Integrate biological replicates from different assay plates

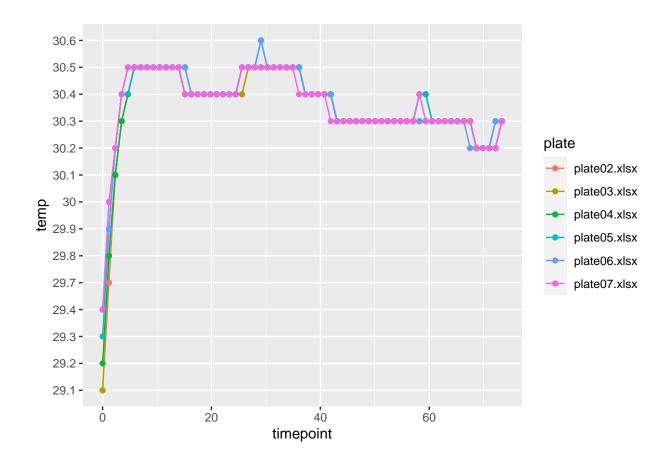
JN

Sat Feb 17 13:43:12 2024

[2] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/input/plate02.xlsx"

[1] "all raw data xlsx files in the directory, absolute paths: "

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## [3] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/input/plate03.xlsx"
## [4] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/input/plate04.xlsx"
## [5] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/input/plate05.xlsx"
## [6] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/input/plate06.xlsx"
## [7] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/input/plate07.xlsx"
## [1] "all raw data xlsx files in the directory: "
## [2] "plate02.xlsx"
## [3] "plate03.xlsx"
## [4] "plate04.xlsx"
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## [6] "plate06.xlsx"
## [7] "plate07.xlsx"
## [1] "all corresponding RData files in the directory: "
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## [3] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/output/plate_data_plate03.xlsx.RDat
## [4] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/output/plate_data_plate04.xlsx.RDat
## [5] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/output/plate_data_plate05.xlsx.RDat
## [6] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/output/plate_data_plate06.xlsx.RDat
## [7] "C:/Users/jahn_/Desktop/analyses/broth_demonstration_dataset/output/plate_data_plate07.xlsx.RDat
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## [1] 768
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[1] 768

[1] 768

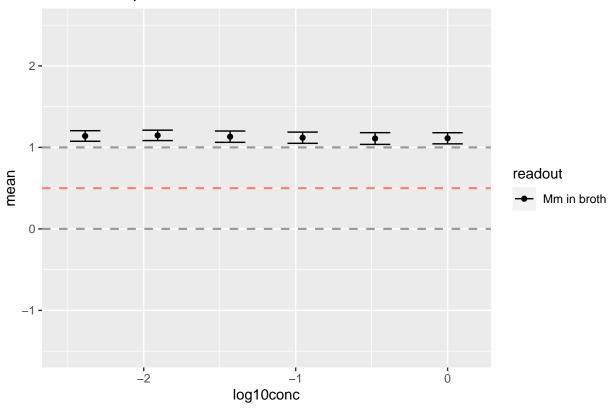
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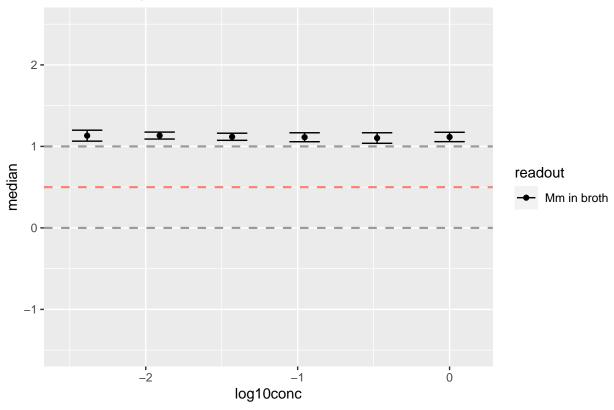
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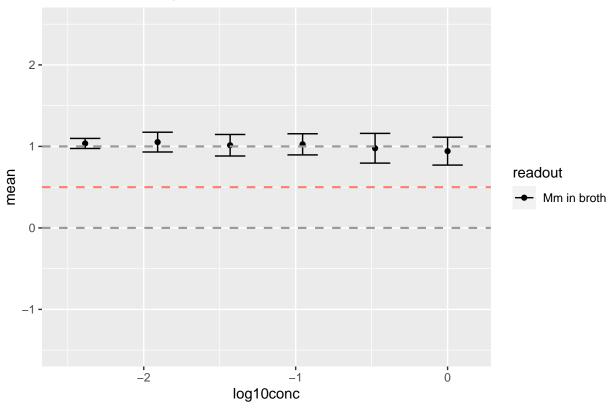
-: dose response curve, mean & sd



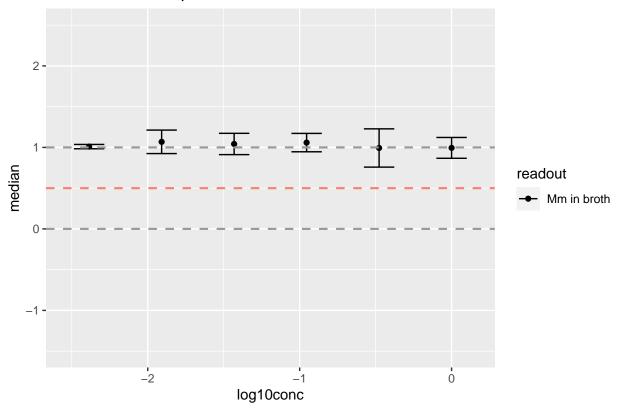
-: dose response curve, median & mad



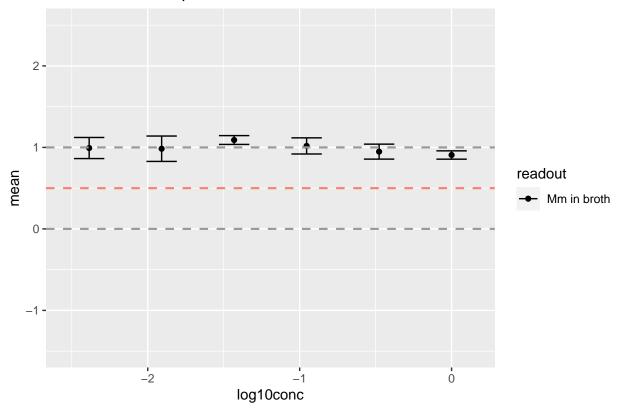
 Ak_07 : dose response curve, mean & sd

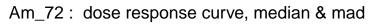


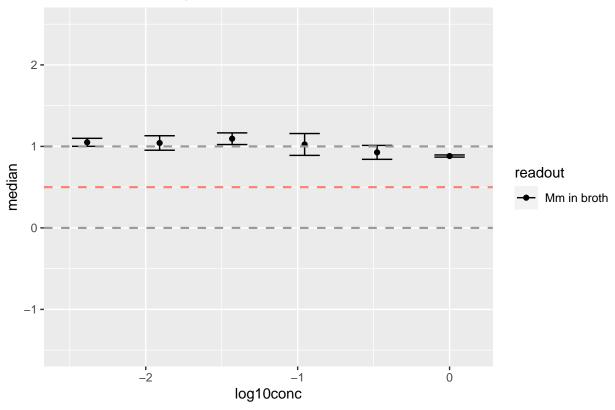
Ak_07: dose response curve, median & mad



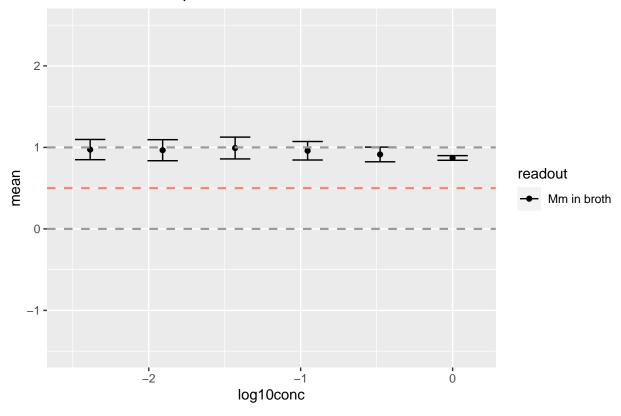
Am_72 : dose response curve, mean & sd

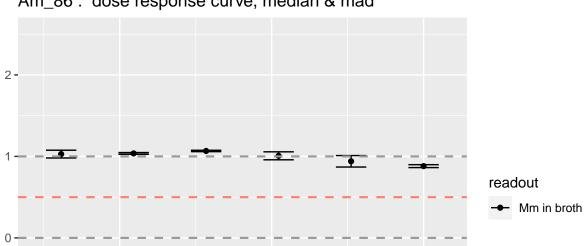






Am_86 : dose response curve, mean & sd





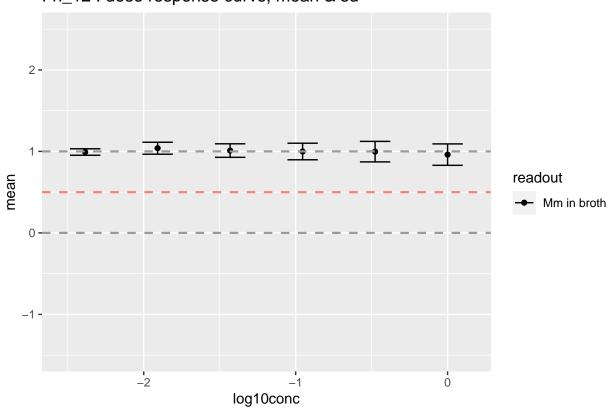
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Am_86: dose response curve, median & mad

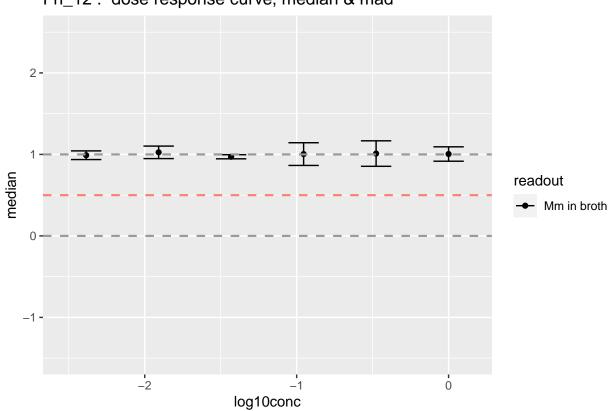
median

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-2

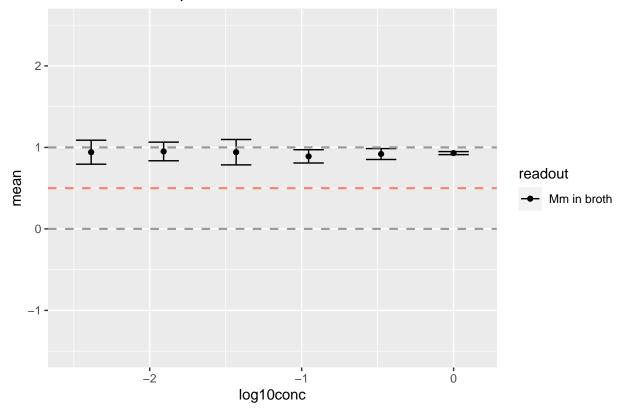


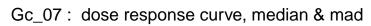
Fh_12 : dose response curve, mean & sd

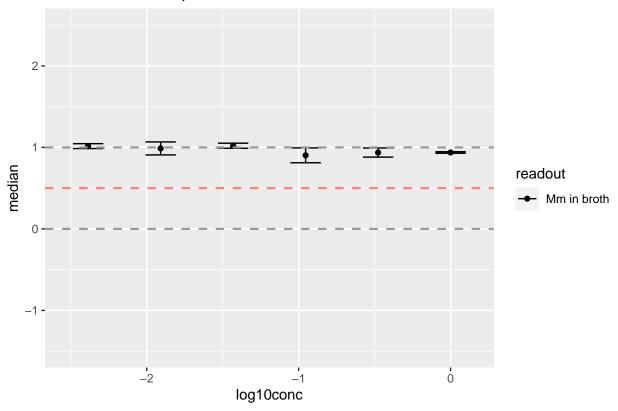


Fh_12: dose response curve, median & mad

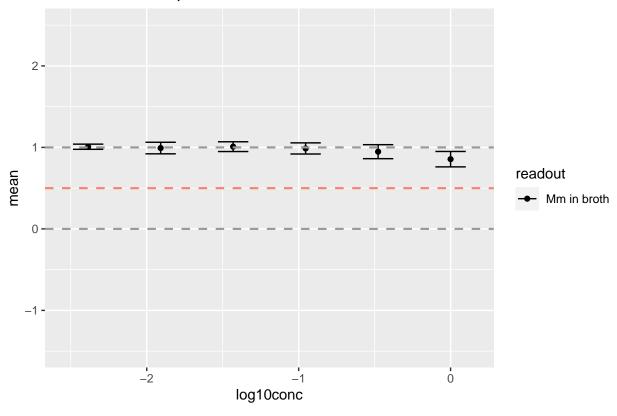
Gc_07 : dose response curve, mean & sd

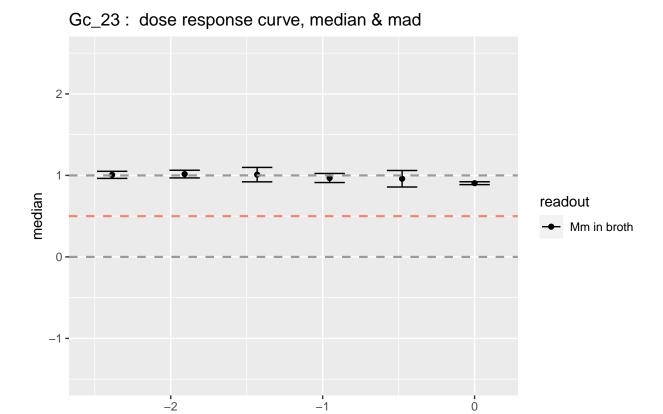






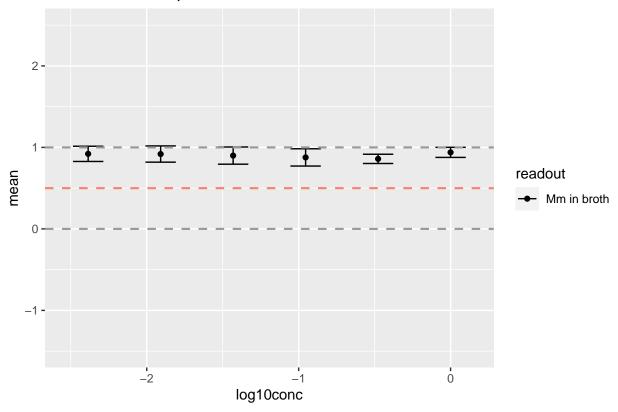
Gc_23 : dose response curve, mean & sd

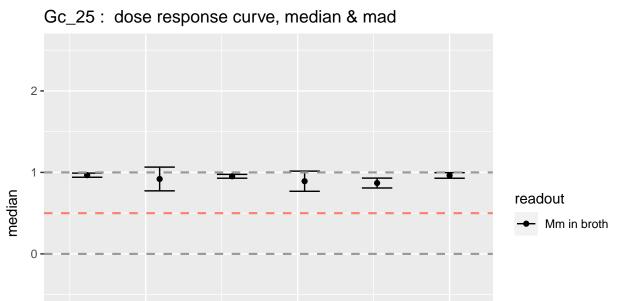




16

Gc_25 : dose response curve, mean & sd

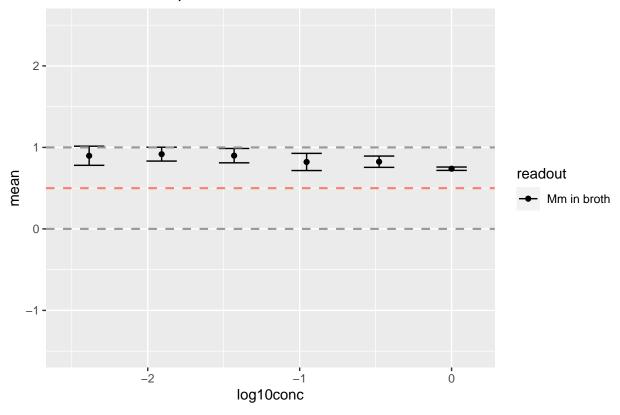


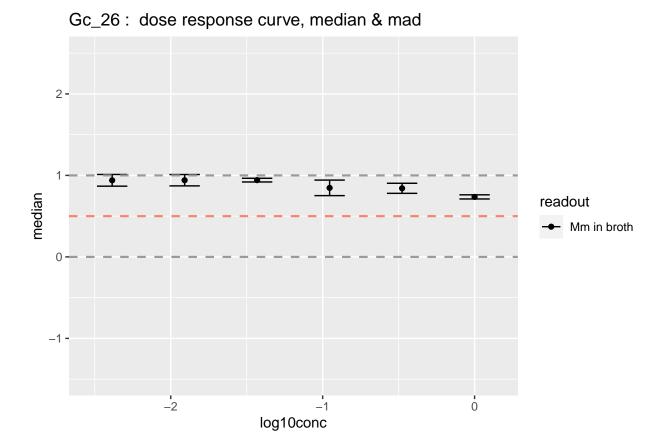


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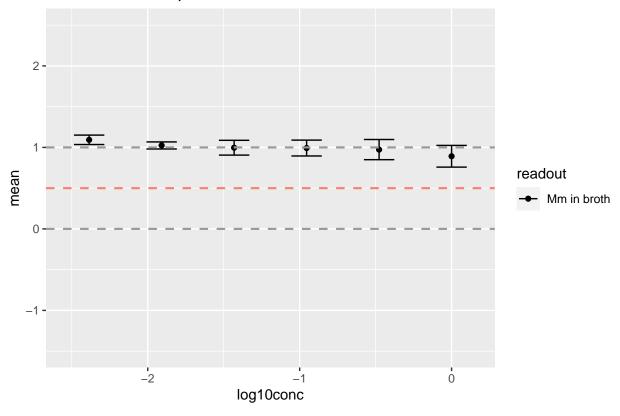
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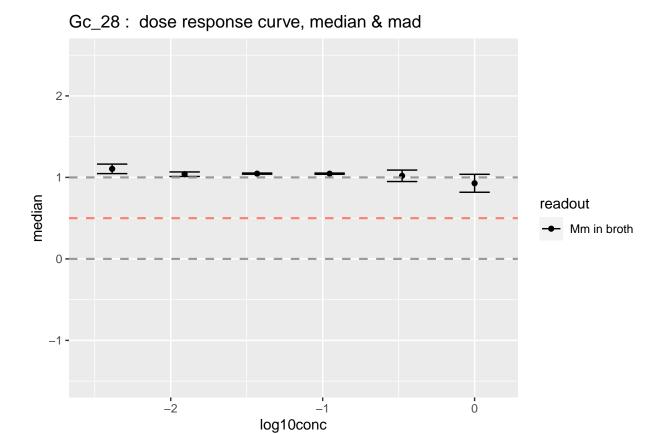




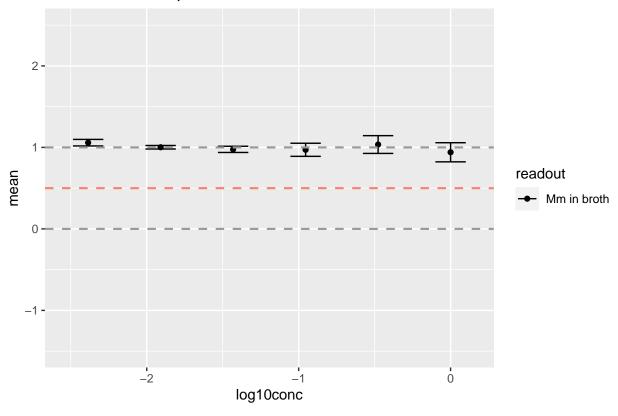


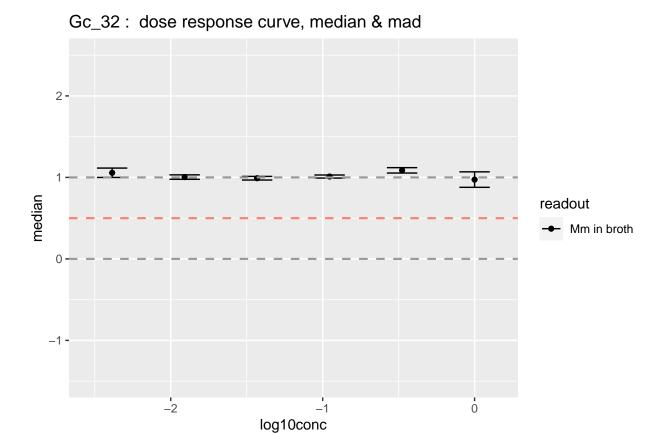
Gc_28 : dose response curve, mean & sd



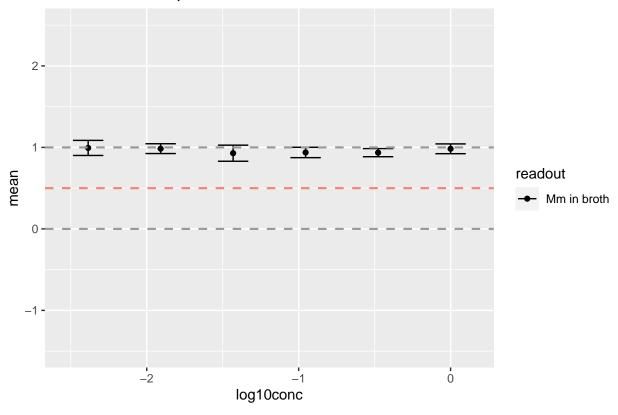


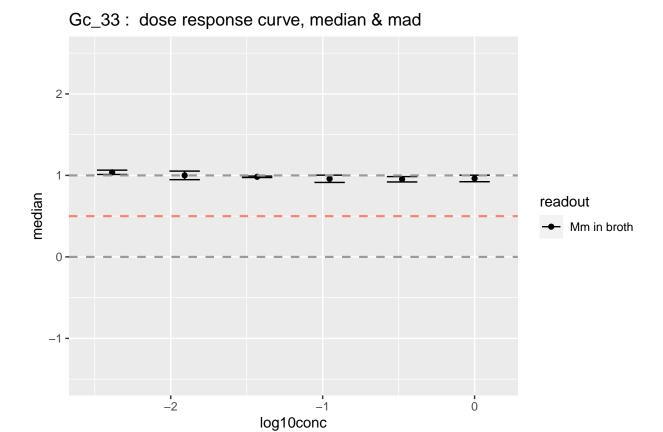
Gc_32 : dose response curve, mean & sd



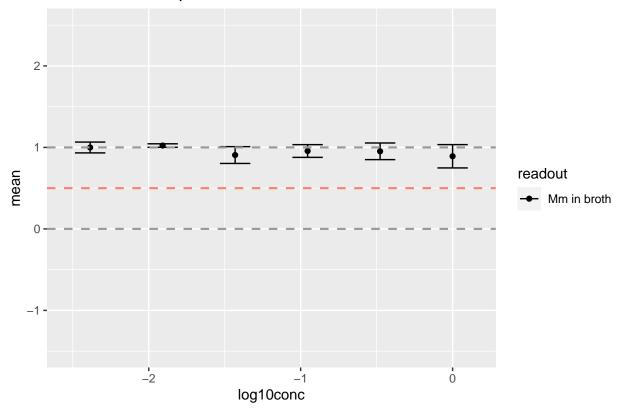


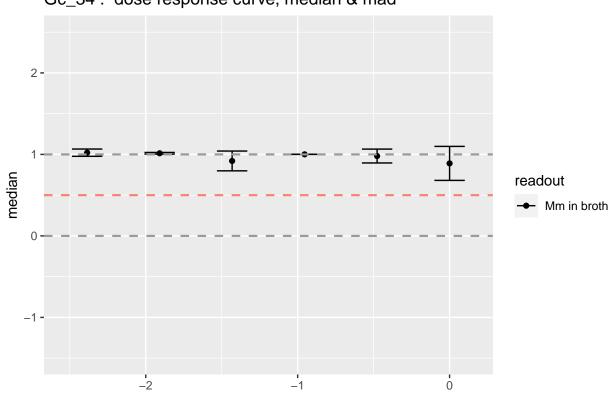






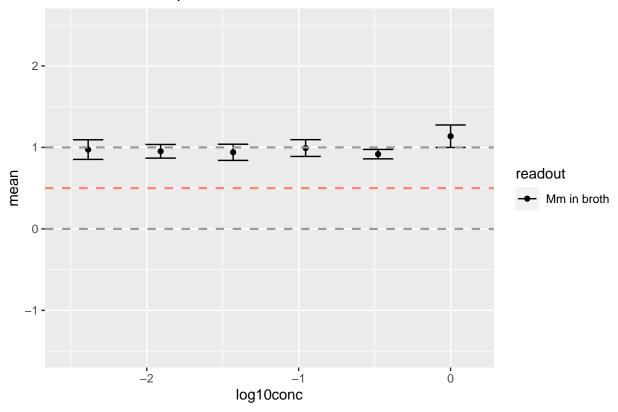
Gc_34 : dose response curve, mean & sd

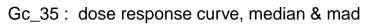


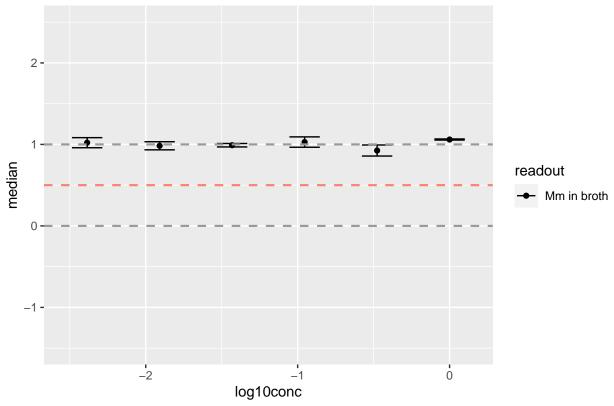


Gc_34: dose response curve, median & mad

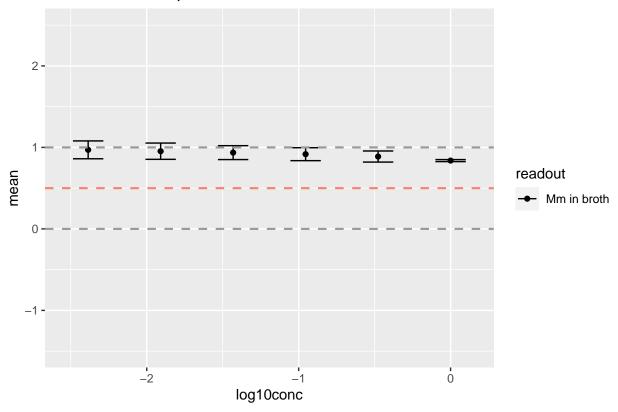
Gc_35 : dose response curve, mean & sd

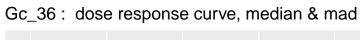


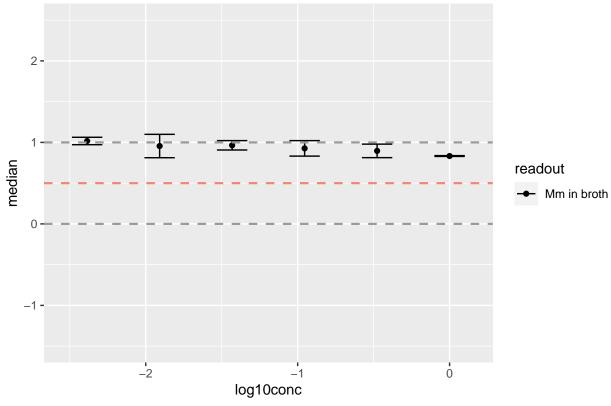




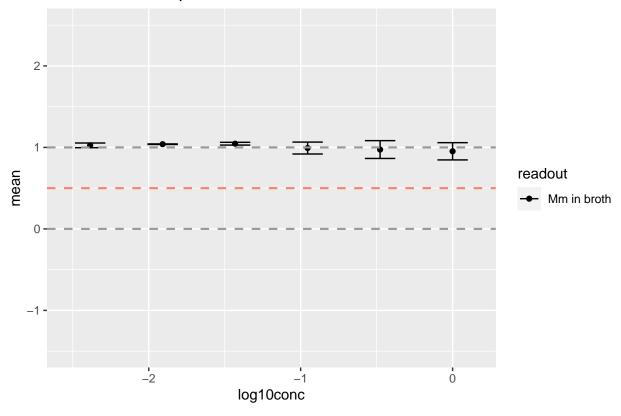
Gc_36 : dose response curve, mean & sd

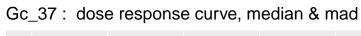


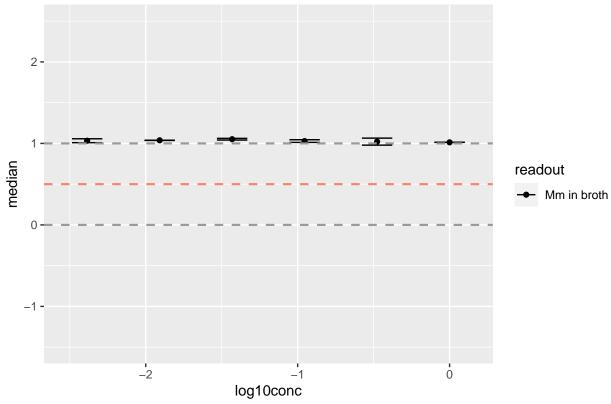




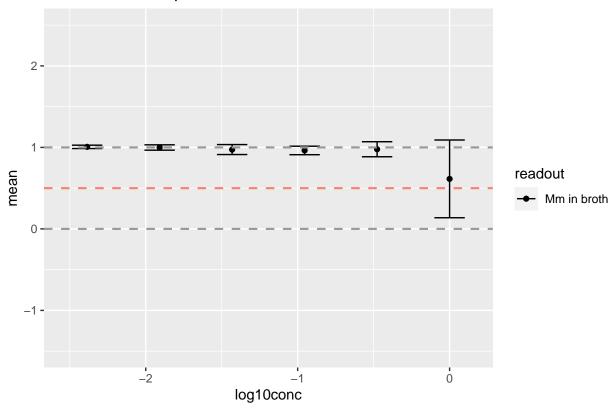
Gc_37 : dose response curve, mean & sd

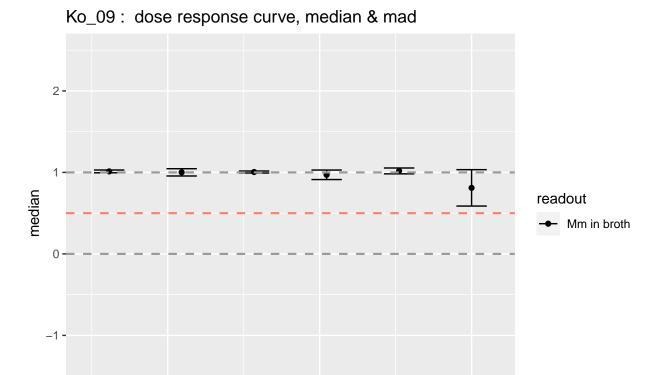






Ko_09 : dose response curve, mean & sd

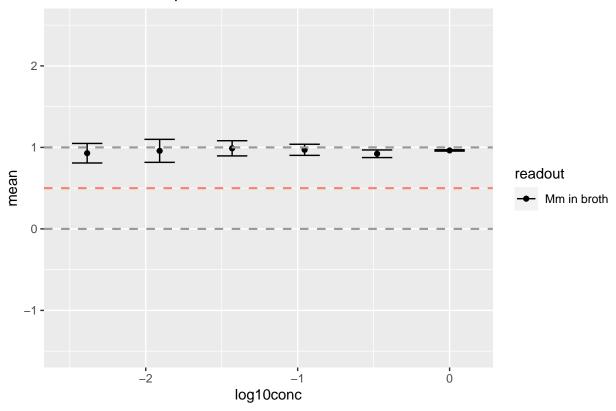


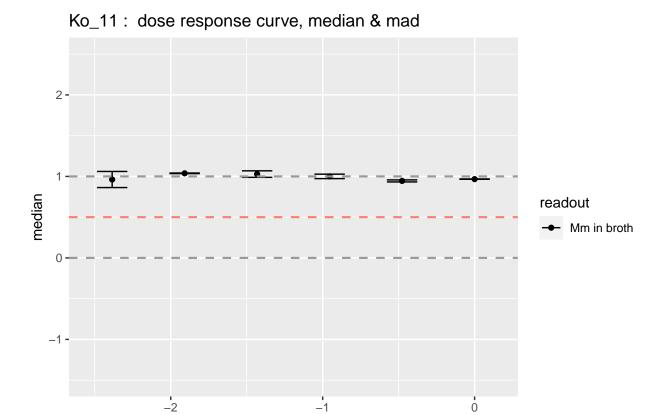


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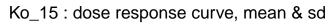
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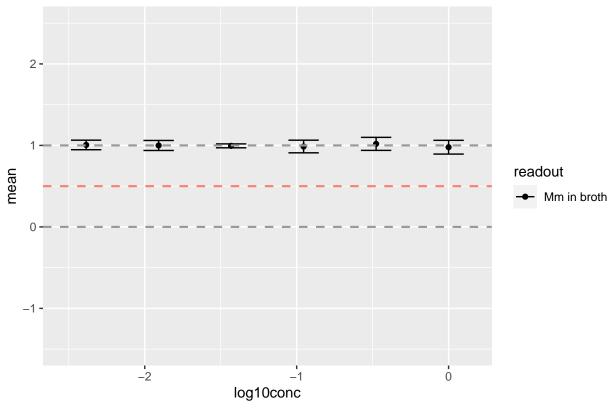


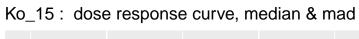


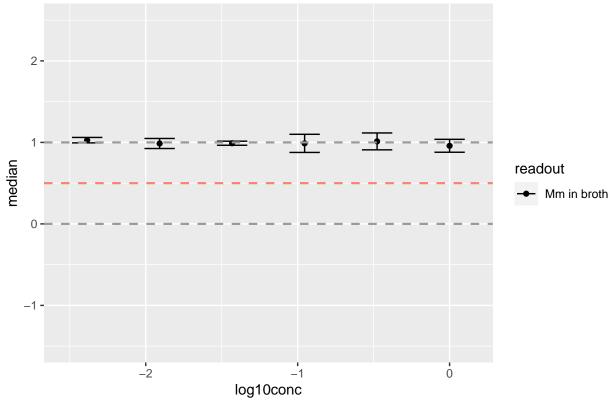


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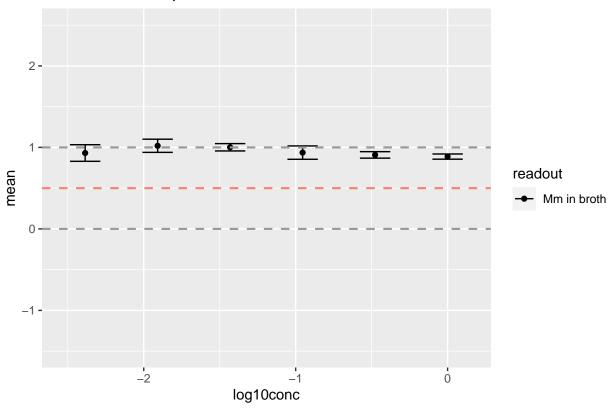


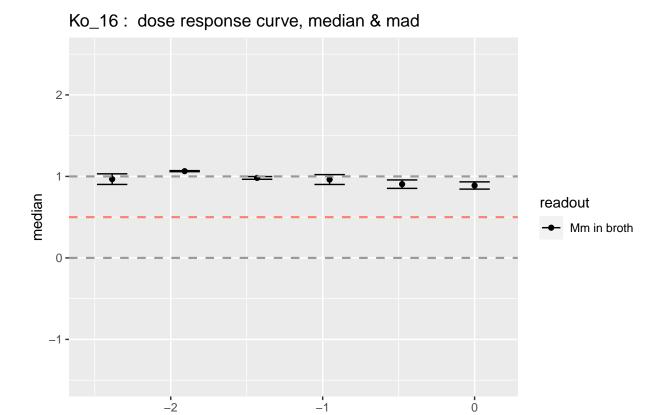






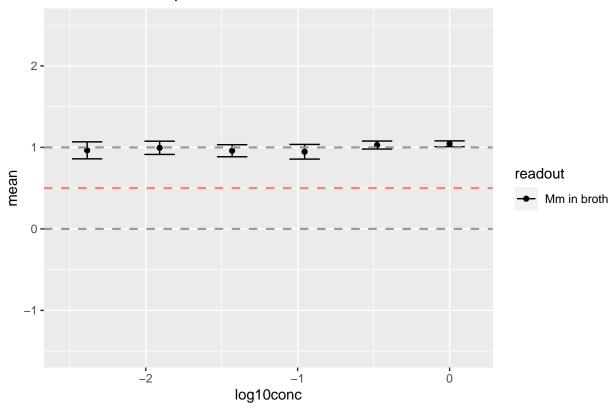
Ko_16 : dose response curve, mean & sd

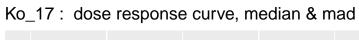


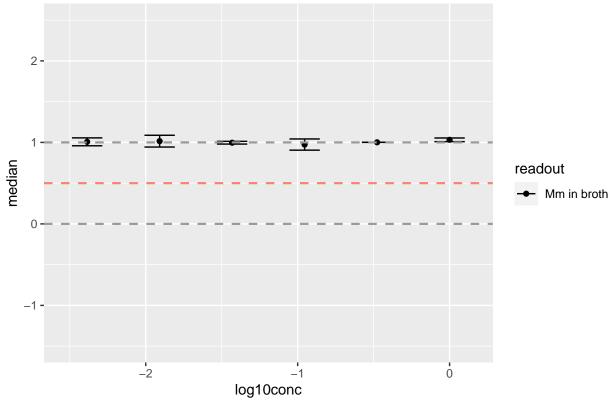


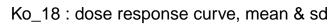
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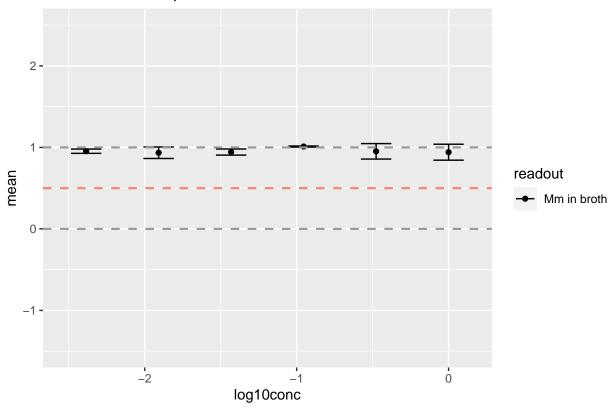


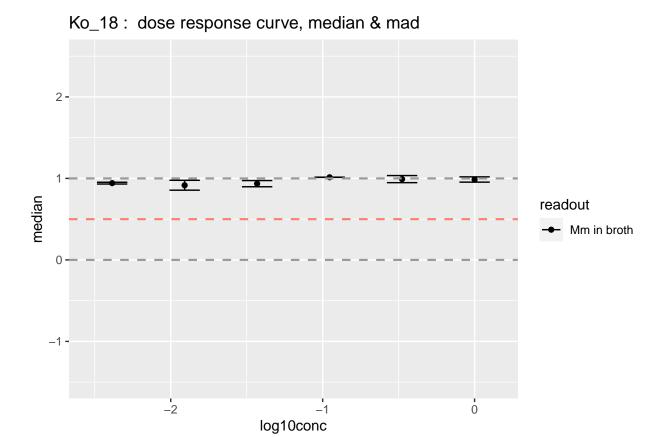




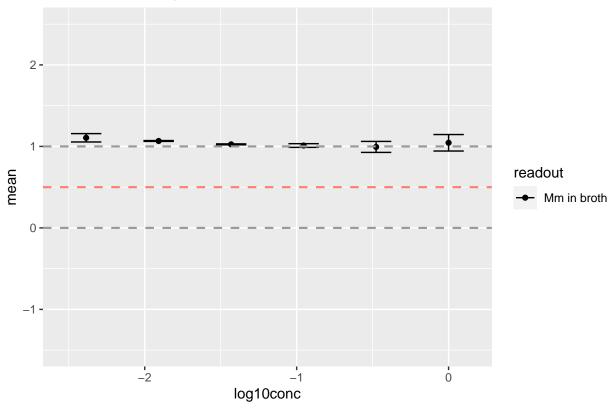


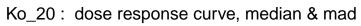


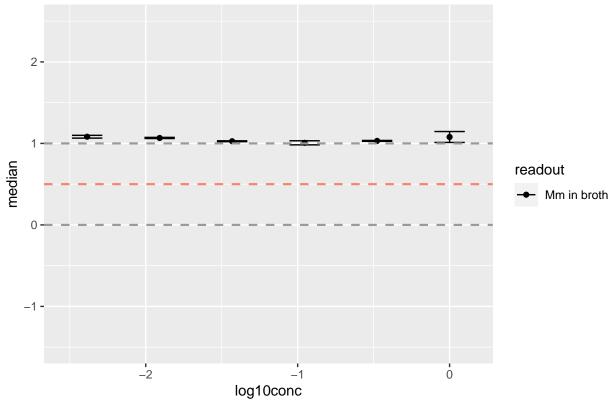




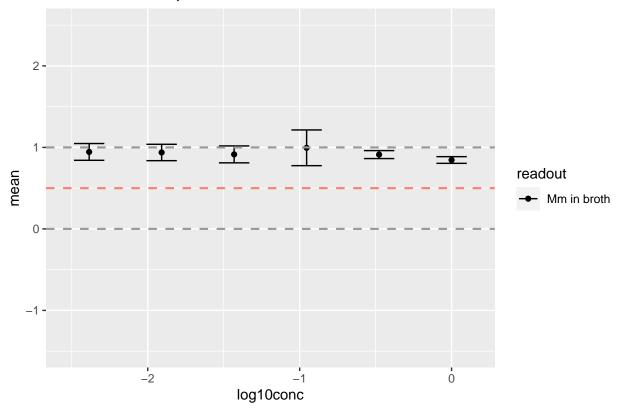
Ko_20 : dose response curve, mean & sd

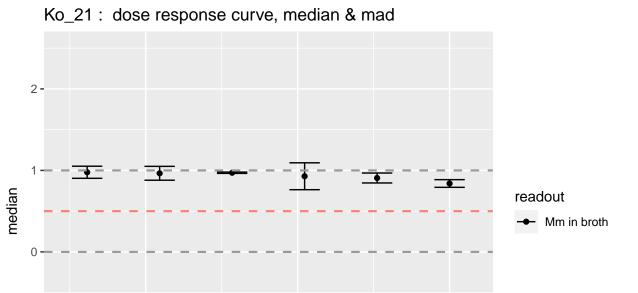






Ko_21 : dose response curve, mean & sd



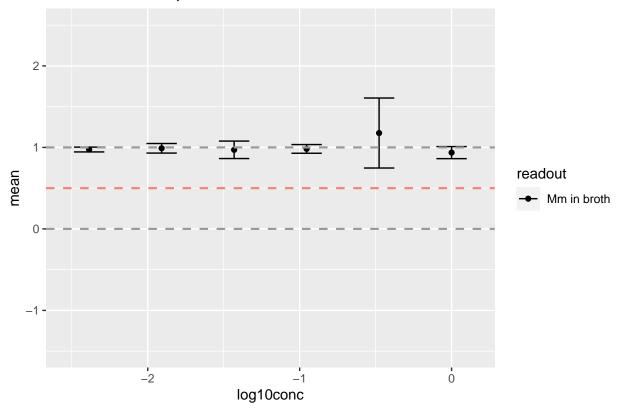


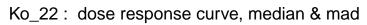
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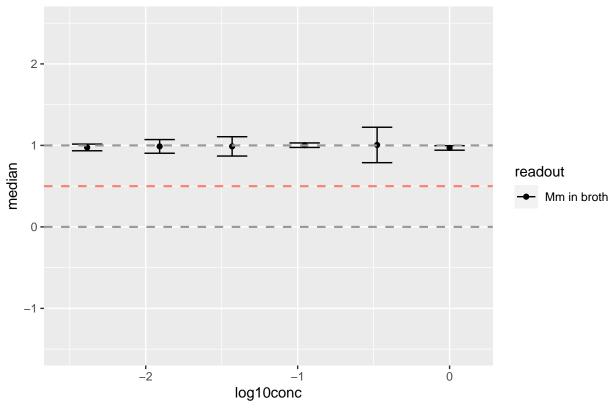
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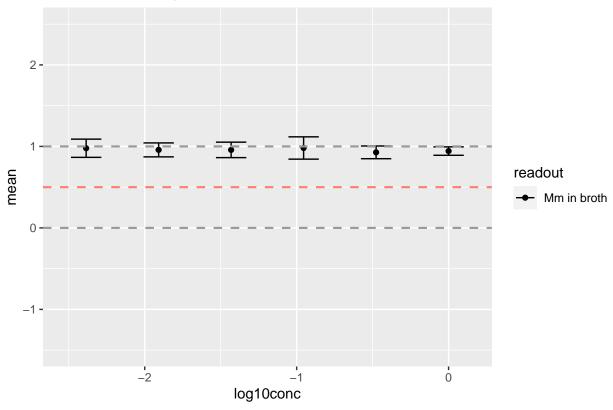
Ko_22 : dose response curve, mean & sd

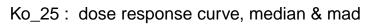


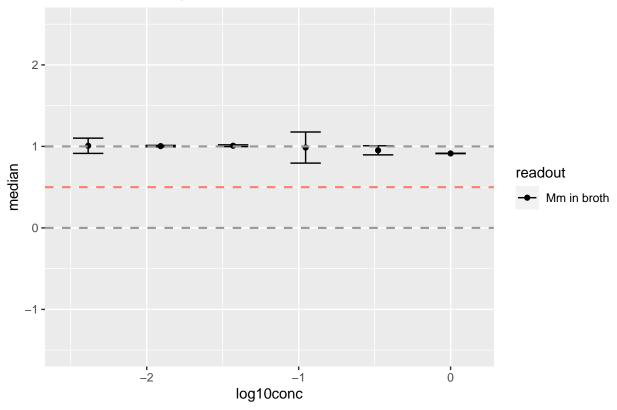


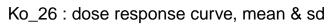


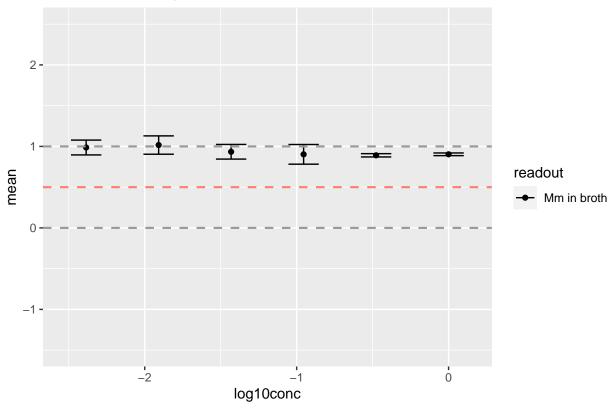
Ko_25 : dose response curve, mean & sd



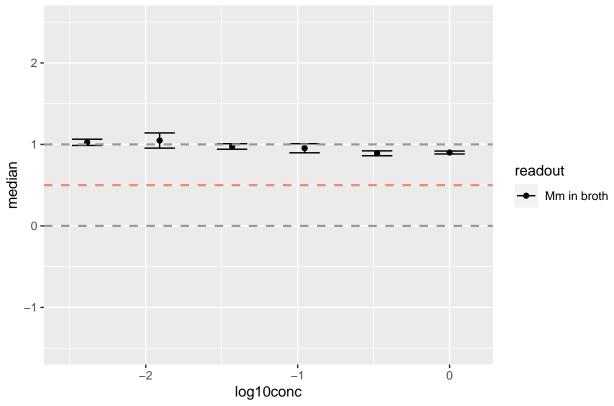




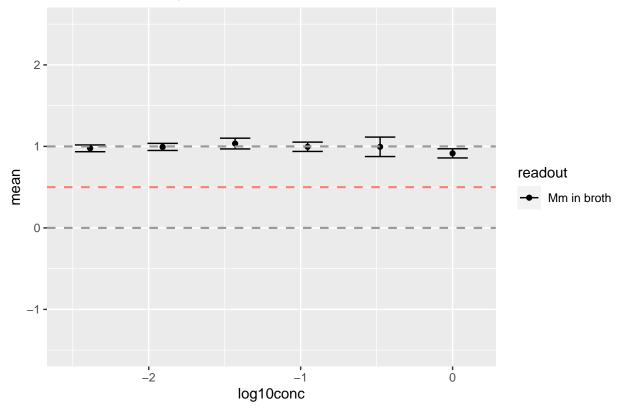


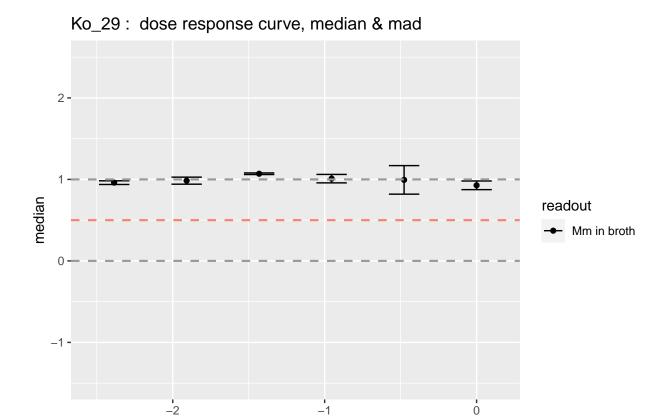






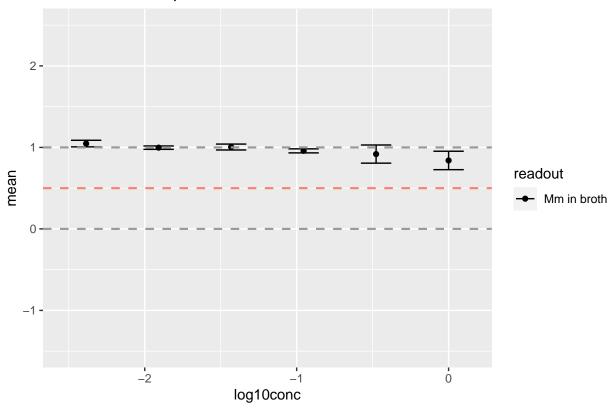
Ko_29 : dose response curve, mean & sd

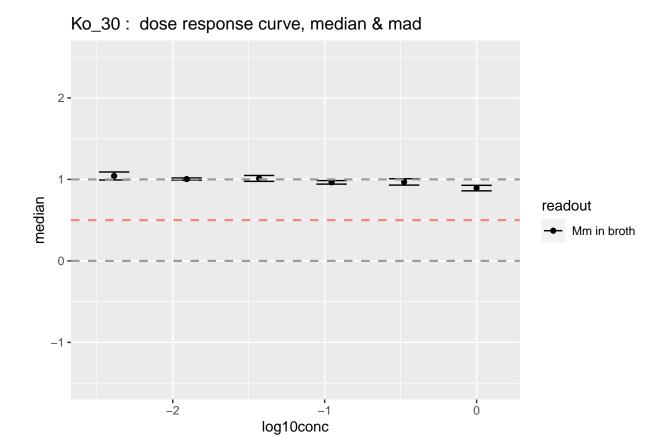




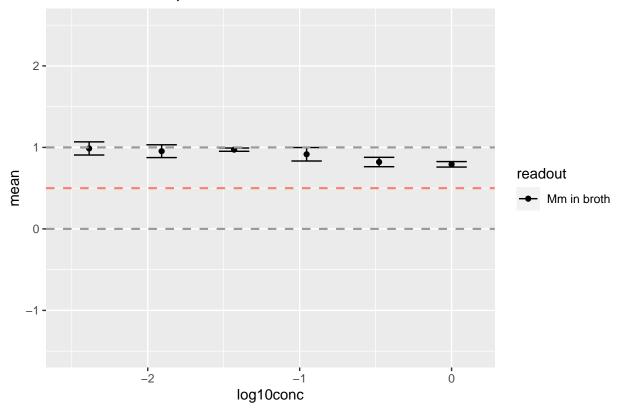
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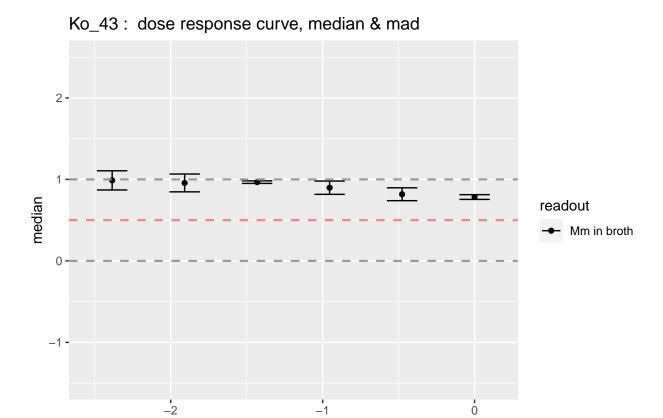






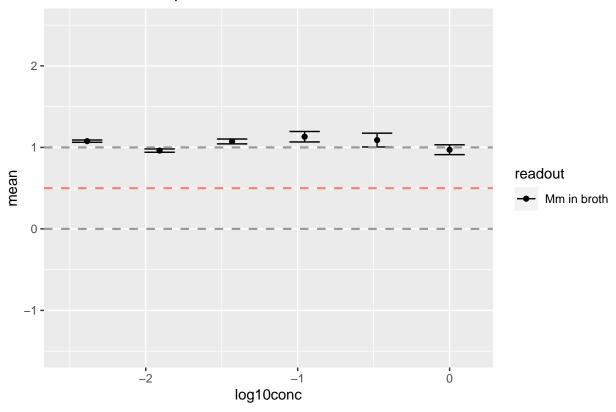
Ko_43 : dose response curve, mean & sd

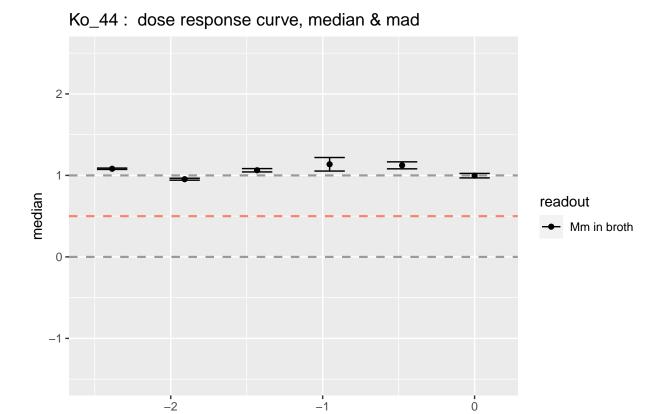




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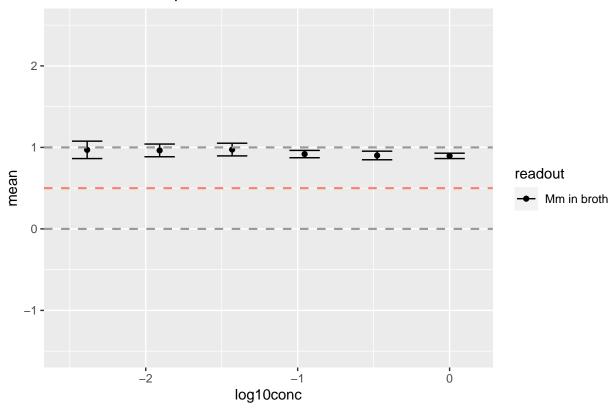
Ko_44 : dose response curve, mean & sd

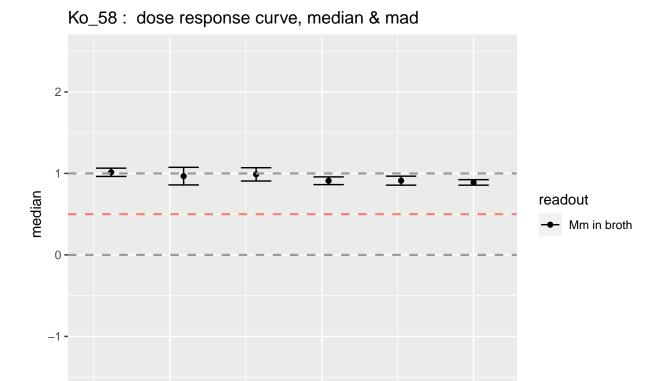




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Ko_58 : dose response curve, mean & sd

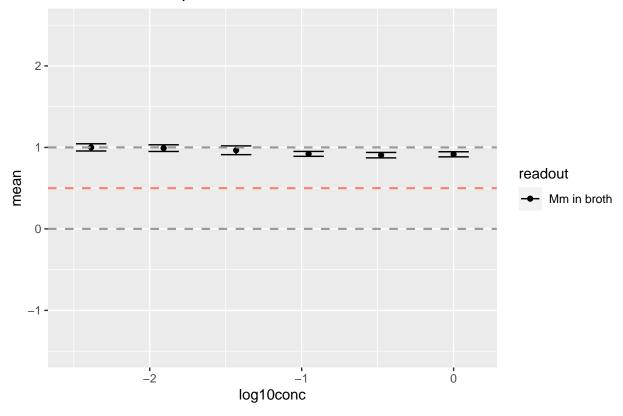


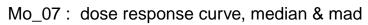


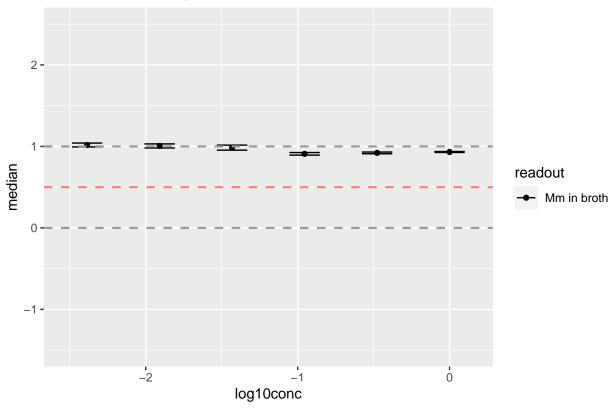
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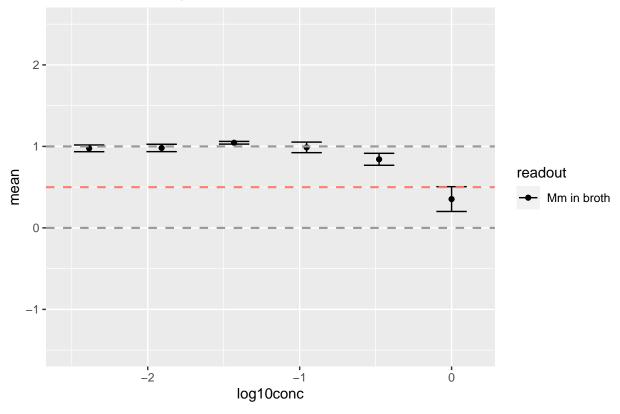
Mo_07 : dose response curve, mean & sd

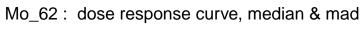


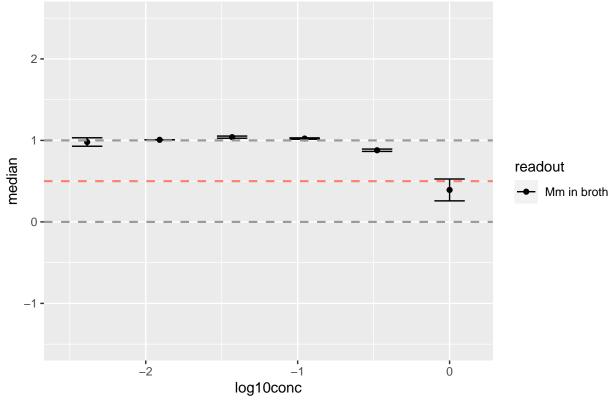




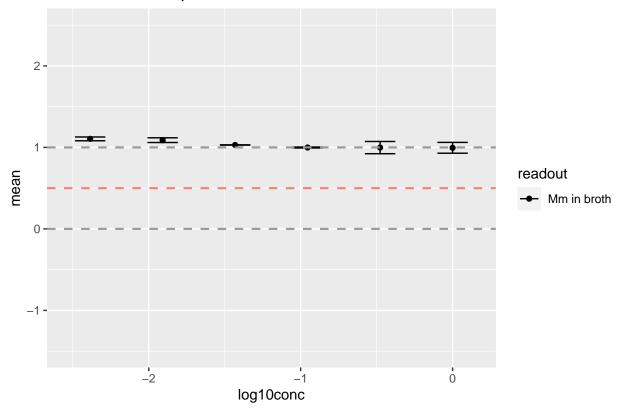
Mo_62 : dose response curve, mean & sd



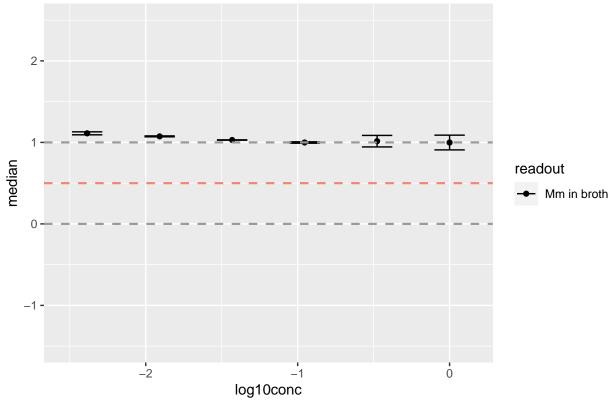




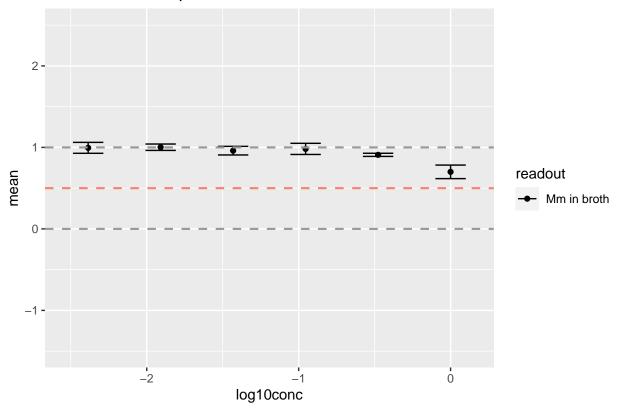
Mo_63 : dose response curve, mean & sd

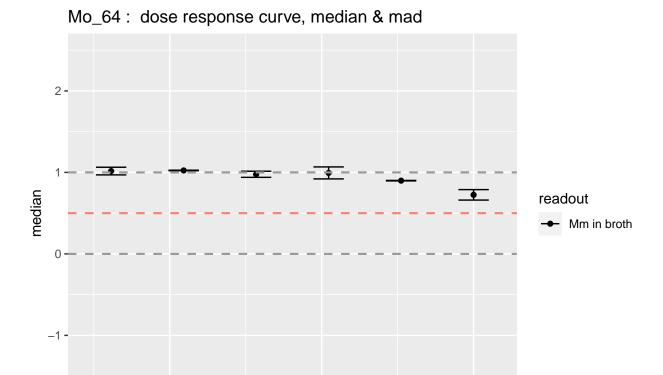


 Mo_63 : dose response curve, median & mad



Mo_64 : dose response curve, mean & sd

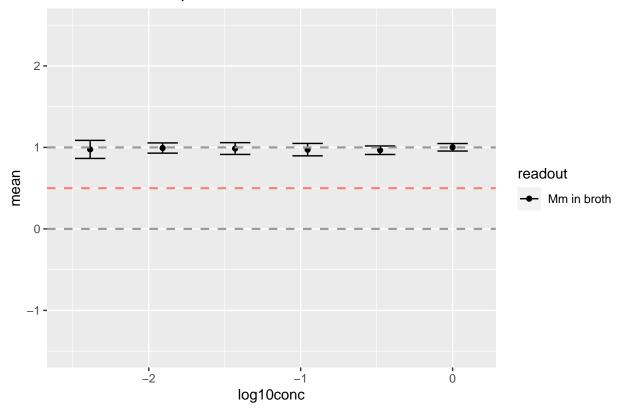




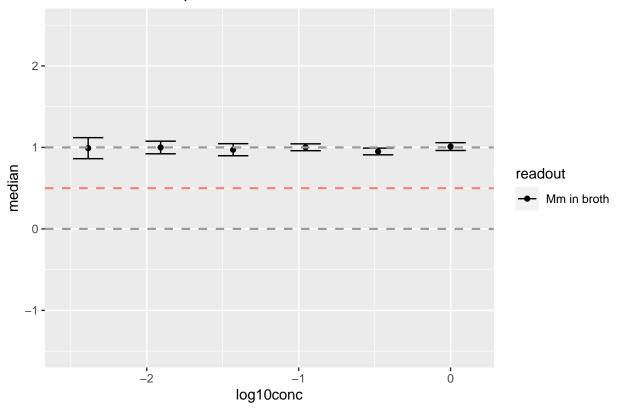
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74

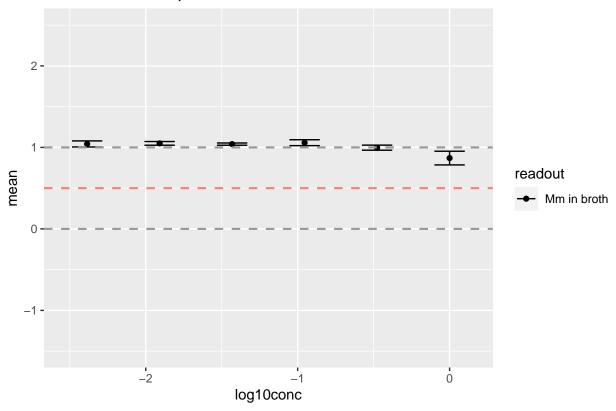
Mo_65 : dose response curve, mean & sd

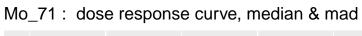


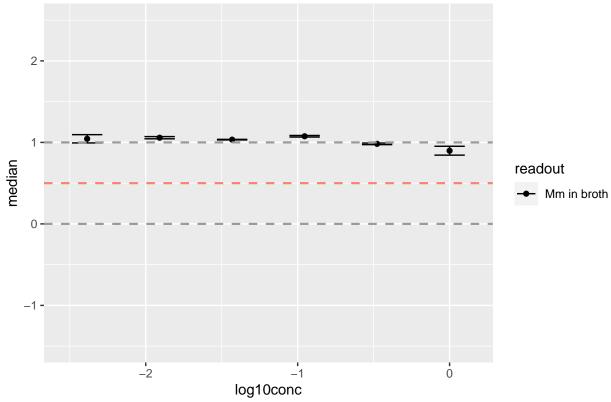
Mo_65: dose response curve, median & mad



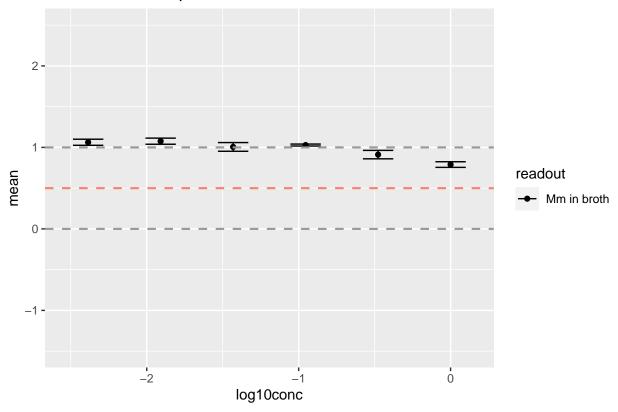


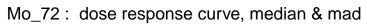


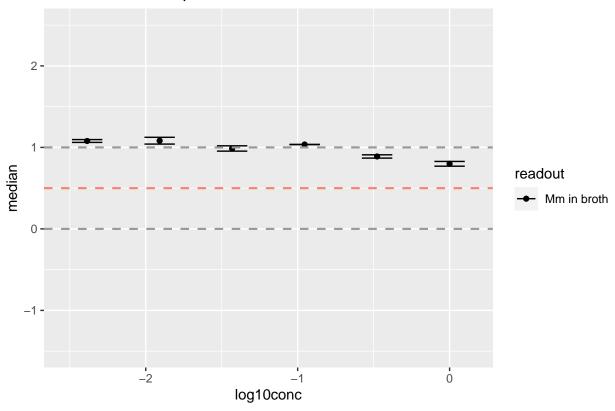




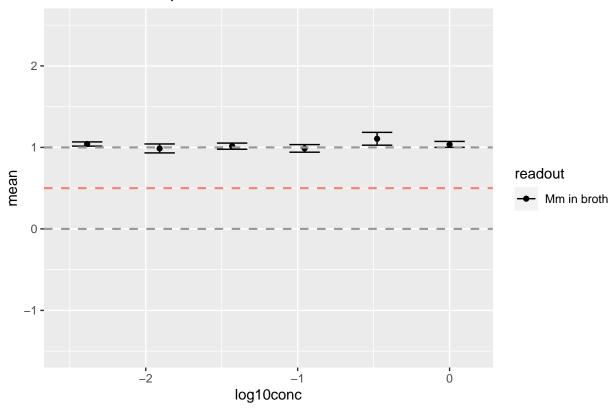
Mo_72 : dose response curve, mean & sd

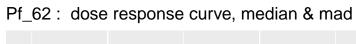


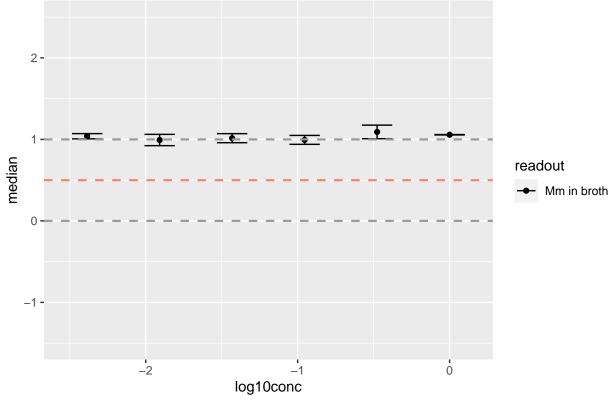


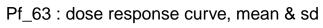


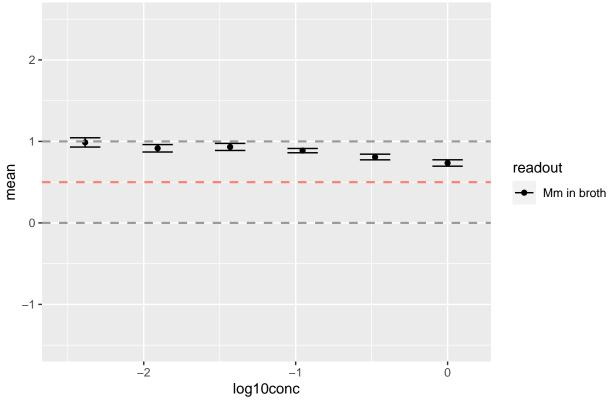


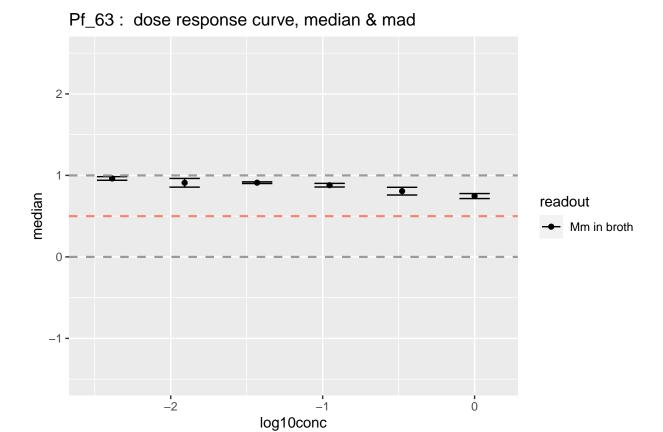




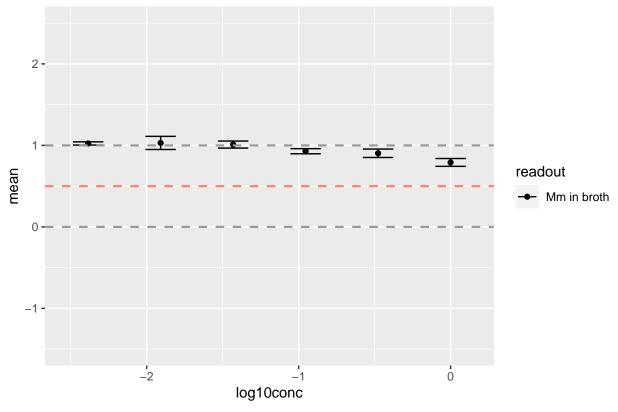


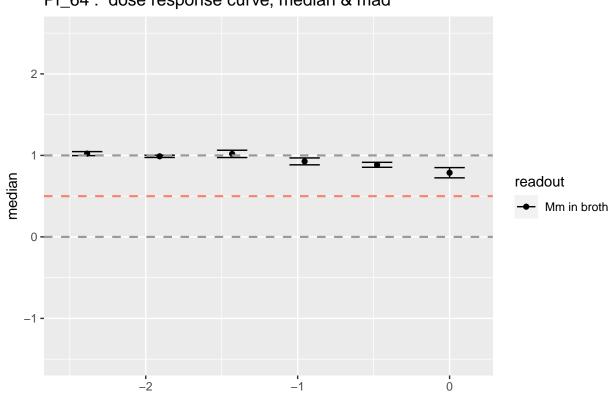




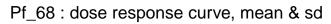


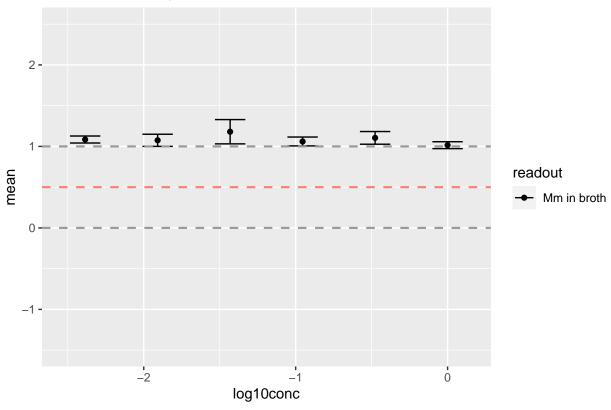






Pf_64: dose response curve, median & mad

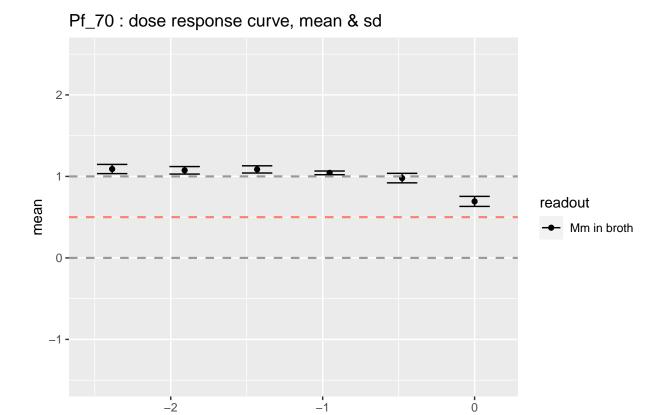




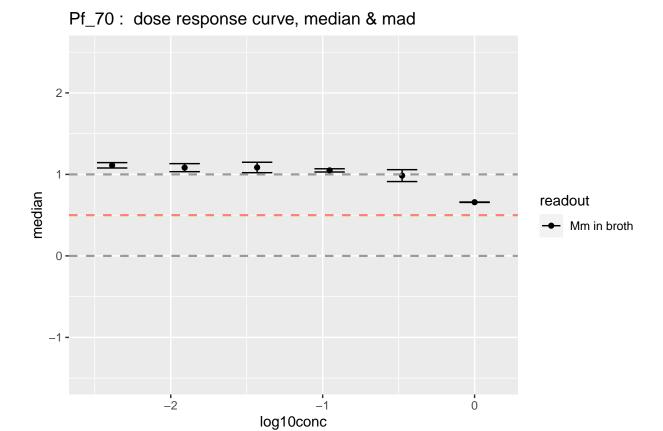


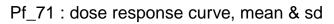
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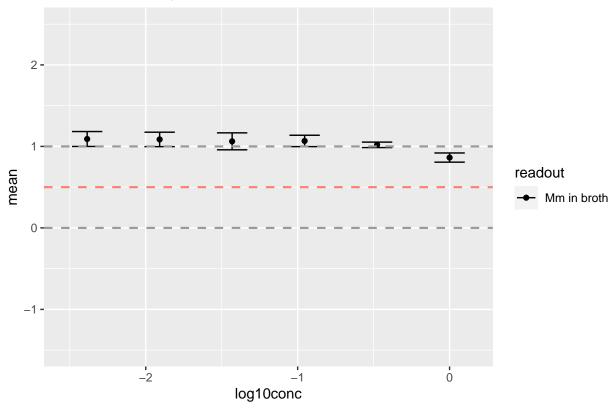
88

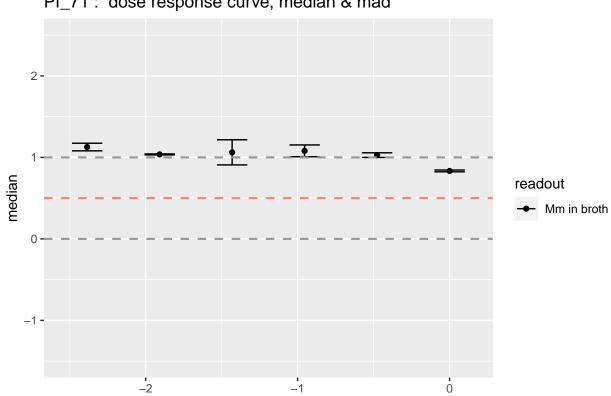


89

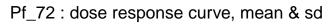


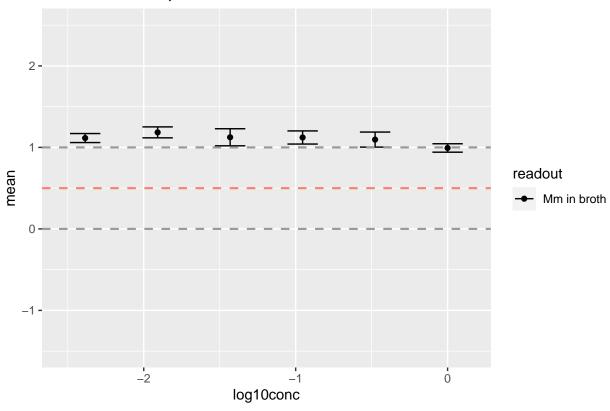


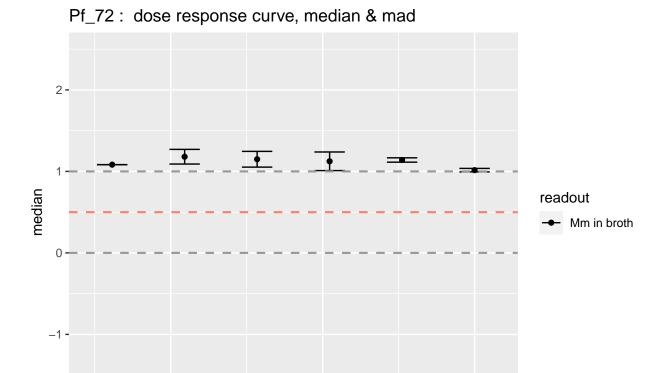




Pf_71: dose response curve, median & mad

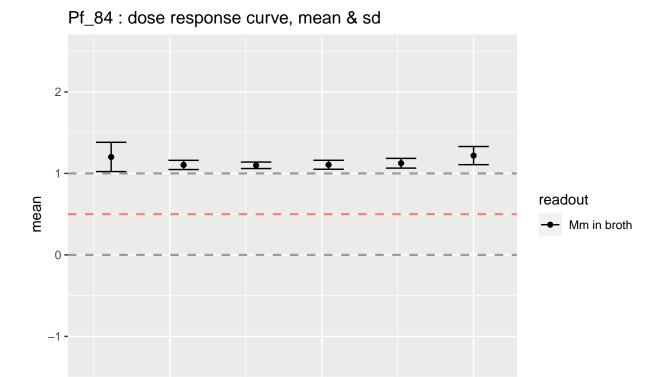






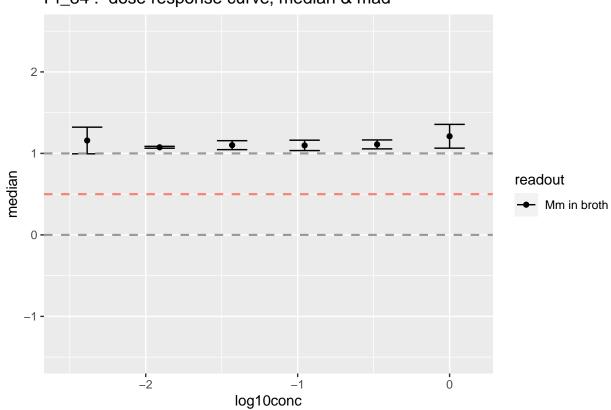
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94

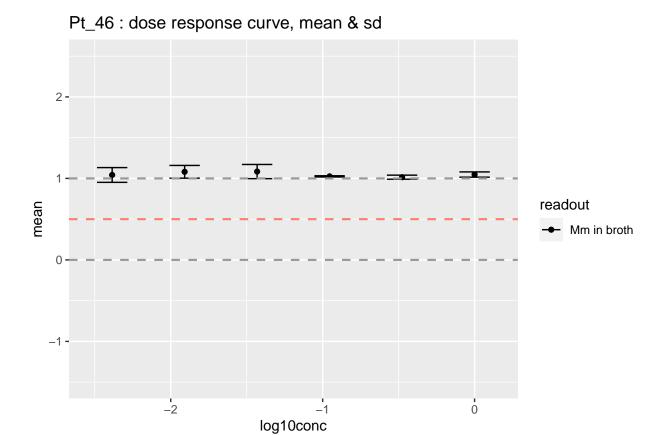


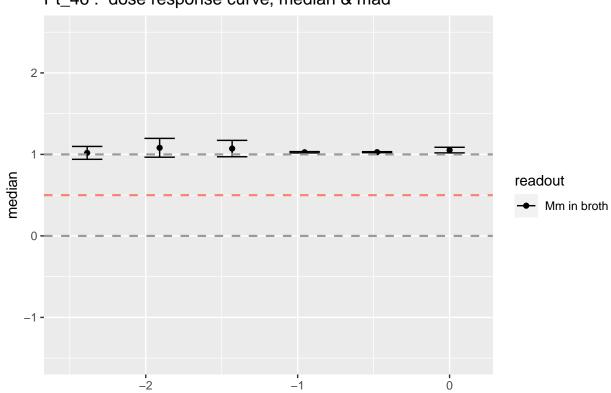
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95

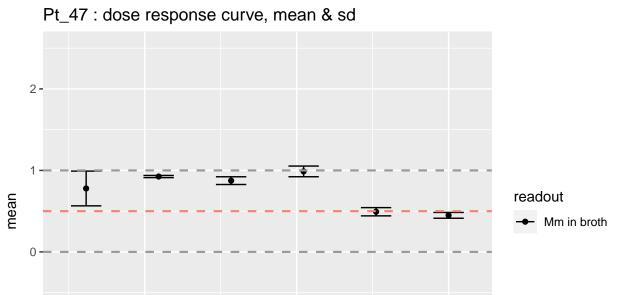


Pf_84: dose response curve, median & mad



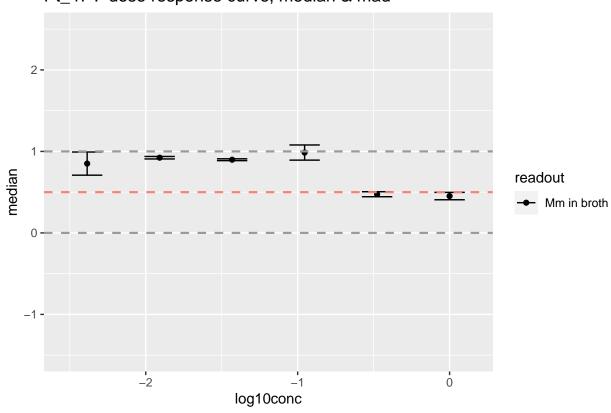


Pt_46: dose response curve, median & mad

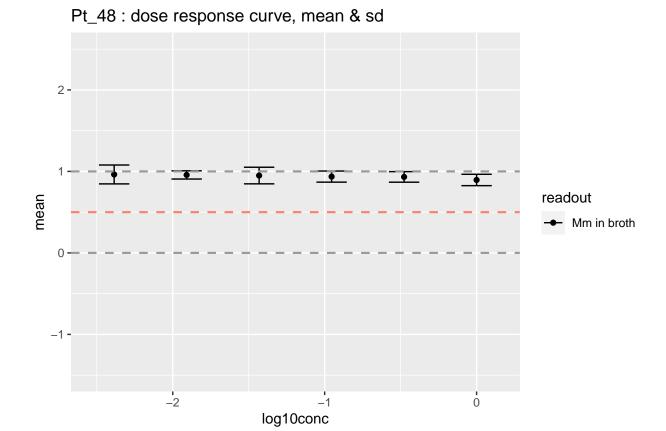


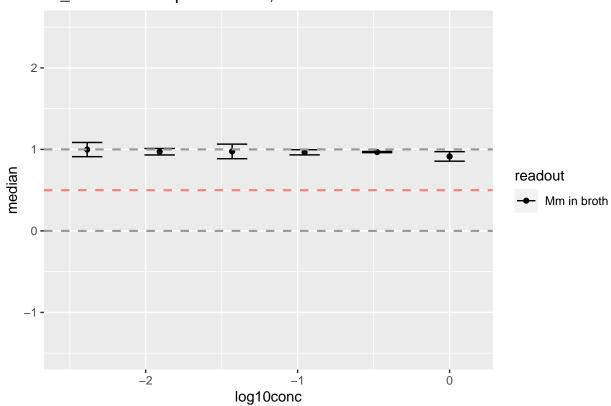
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_1 **-**



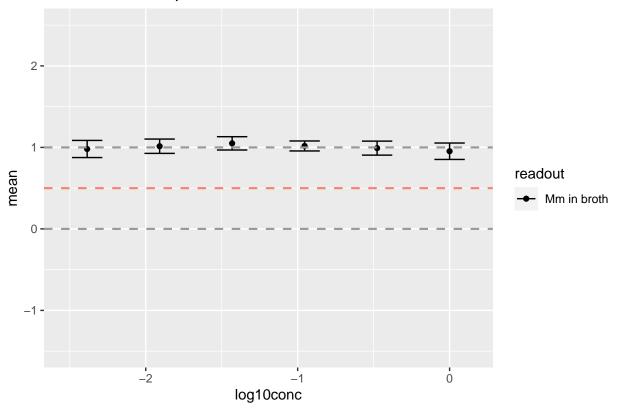
Pt_47: dose response curve, median & mad

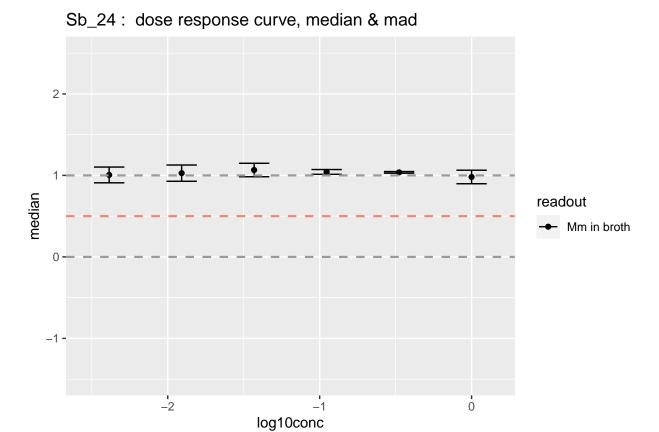




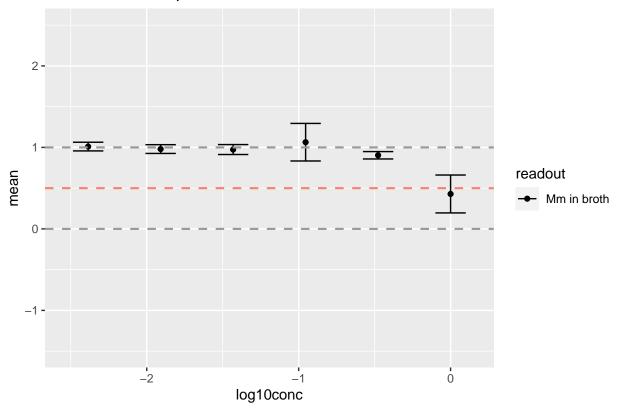
Pt_48: dose response curve, median & mad

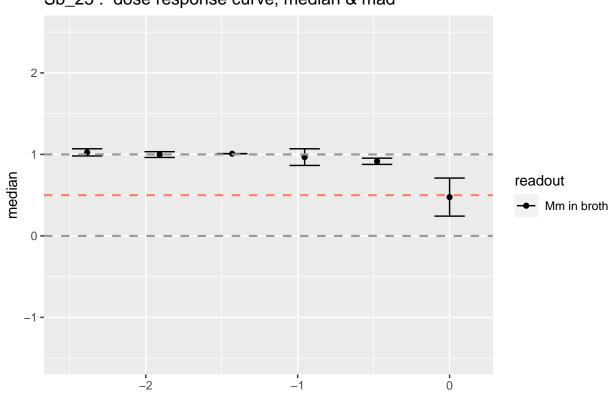
Sb_24 : dose response curve, mean & sd





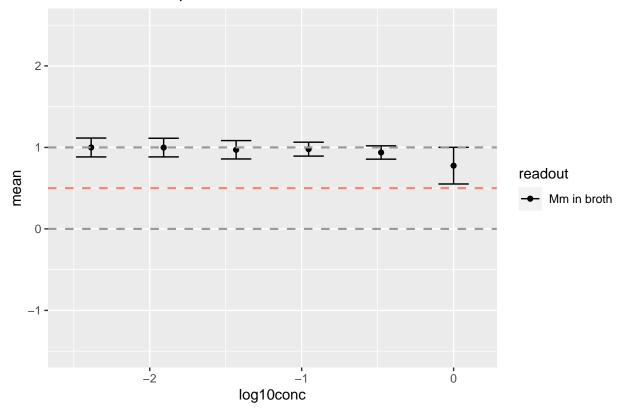
Sb_25 : dose response curve, mean & sd

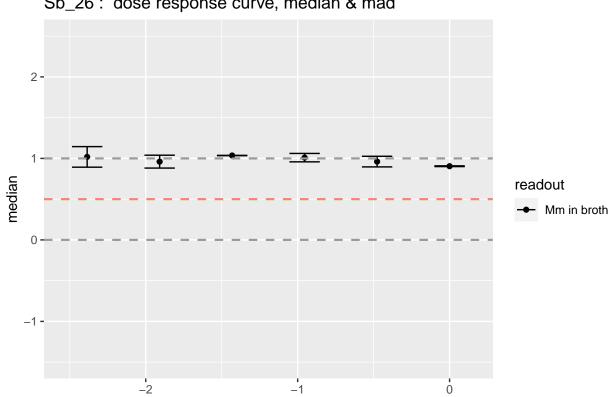




Sb_25: dose response curve, median & mad

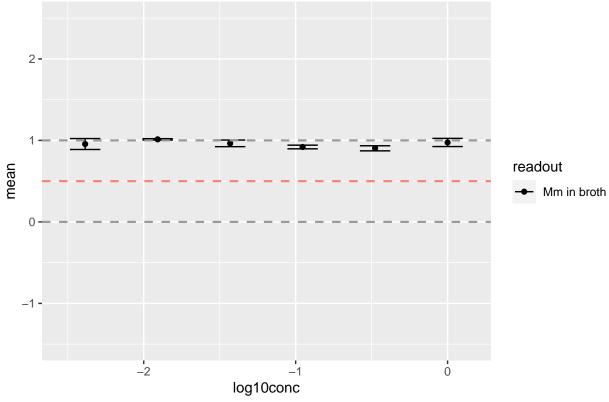
Sb_26 : dose response curve, mean & sd

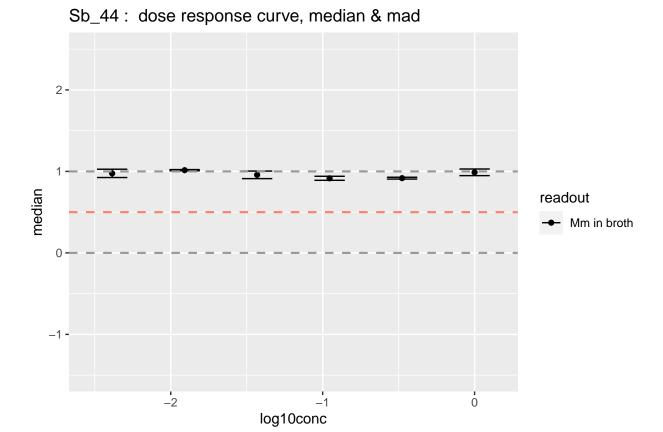


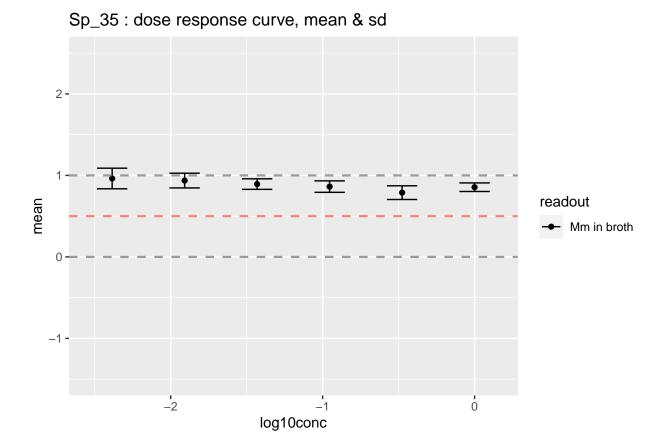


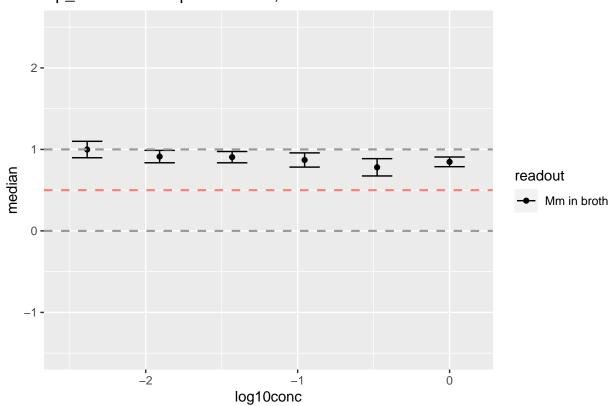
Sb_26: dose response curve, median & mad

Sb_44 : dose response curve, mean & sd

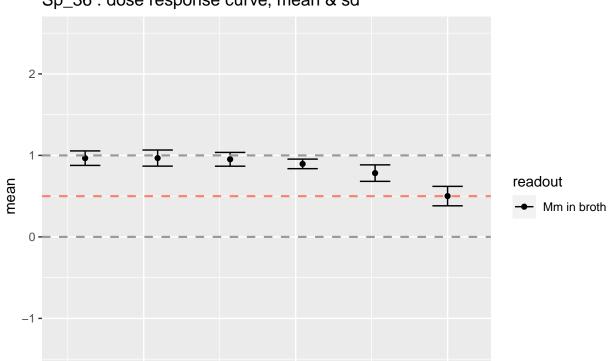






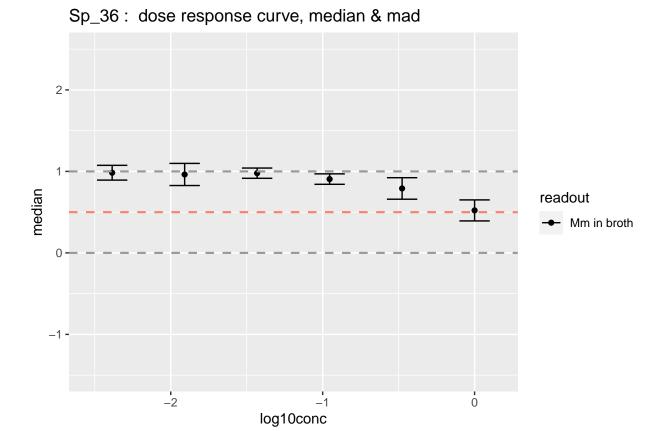


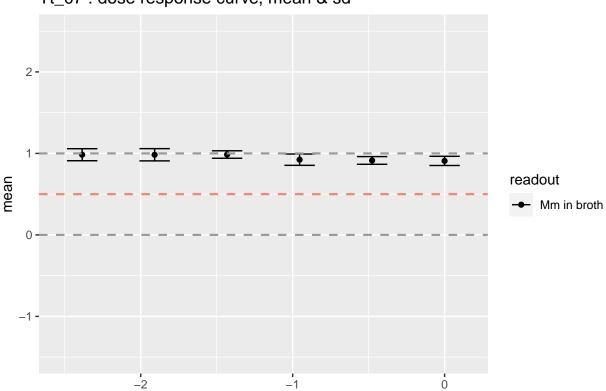
Sp_35: dose response curve, median & mad



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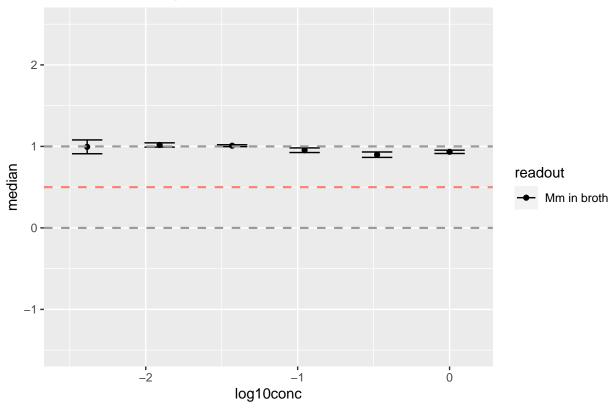
 $\ensuremath{\mathsf{Sp_36}}$: dose response curve, mean & sd





Tt_07 : dose response curve, mean & sd

Tt_07: dose response curve, median & mad



```
## R version 4.3.1 (2023-06-16 ucrt)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19045)
## Matrix products: default
##
##
## locale:
## [1] LC_COLLATE=English_Switzerland.utf8 LC_CTYPE=English_Switzerland.utf8
## [3] LC_MONETARY=English_Switzerland.utf8 LC_NUMERIC=C
## [5] LC_TIME=English_Switzerland.utf8
##
## time zone: Europe/Zurich
## tzcode source: internal
## attached base packages:
## [1] stats
                graphics grDevices utils
                                              datasets methods
                                                                   base
##
## other attached packages:
## [1] reshape2_1.4.4 tidyr_1.3.0
                                     dplyr_1.1.3
                                                    ggplot2_3.4.4 gplots_3.1.3
## [6] readxl_1.4.3 xlsx_0.6.5
##
## loaded via a namespace (and not attached):
##
  [1] gtable_0.3.4
                          compiler_4.3.1
                                             gtools_3.9.4
                                                                 Rcpp_1.0.11
  [5] tidyselect_1.2.0 stringr_1.5.0
                                             bitops_1.0-7
                                                                 scales_1.2.1
##
  [9] yaml_2.3.7
                          fastmap_1.1.1
                                                                 R6_2.5.1
                                             plyr_1.8.9
```

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##	[17]	munsell_0.5.0	pillar_1.9.0	rlang_1.1.1	utf8_1.2.3
##	[21]	stringi_1.7.12	xfun_0.40	caTools_1.18.2	cli_3.6.1
##	[25]	withr_2.5.1	magrittr_2.0.3	digest_0.6.33	grid_4.3.1
##	[29]	rstudioapi_0.15.0	rJava_1.0-6	lifecycle_1.0.3	vctrs_0.6.3
##	[33]	${\tt KernSmooth_2.23-21}$	evaluate_0.22	glue_1.6.2	farver_2.1.1
##	[37]	cellranger_1.1.0	xlsxjars_0.6.1	fansi_1.0.5	<pre>colorspace_2.1-0</pre>
##	[41]	rmarkdown_2.25	purrr_1.0.2	tools_4.3.1	pkgconfig_2.0.3
##	[45]	htmltools_0.5.6.1			