

# Summary of biological replicates

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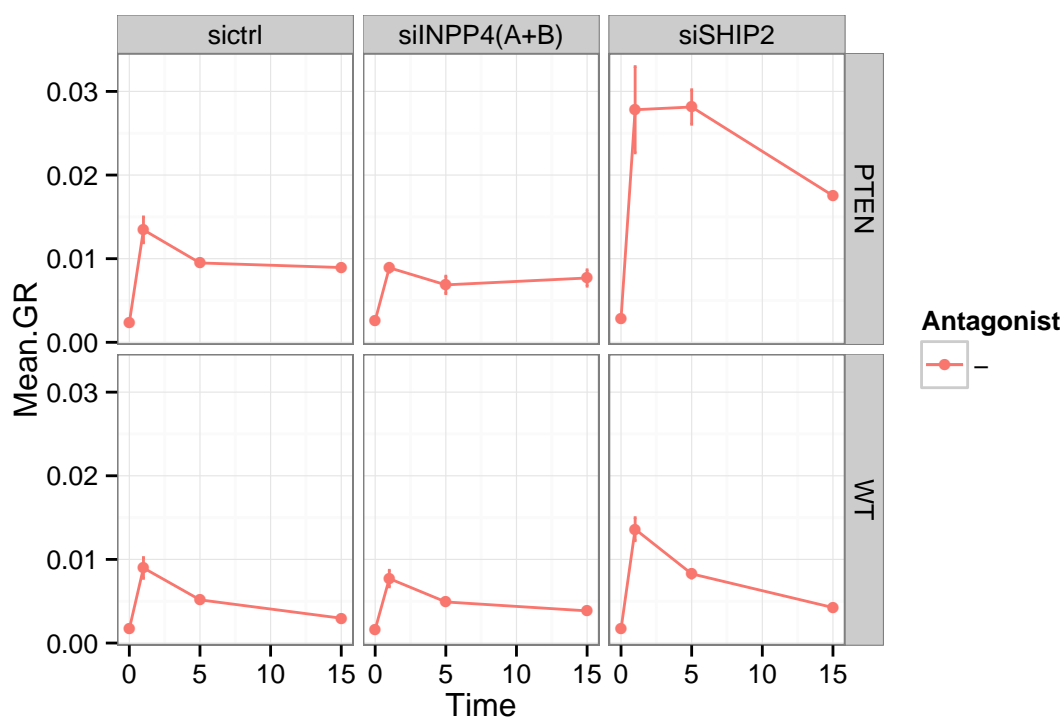
*18 August 2015*

In this report I combine all available data in one file. The purpose is to overview the data and to understand what replicates to use in the modeling project.

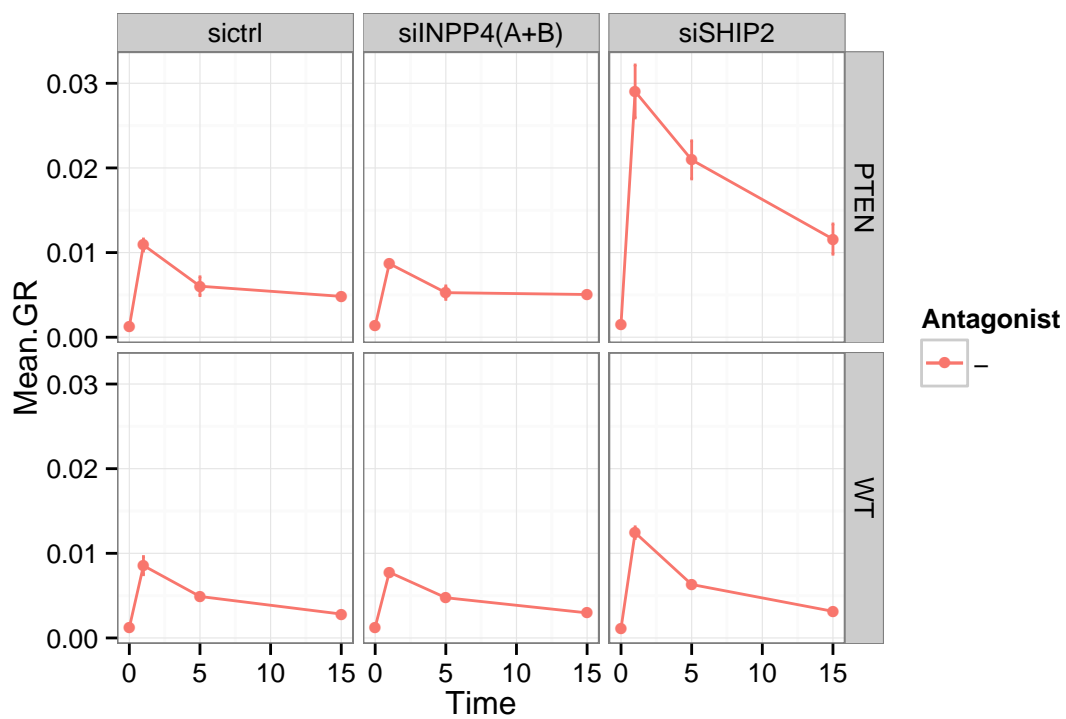
I add plots of each species splitted by file names.

## P3\_4

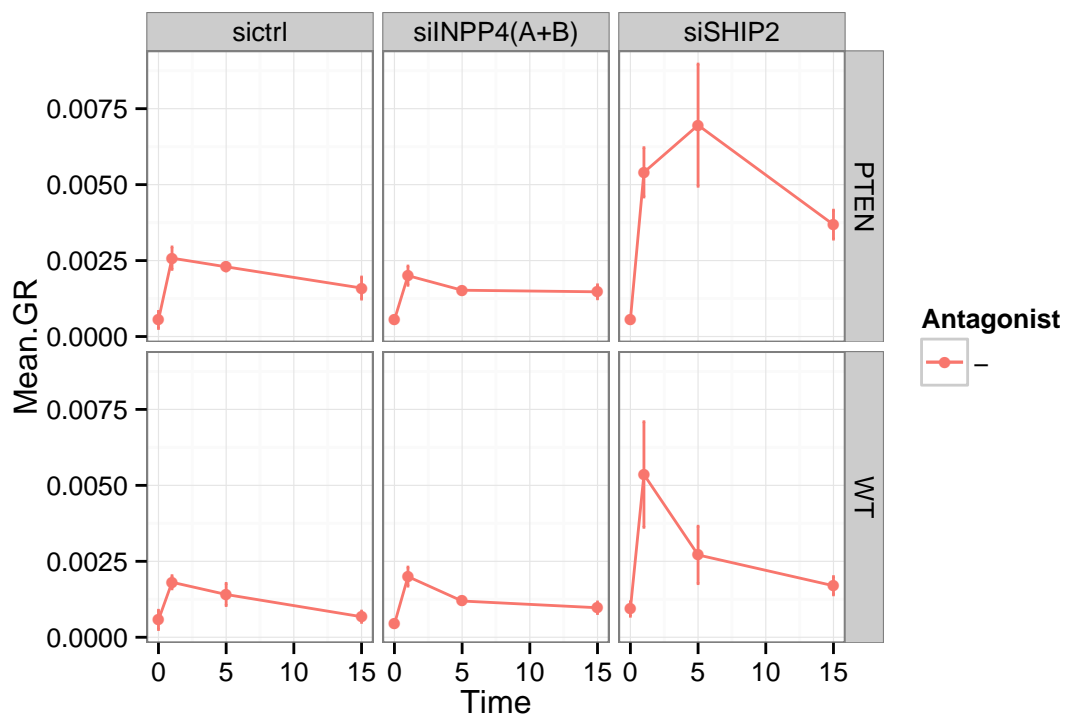
2014\_09\_05B\_c4\_FULLn1



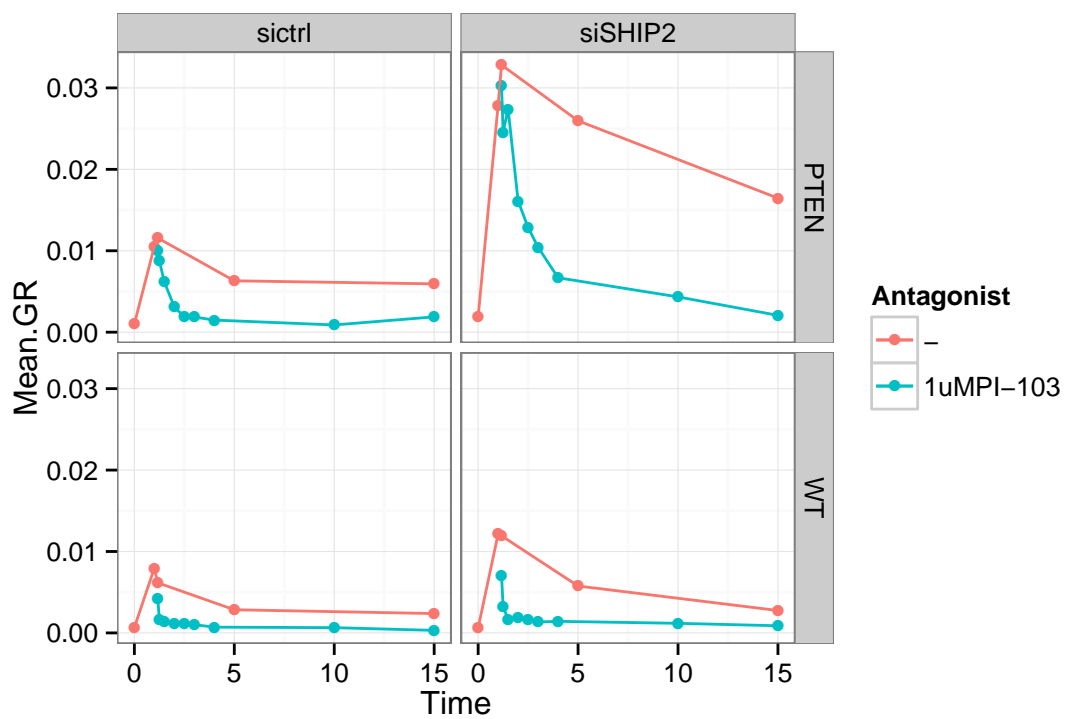
2014\_09\_19B\_c4\_FULLn2



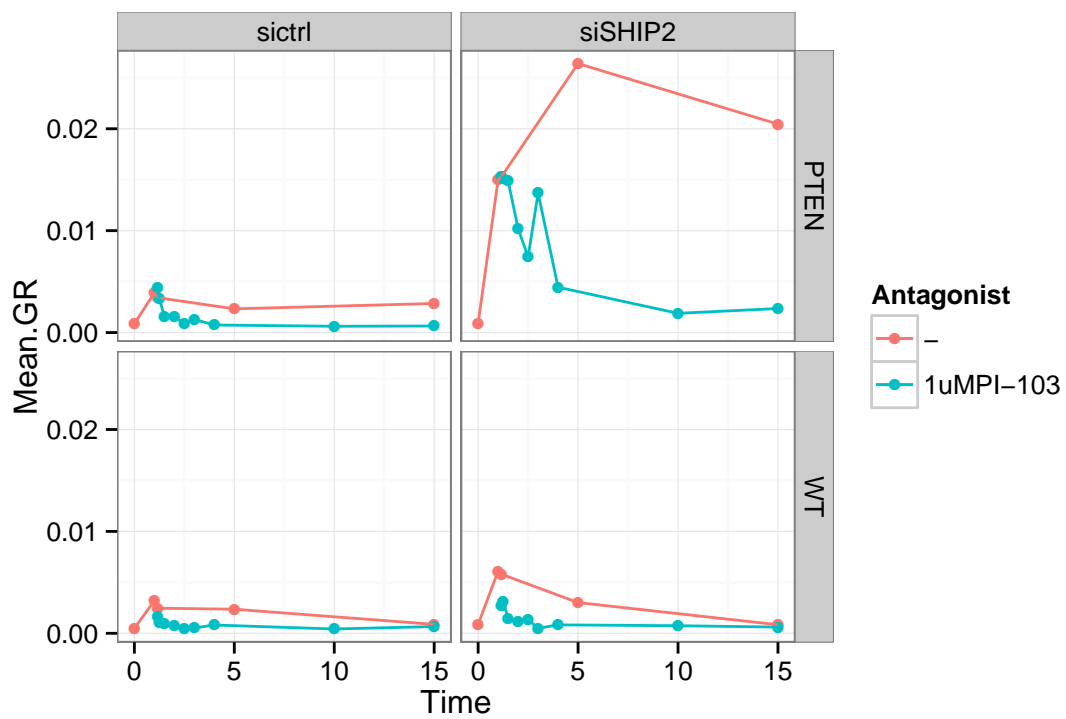
2015\_02\_23B\_c4\_FULLLn3



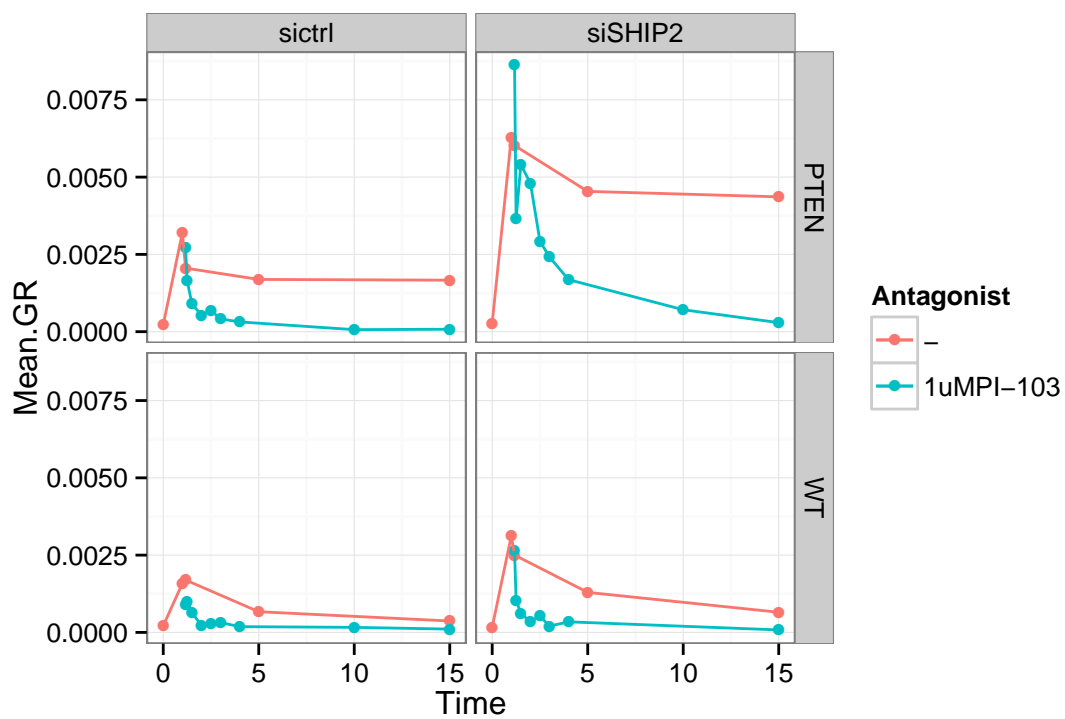
2014\_12\_18A\_c4\_full



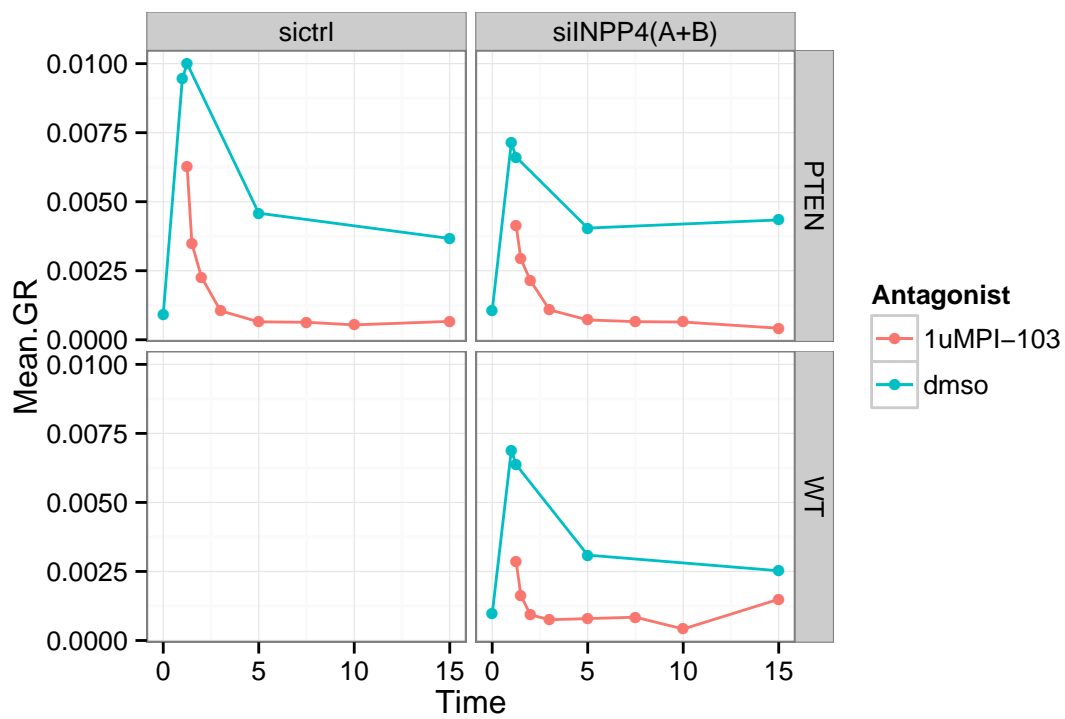
2015\_02\_06B\_c4\_pten\_ship2\_ptenship2\_n2



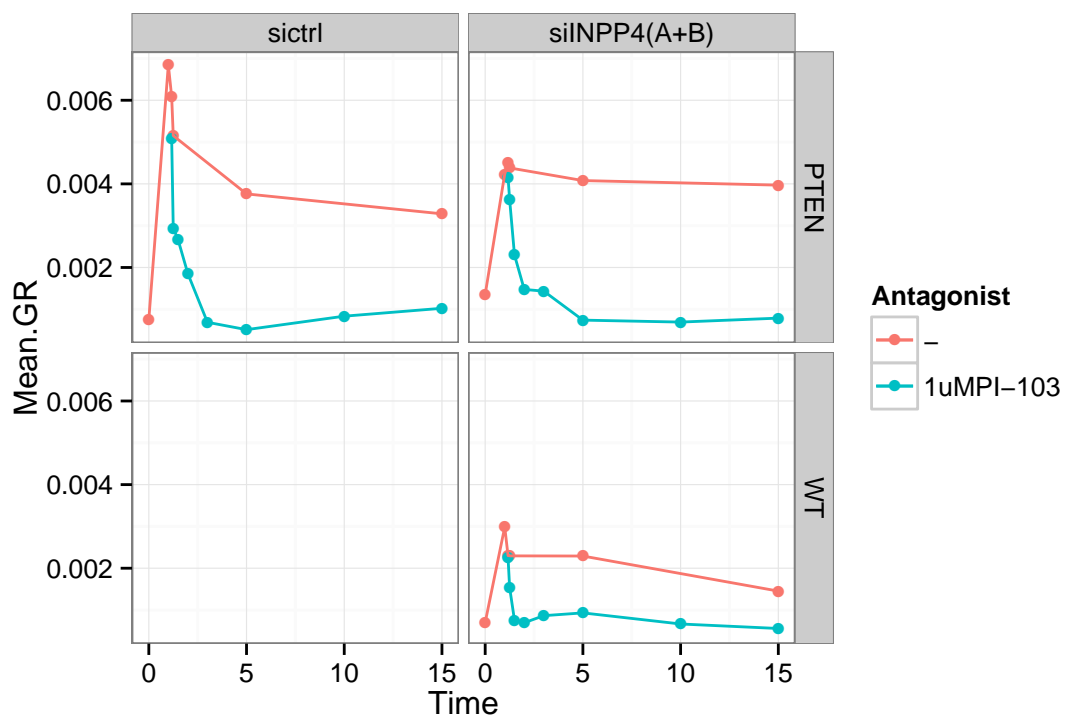
2015\_04\_04A\_c4\_wt\_ship2\_pten\_ptenship2\_N3



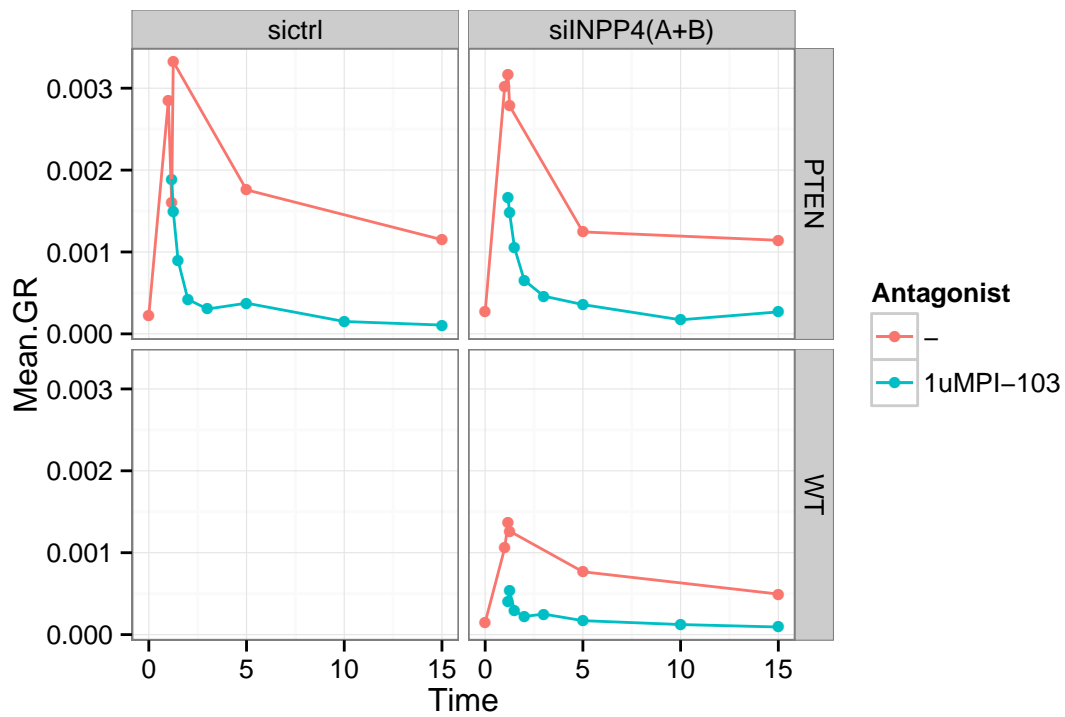
2014\_12\_12A\_c4\_wt\_ab\_pten\_tc



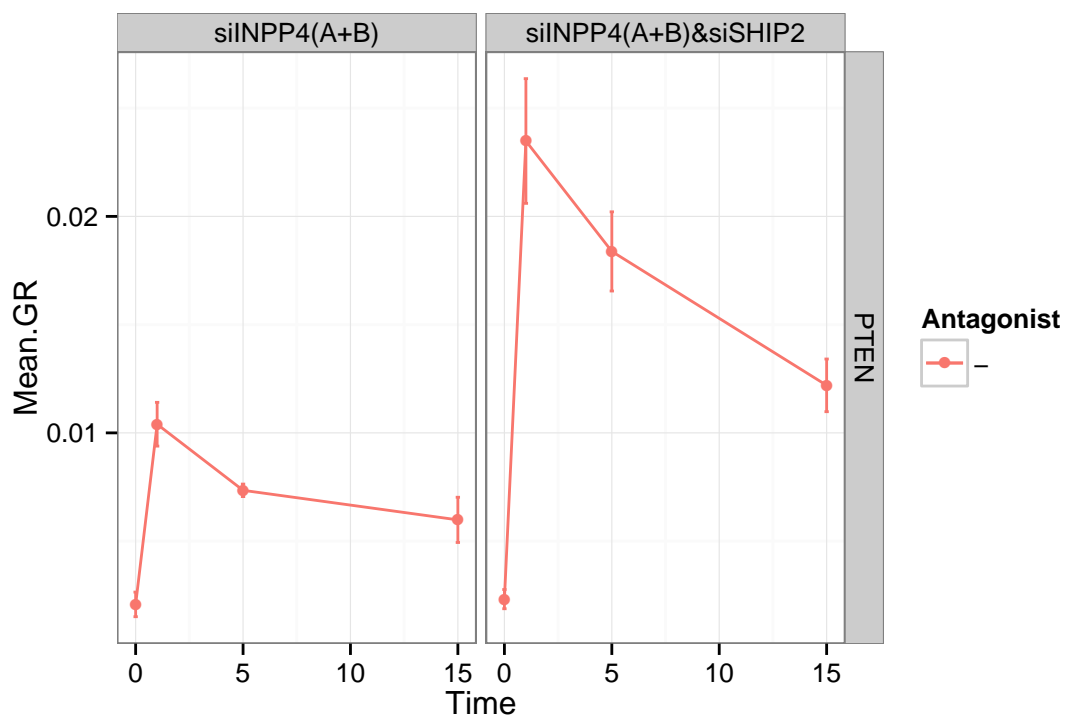
2015\_02\_14B\_c4\_pten\_ab\_pab\_tcN2\_pabVSpabsN2



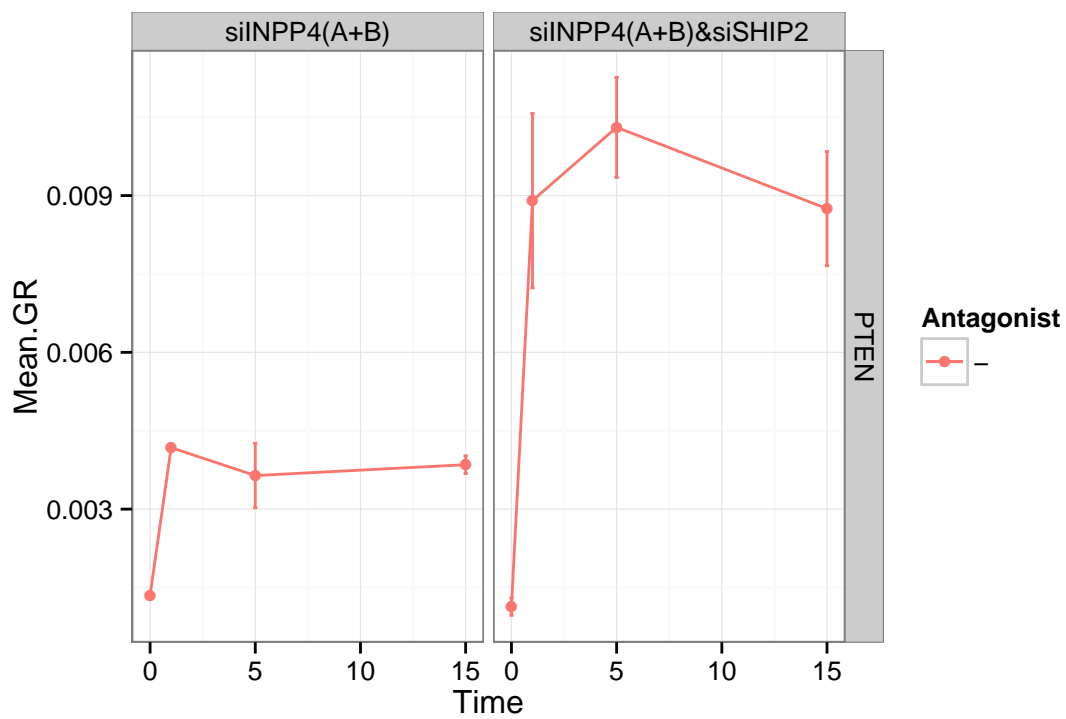
2015\_03\_06A\_c4\_pi34p2Trans\_n3



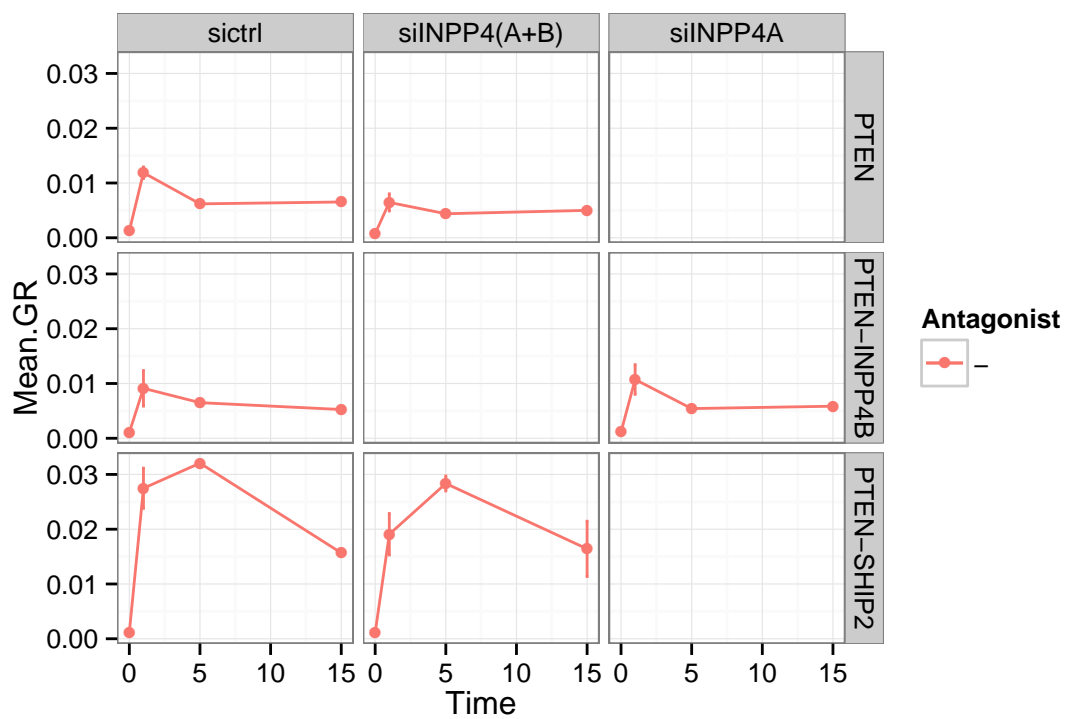
2014\_10\_10A\_c4\_pabVSpabs



2015\_02\_14B\_c4\_pabVSpabsN2

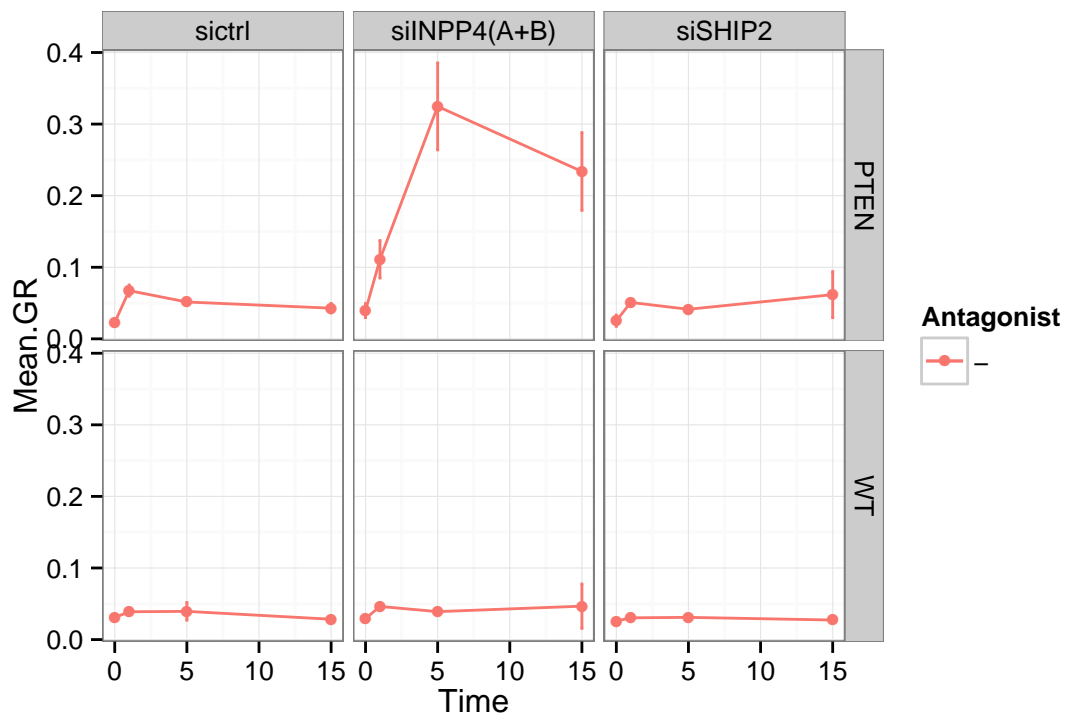


2015\_08\_06A\_c4\_PTEN\_crispr\_n1

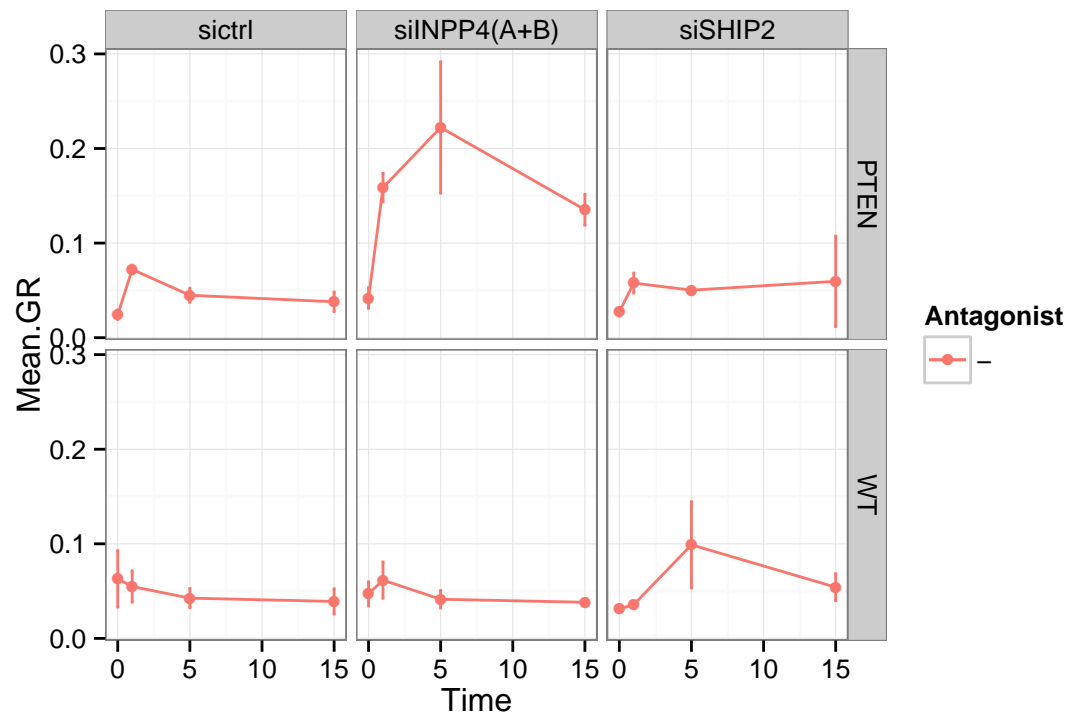


## SA\_P2\_aldehyde\_34

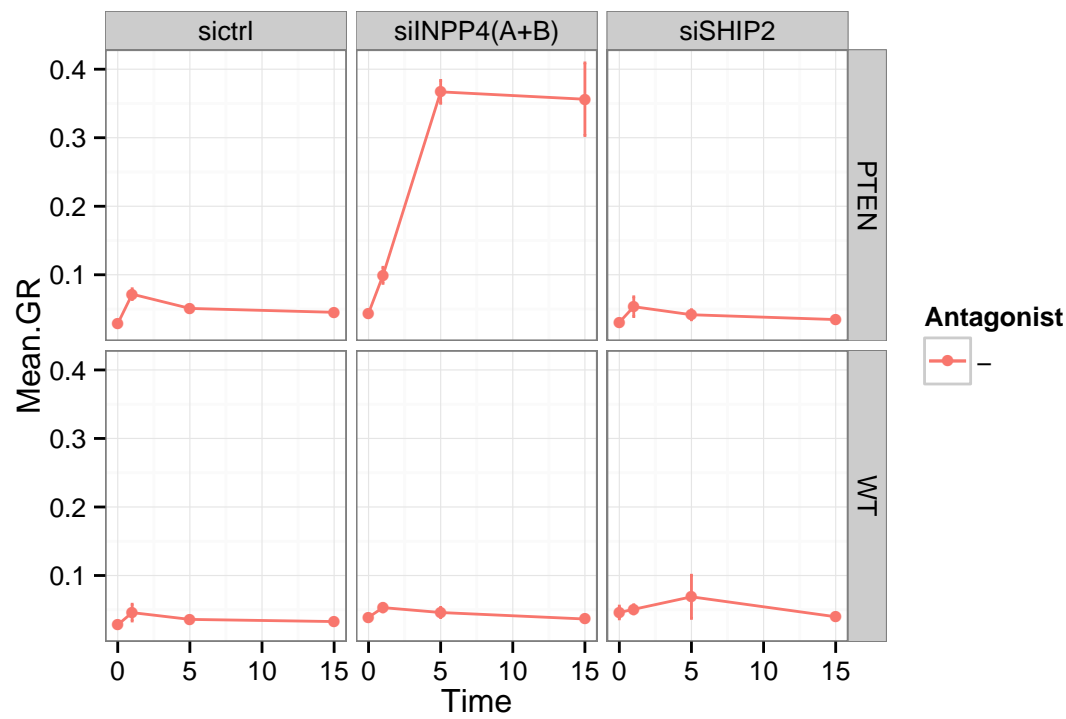
2014\_09\_05A\_c18\_FULLLn1



2014\_09\_19A\_c18\_FULLLn2

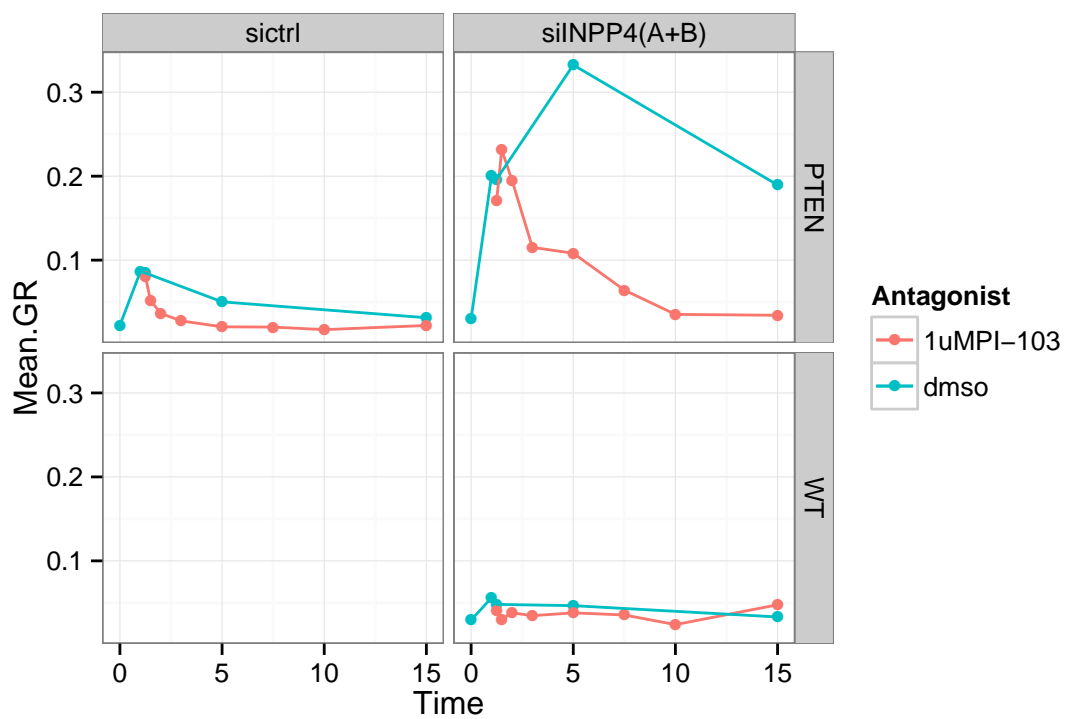


2015\_02\_23A\_c18\_FULLLn3

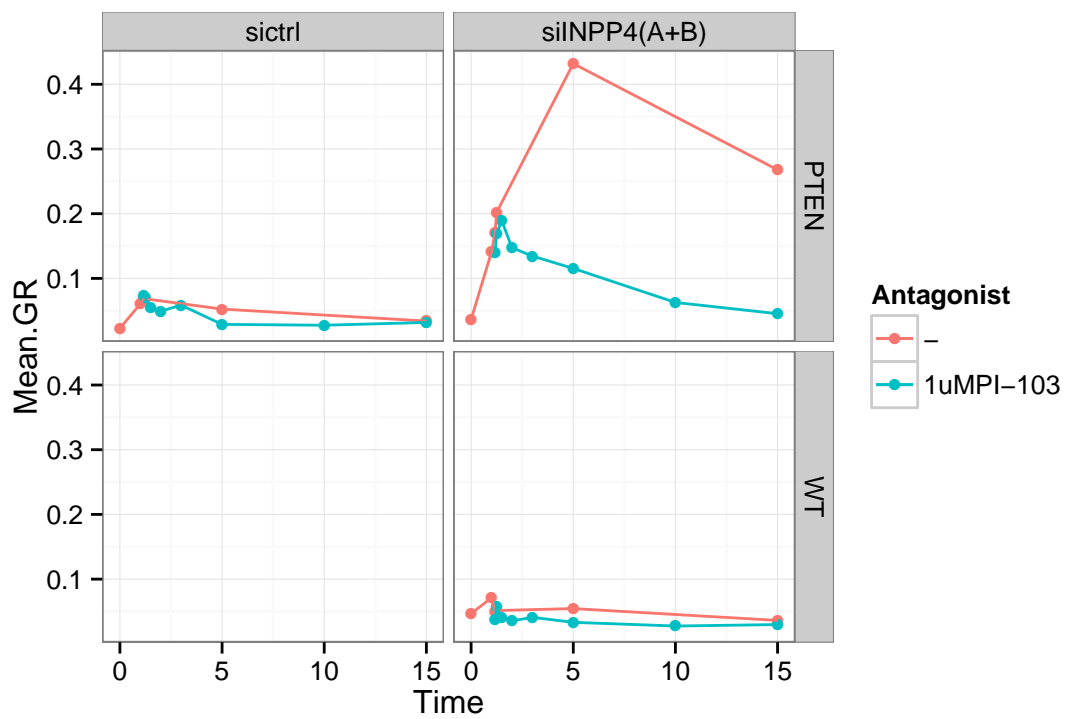


2014\_12\_12B\_c18\_wt\_ab\_pten\_tc

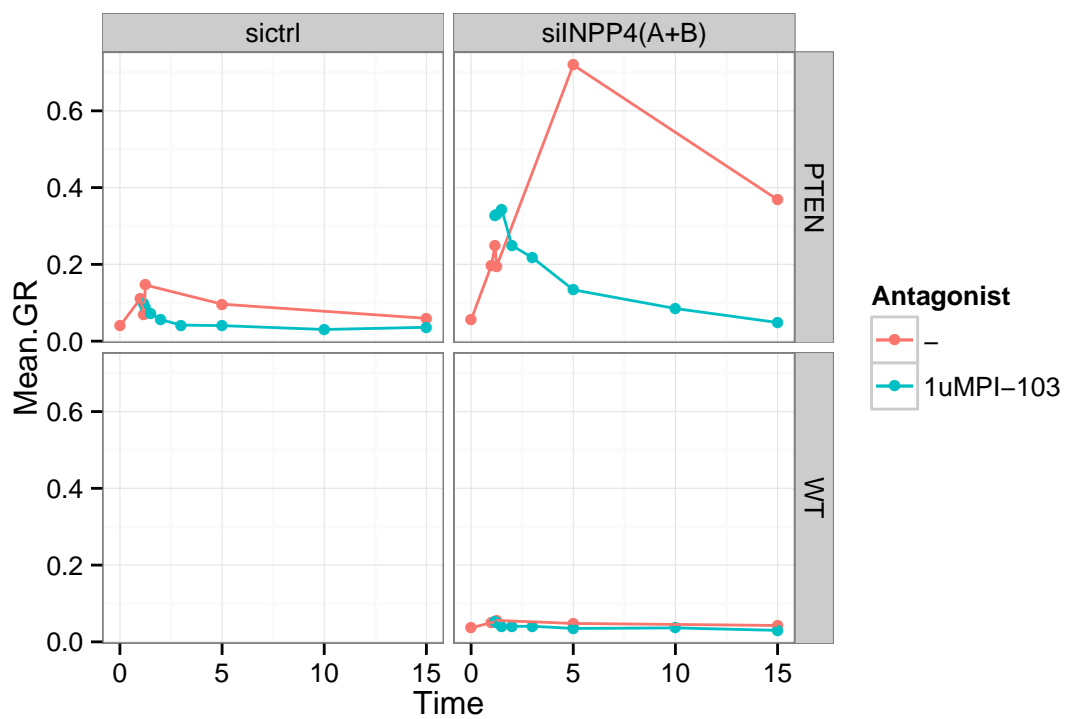




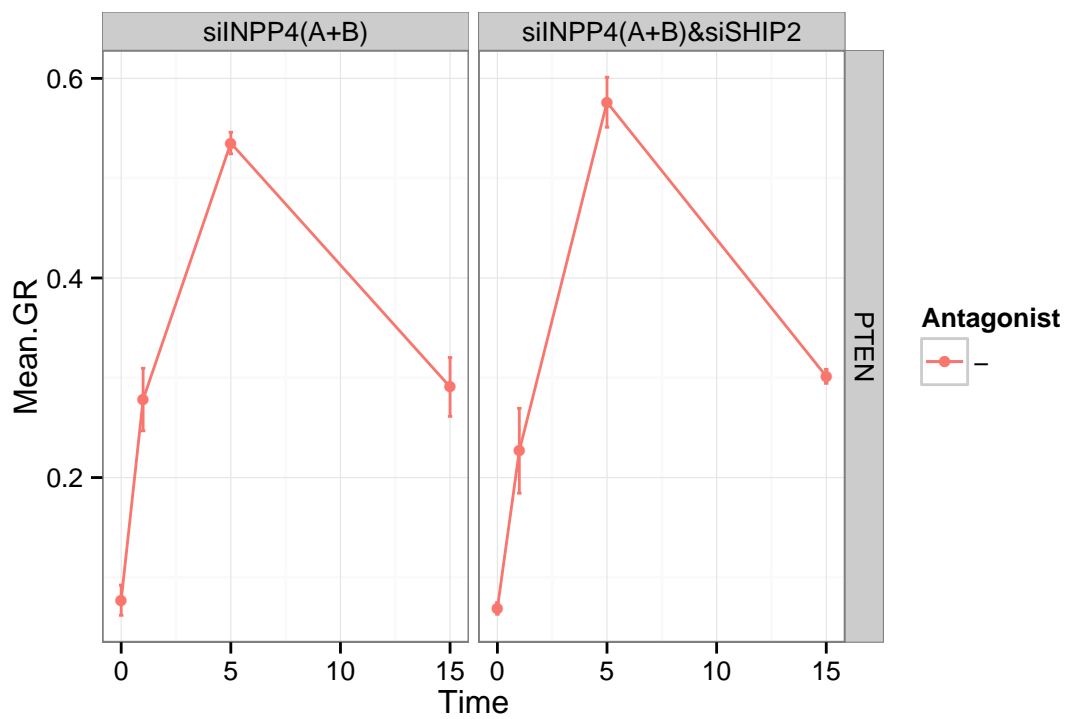
2015\_02\_14A\_c18\_pten\_ab\_pab\_tcN2\_pabVSpabsN2



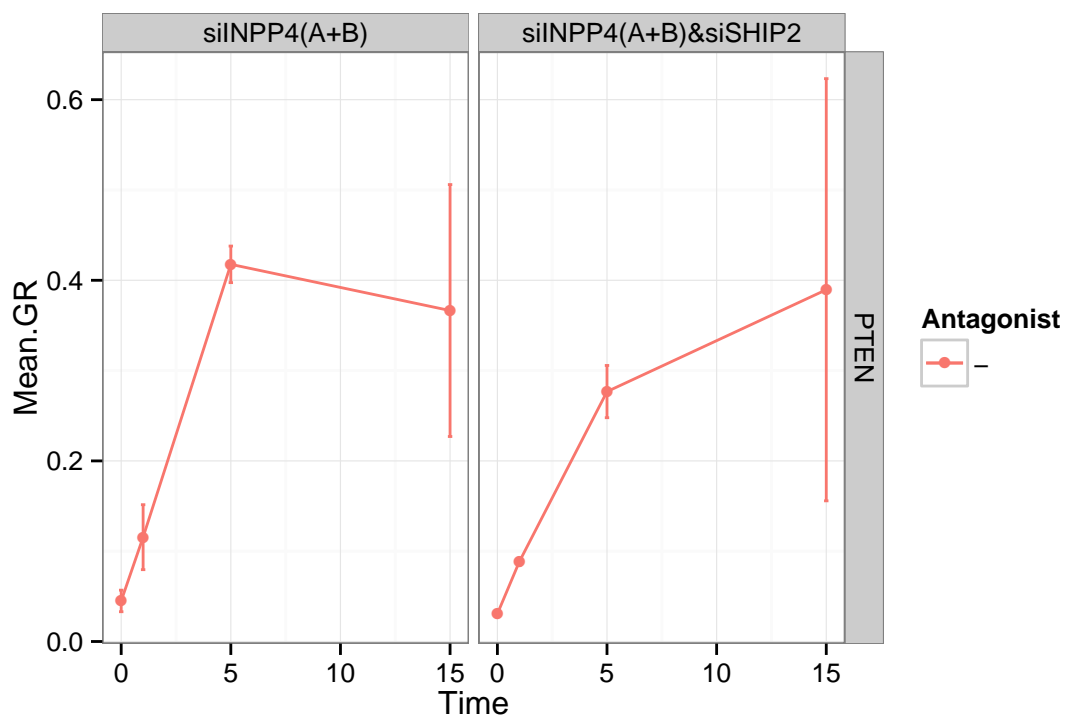
2015\_03\_06B\_c18\_pi34p2Trans\_n3



2014\_10\_10B\_c18\_pabVSpabs



2015\_02\_14A\_c18\_pabVSpabsN2



2015\_08\_06B\_c18\_PTEN\_crispr\_n1

