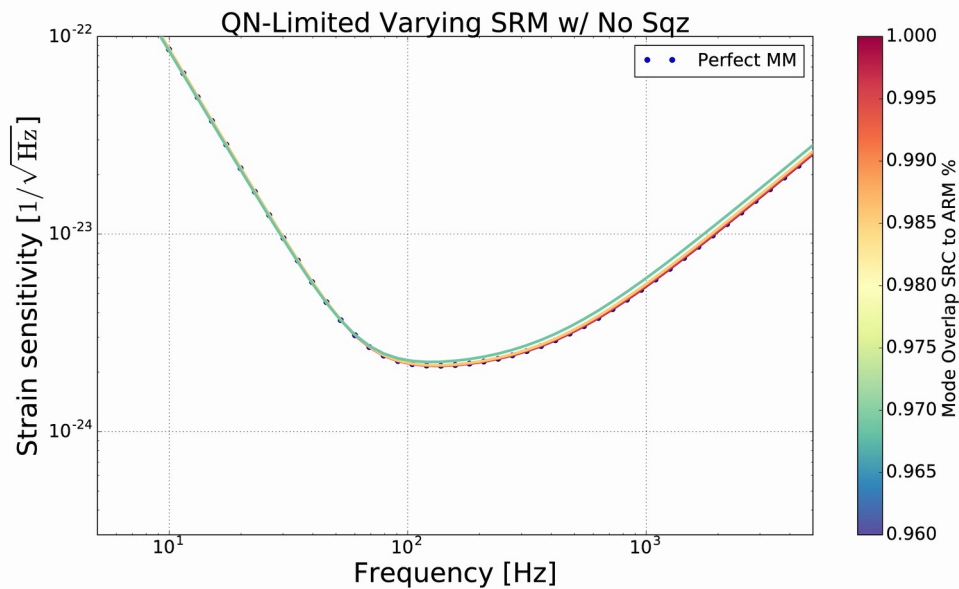
# Sensitivity Outputs

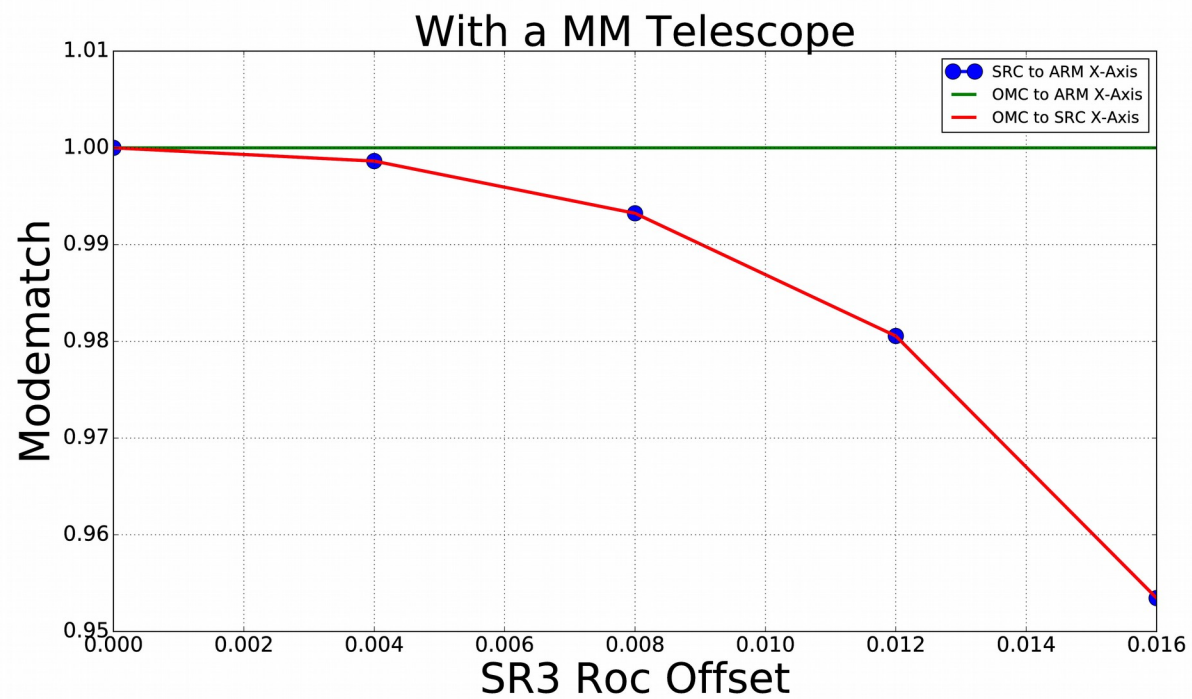
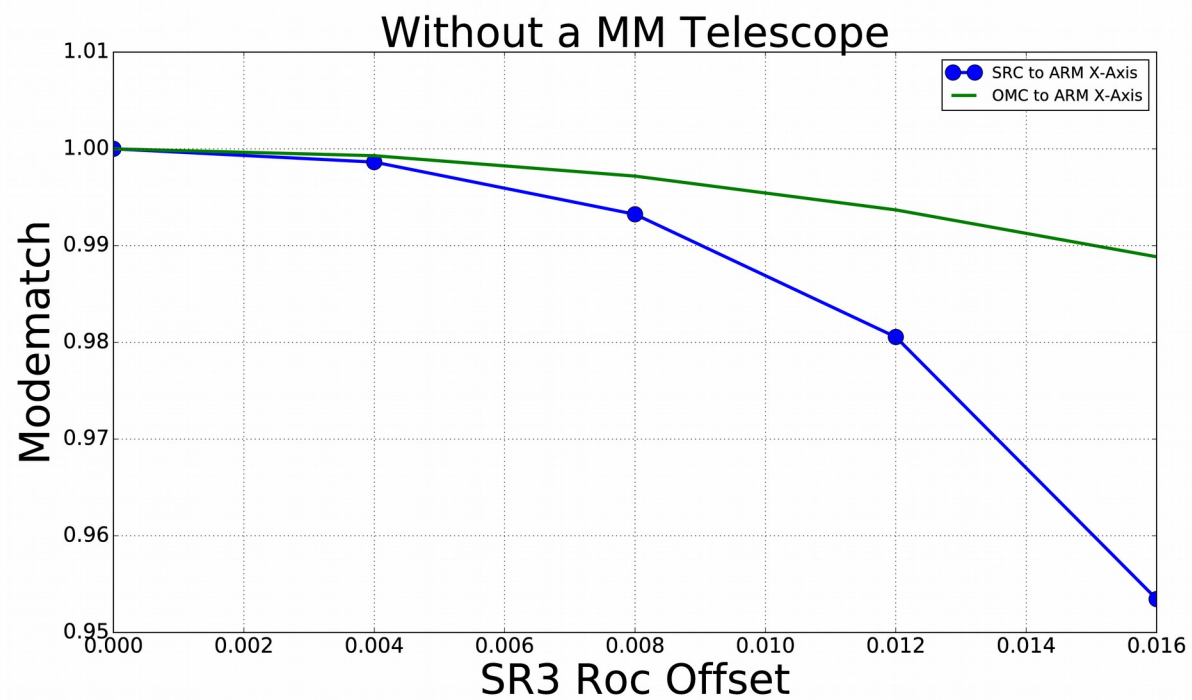
- Changing SR3 curvature by up to 2cm
  - It's clear to see that that mode-mismatching SR3 hurts us, especially in the Squeezed IFO



# Sensitivity Outputs

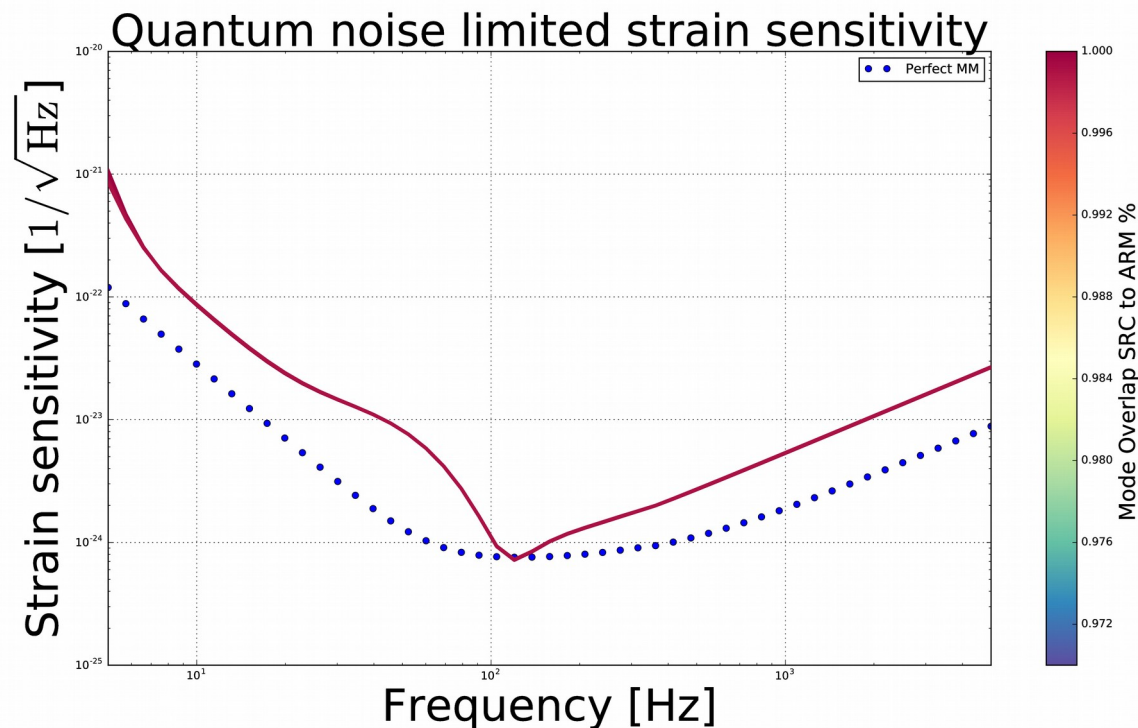
- Changing SRM curvature by up to 50cm (this is huge!)
  - Not terrible for either squeezed or non-squeezed.





# Checking Ourselves

- Setting the SRM Trans to 1:
  - We should see no effects when varying the curvature as long as we include sufficient higher order modes.

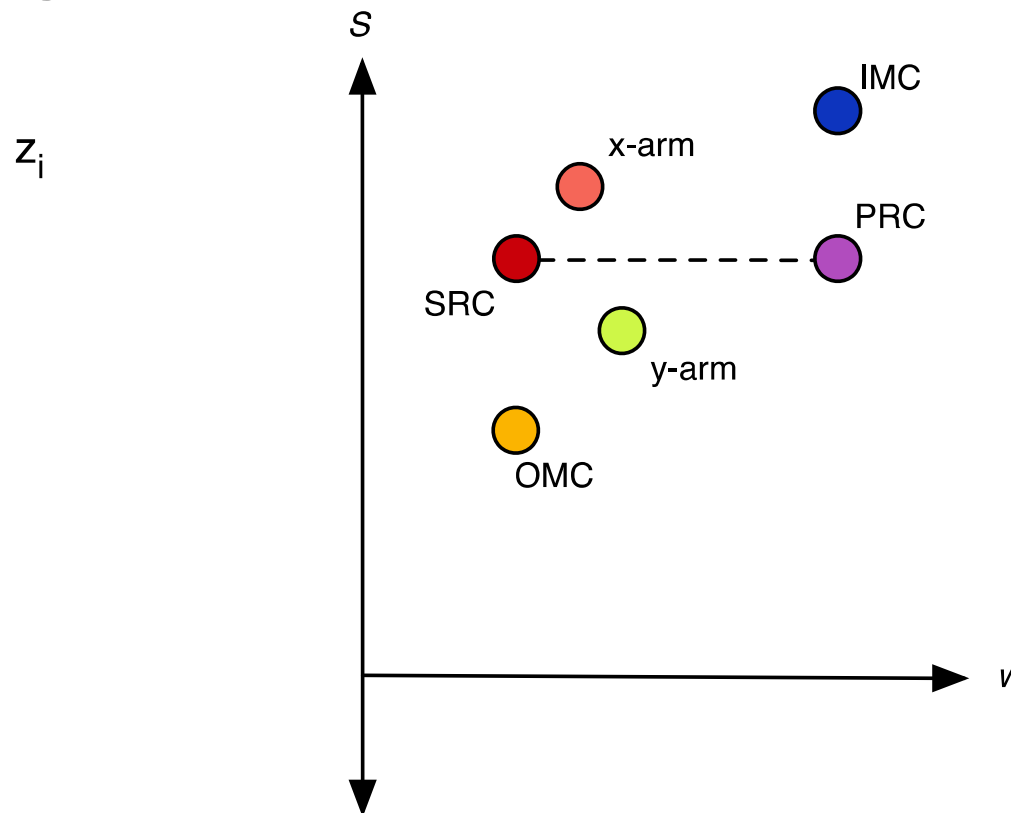




# WS Space

# Next Steps

- Look to see how mode-mismatch affects other cavities and the quoy phase relation between actuators:



# Extra Slides: Finesse+Pykat

- Advantages
  - A full aLIGO Fourier domain analyses
  - Output a QM-limited sensitivity
  - You can add optics as needed for design purposes
- Disadvantages
  - Complexity (multiple cavities & a lot of code)
  - Need careful consideration of higher order modes
    - The larger the mismatches, the more coupling to HOMs, the more computationally expensive it is to run this analysis.