

gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneAA 40.0 ng/well 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.07 48.0 h=-0.08 66.0 h=-0.07 time (h)

1.5

1.0

0.5

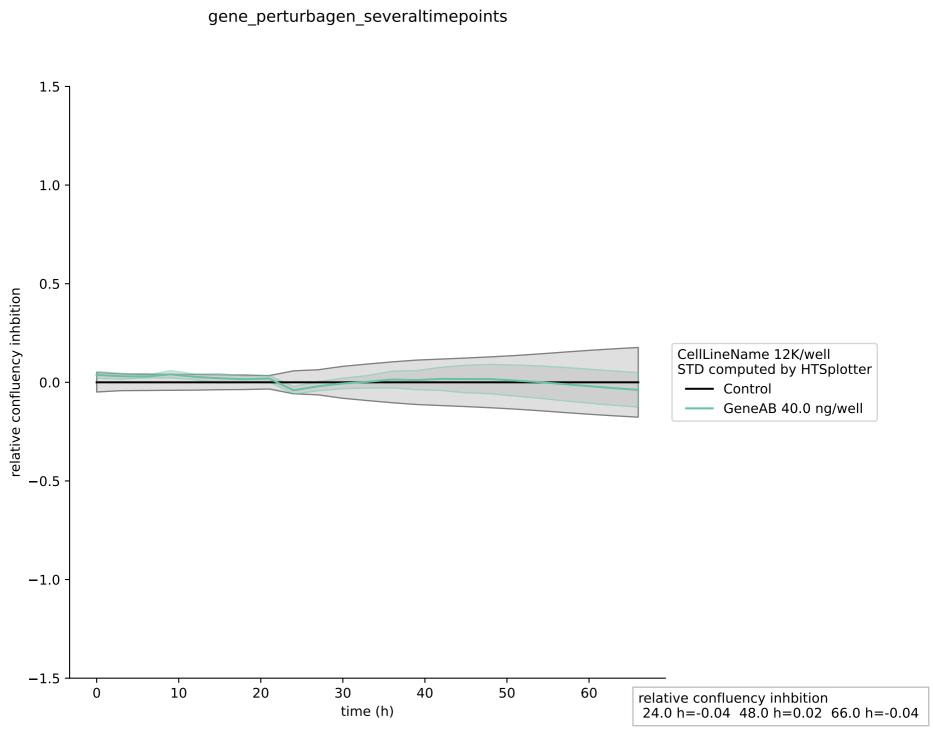
0.0

-0.5

-1.0

-1.5

0



gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneAC 40.0 ng/well 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.09 48.0 h=-0.14 66.0 h=-0.18 time (h)

1.5

1.0

0.5

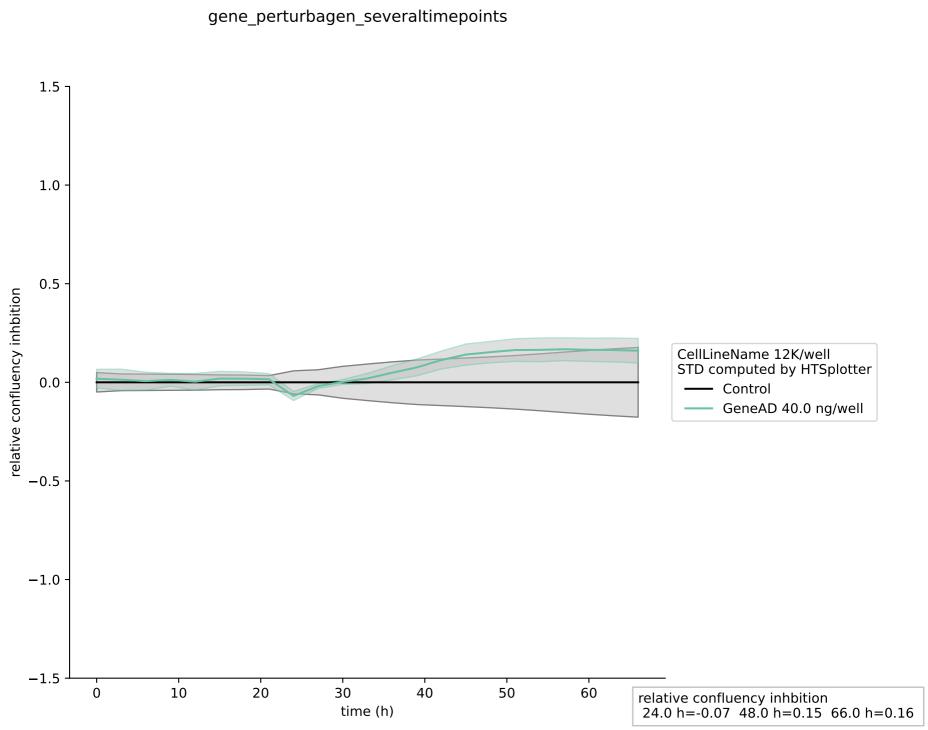
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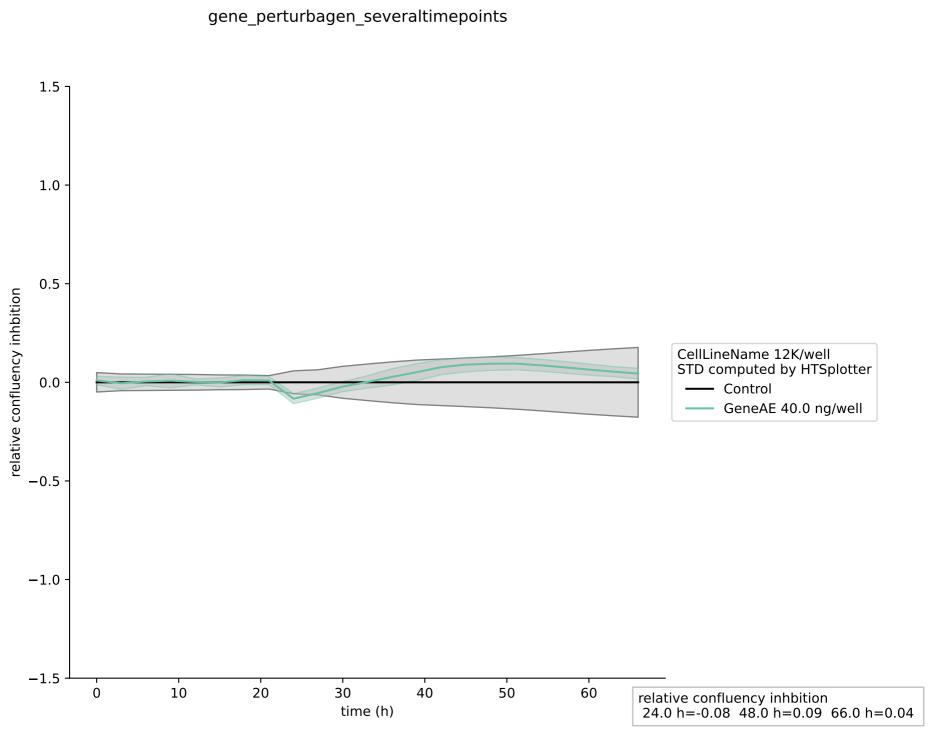
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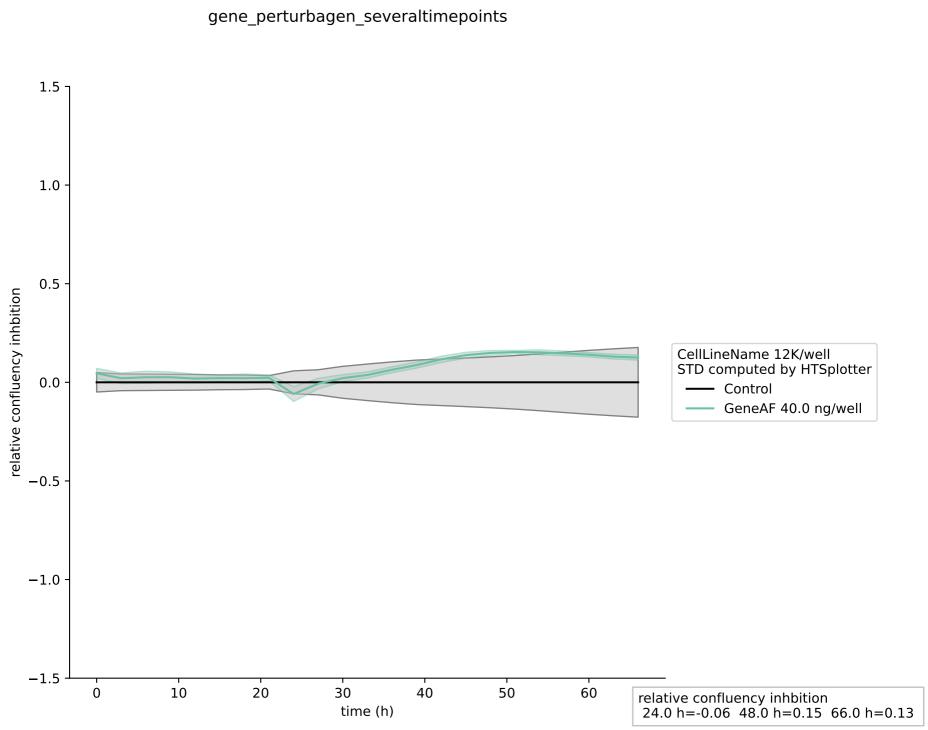
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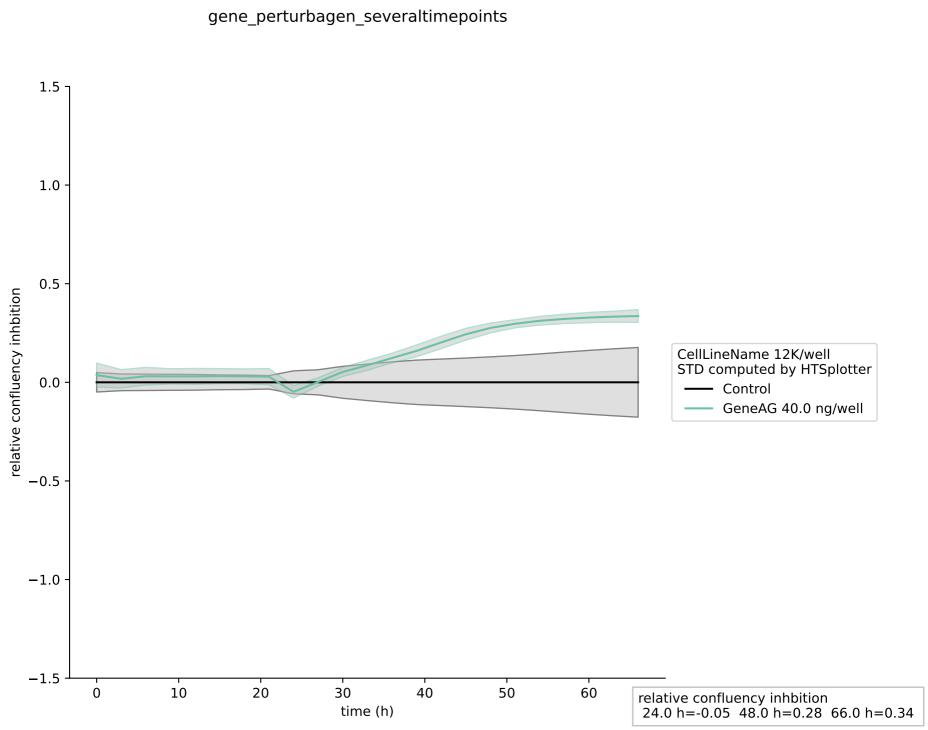
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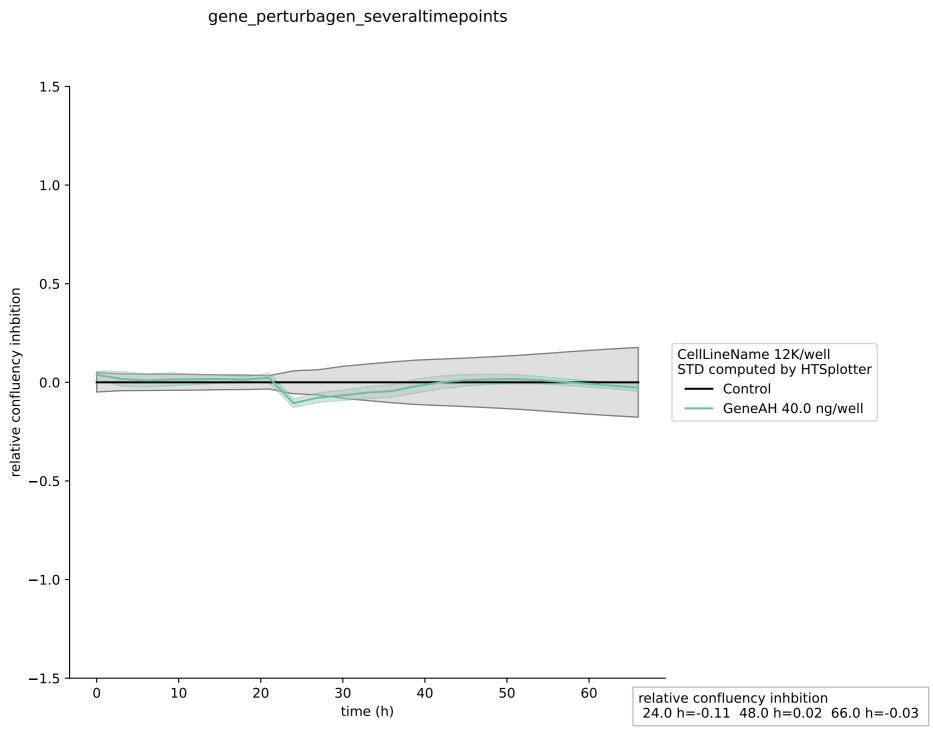
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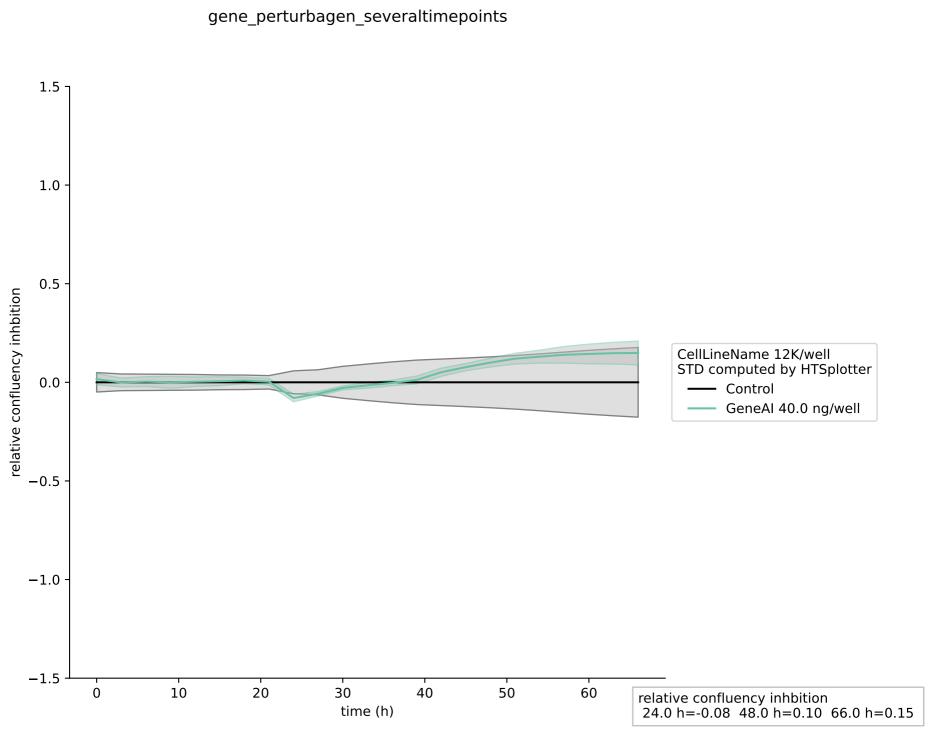


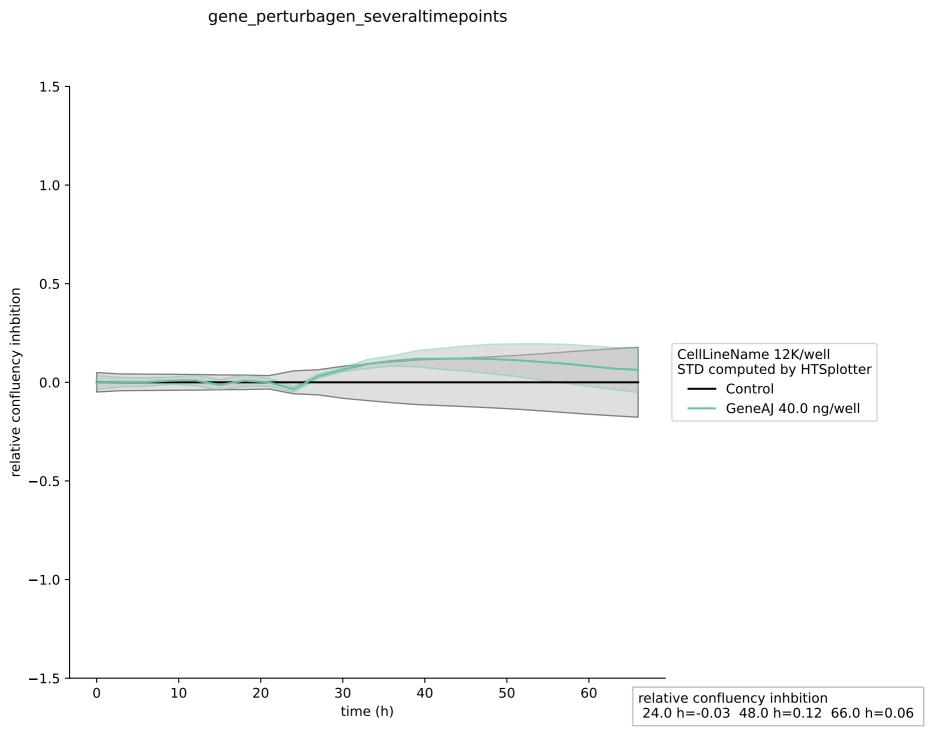


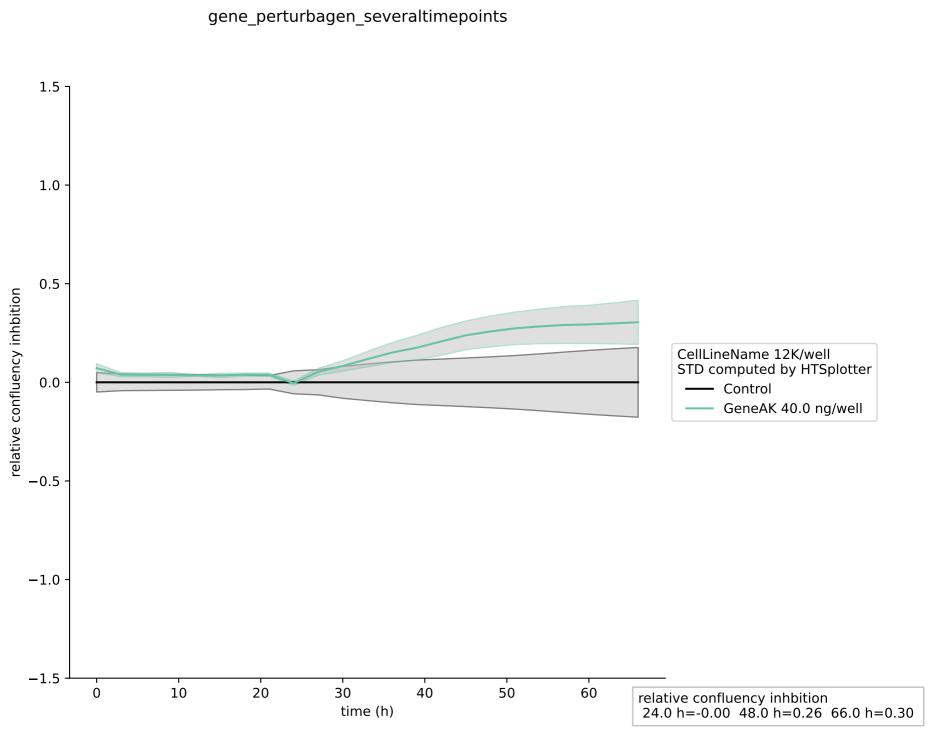


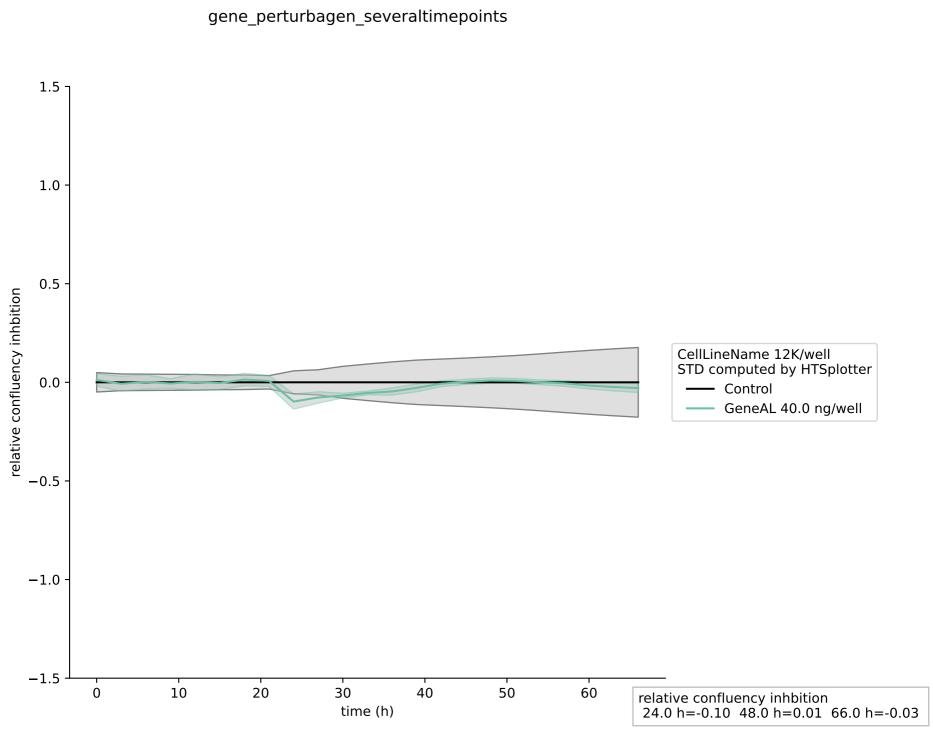




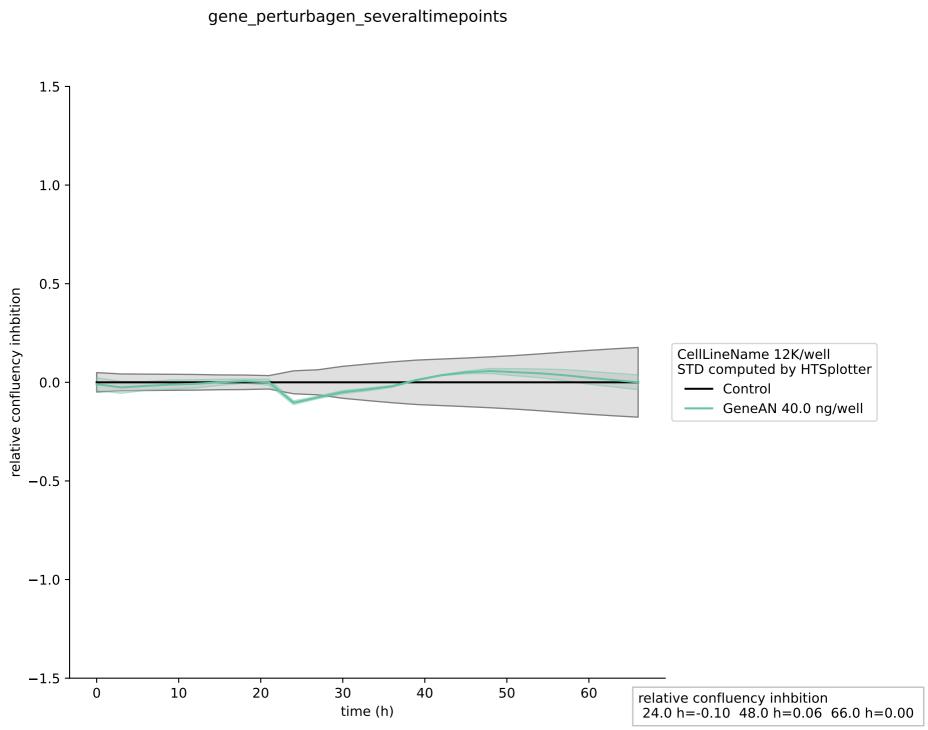


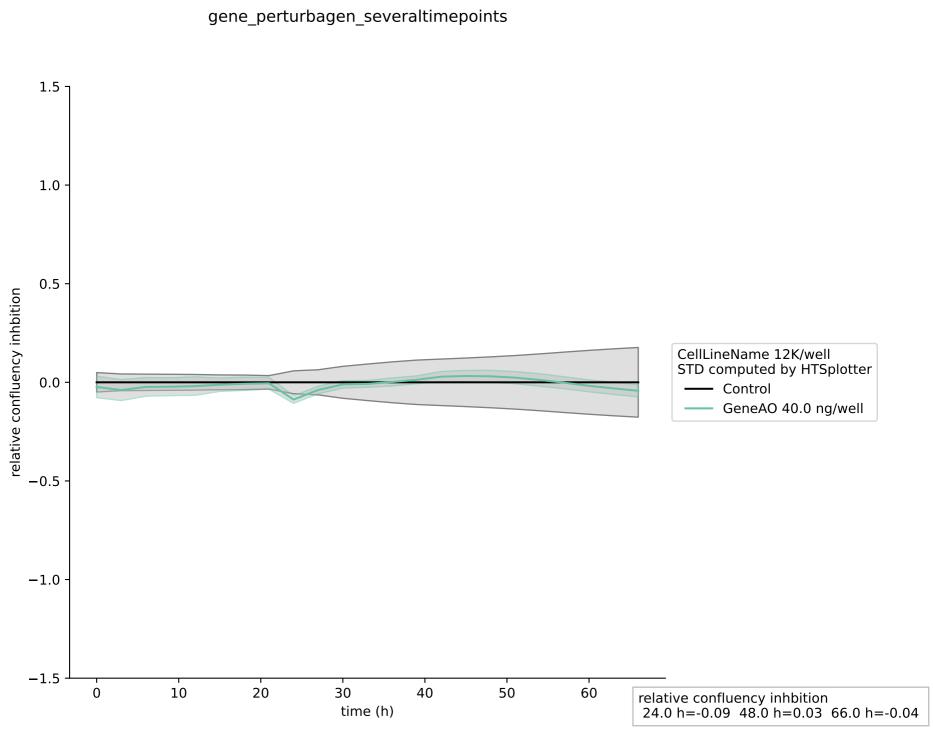


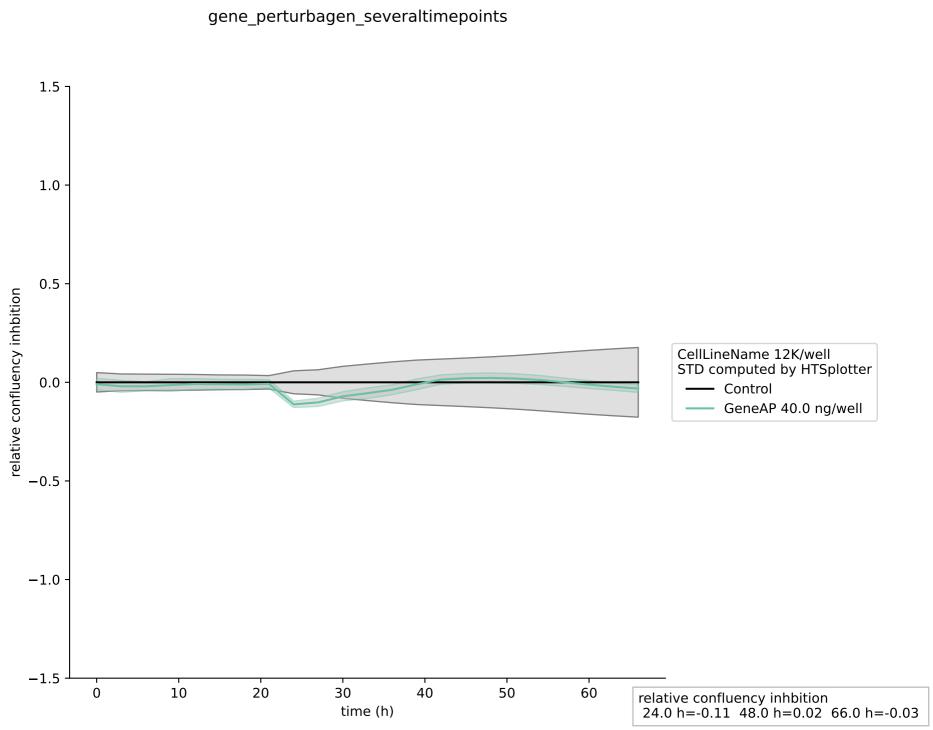


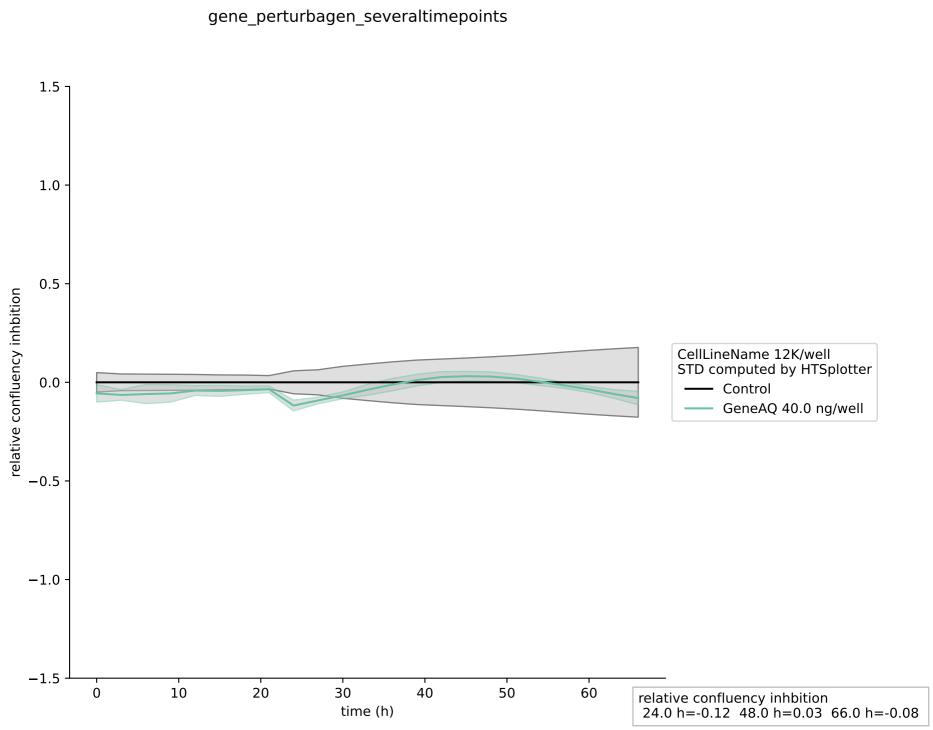


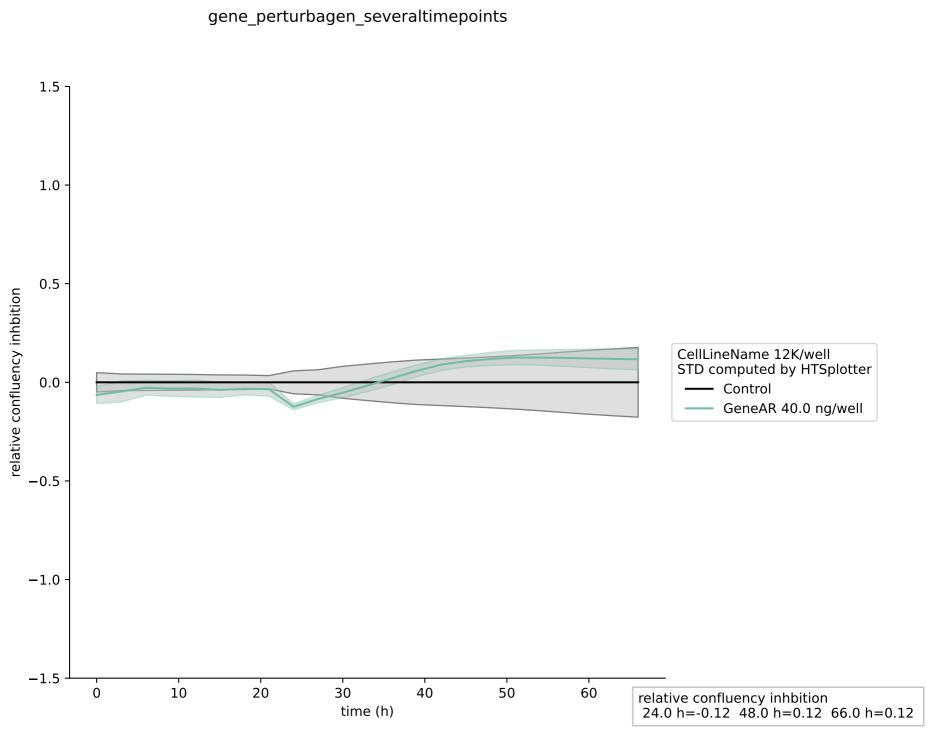
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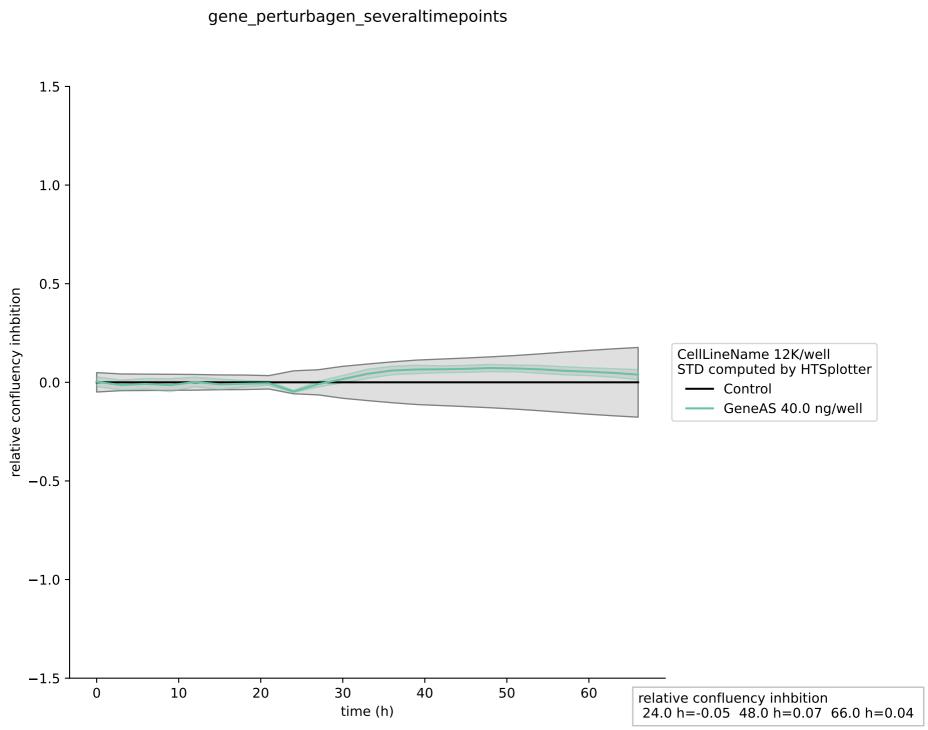




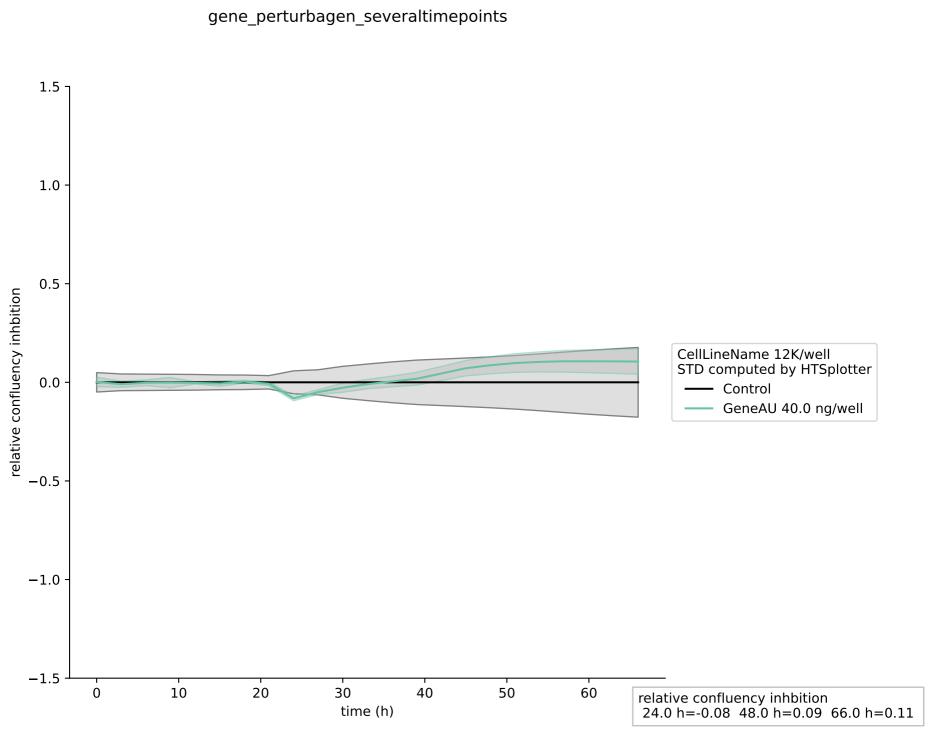


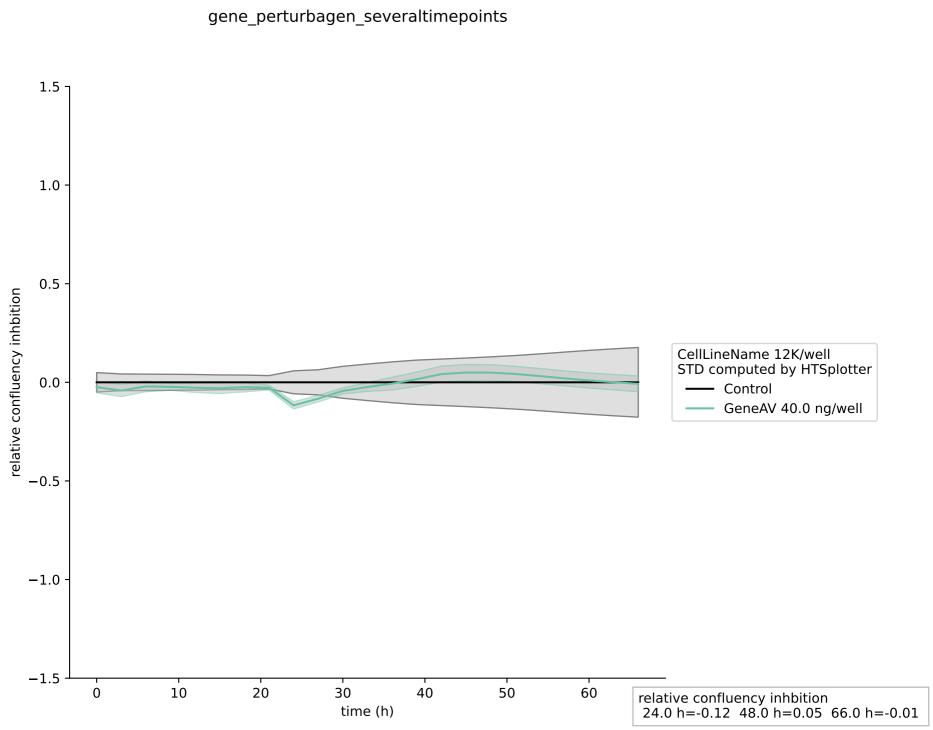


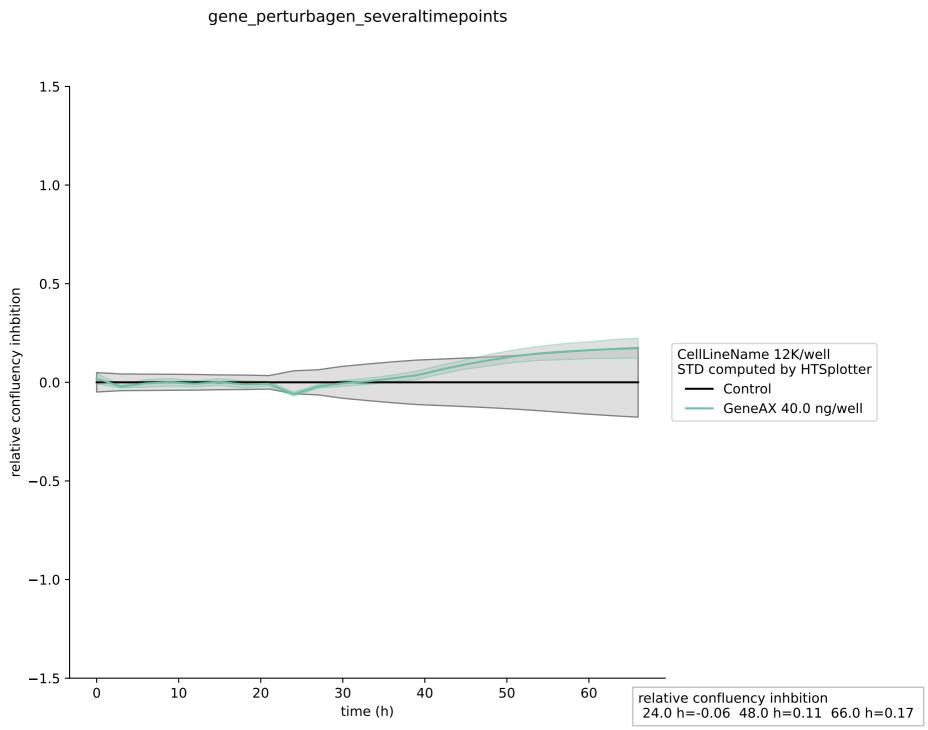


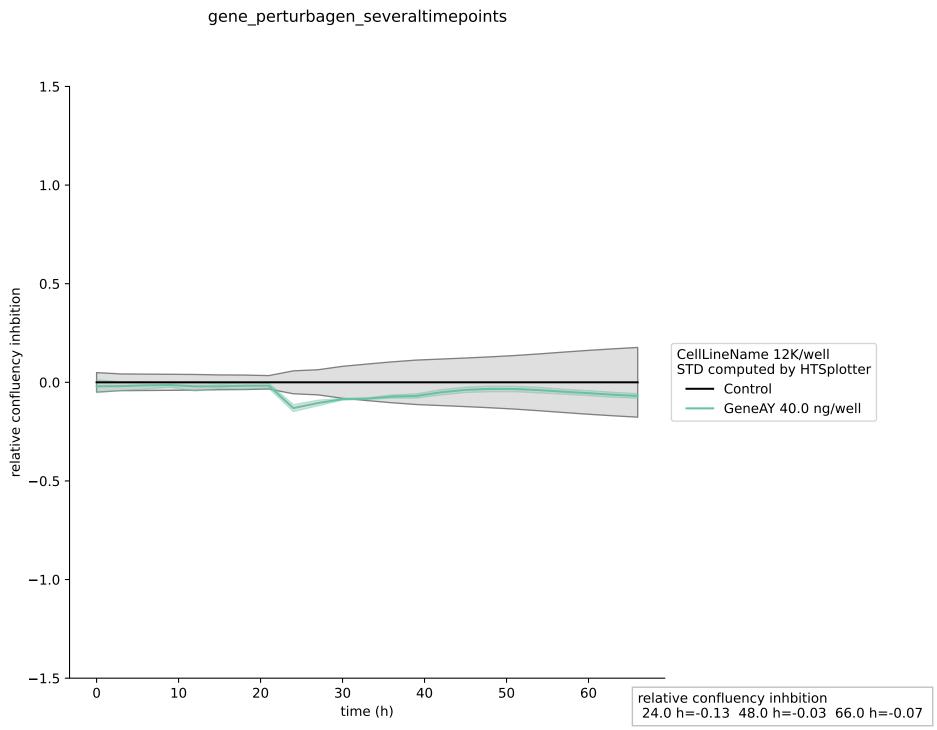


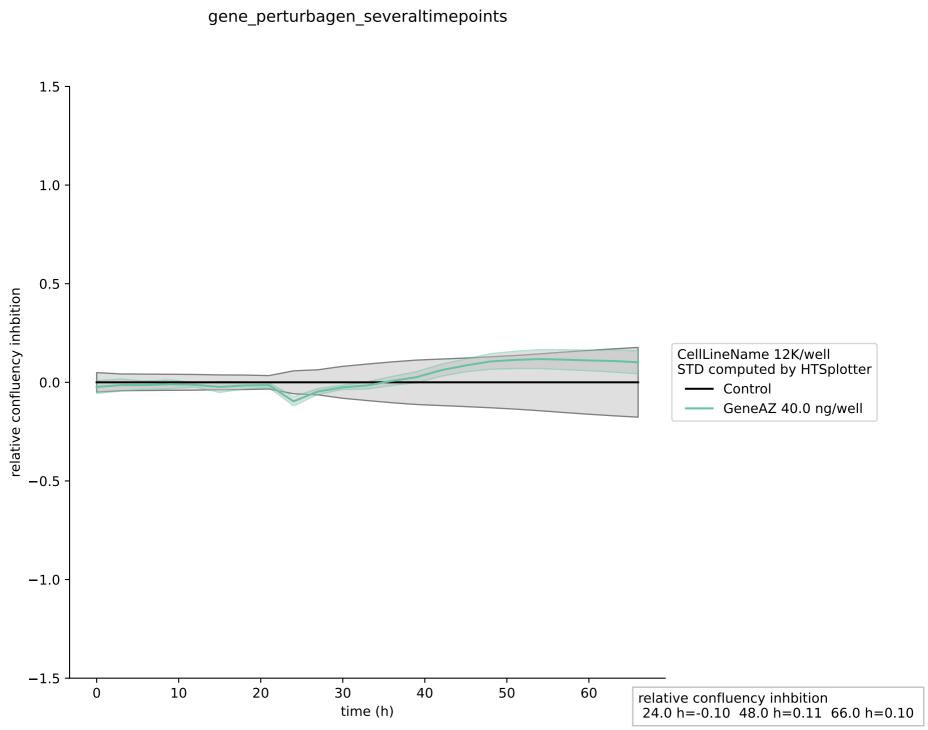
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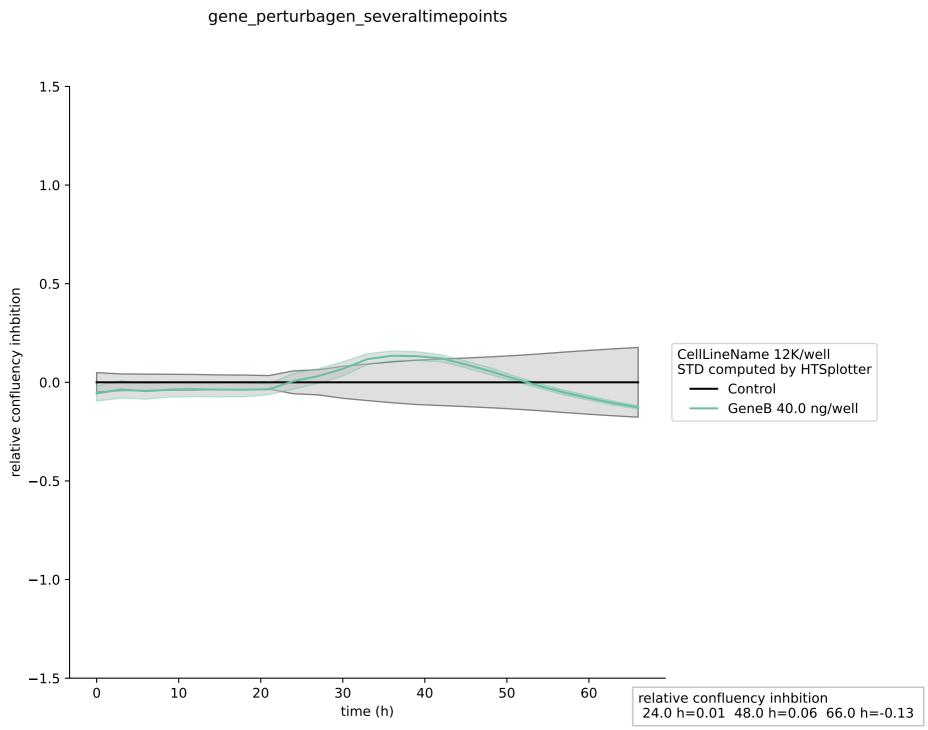




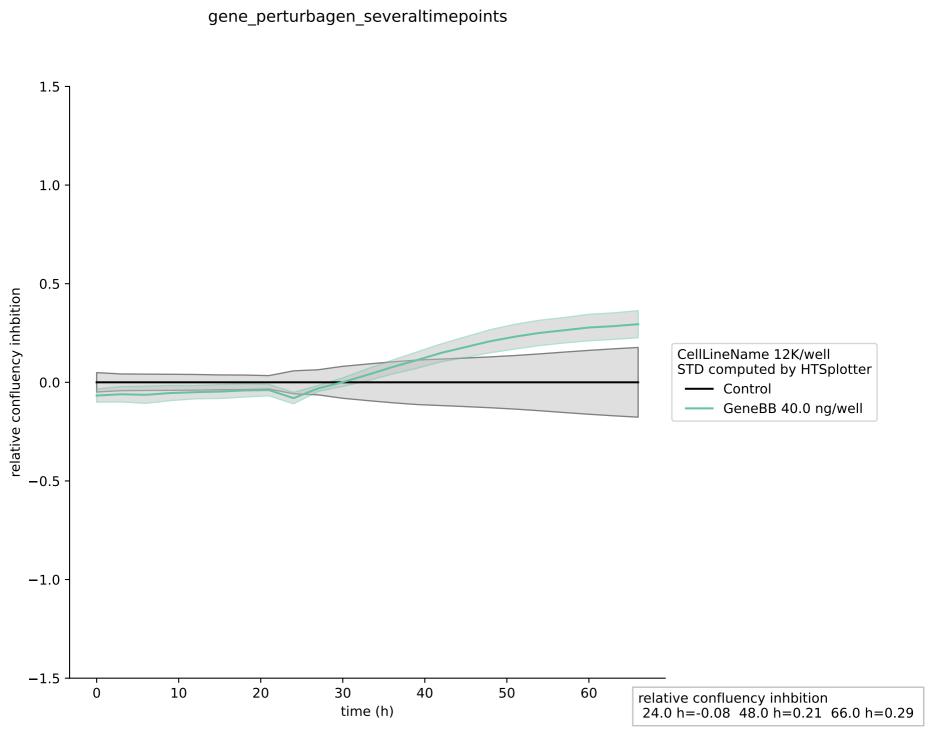


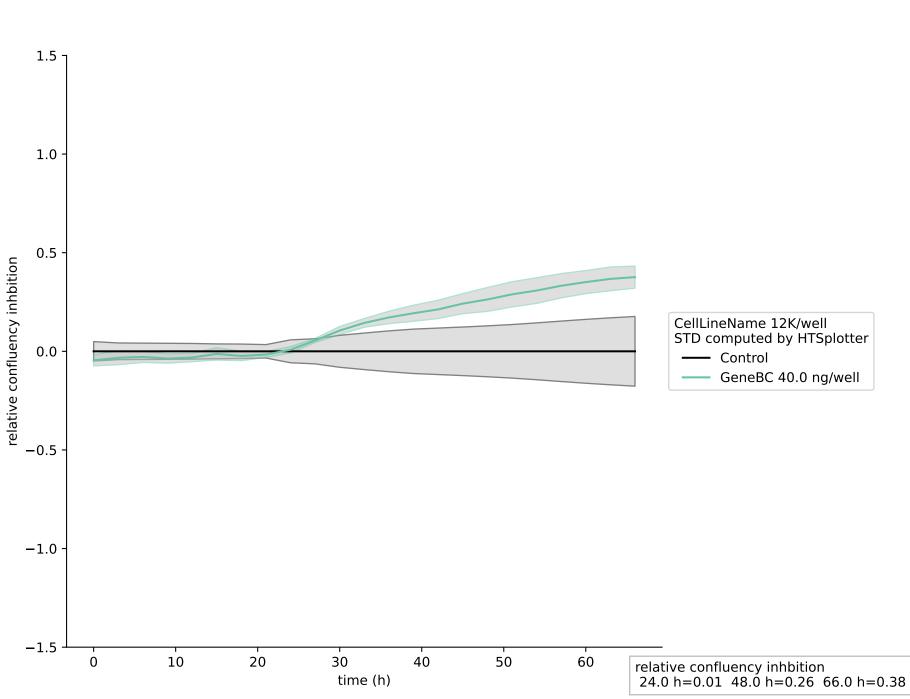




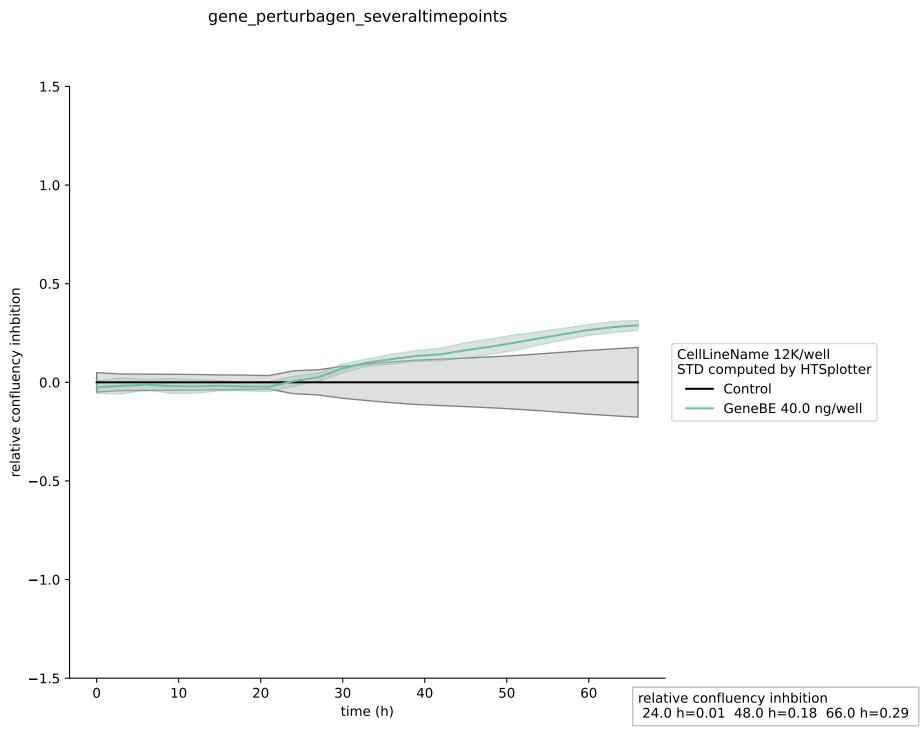


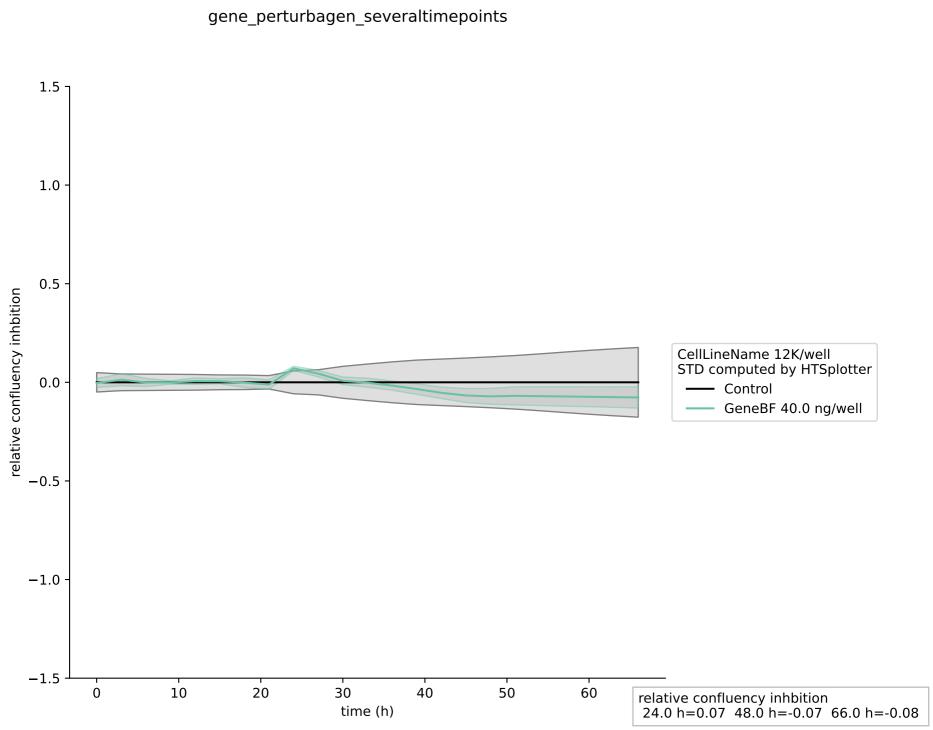
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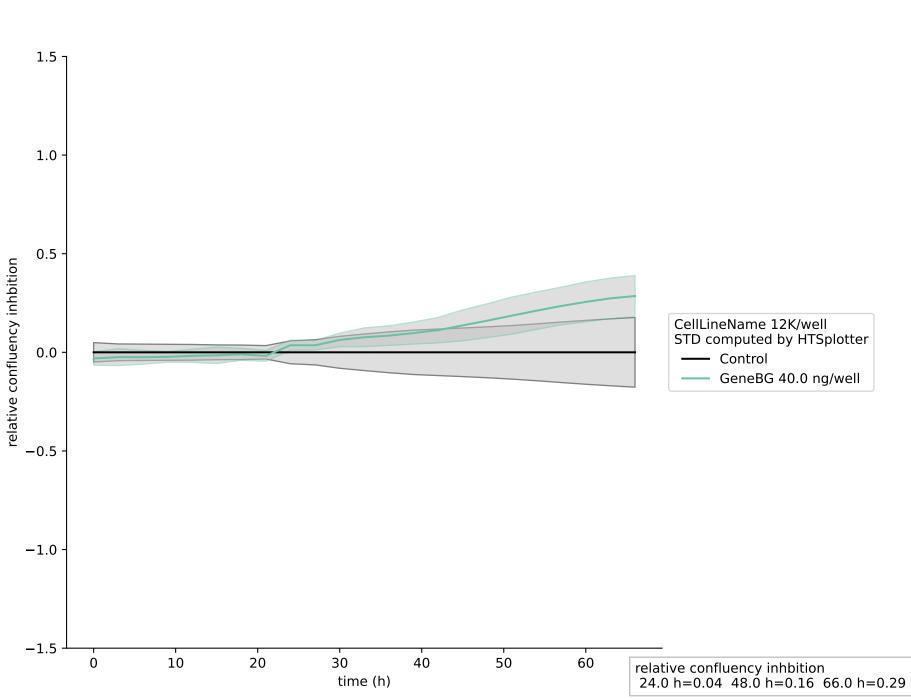




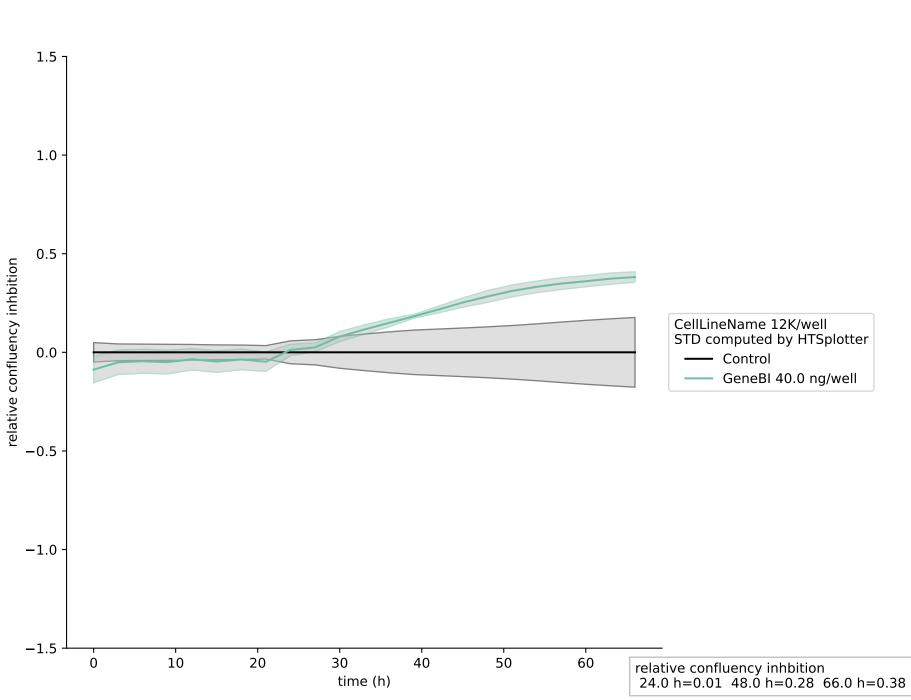
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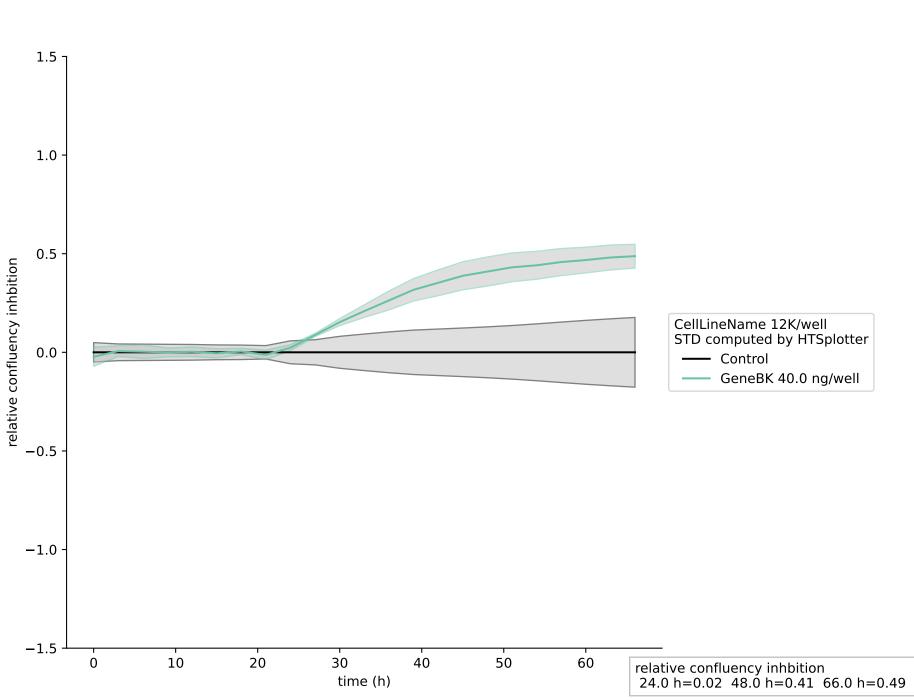




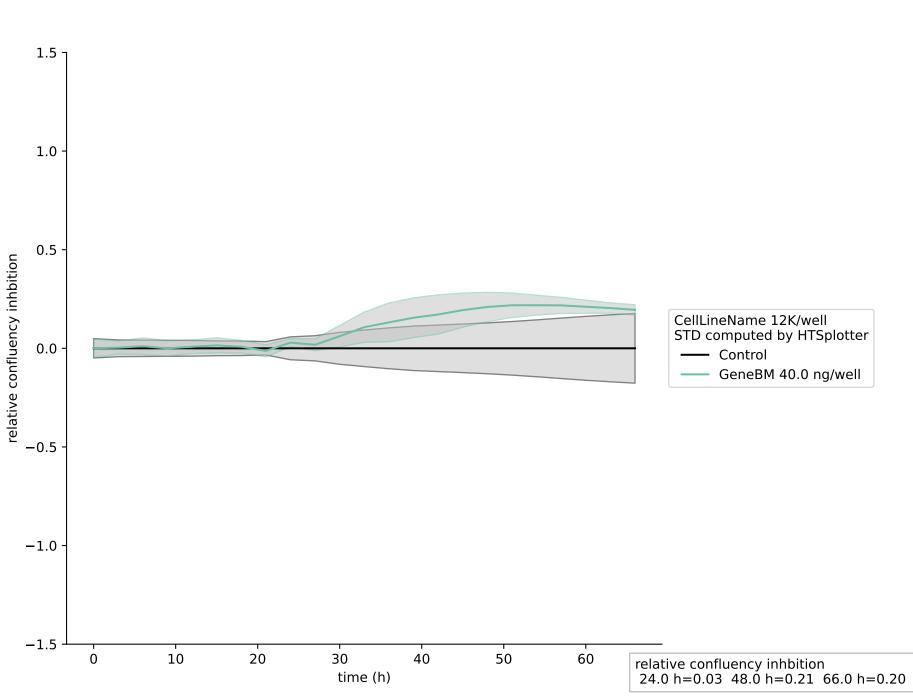
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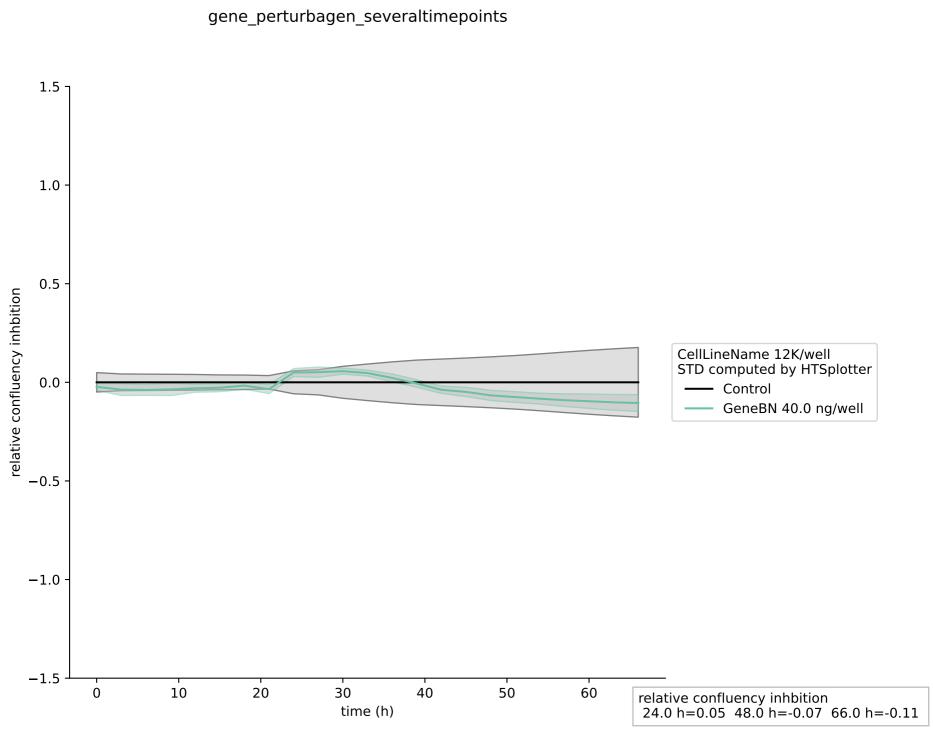


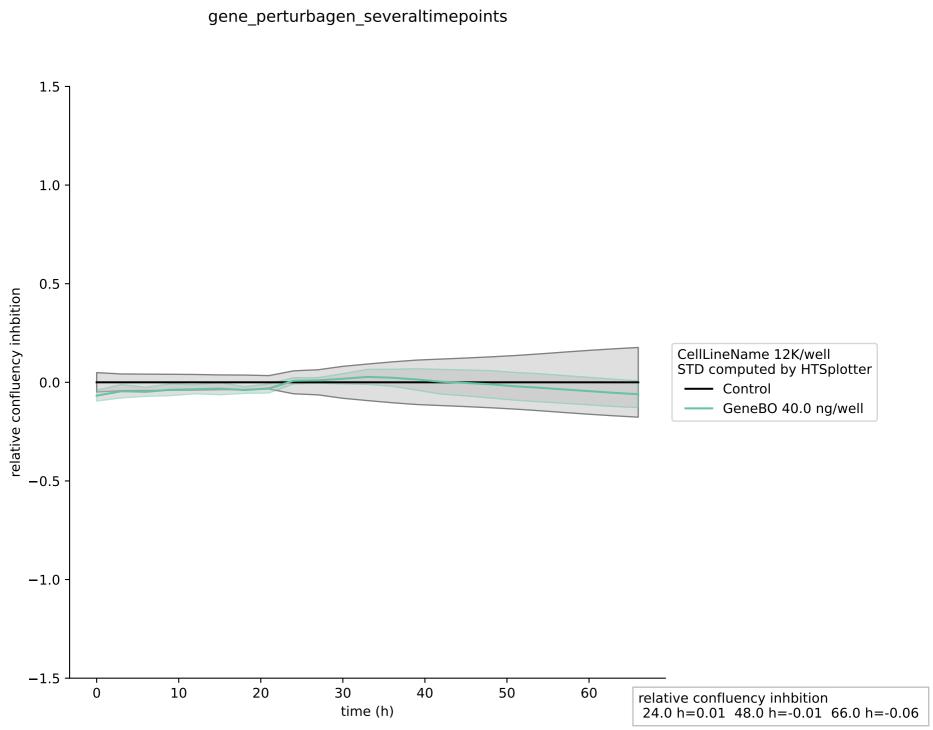
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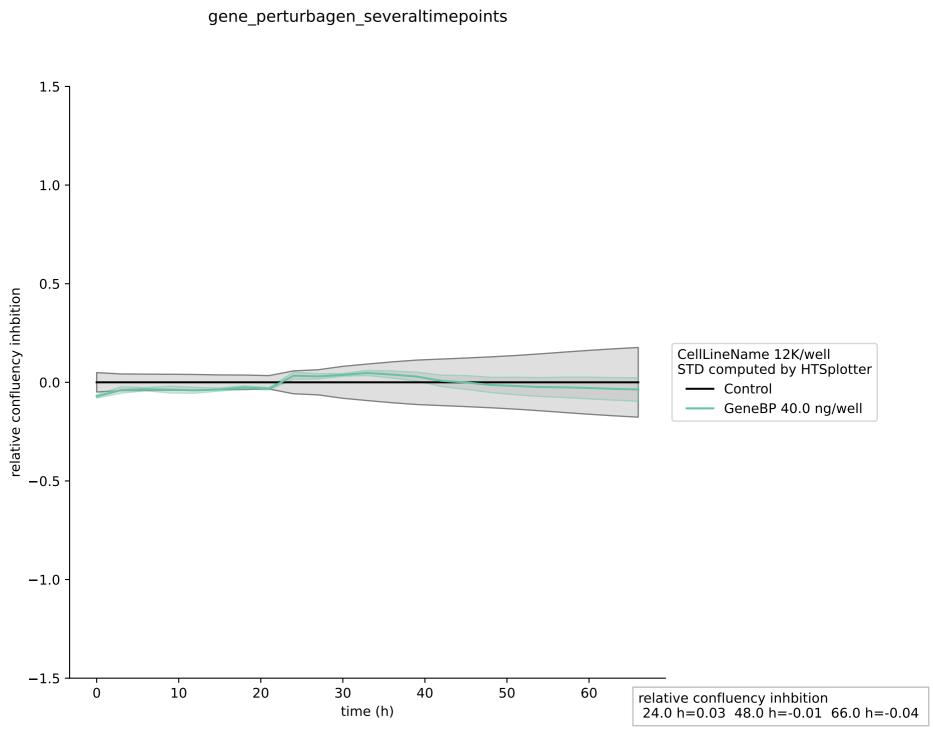


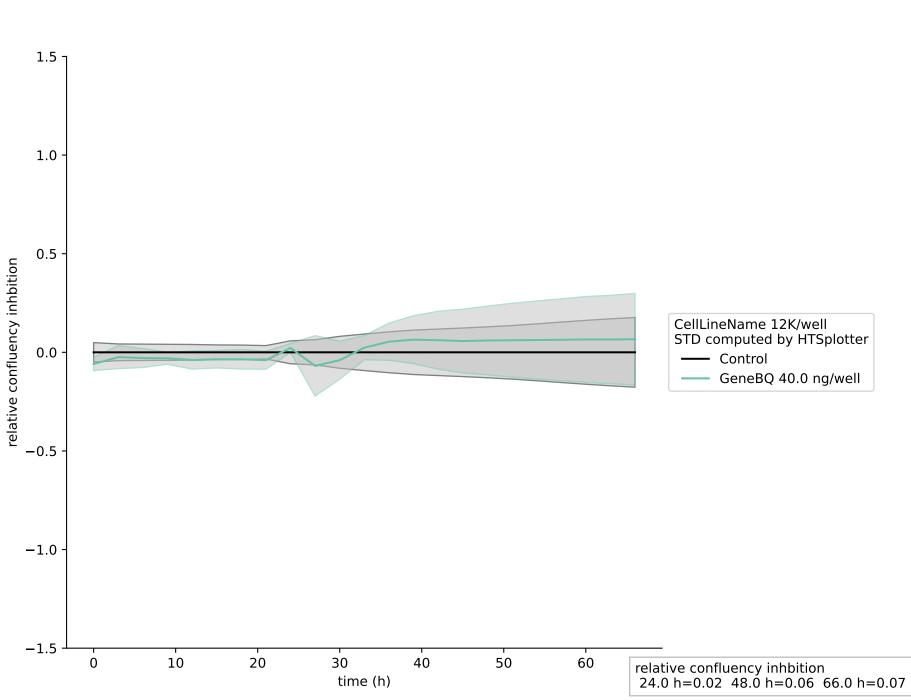
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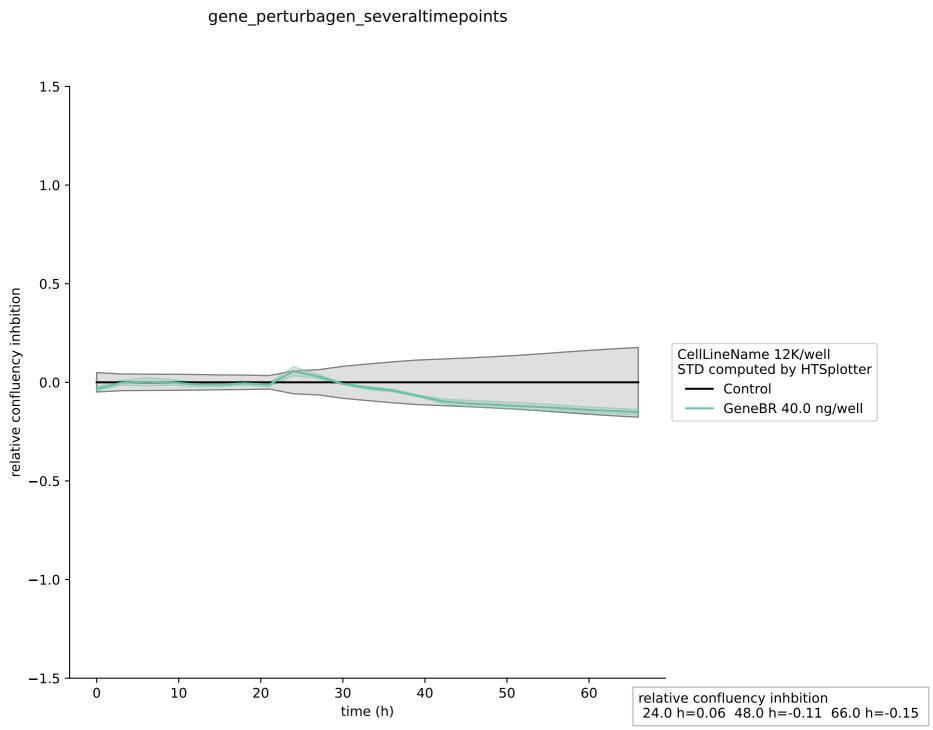




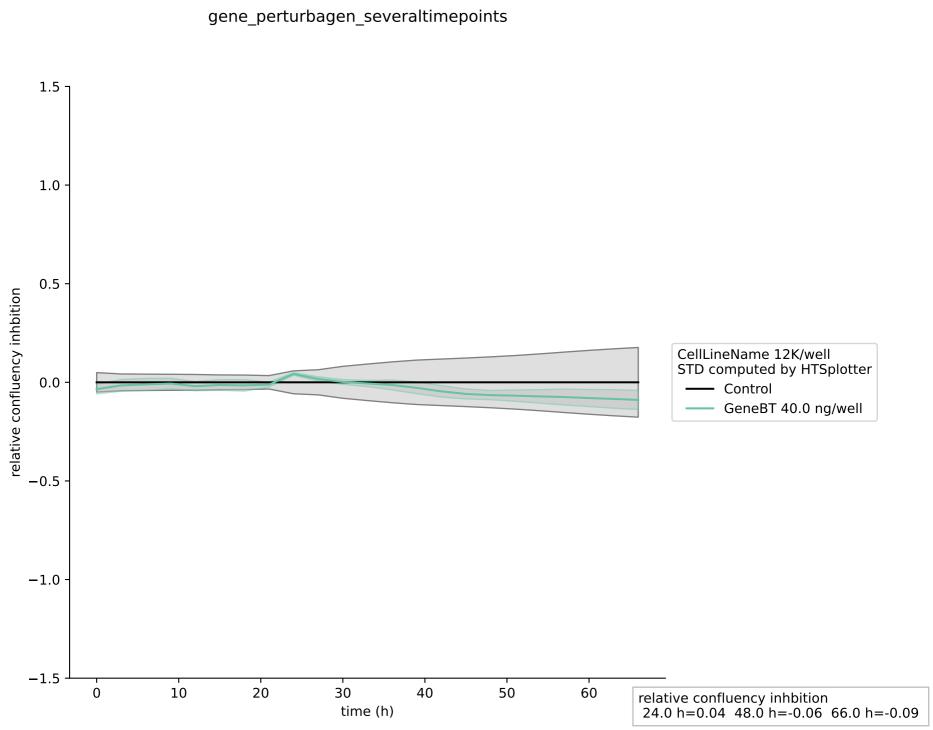


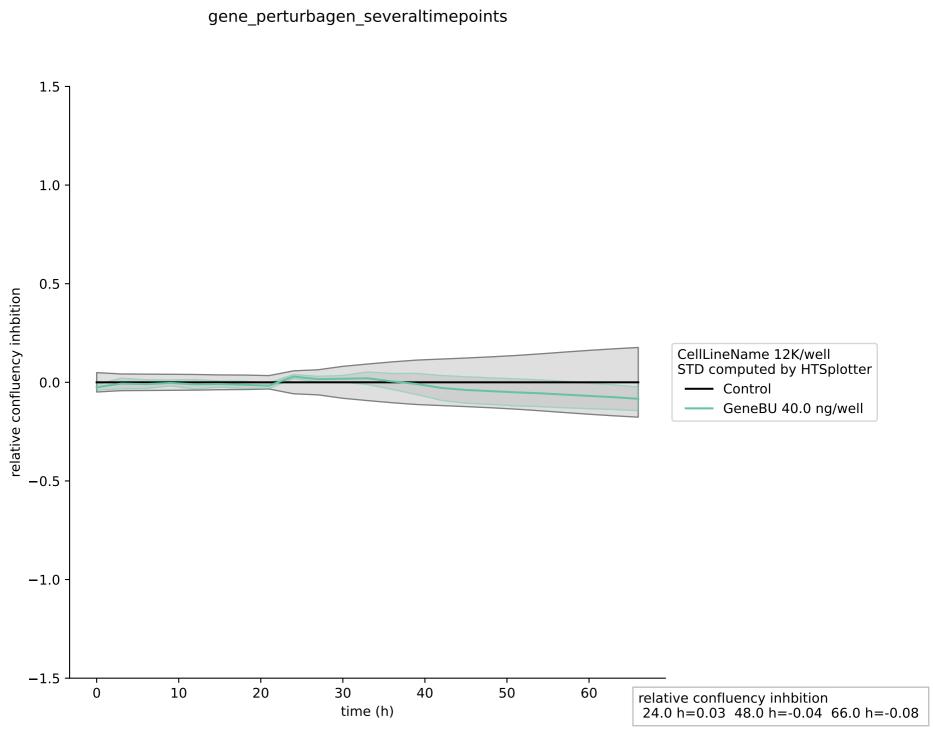


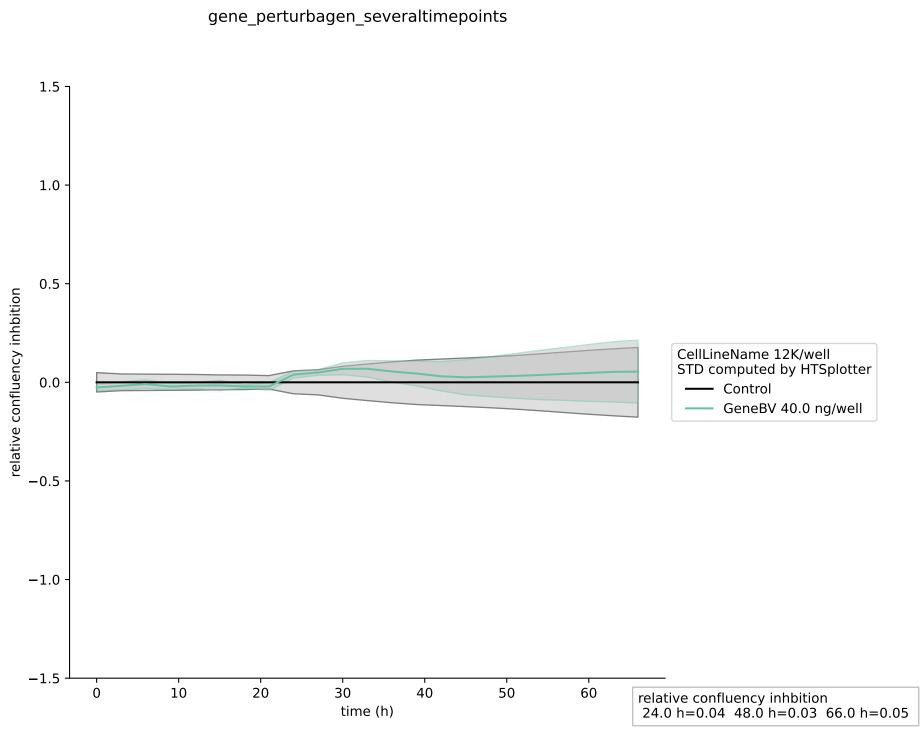


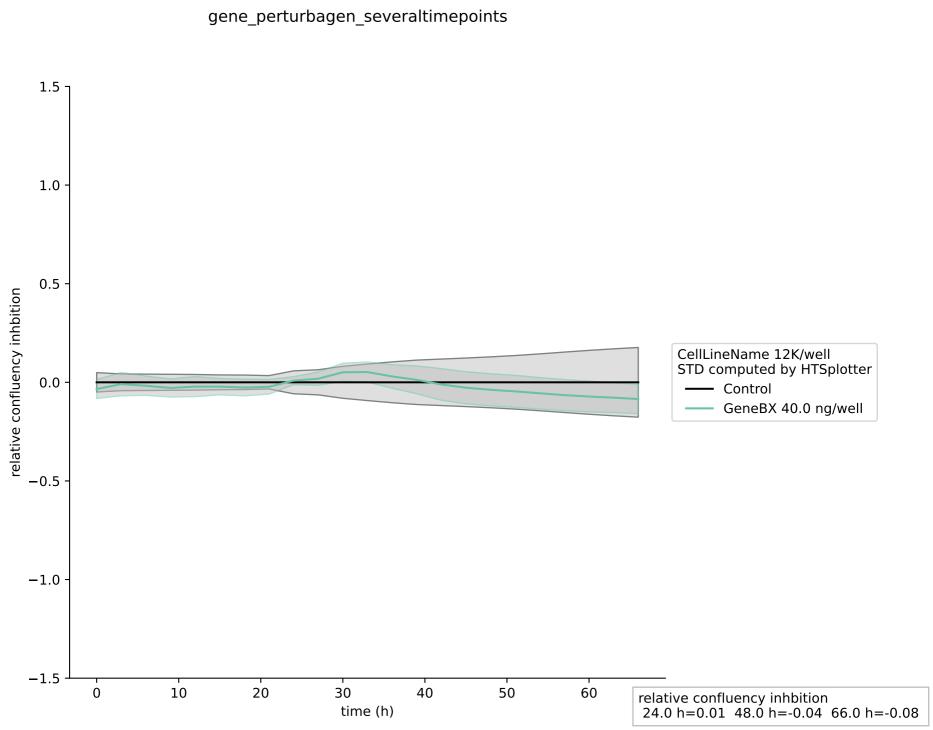


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gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneBY 40.0 ng/well 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.00 48.0 h=-0.04 66.0 h=-0.02 time (h)

1.5

1.0

0.5

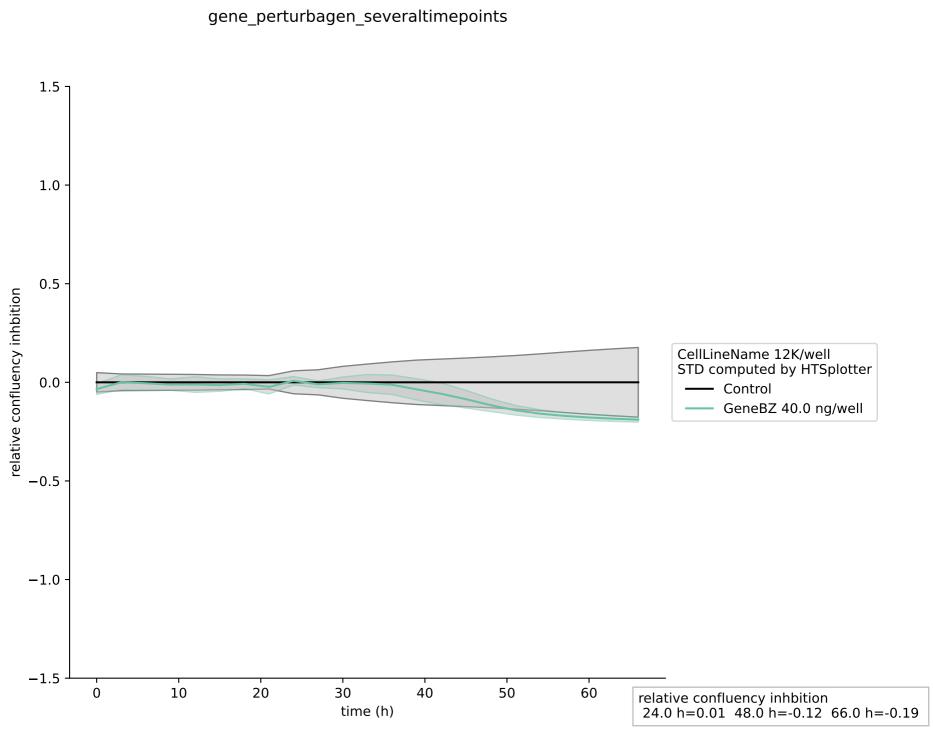
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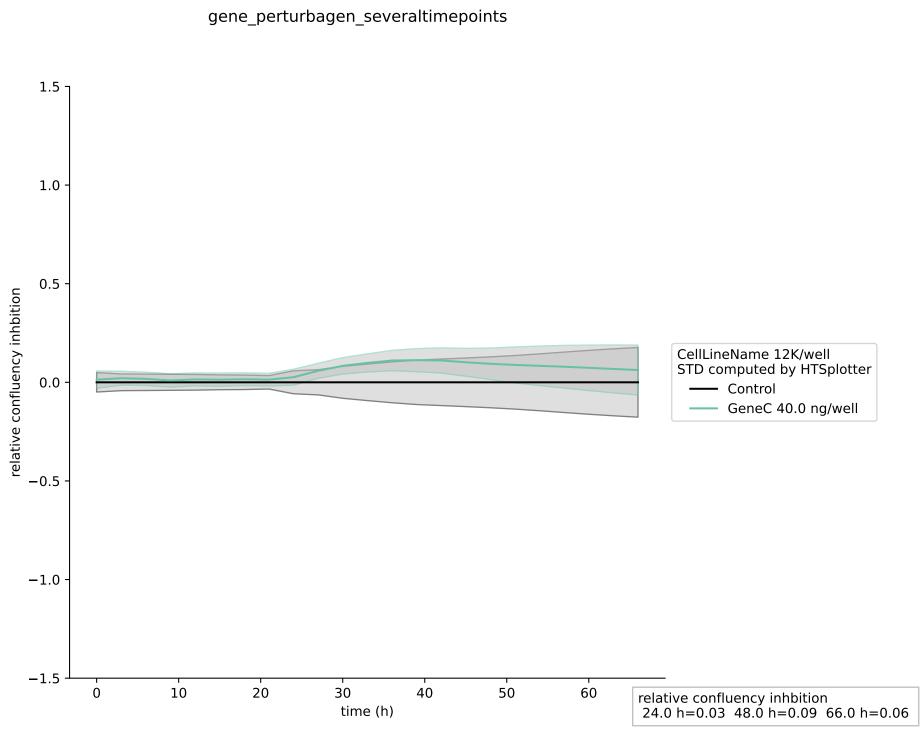
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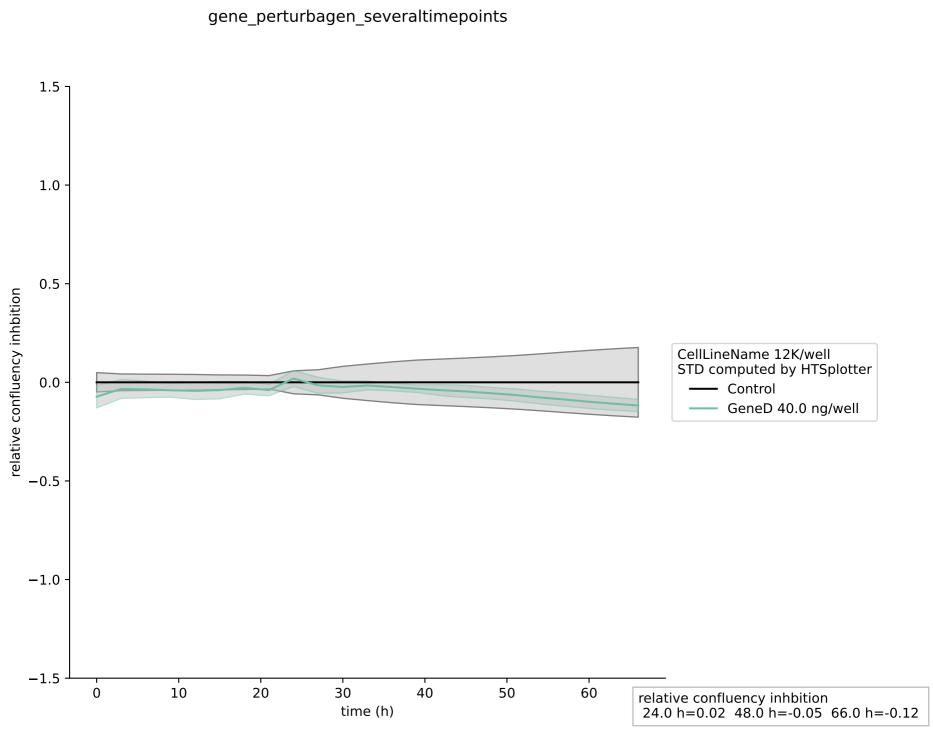
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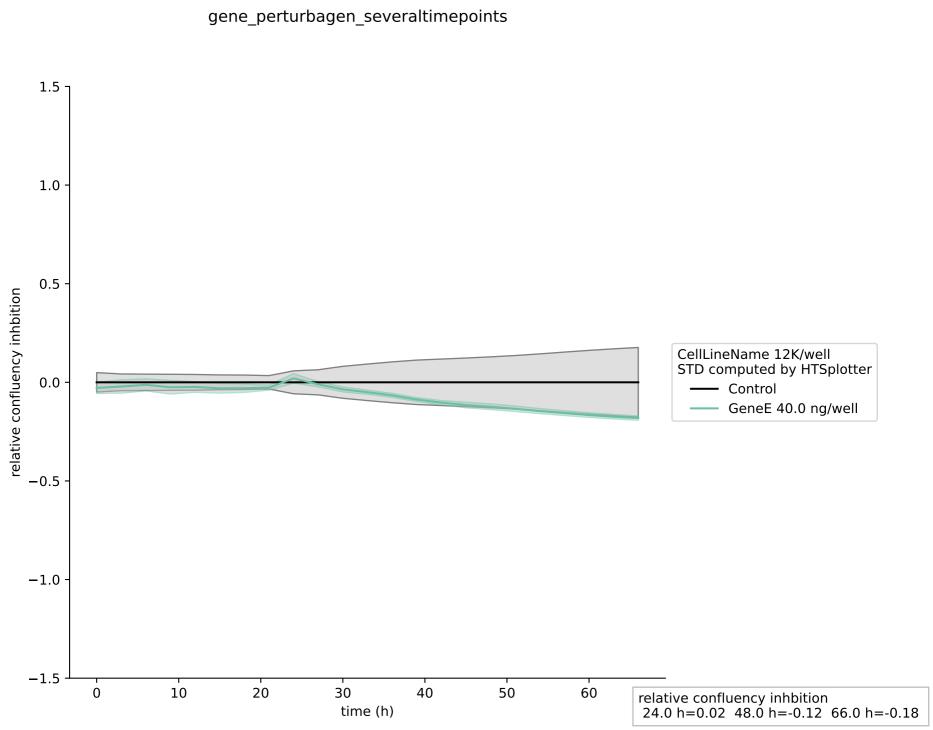
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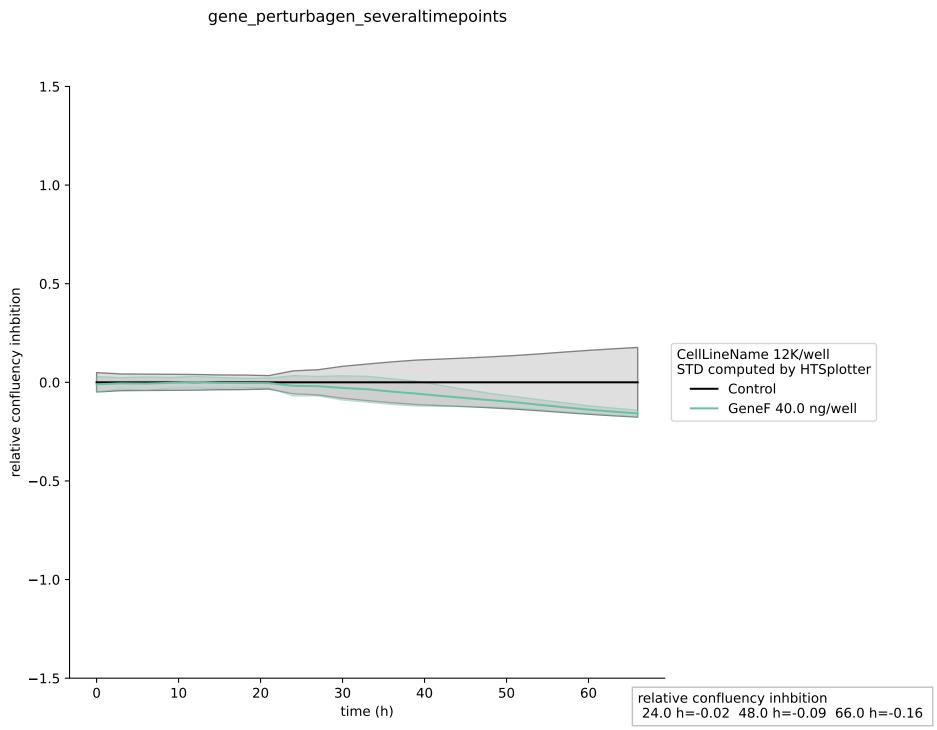




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gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneG 40.0 ng/well 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.01 48.0 h=-0.06 66.0 h=-0.15 time (h)

1.5

1.0

0.5

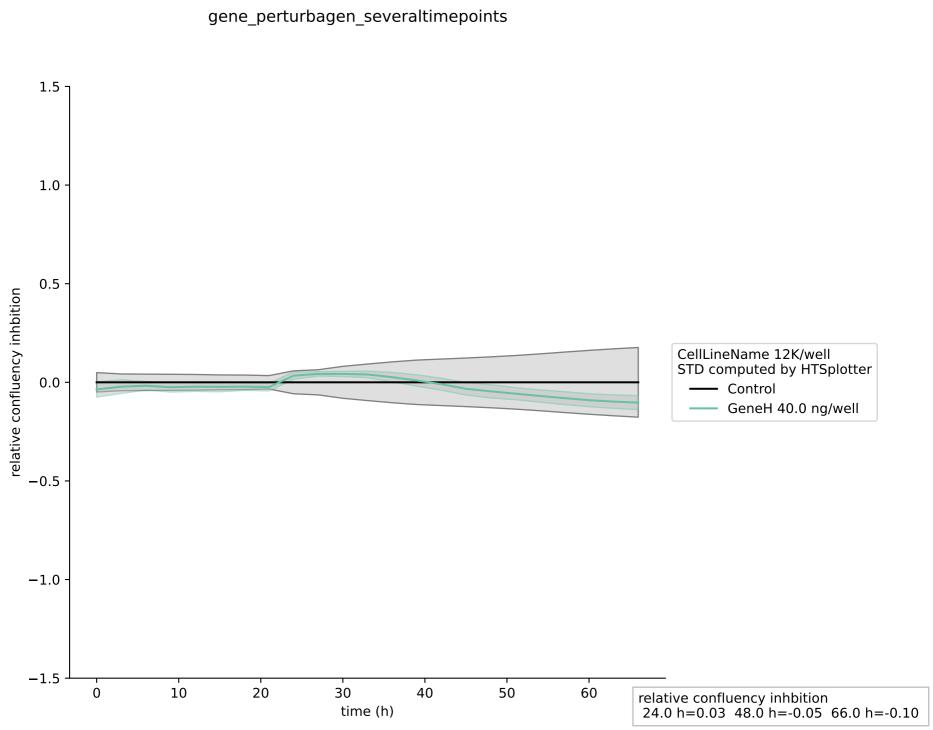
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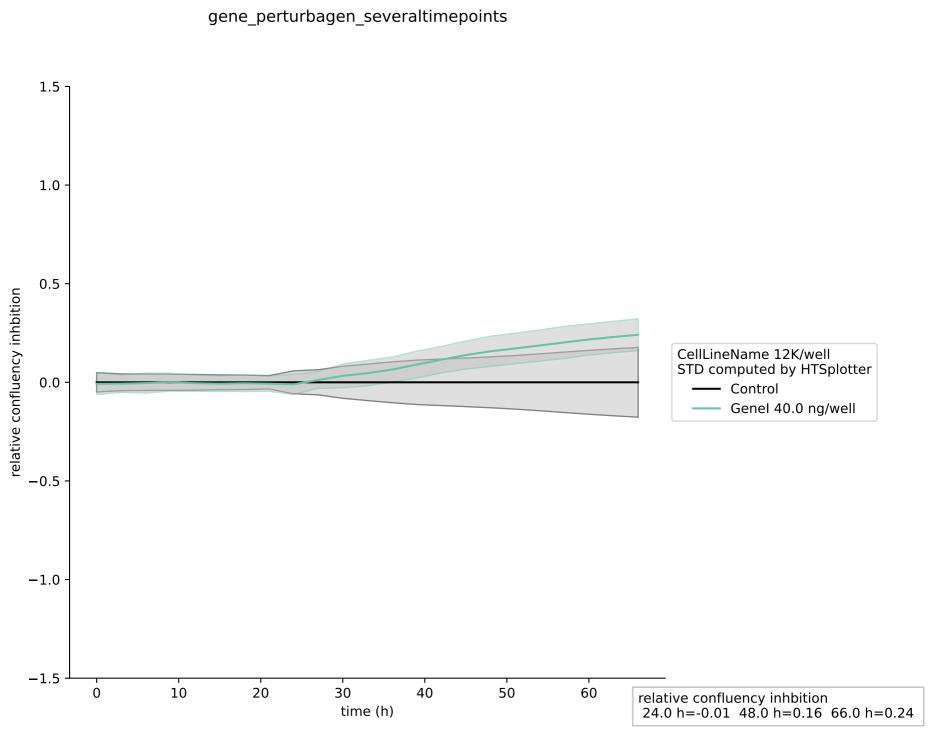
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-1.0

-1.5

0





gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneJ 40.0 ng/well 10 20 30 40 50 60 relative confluency inhbition 24.0 h=0.04 48.0 h=0.12 66.0 h=0.05

time (h)

1.5

1.0

0.5

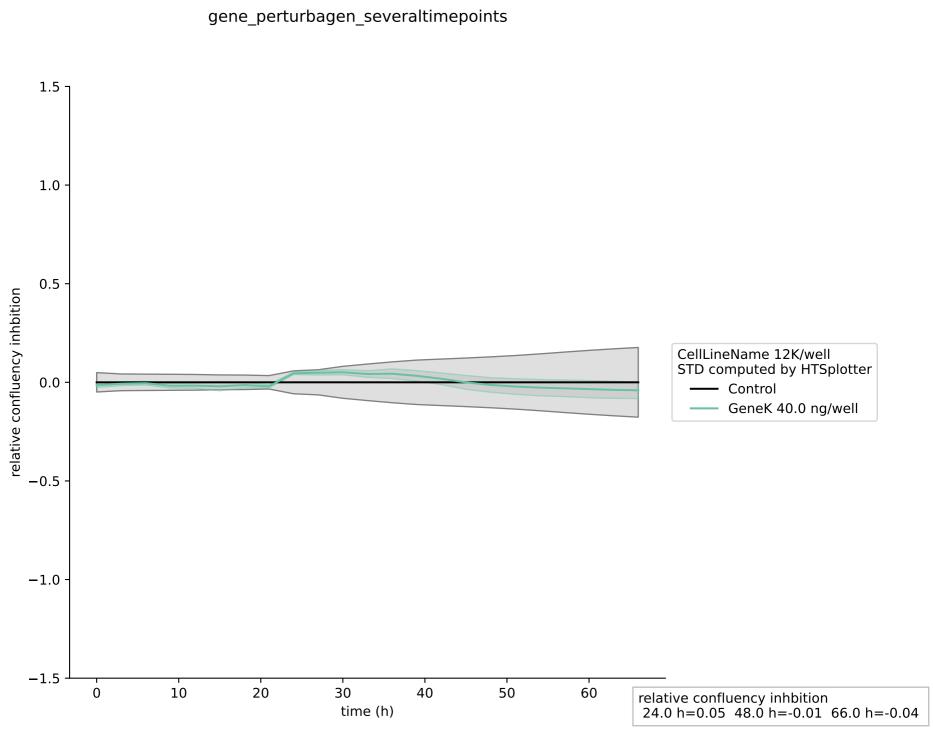
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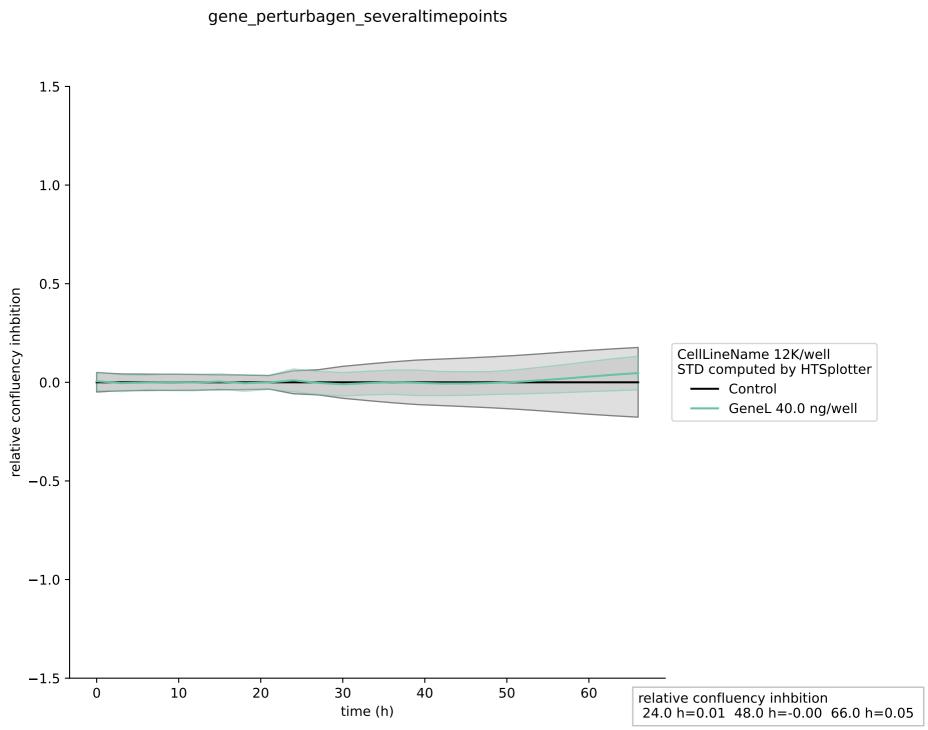
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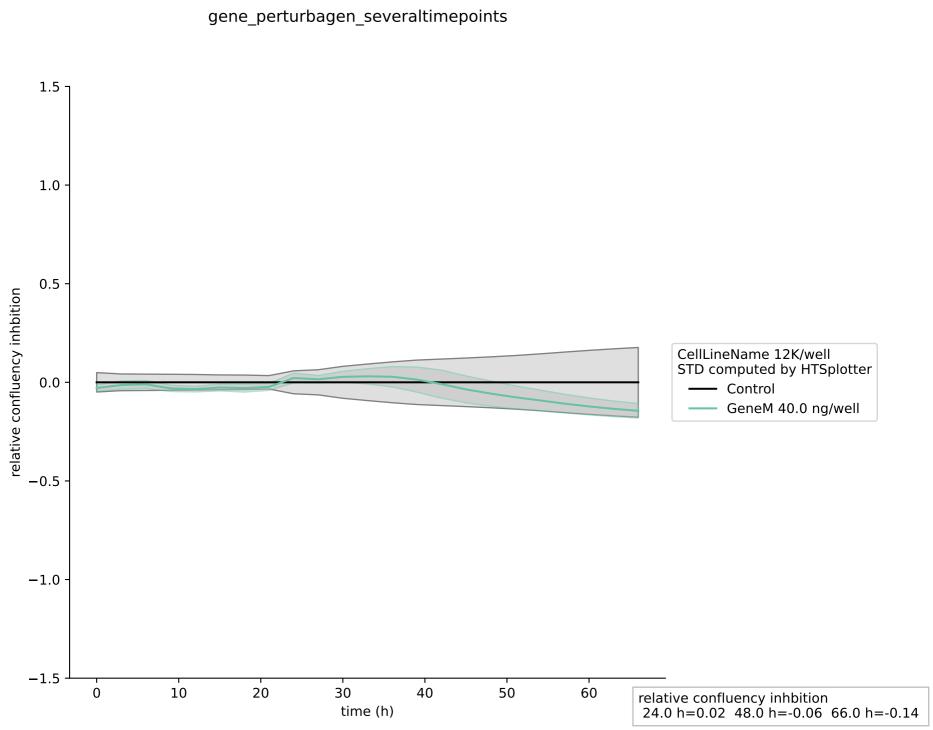
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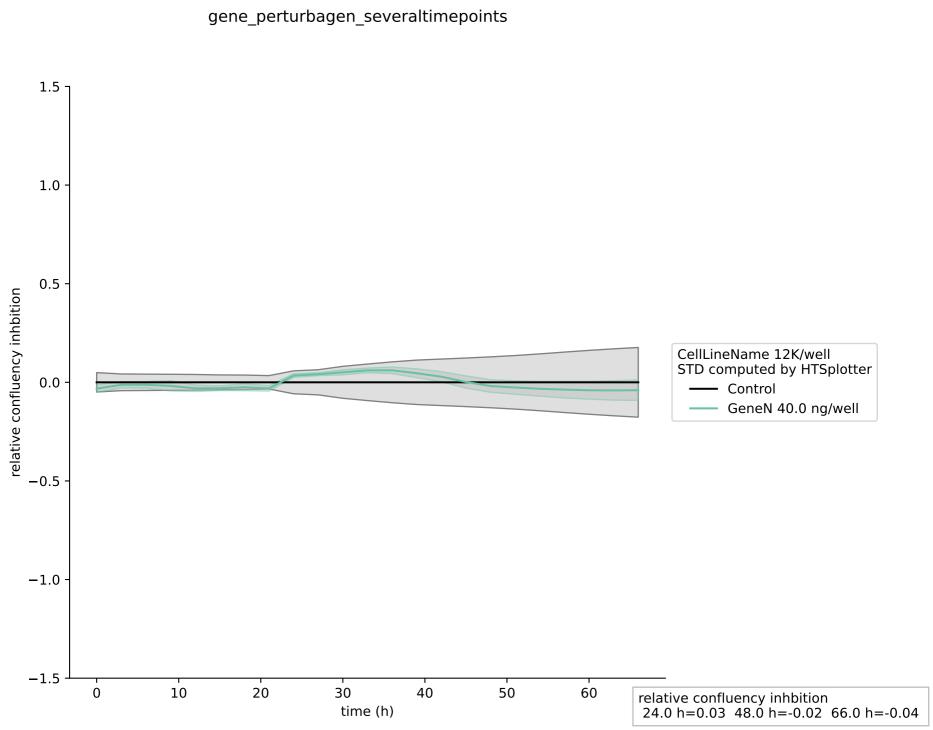
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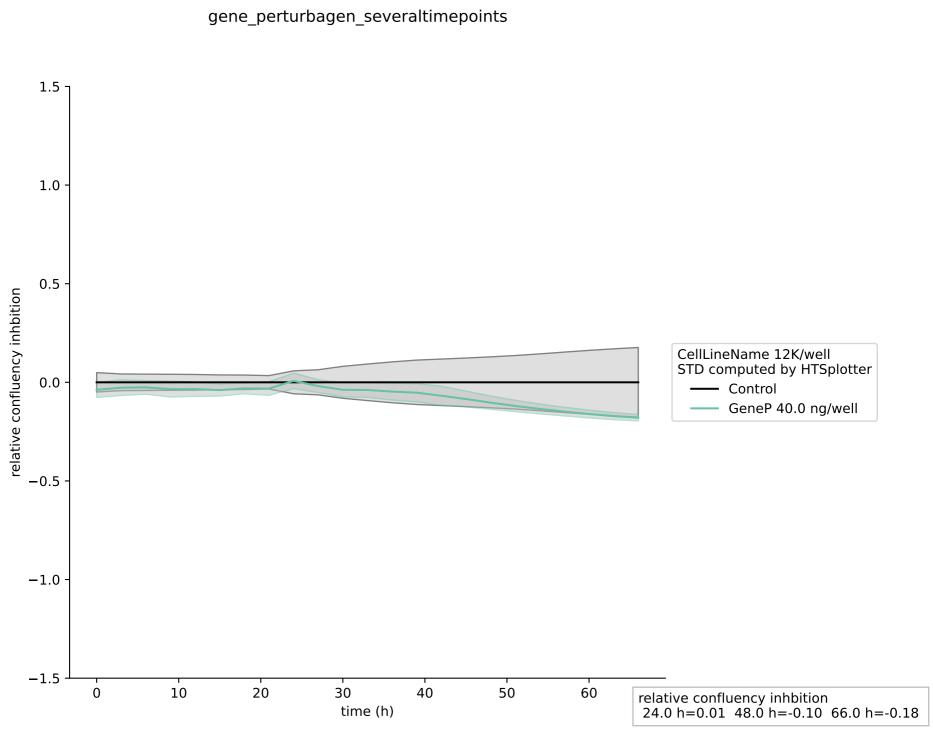








gene\_perturbagen\_severaltimepoints 1.5 1.0 0.5 relative confluency inhbition CellLineName 12K/well STD computed by HTSplotter 0.0 Control GeneO 40.0 ng/well -0.5-1.0-1.50 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.01 48.0 h=-0.16 66.0 h=-0.19 time (h)



gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneQ 40.0 ng/well

1.5

1.0

0.5

0.0

-0.5

-1.0

-1.5

0

10

20

30

time (h)

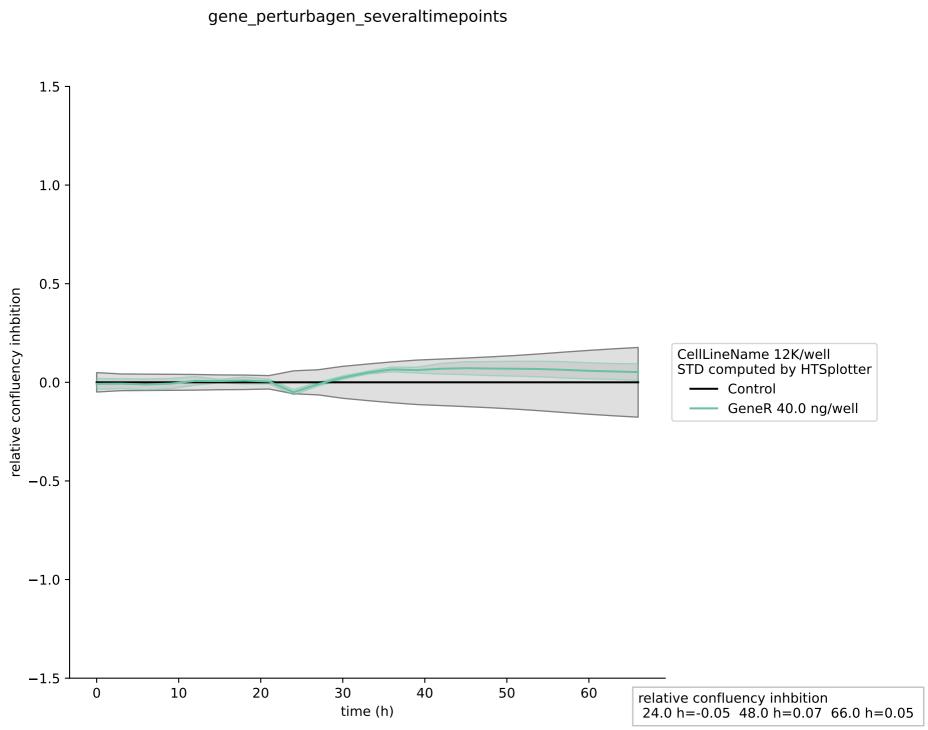
40

50

60

relative confluency inhbition 24.0 h=0.02 48.0 h=0.13 66.0 h=0.15

relative confluency inhbition



gene\_perturbagen\_severaltimepoints 1.5 1.0 0.5 relative confluency inhbition CellLineName 12K/well STD computed by HTSplotter 0.0 Control GeneS 40.0 ng/well -0.5-1.0-1.50 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.04 48.0 h=-0.10 66.0 h=-0.13 time (h)

gene\_perturbagen\_severaltimepoints 1.5 1.0 0.5 relative confluency inhbition CellLineName 12K/well STD computed by HTSplotter 0.0 Control GeneT 40.0 ng/well -0.5-1.0-1.50 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.05 48.0 h=-0.04 66.0 h=-0.08 time (h)

gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneU 40.0 ng/well 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.05 48.0 h=-0.08 66.0 h=-0.11

time (h)

1.5

1.0

0.5

0.0

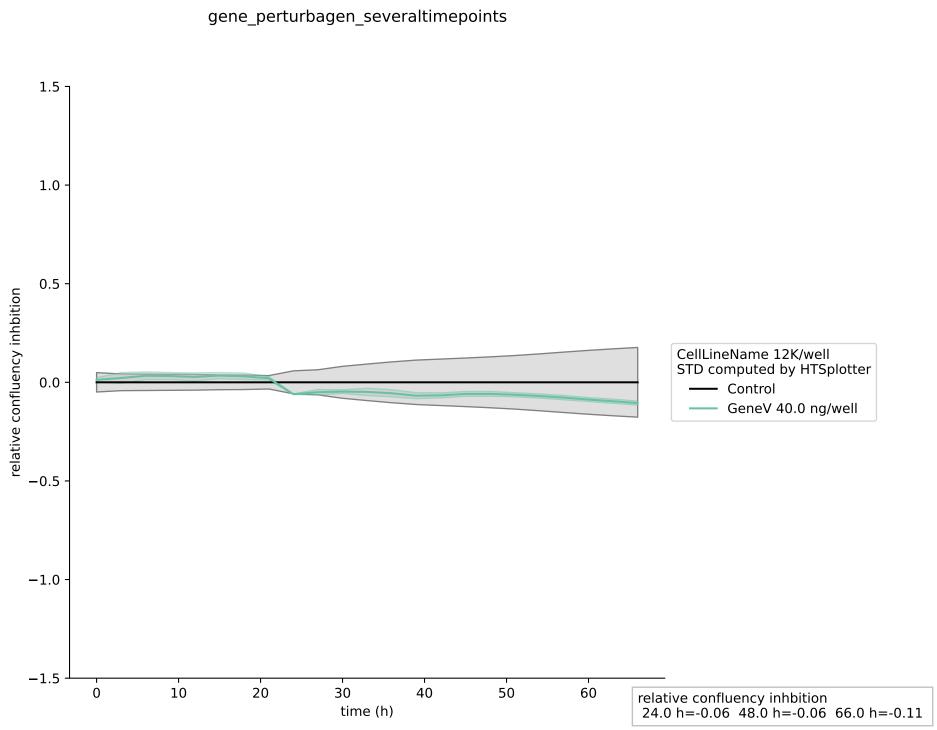
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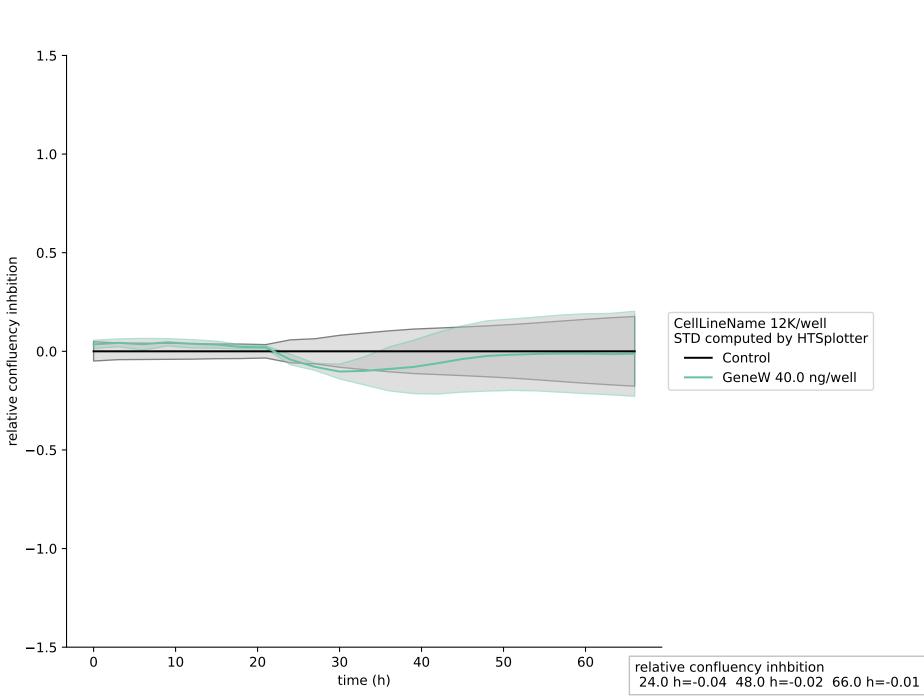
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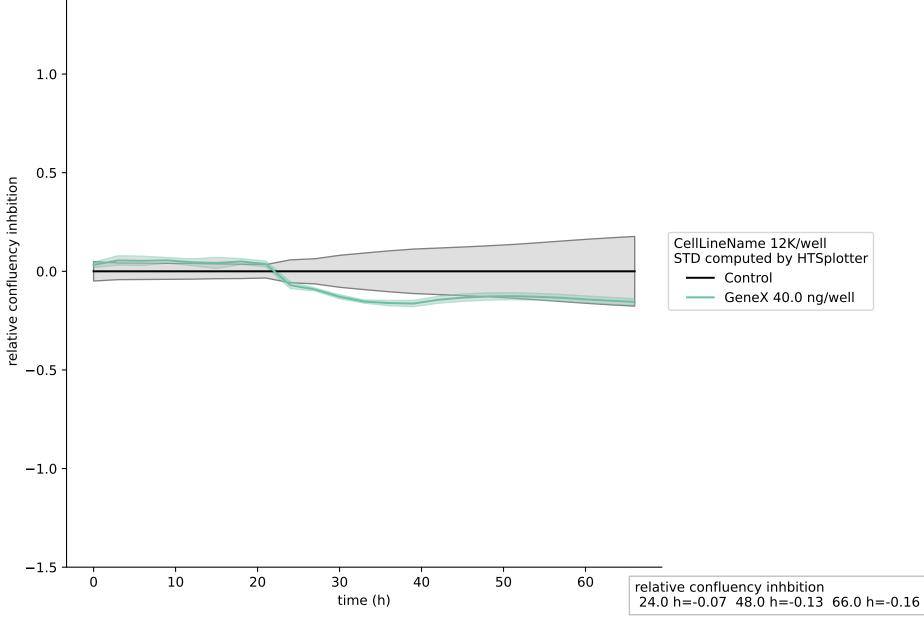
relative confluency inhbition



gene\_perturbagen\_severaltimepoints



gene\_perturbagen\_severaltimepoints CellLineName 12K/well STD computed by HTSplotter Control GeneX 40.0 ng/well



1.5

gene\_perturbagen\_severaltimepoints 1.5 1.0 0.5 CellLineName 12K/well STD computed by HTSplotter 0.0 Control GeneY 40.0 ng/well -0.5-1.0-1.50 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.02 48.0 h=-0.08 66.0 h=-0.12 time (h)

relative confluency inhbition

gene\_perturbagen\_severaltimepoints 1.5 1.0 0.5 relative confluency inhbition CellLineName 12K/well STD computed by HTSplotter 0.0 Control GeneZ 40.0 ng/well -0.5-1.0-1.50 10 20 30 40 50 60 relative confluency inhbition 24.0 h=-0.06 48.0 h=-0.01 66.0 h=-0.04 time (h)

