

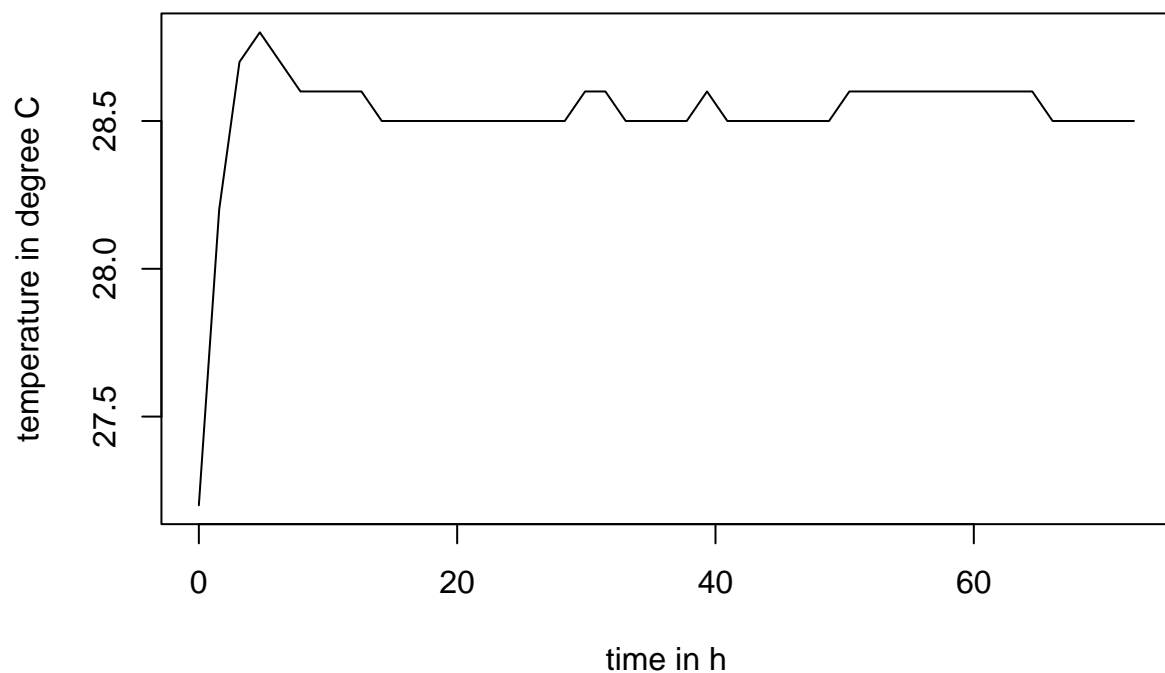
Normalized residual growth for experiment XYZ

JN

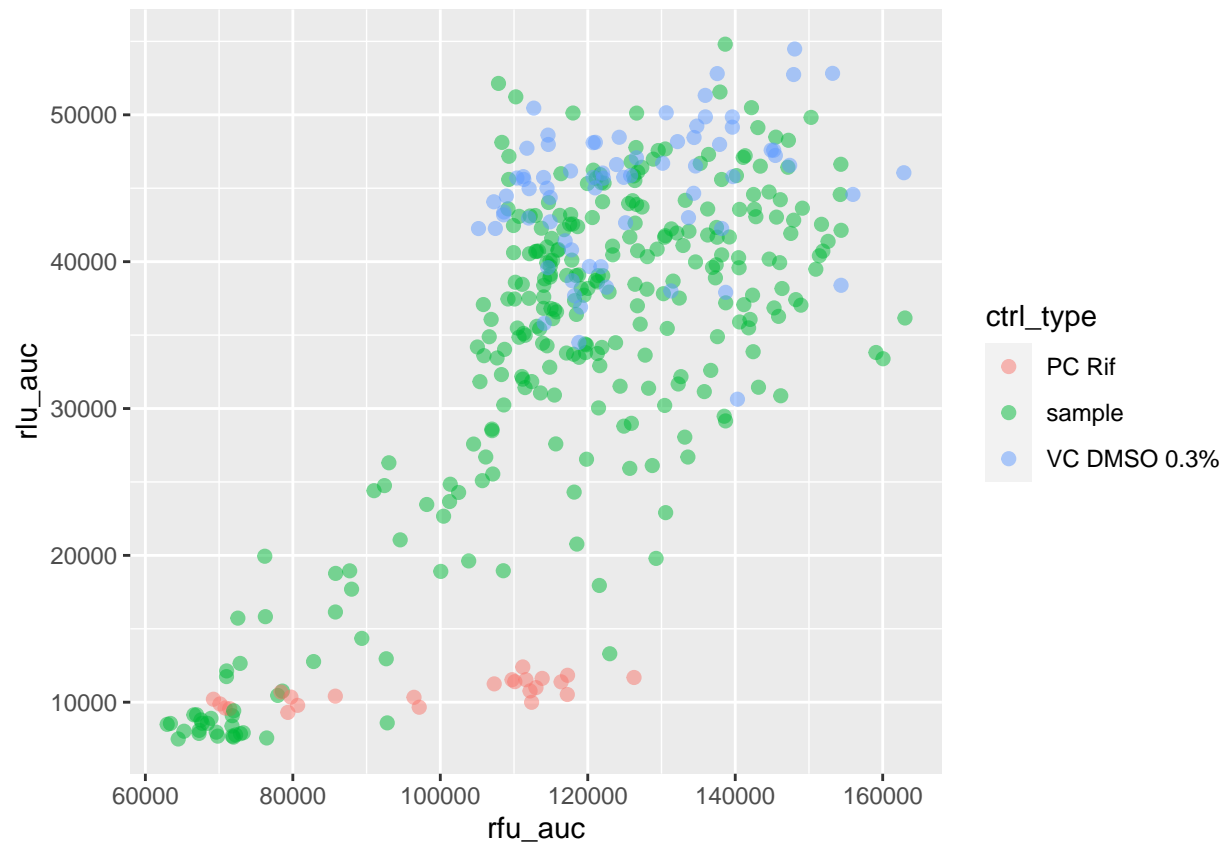
Sat Feb 17 13:20:51 2024

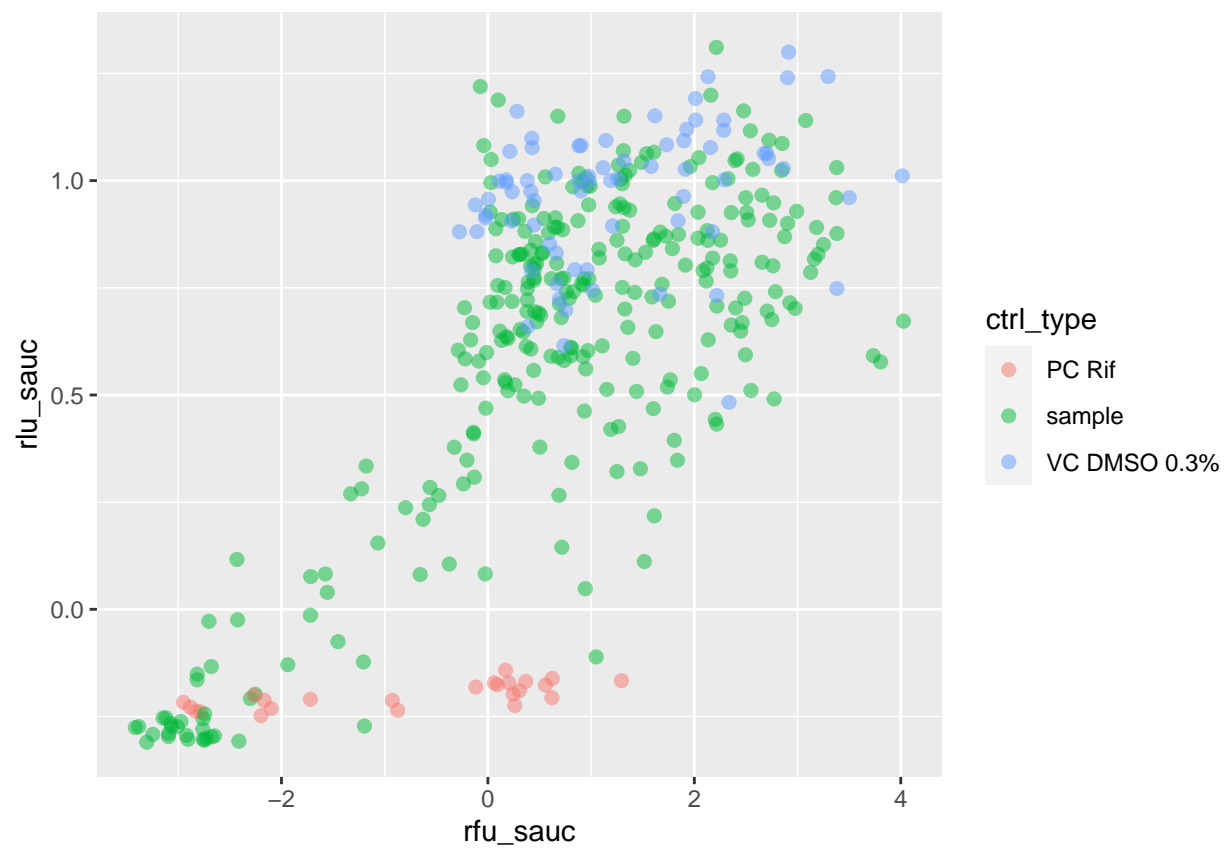
```
## [1] "all files in the directory with the absolute path: "  
## [2] "C:/Users/jahn_/Desktop/analyses/infection_demonstration_dataset/input/plate02.xlsx"  
## [3] "C:/Users/jahn_/Desktop/analyses/infection_demonstration_dataset/input/plate03.xlsx"  
## [4] "C:/Users/jahn_/Desktop/analyses/infection_demonstration_dataset/input/plate04.xlsx"  
## [5] "C:/Users/jahn_/Desktop/analyses/infection_demonstration_dataset/input/plate05.xlsx"  
## [6] "C:/Users/jahn_/Desktop/analyses/infection_demonstration_dataset/input/plate06.xlsx"  
## [7] "C:/Users/jahn_/Desktop/analyses/infection_demonstration_dataset/input/plate07.xlsx"  
  
## [1] "all files in the directory: " "plate02.xlsx"  
## [3] "plate03.xlsx"                "plate04.xlsx"  
## [5] "plate05.xlsx"                "plate06.xlsx"  
## [7] "plate07.xlsx"  
  
## [1] "analysis for plate02.xlsx"
```

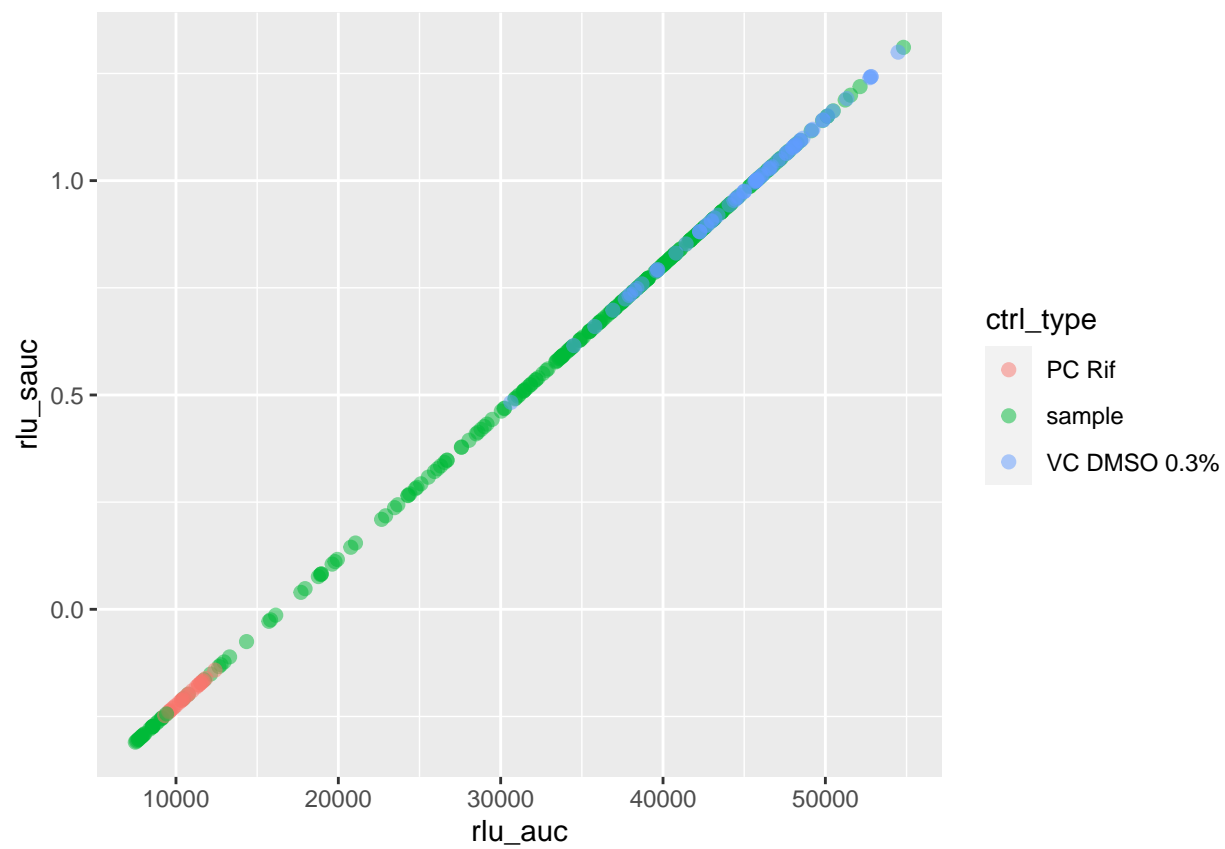
Temperature: plate 1

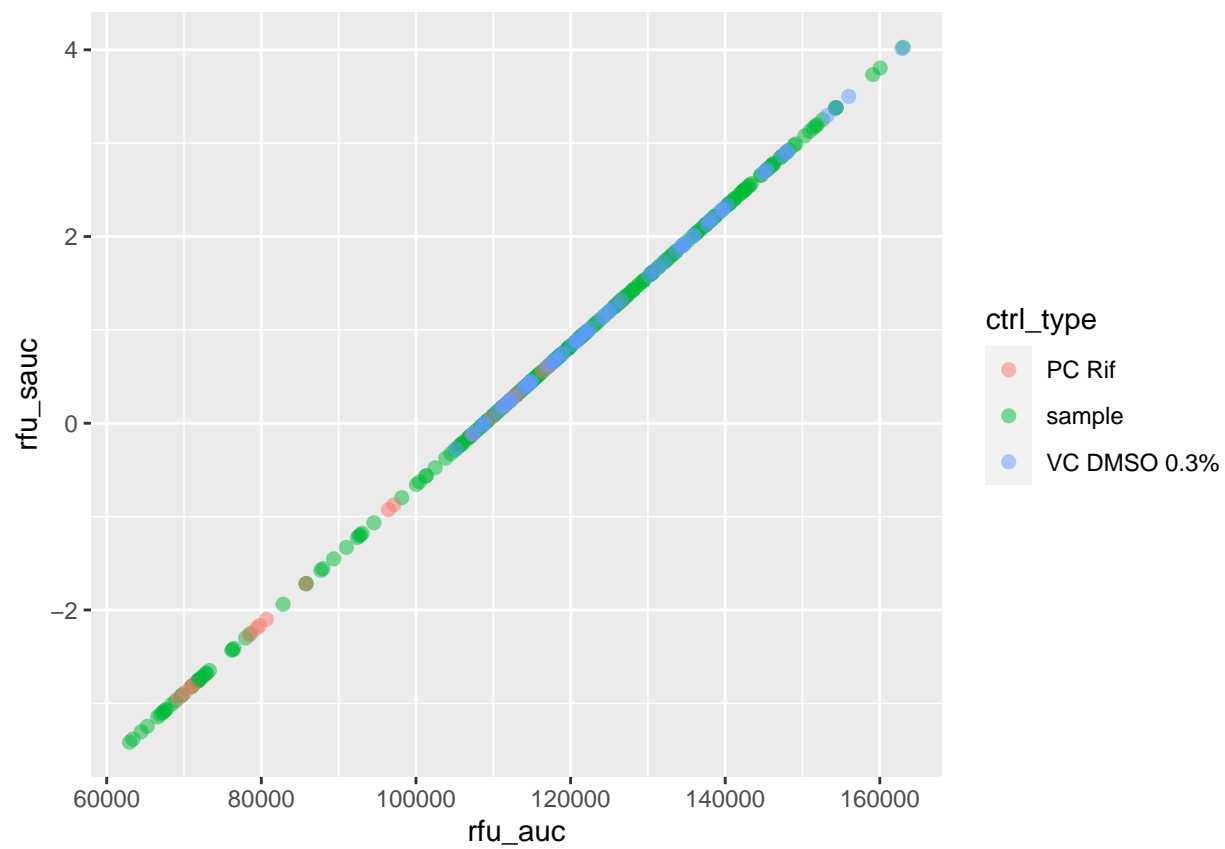


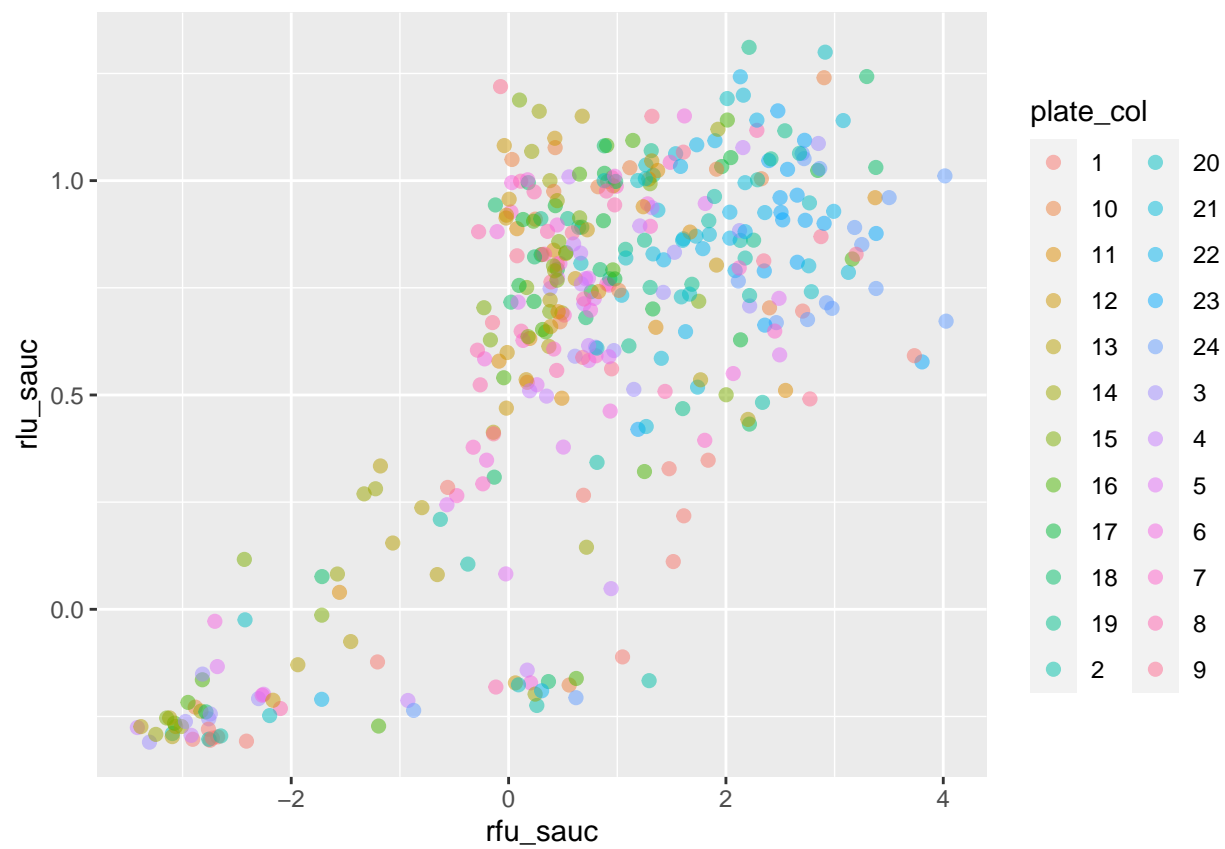
```
## [1] "VC DMSO 0.3% Robust z'-factor of rlu for plate plate02.xlsx, biorep 1 : "  
## [1] 0.57
```

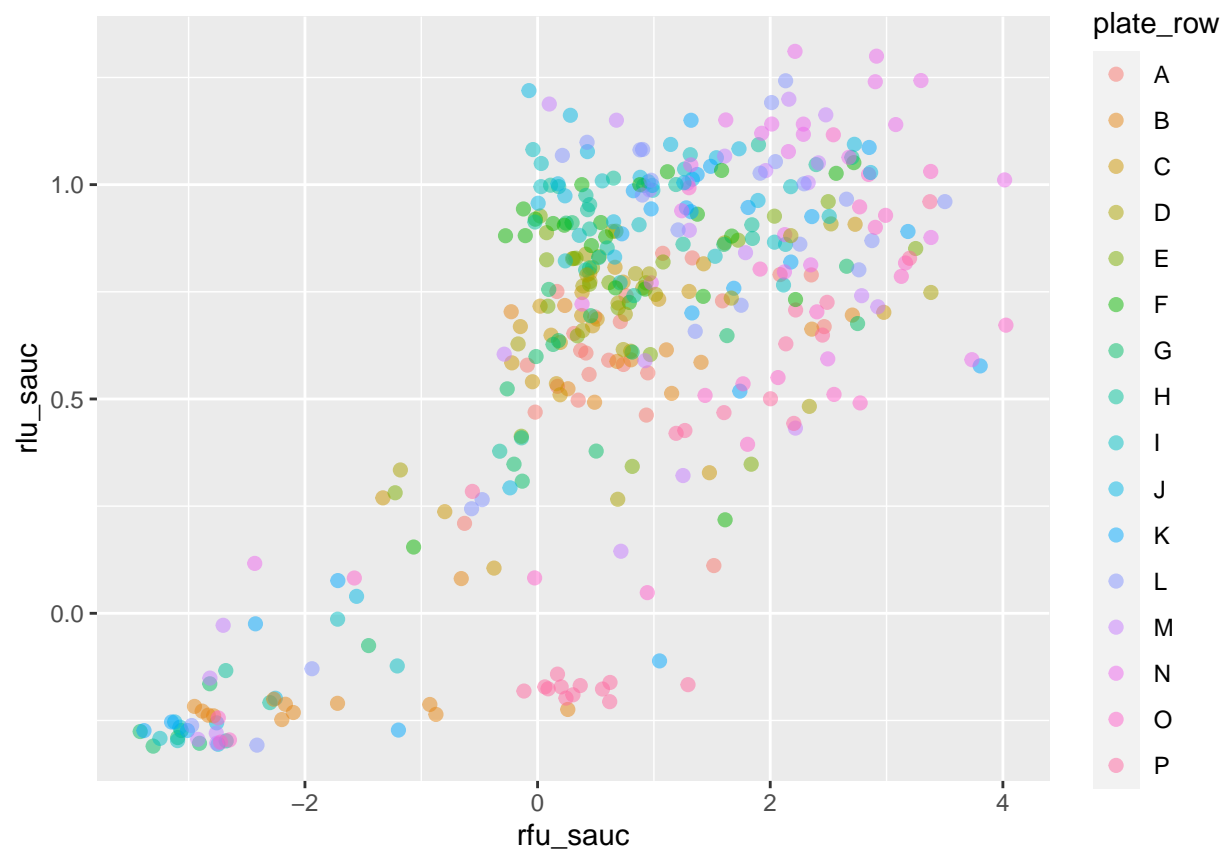




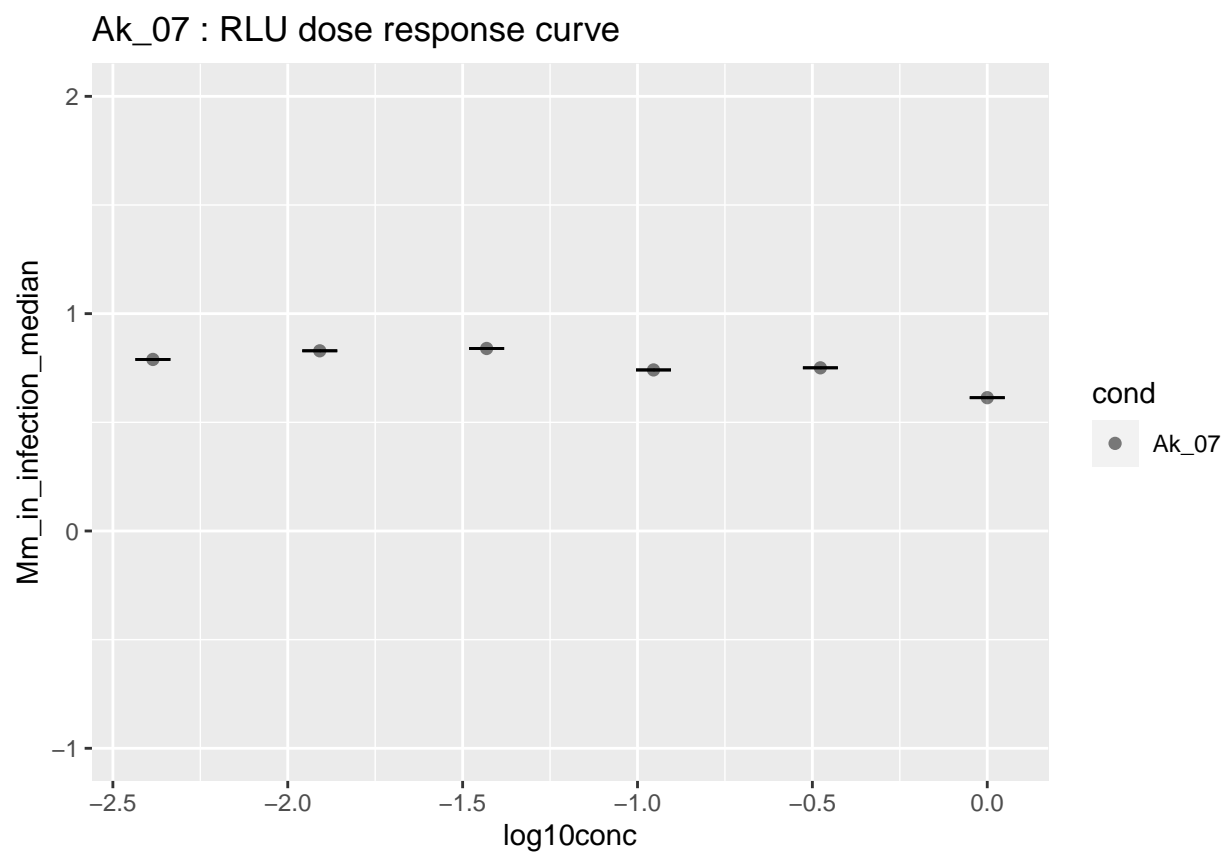


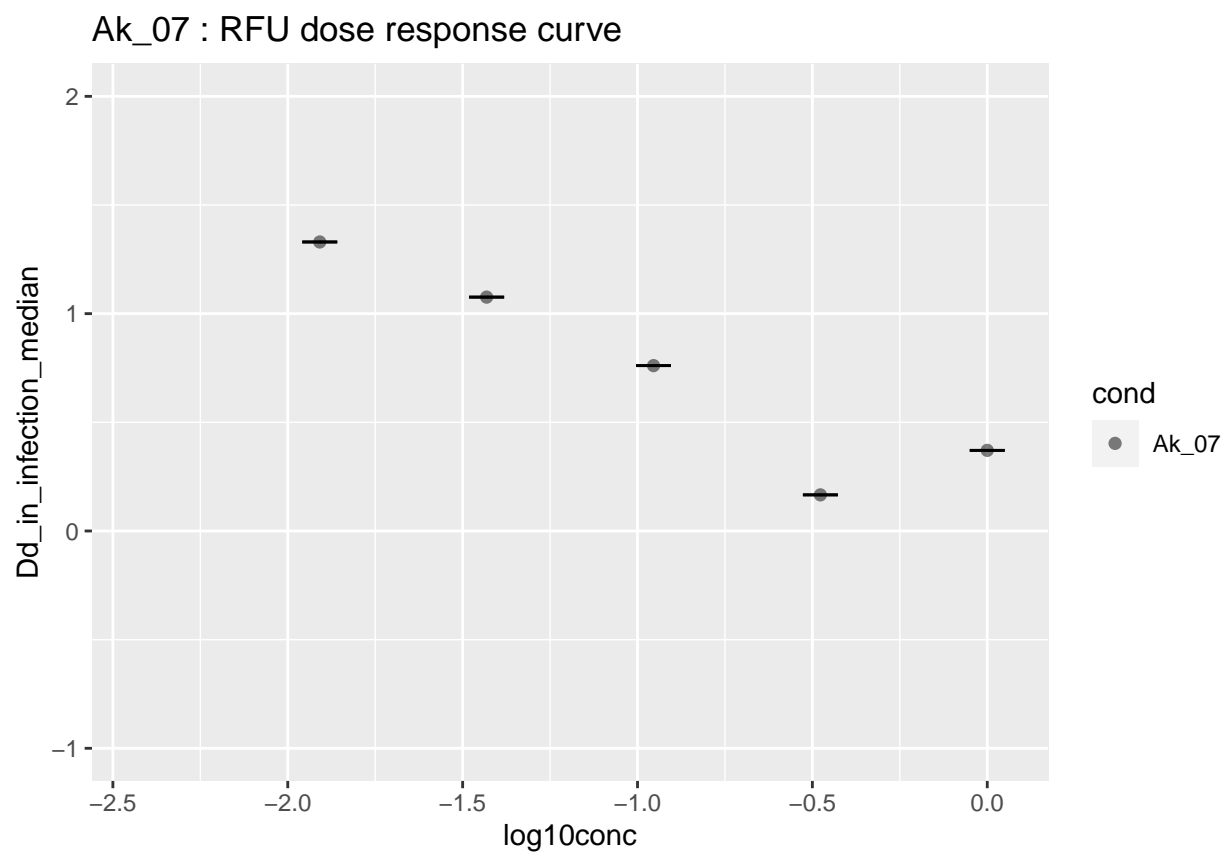


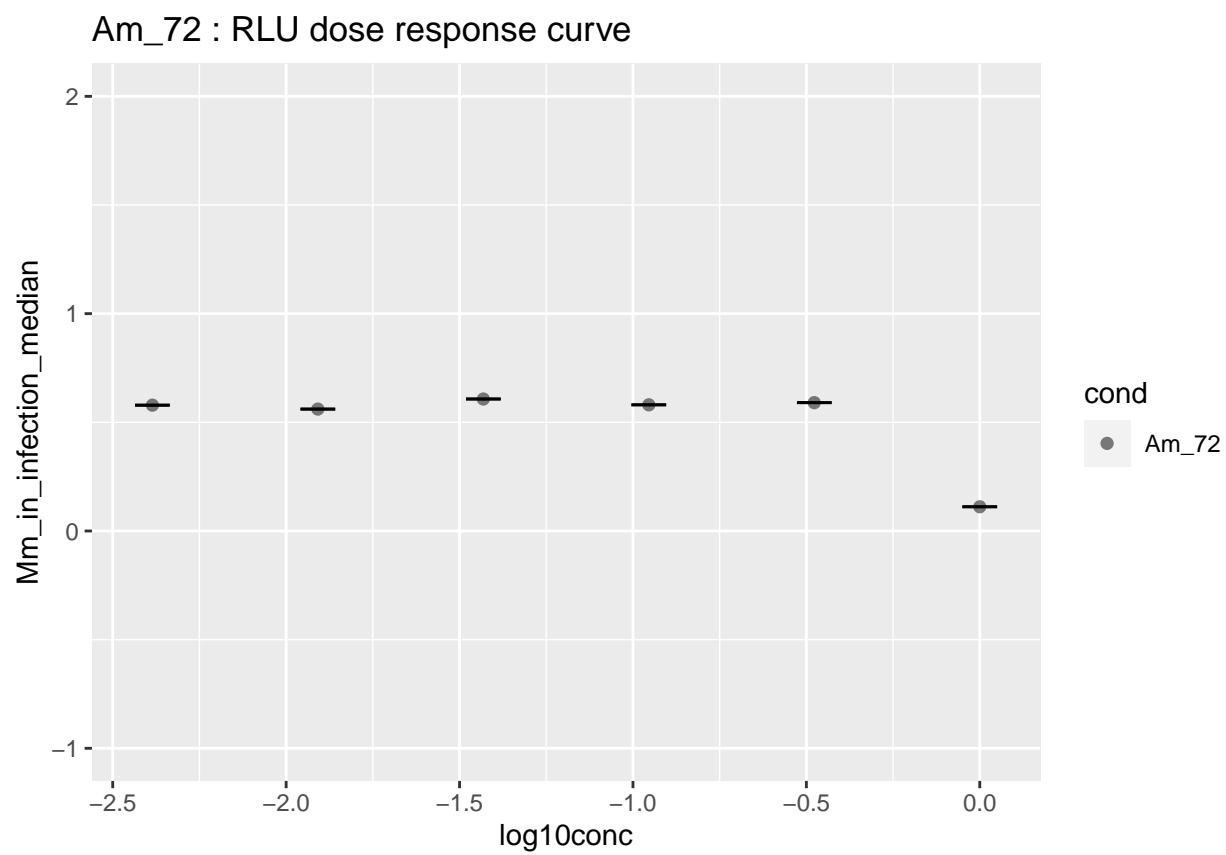


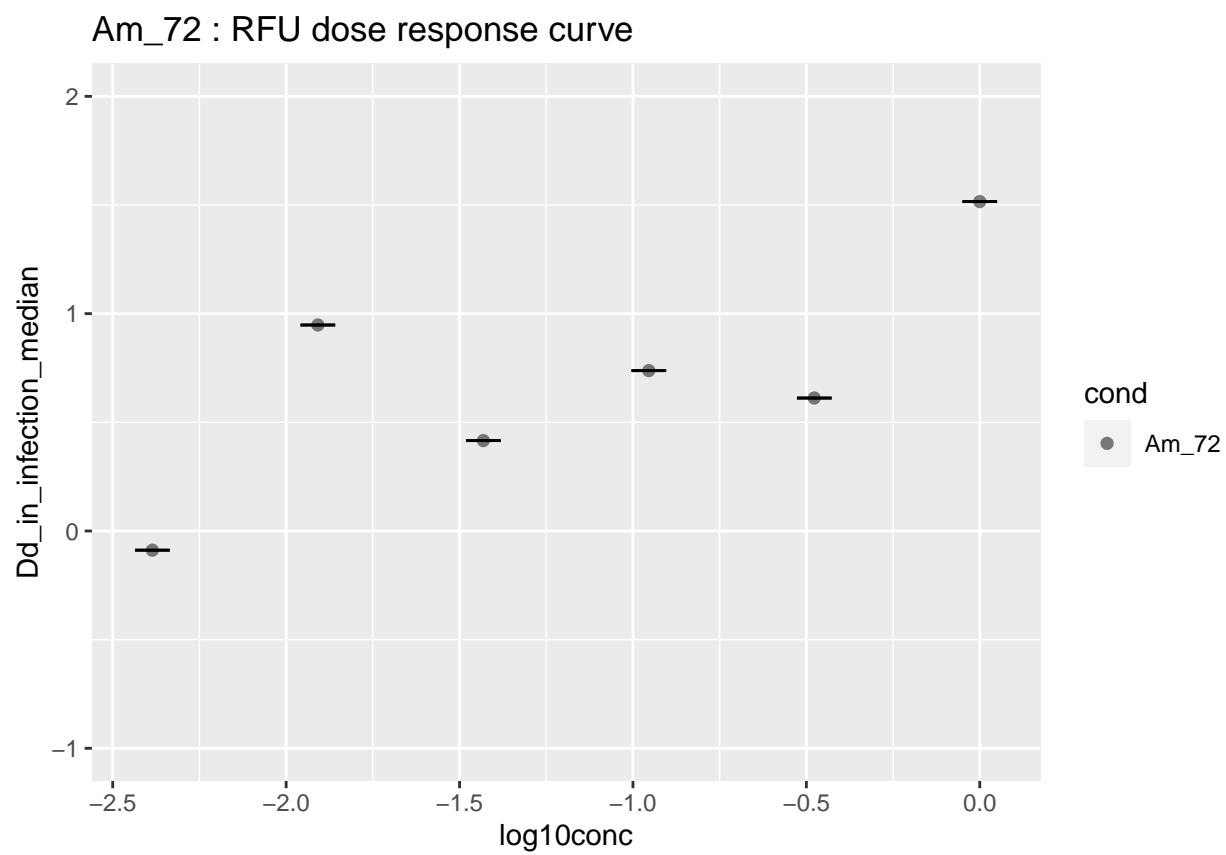


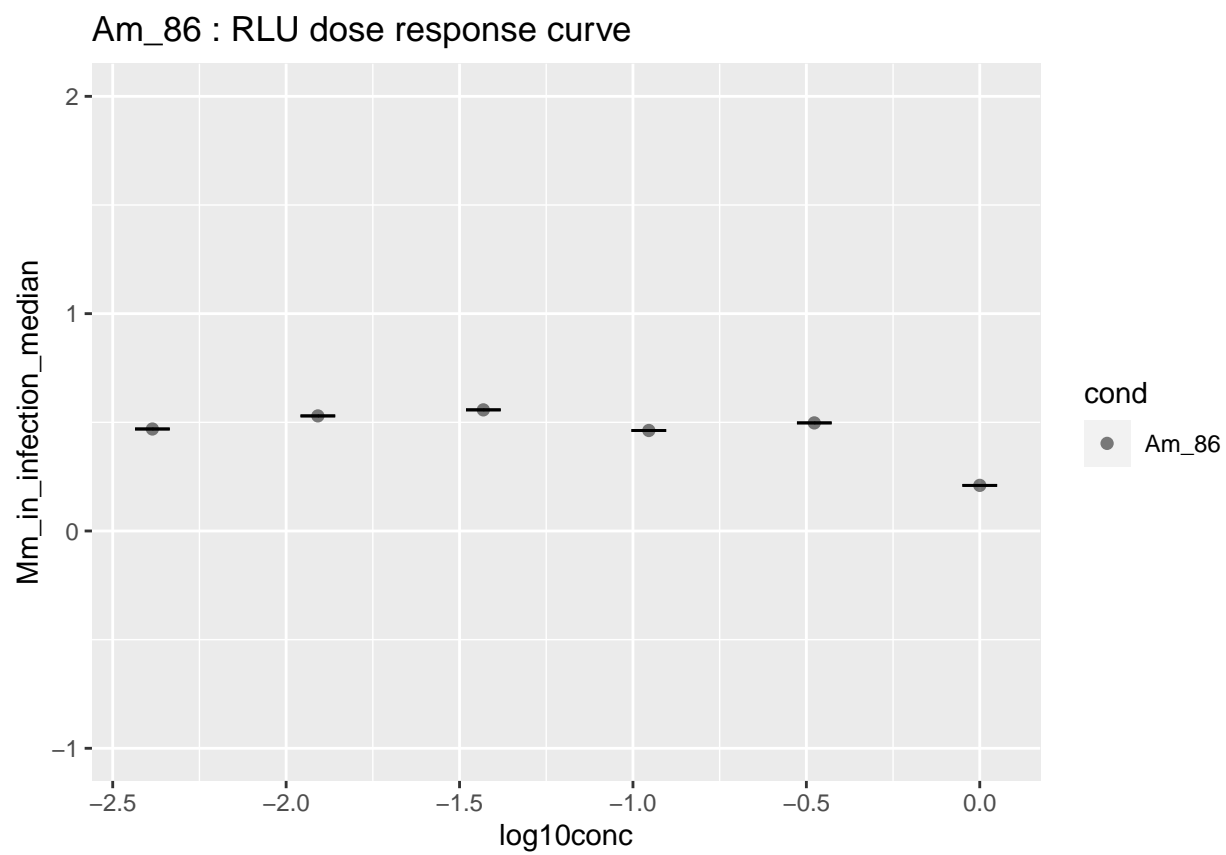
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## [1] "Dose response curves over all bioreps within this plate"
```

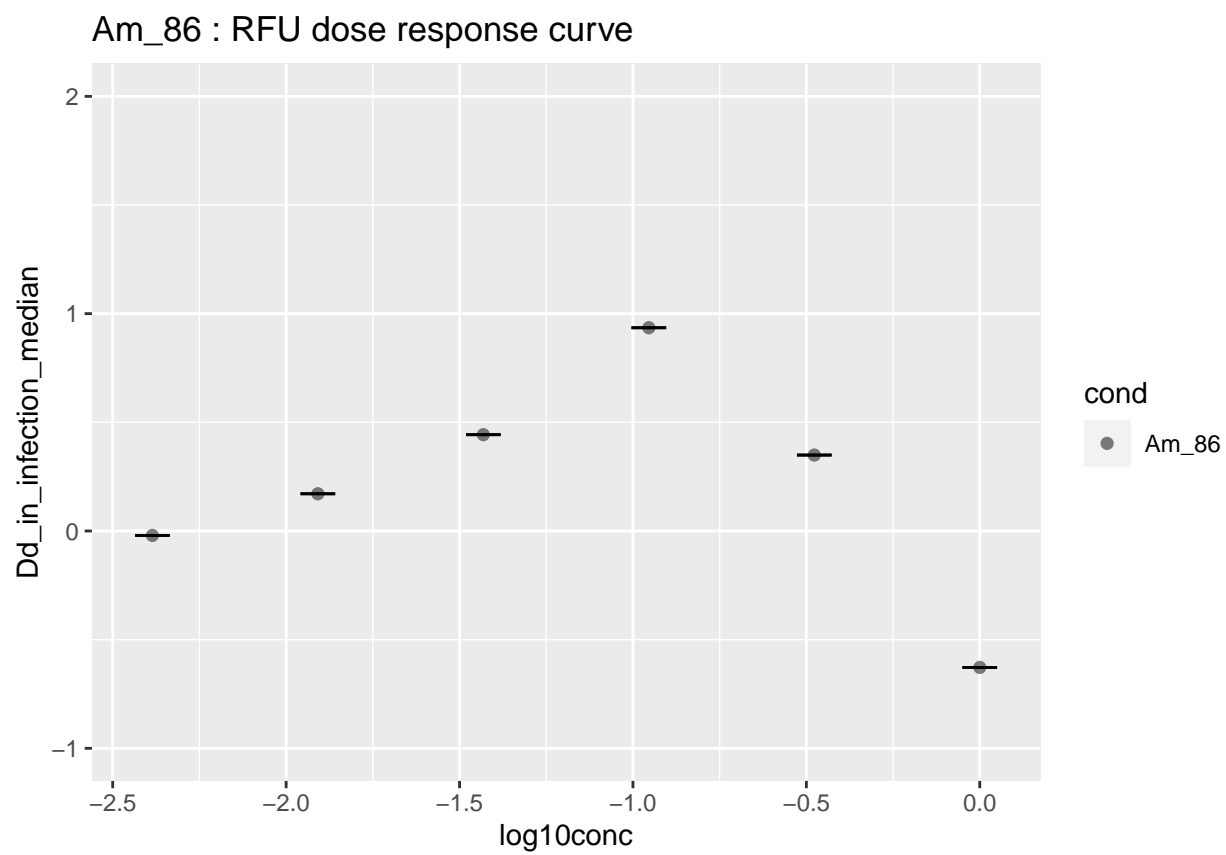


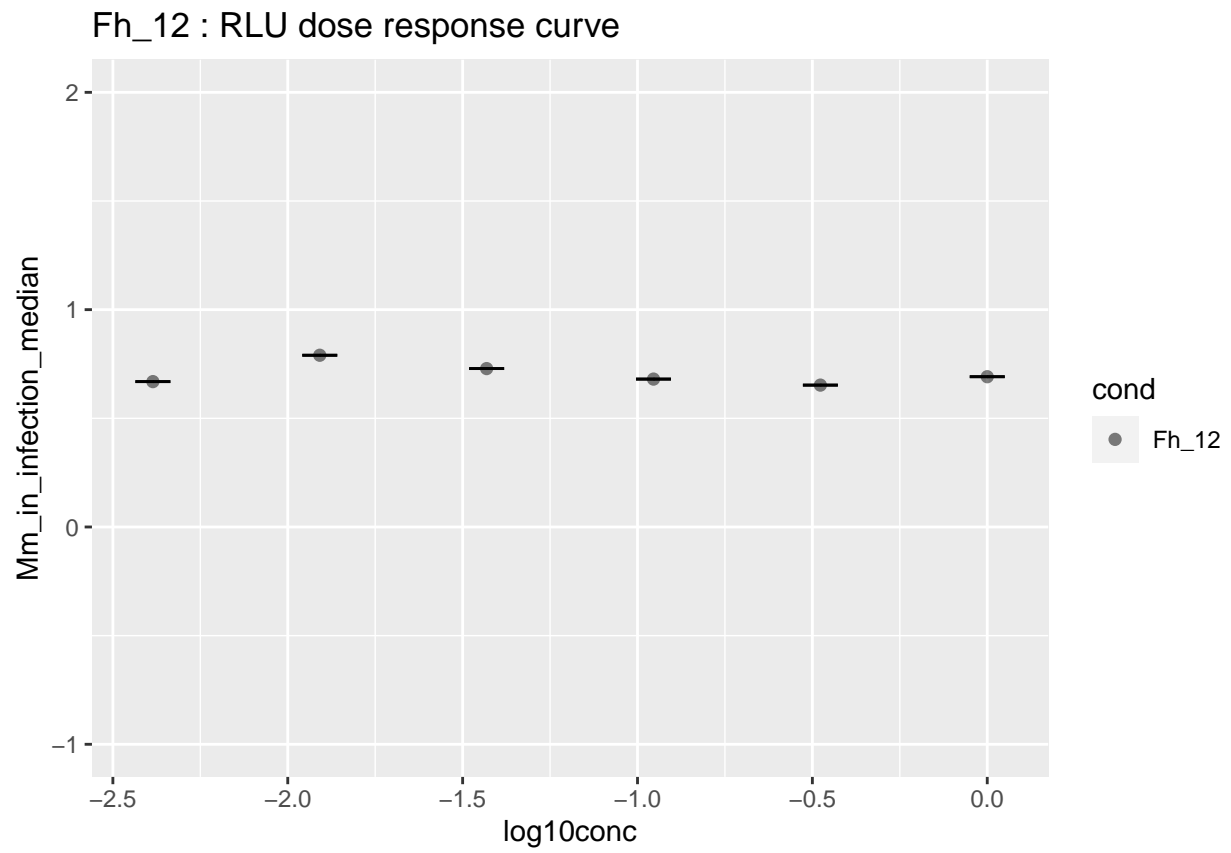


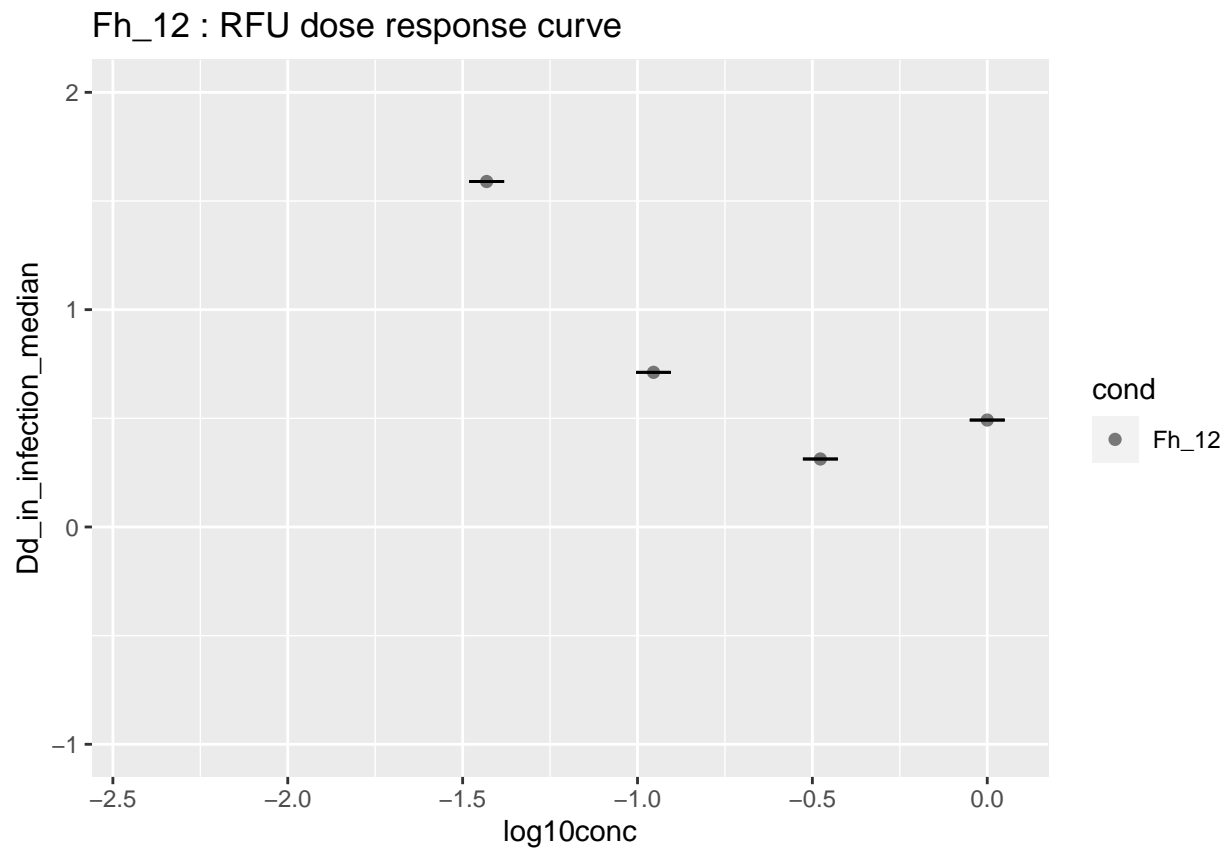


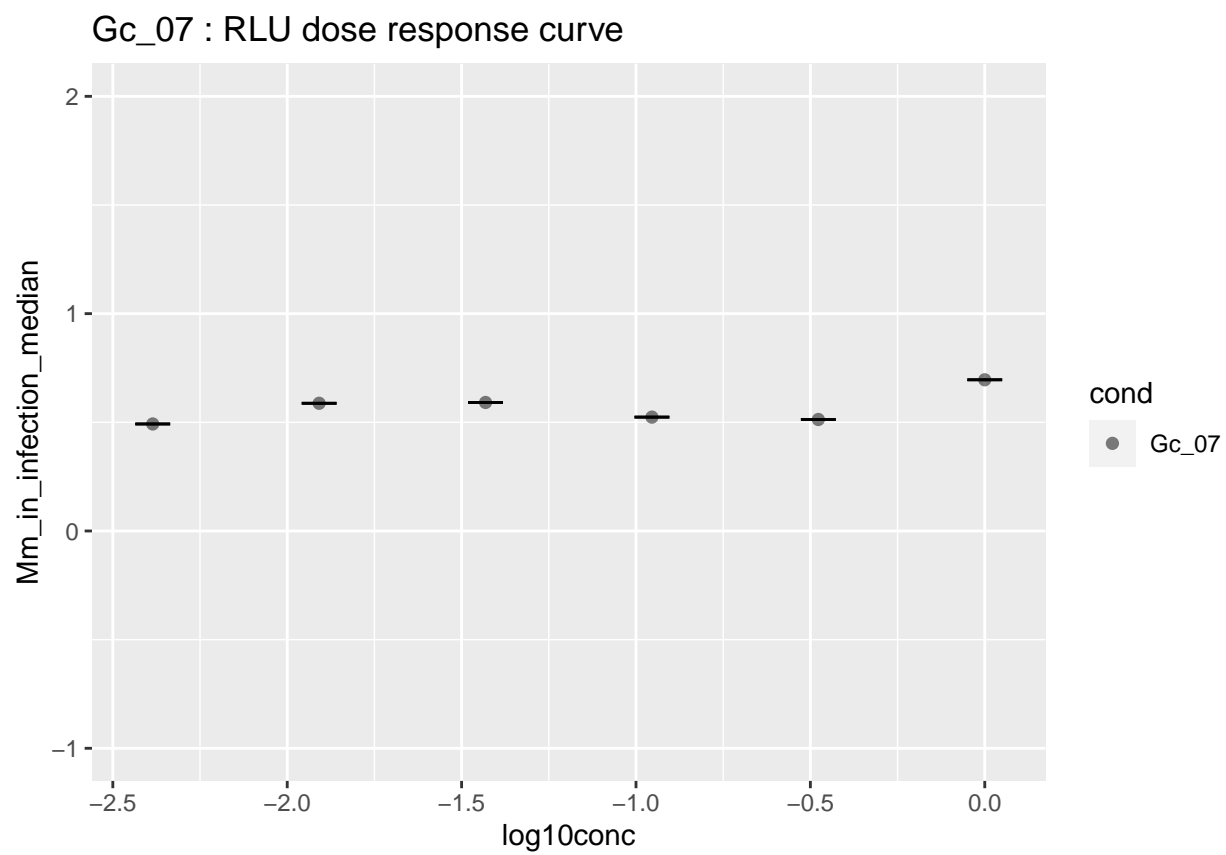


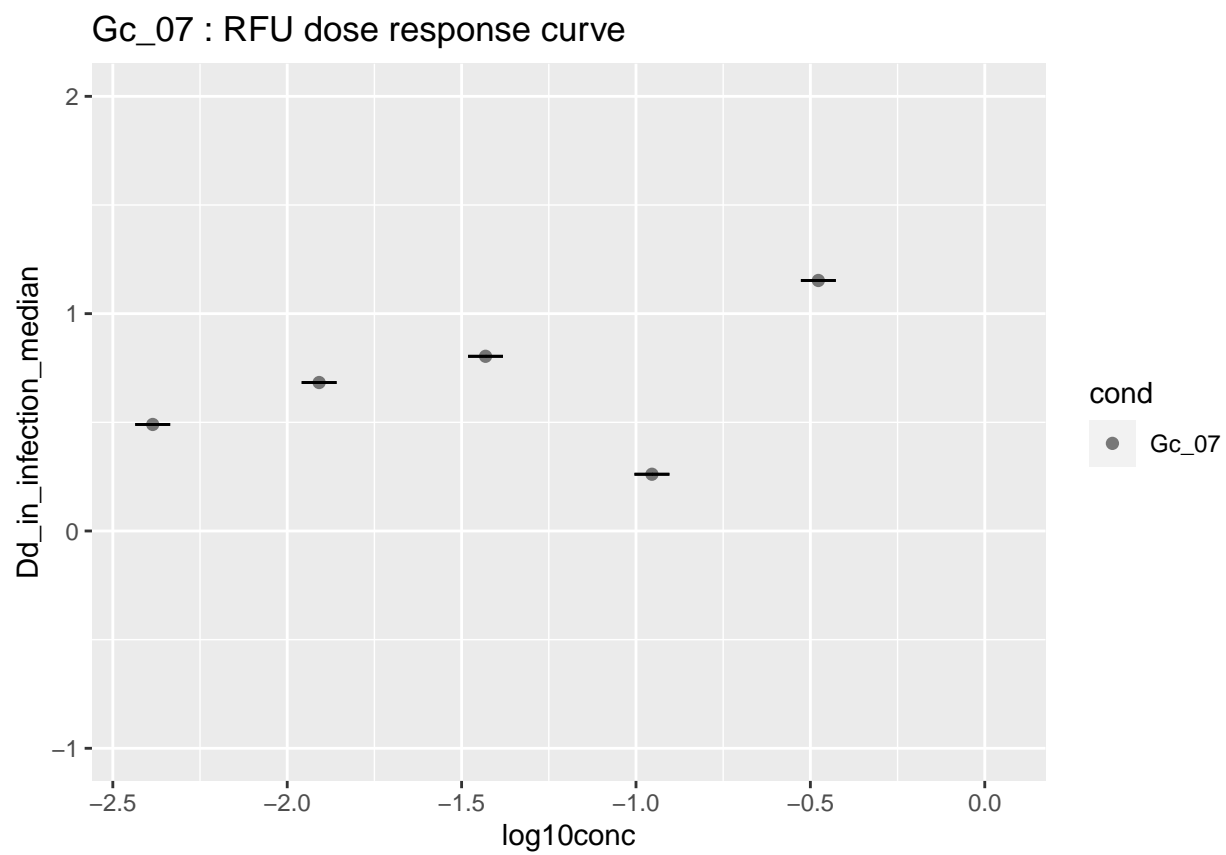


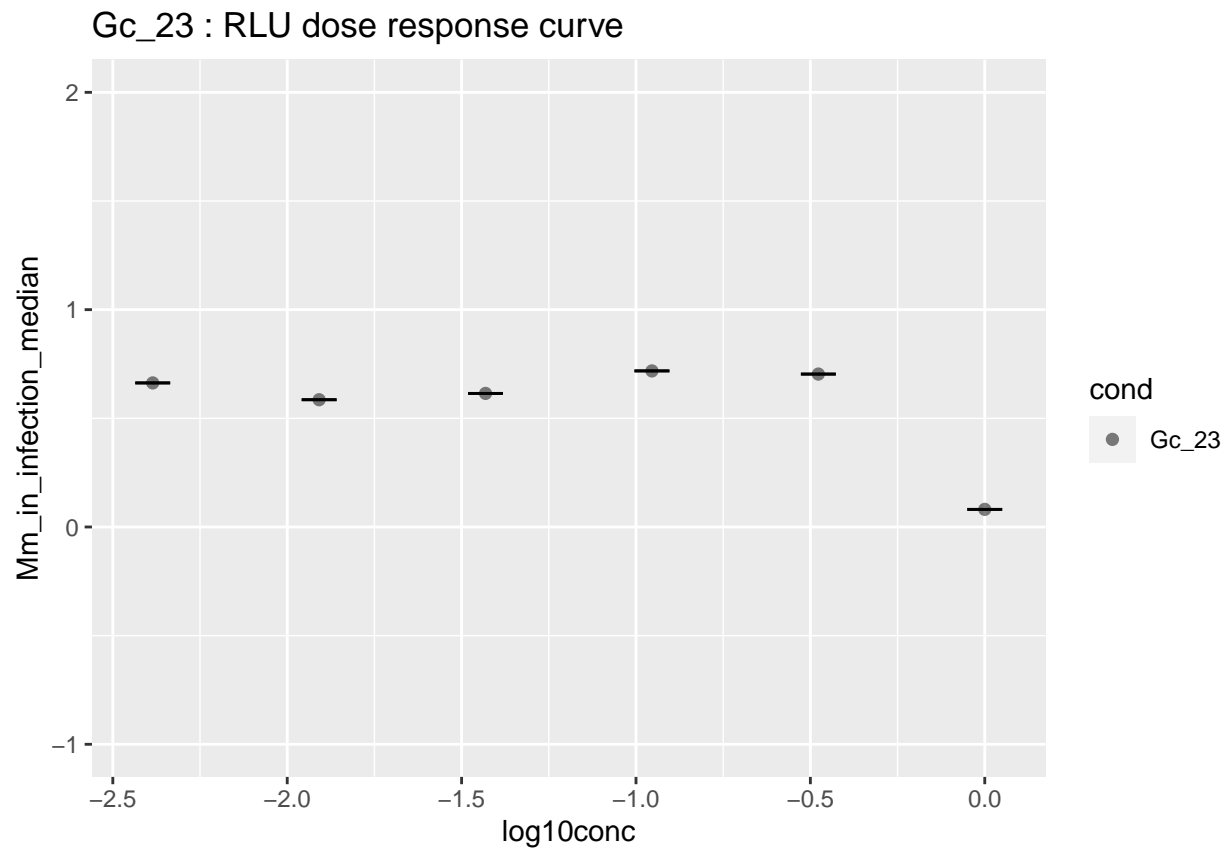


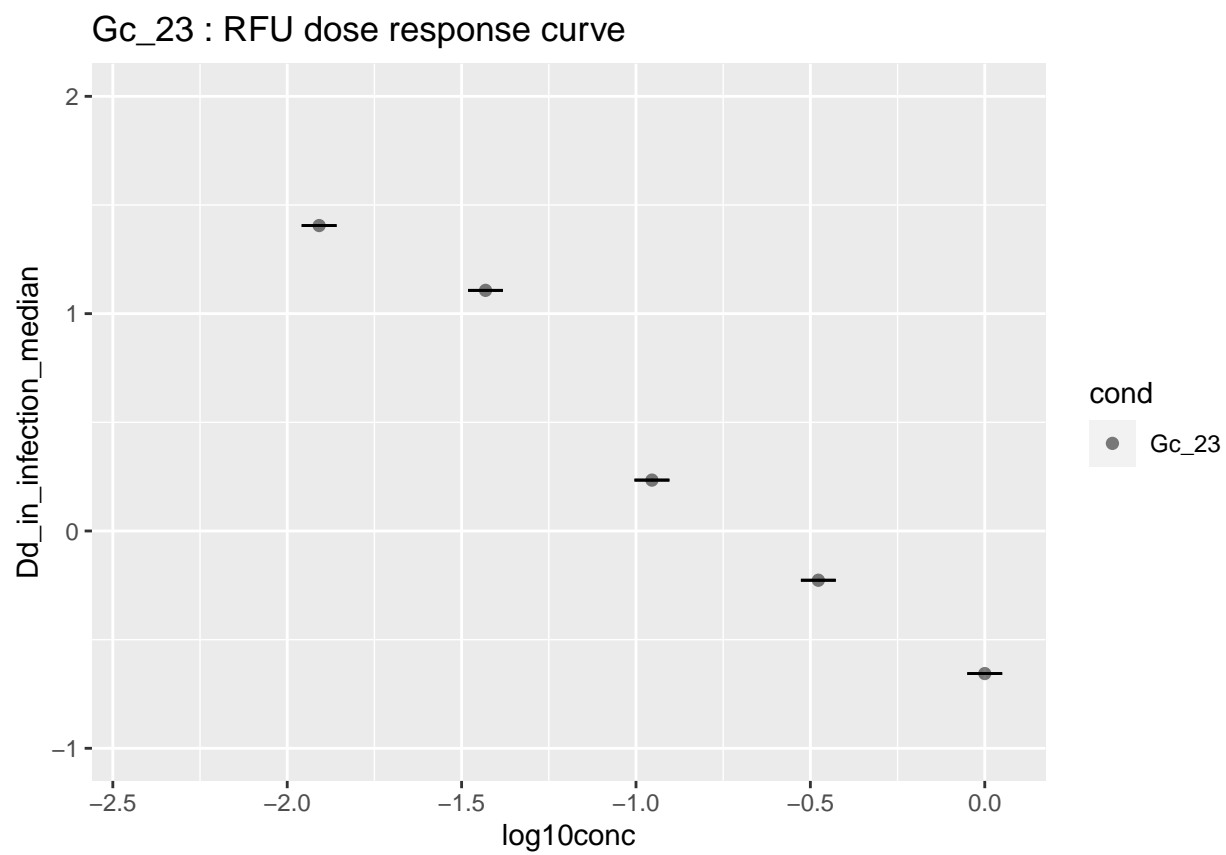


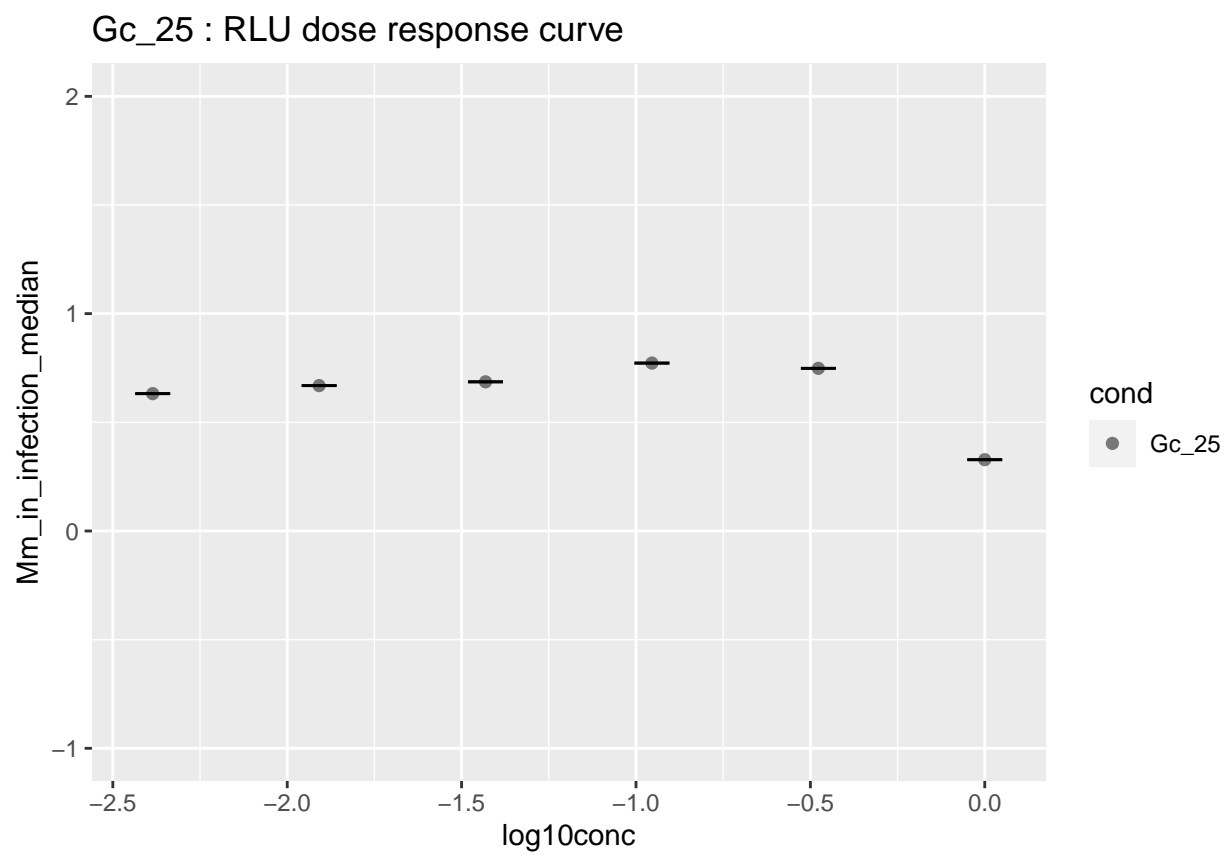


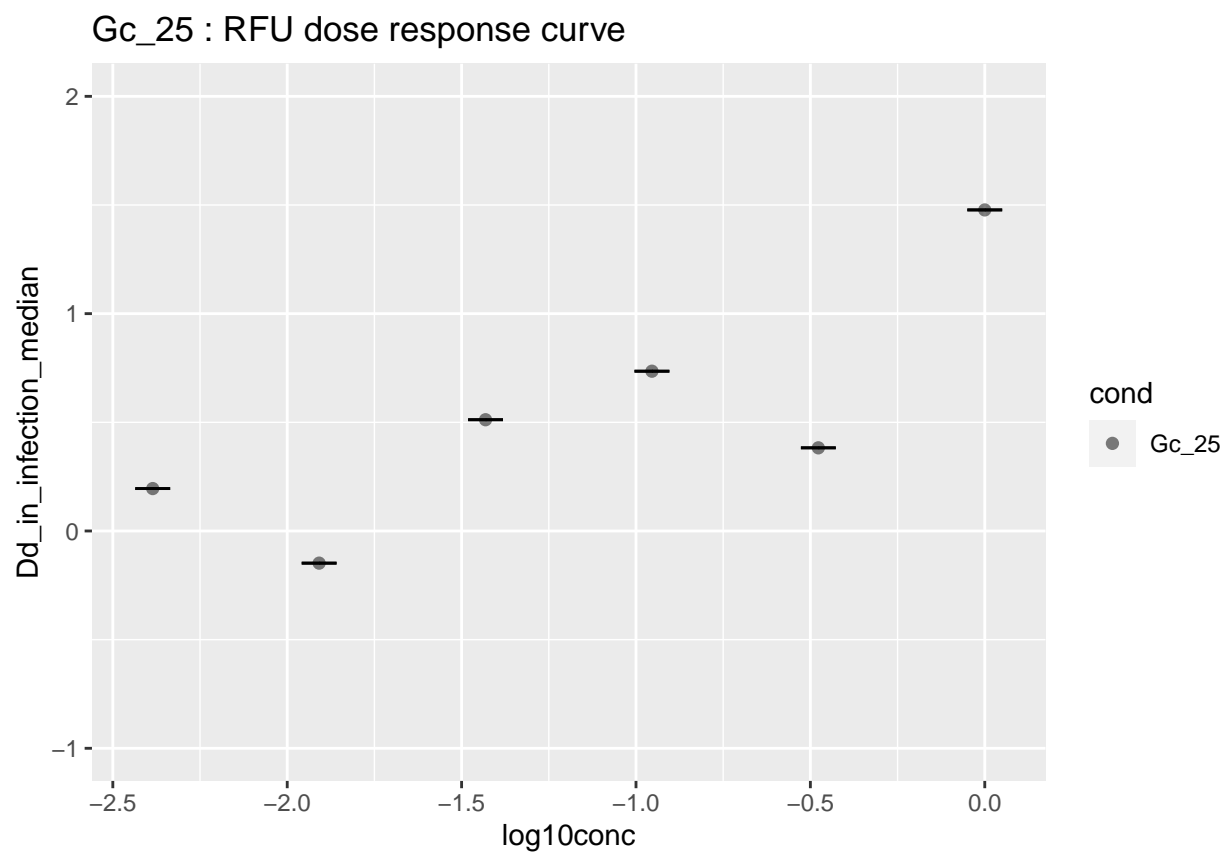


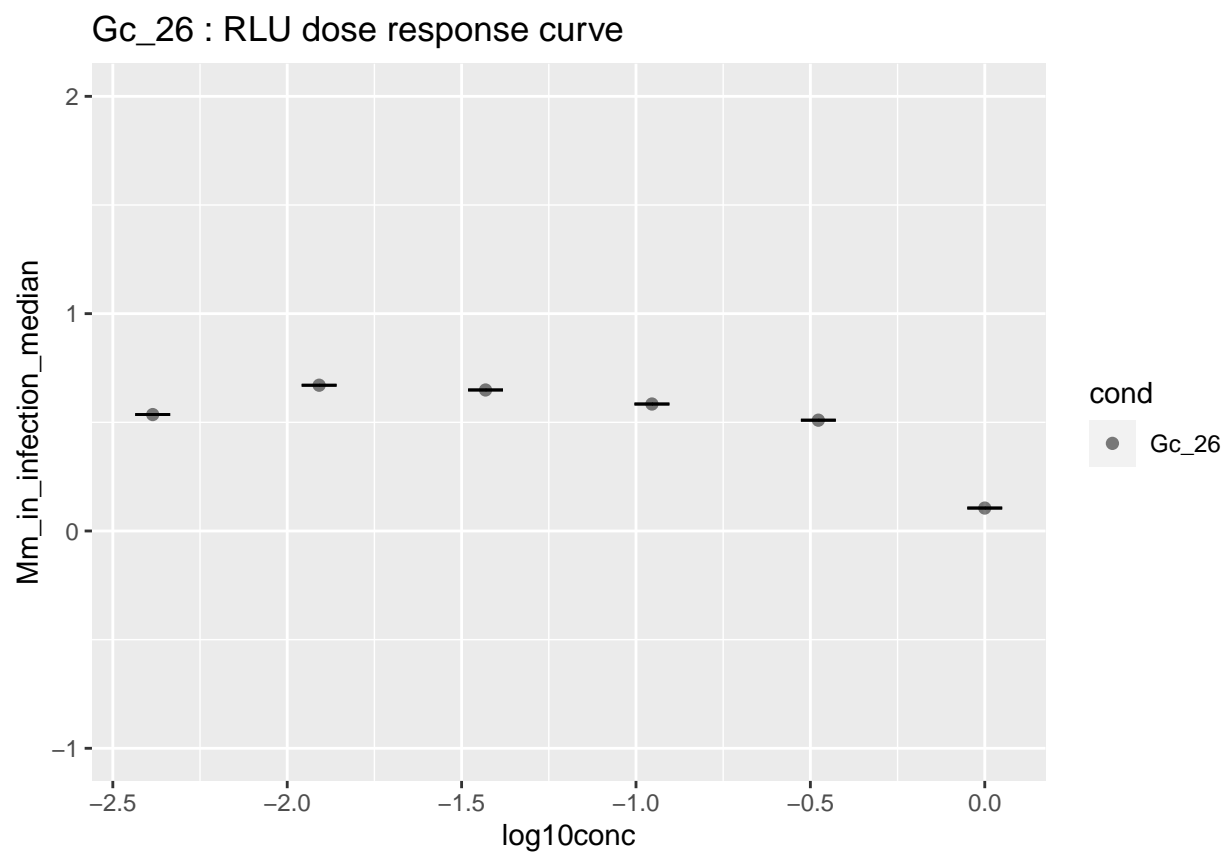


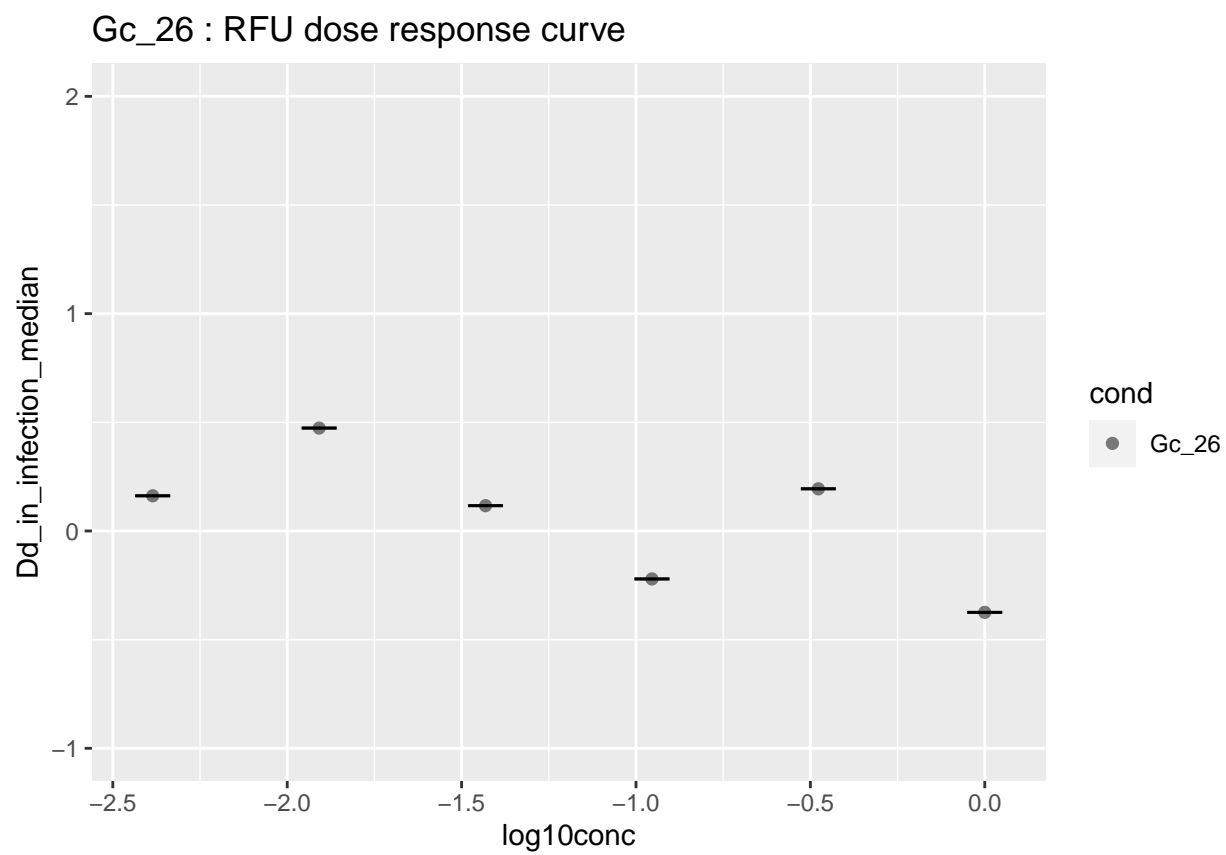


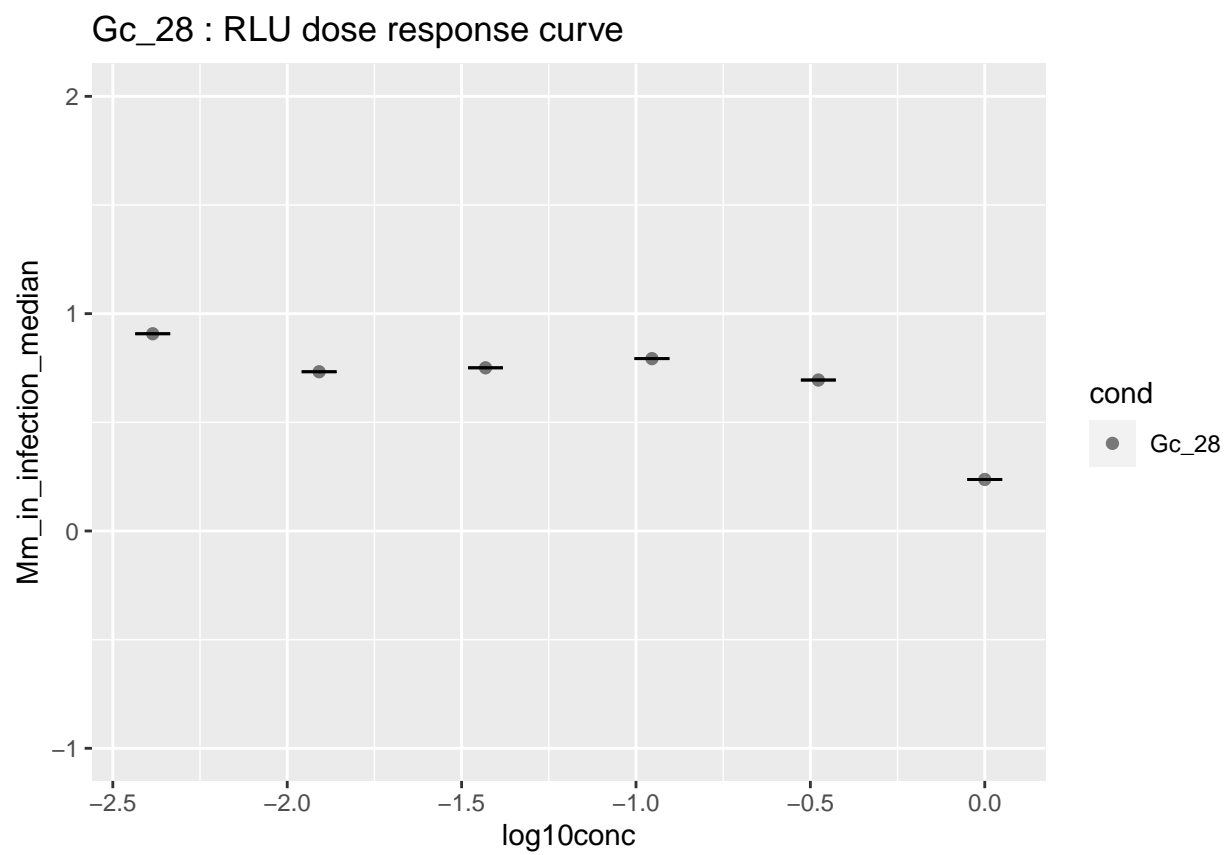


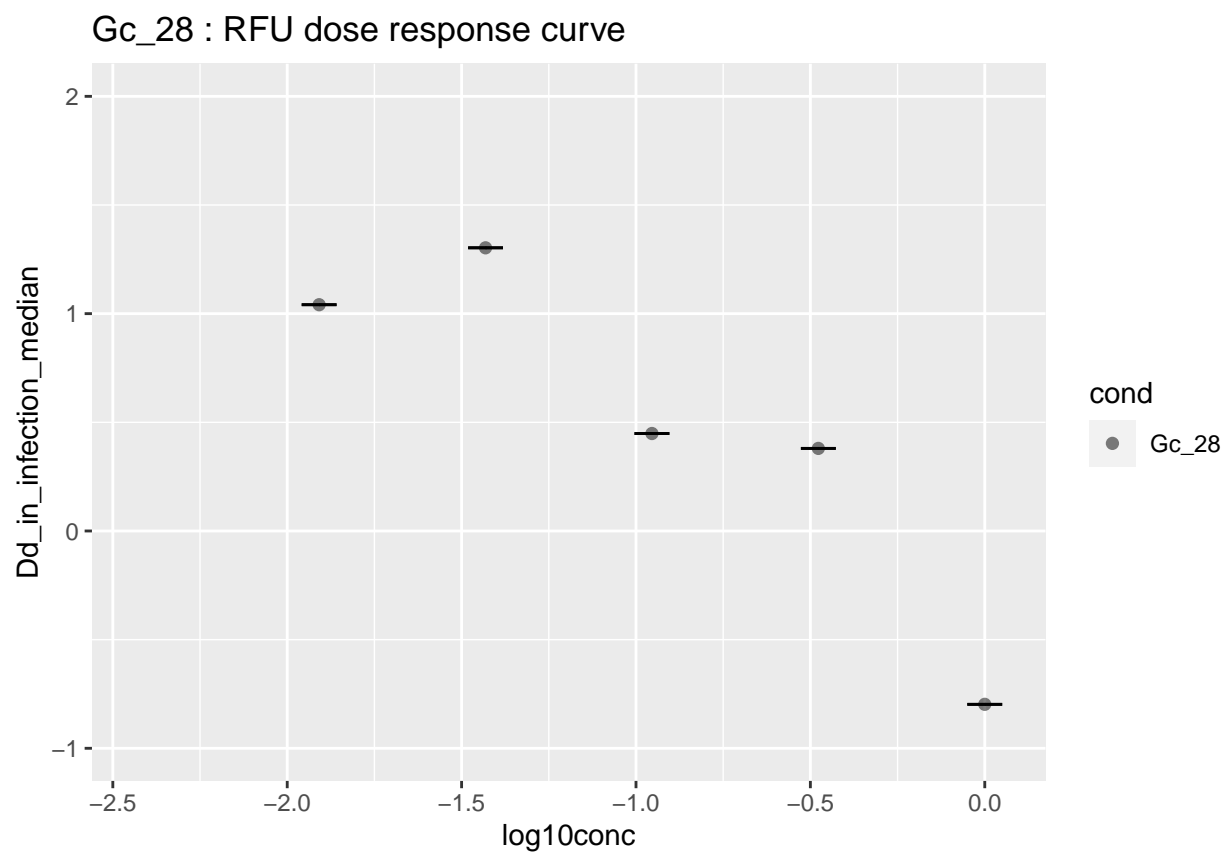


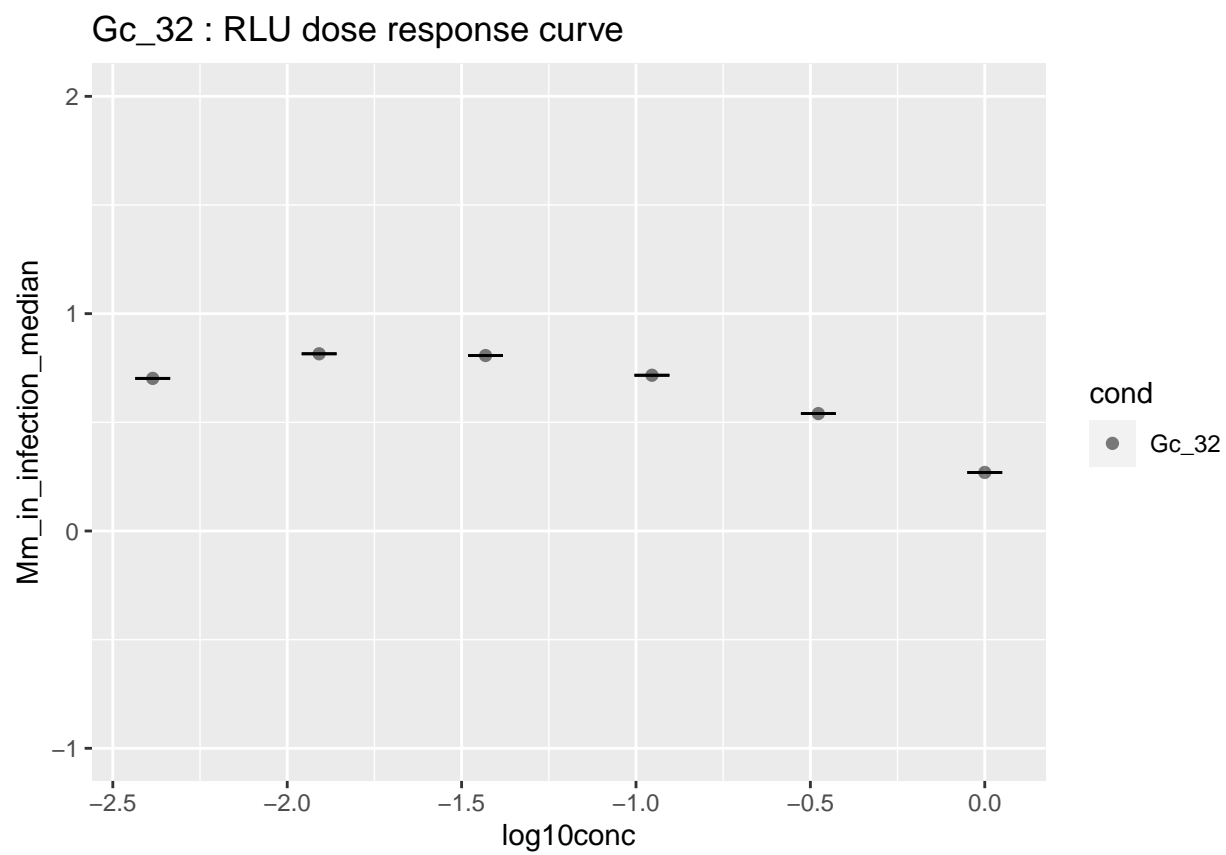




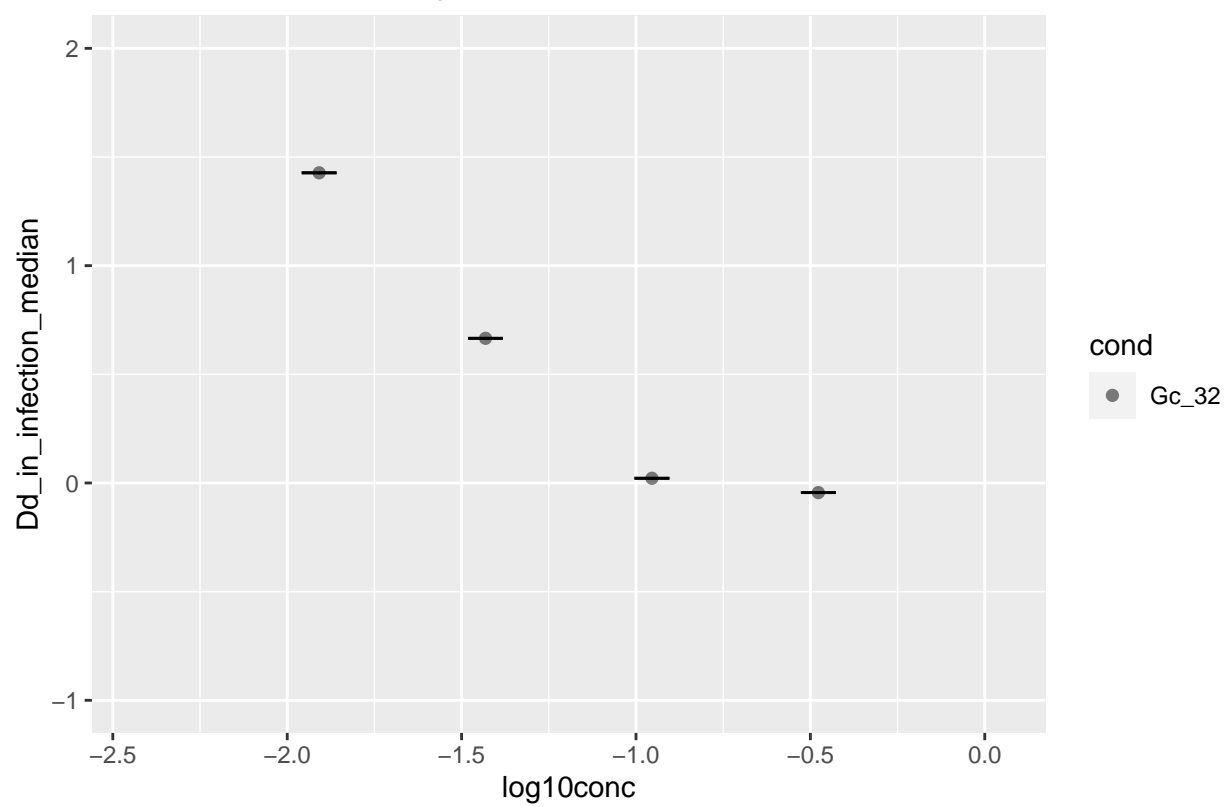


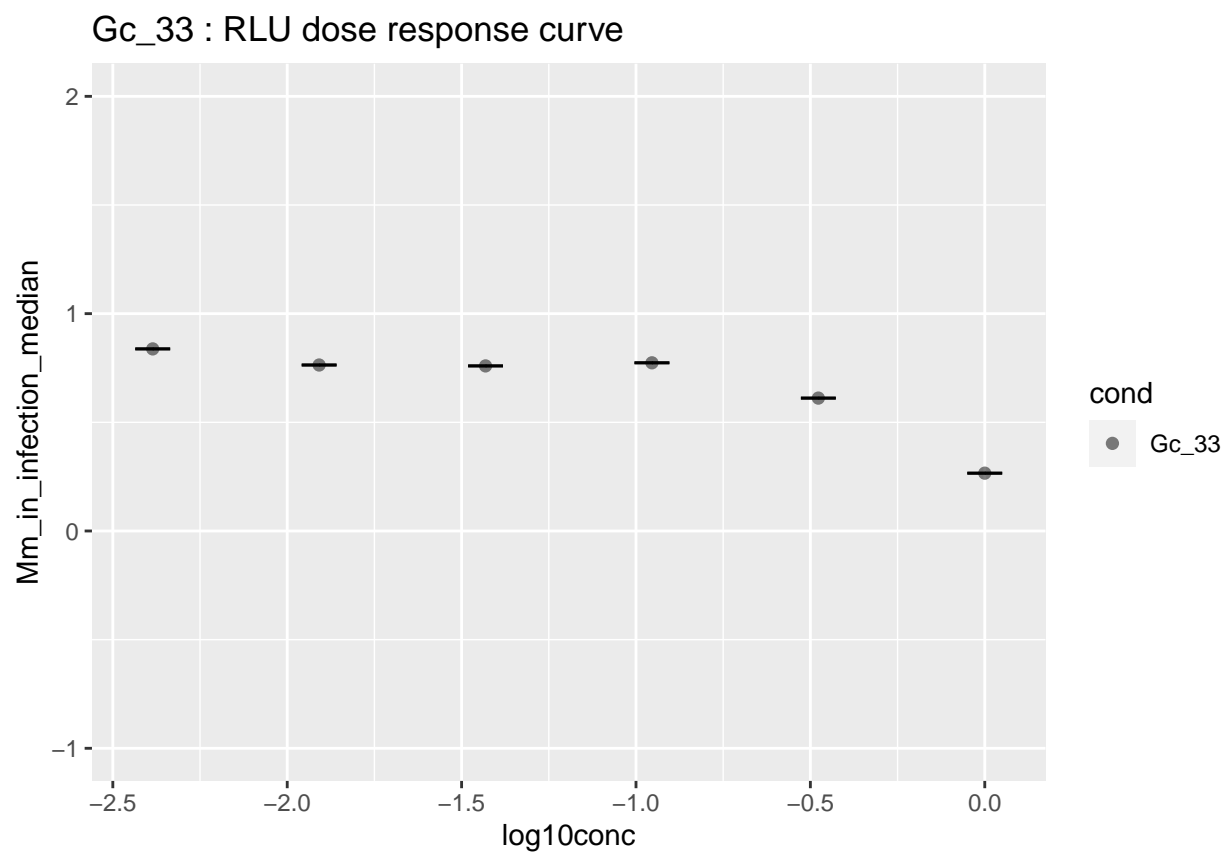


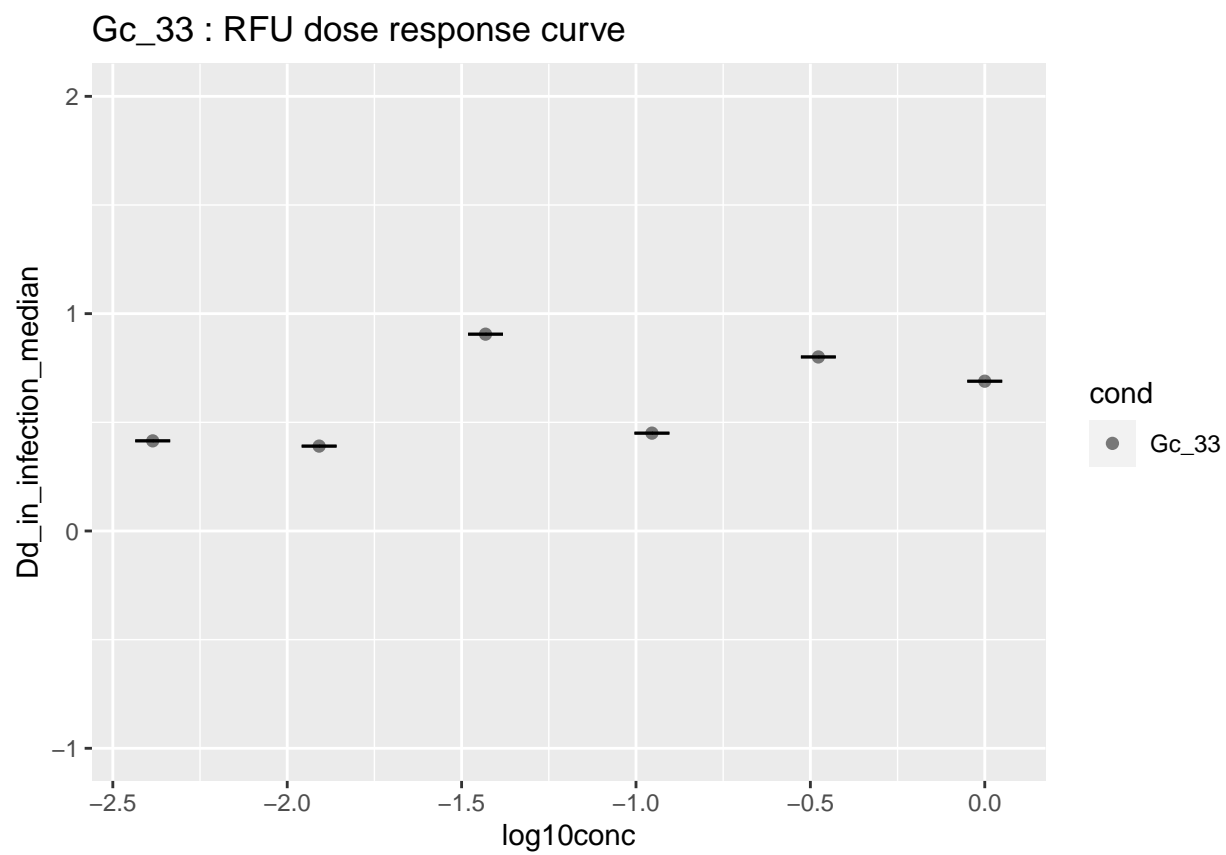


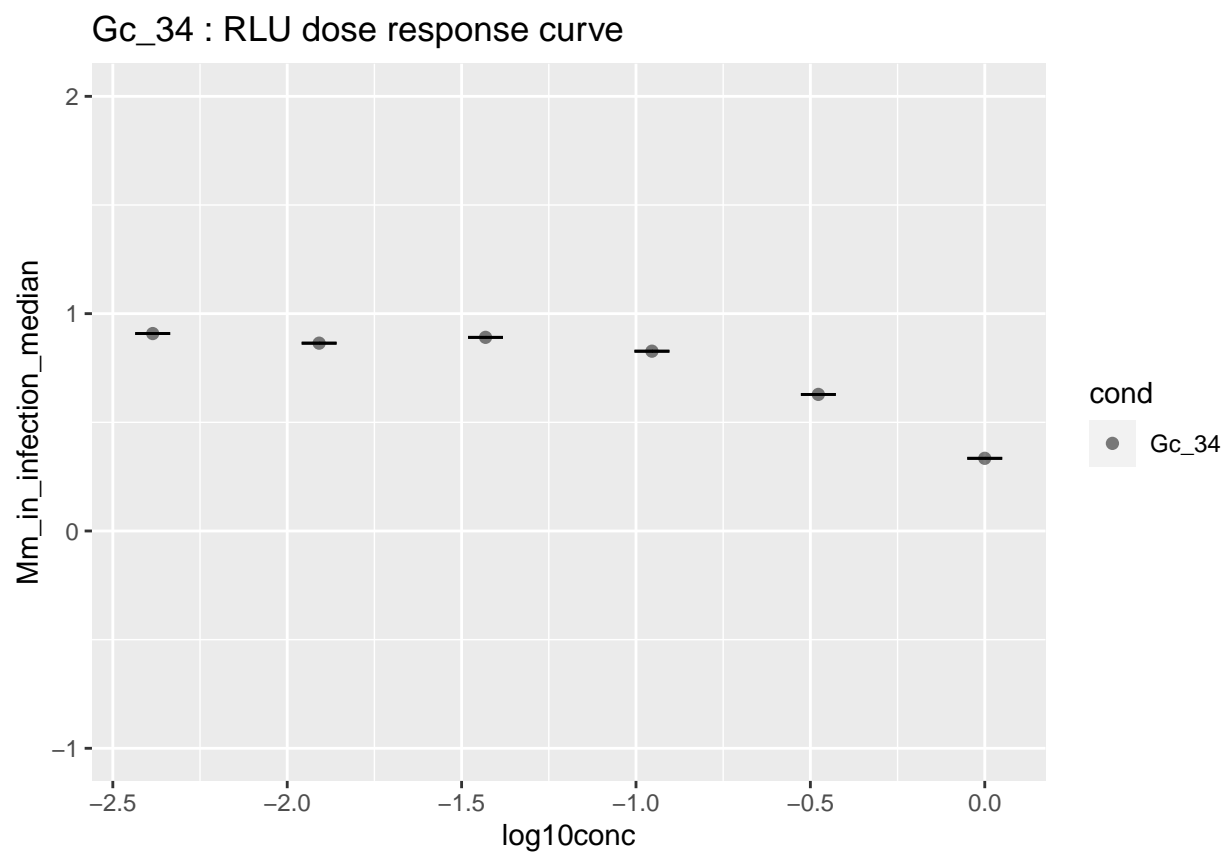


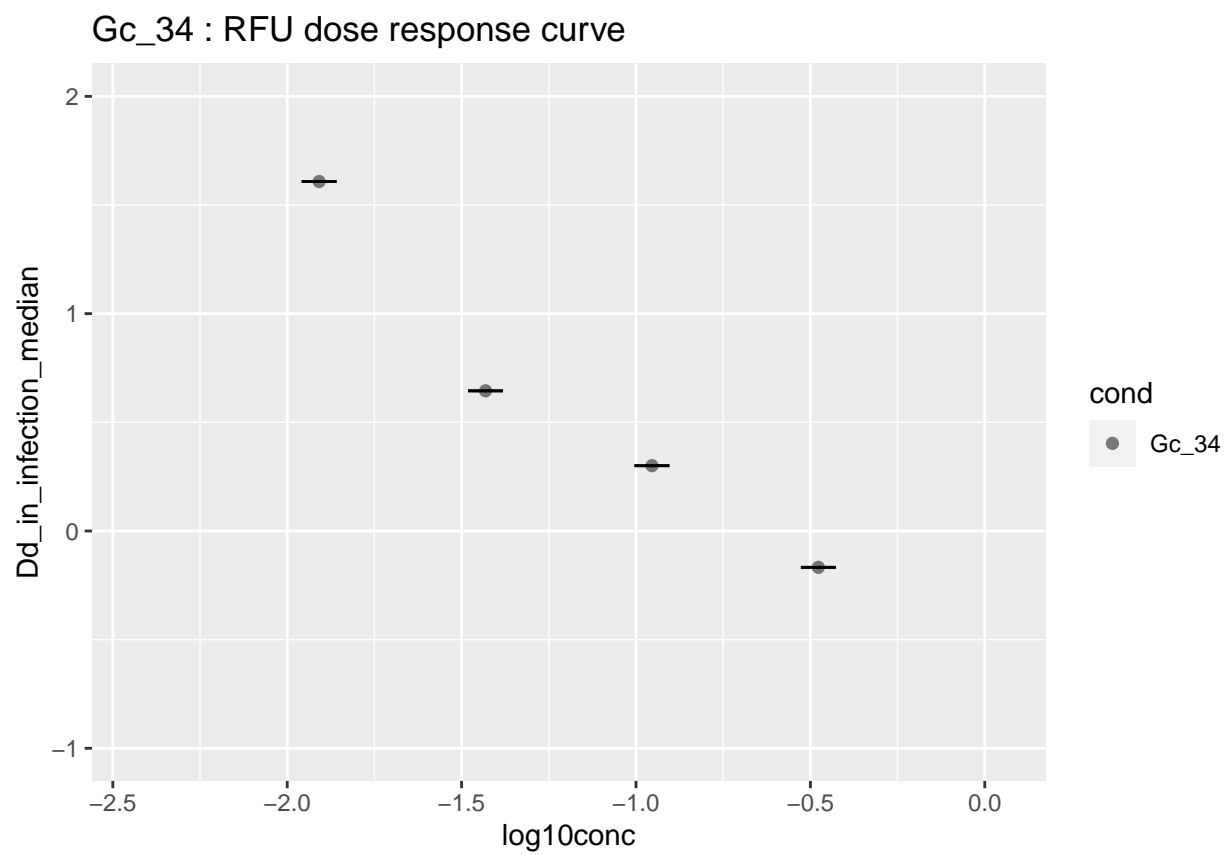
Gc_32 : RFU dose response curve

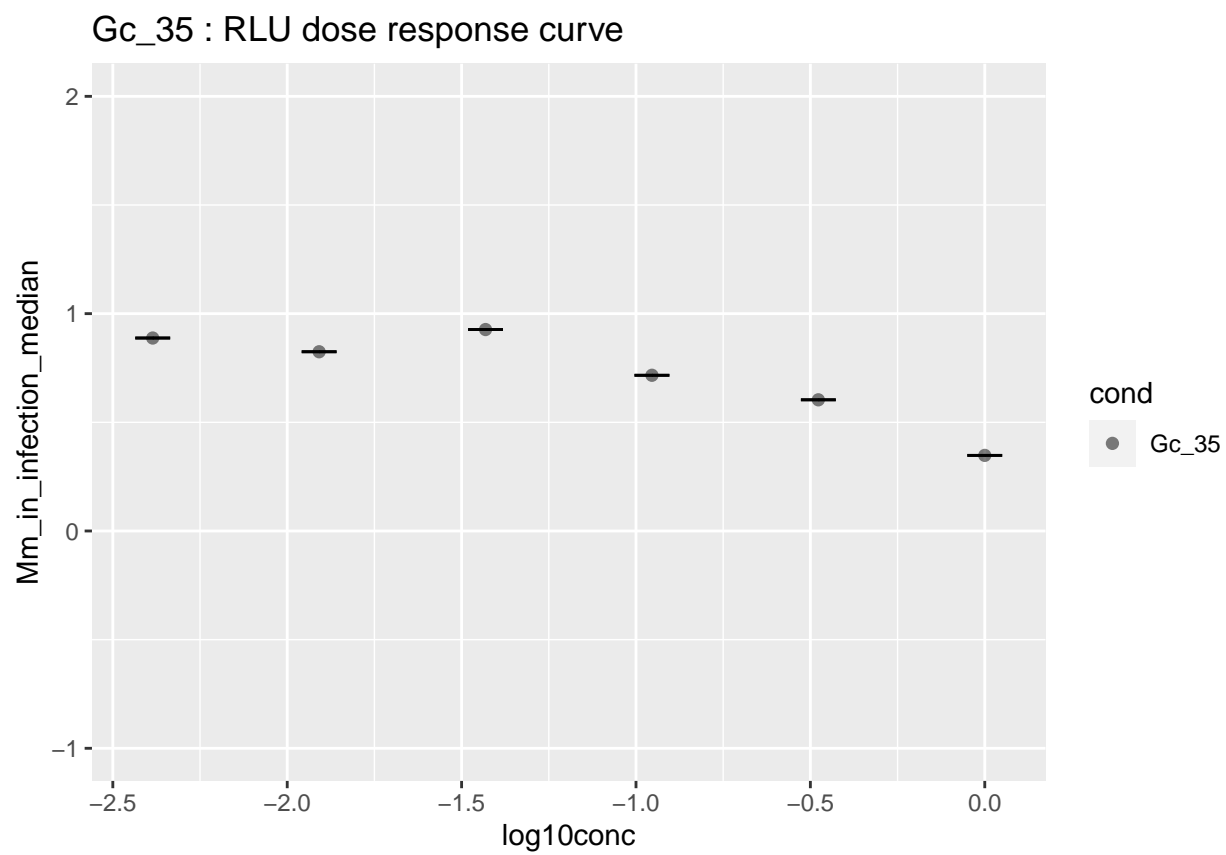


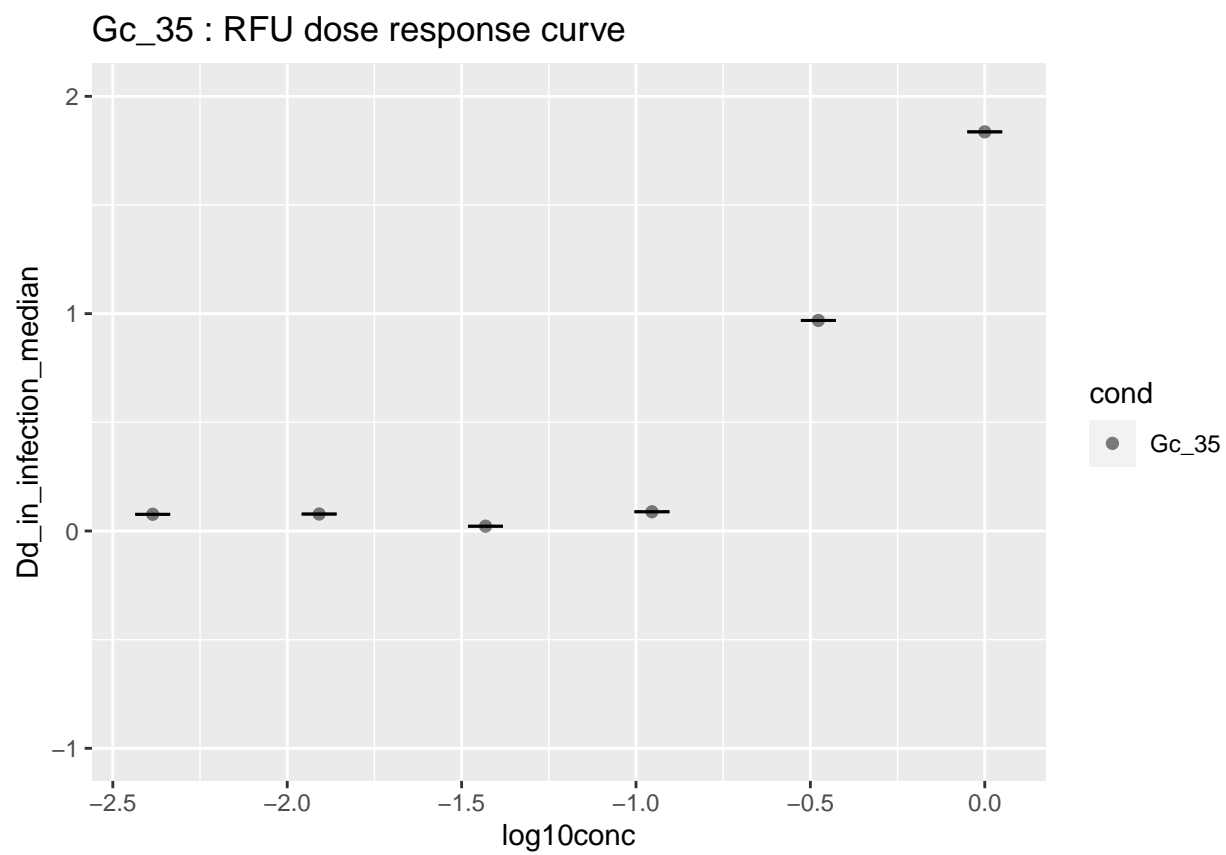


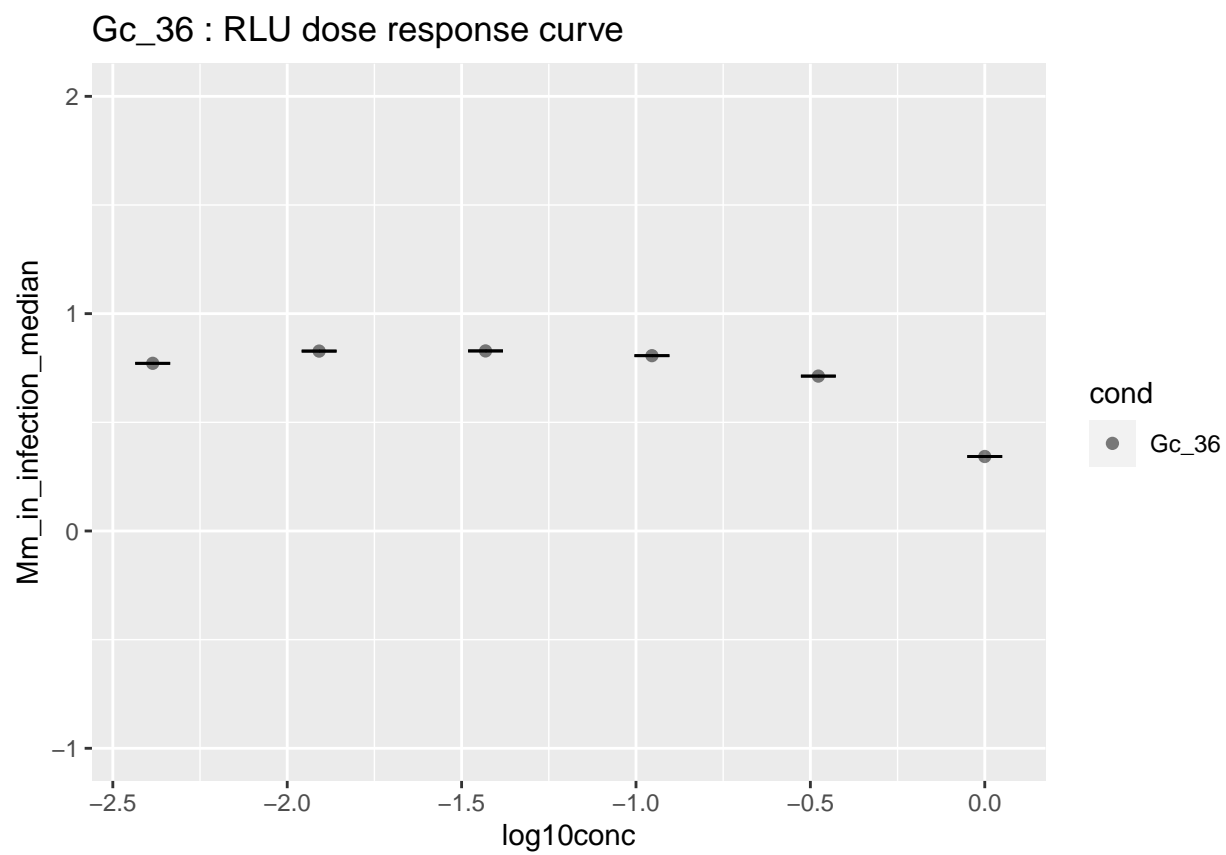


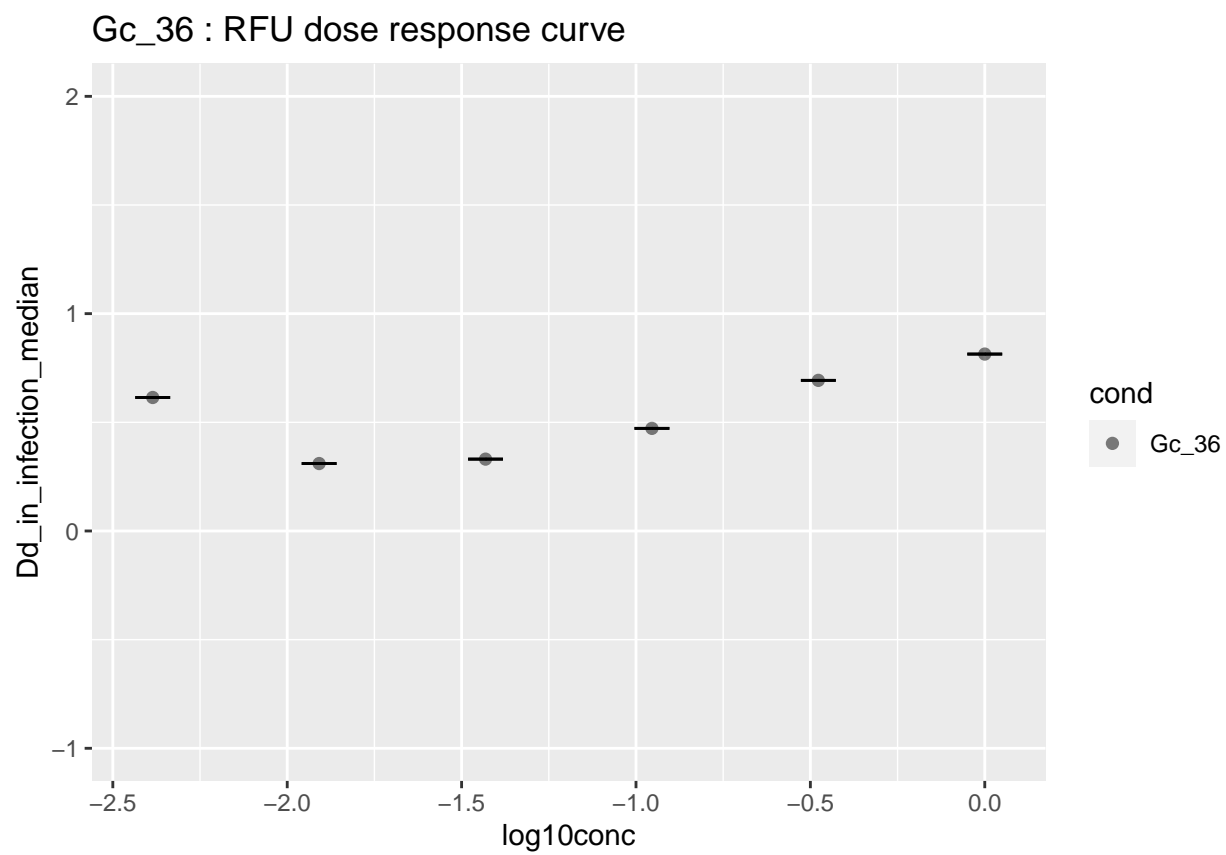


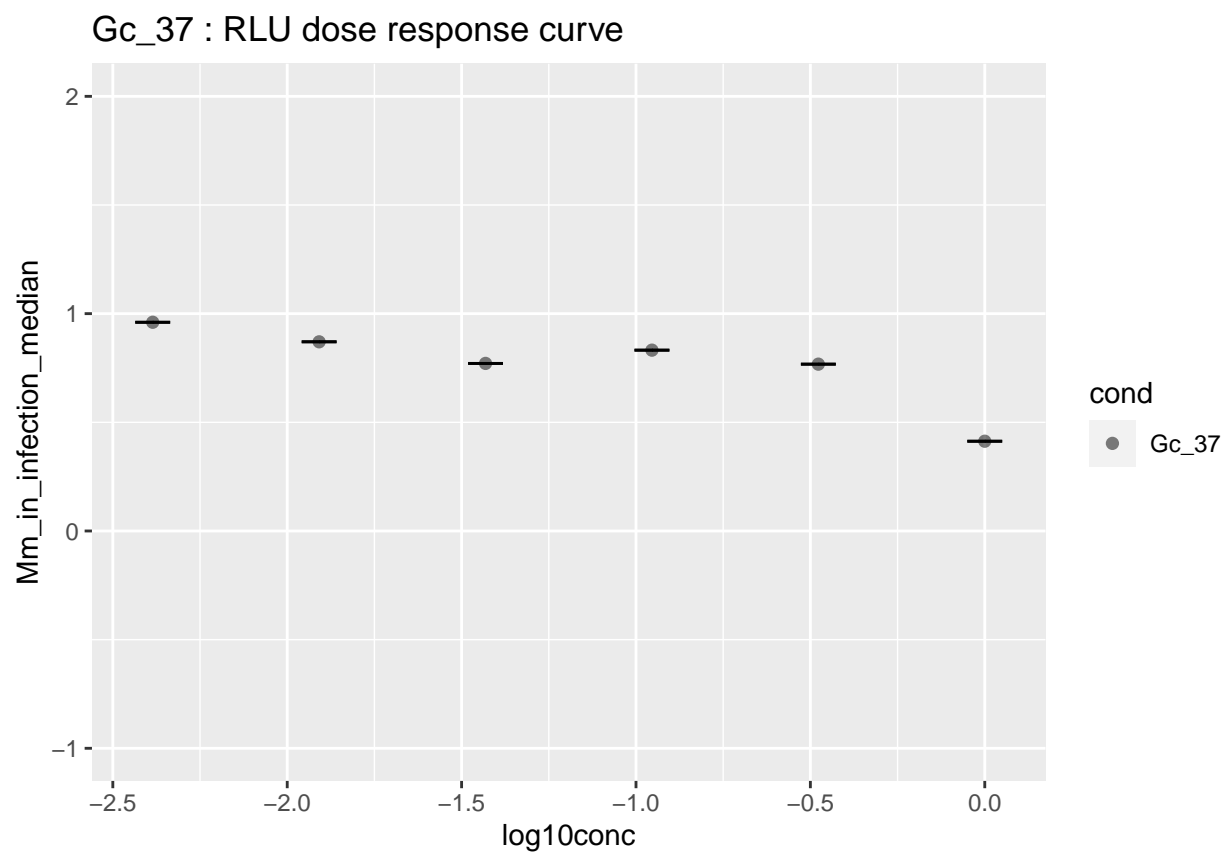


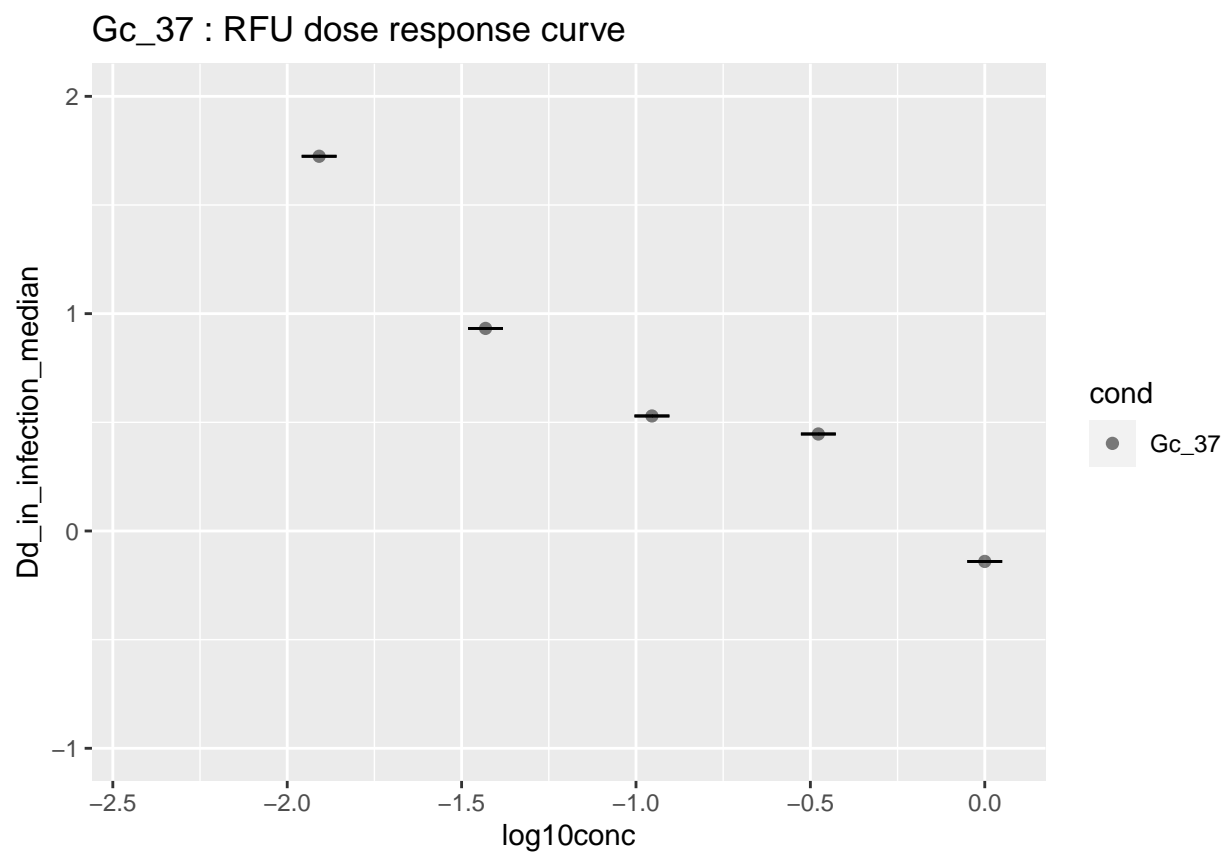


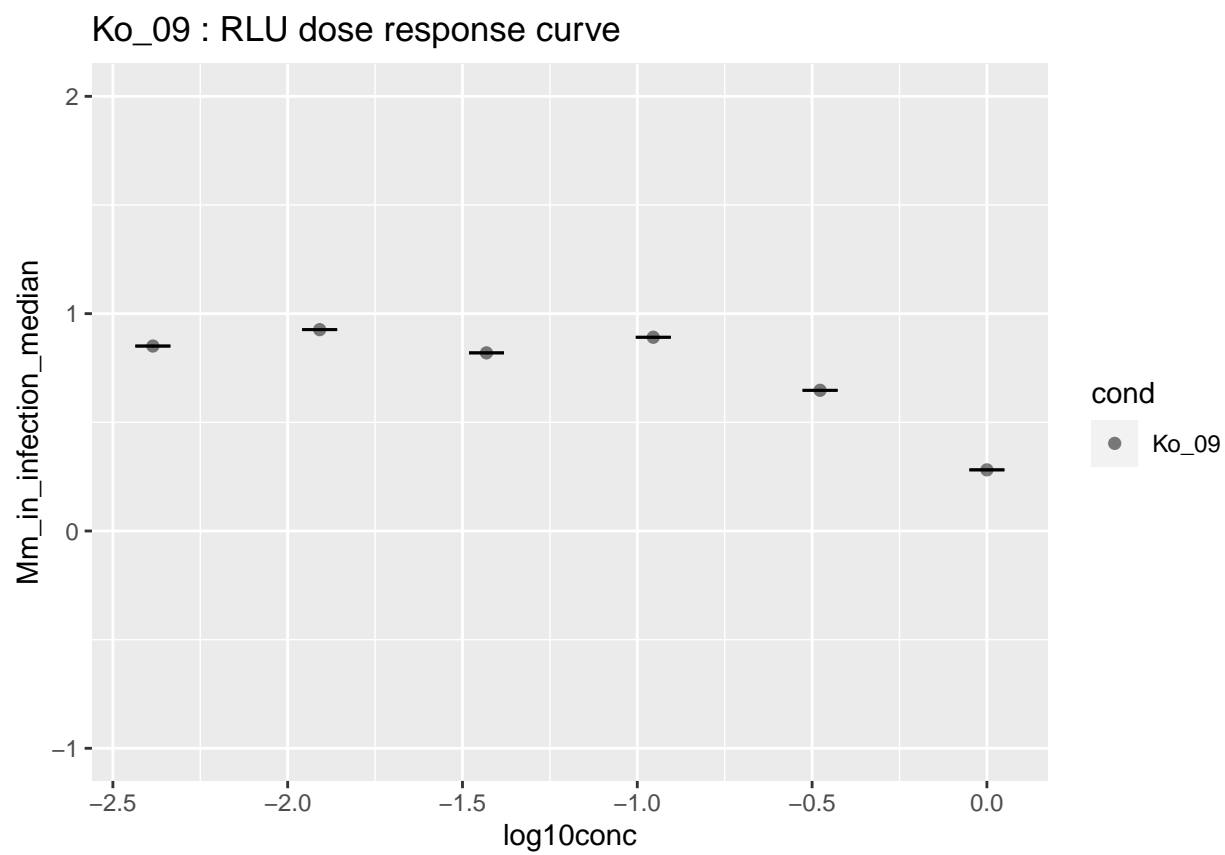


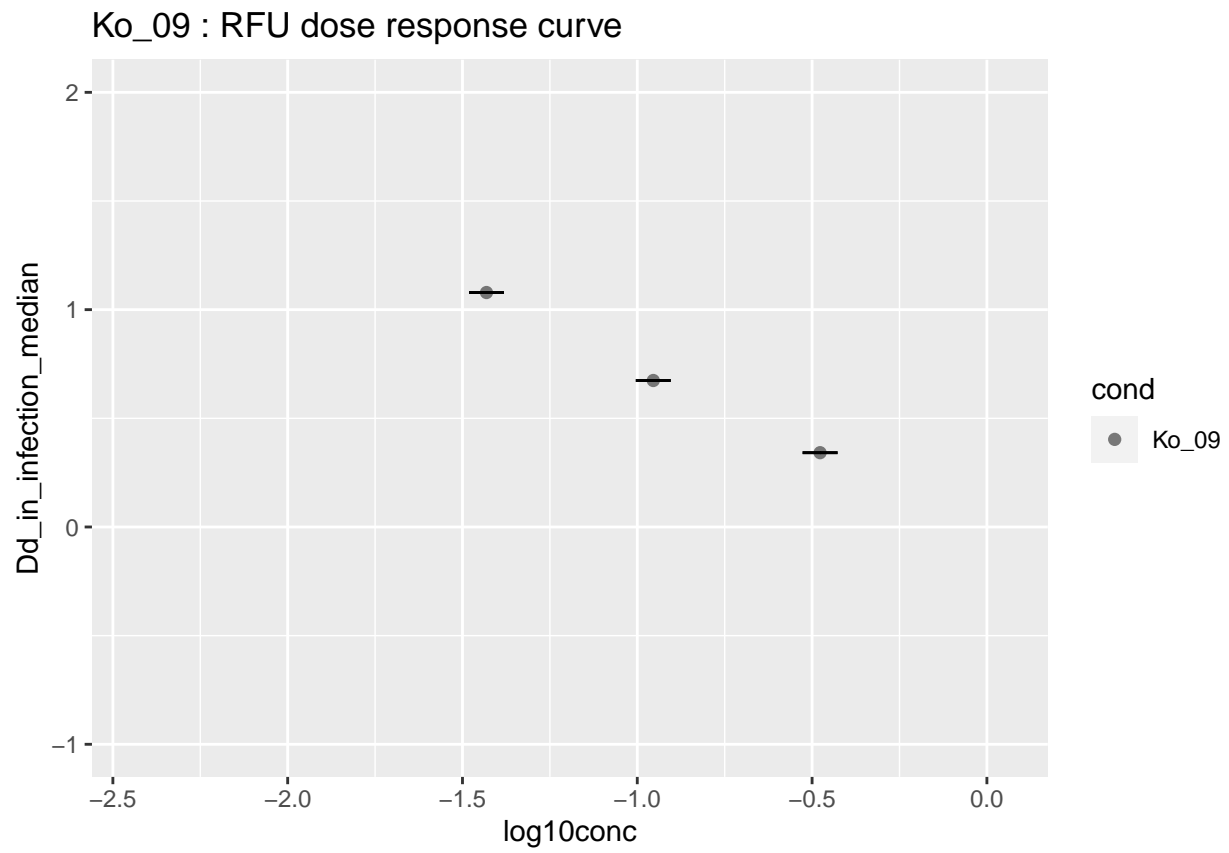


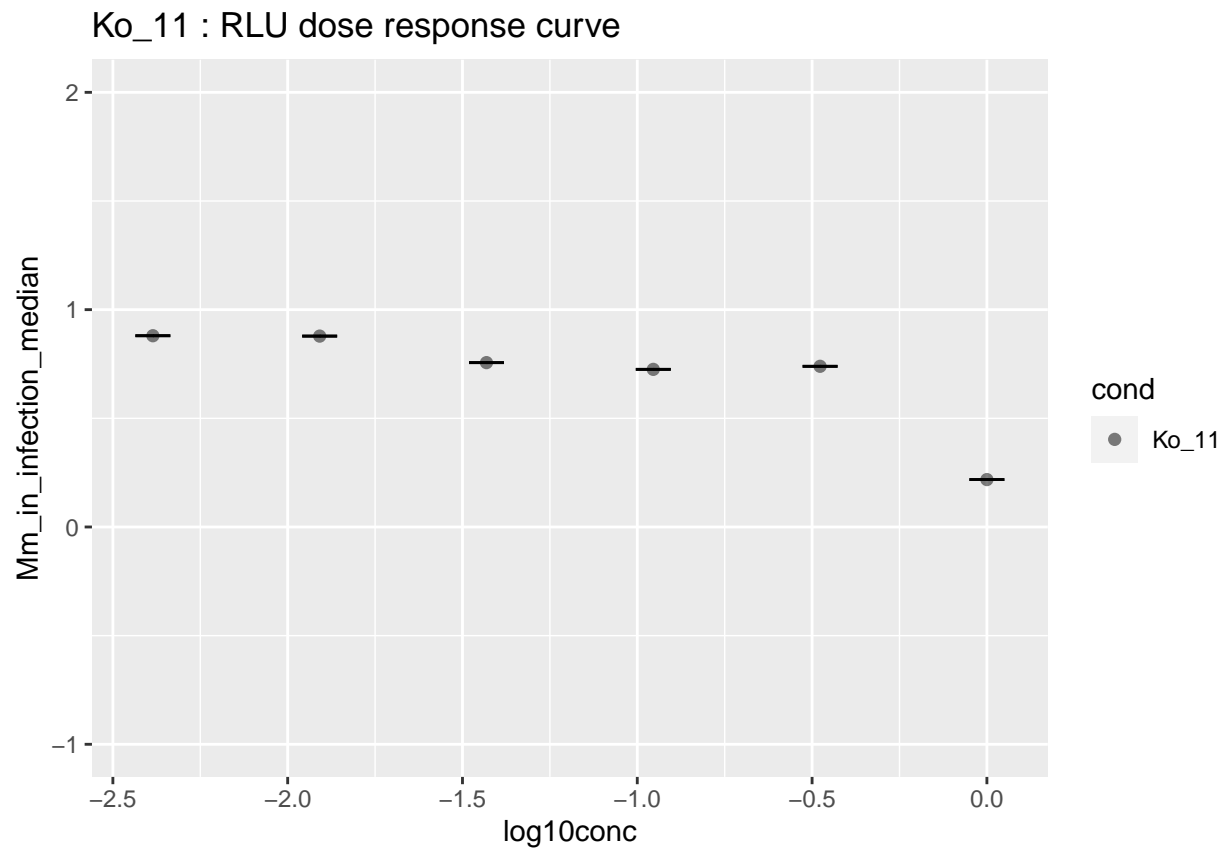


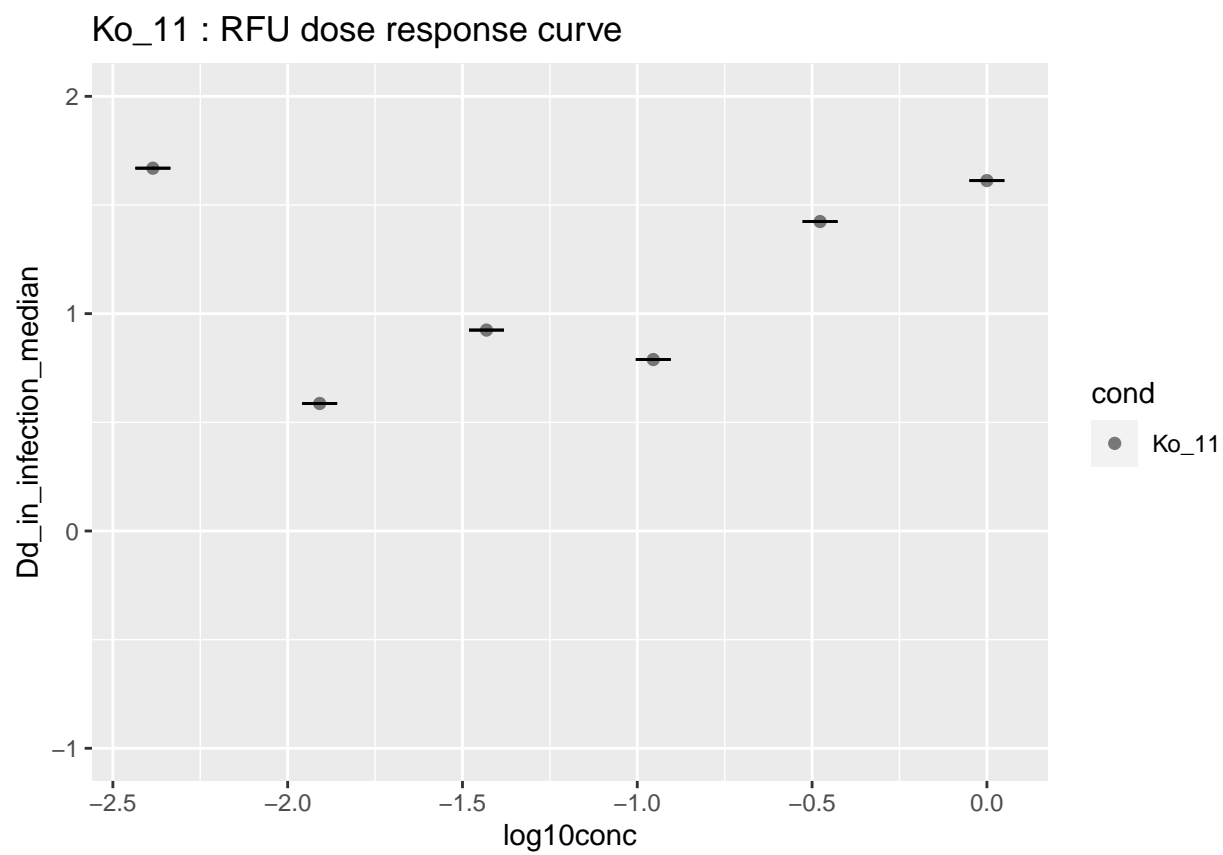


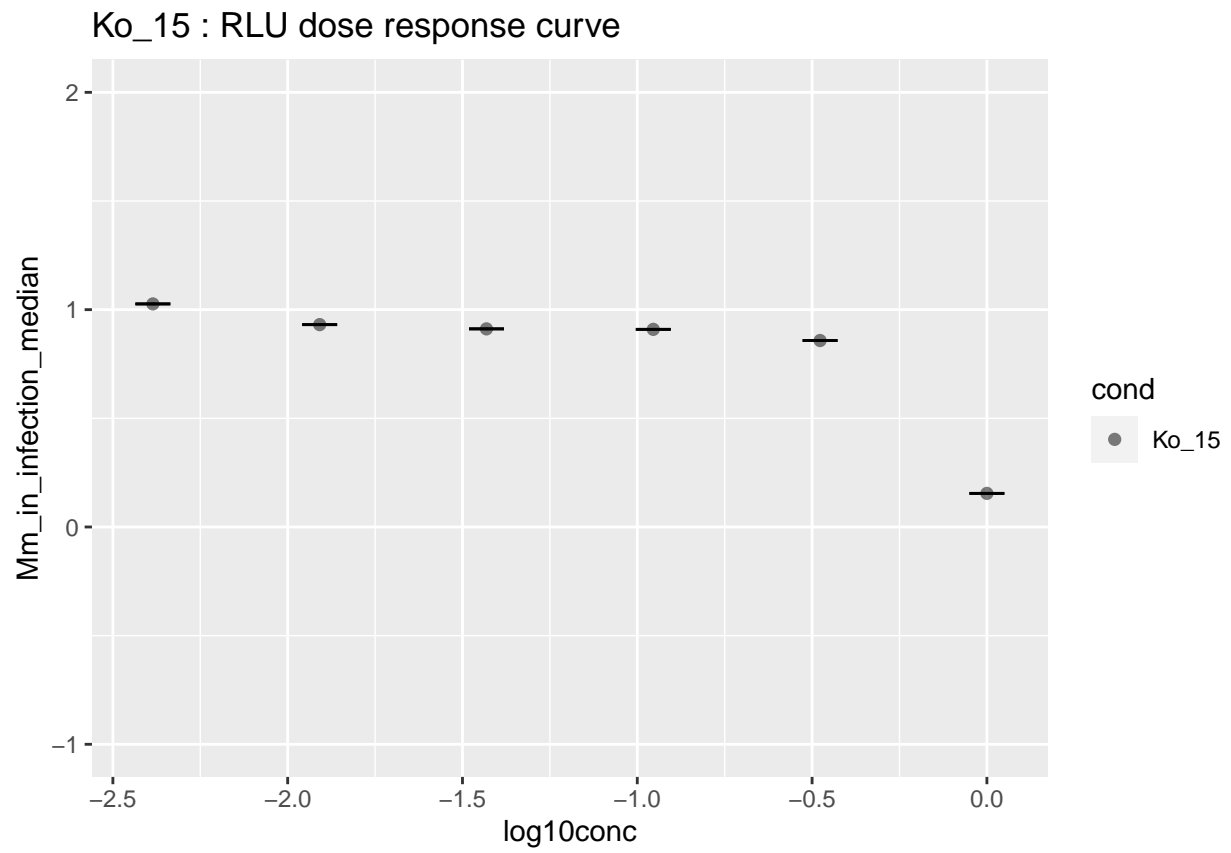


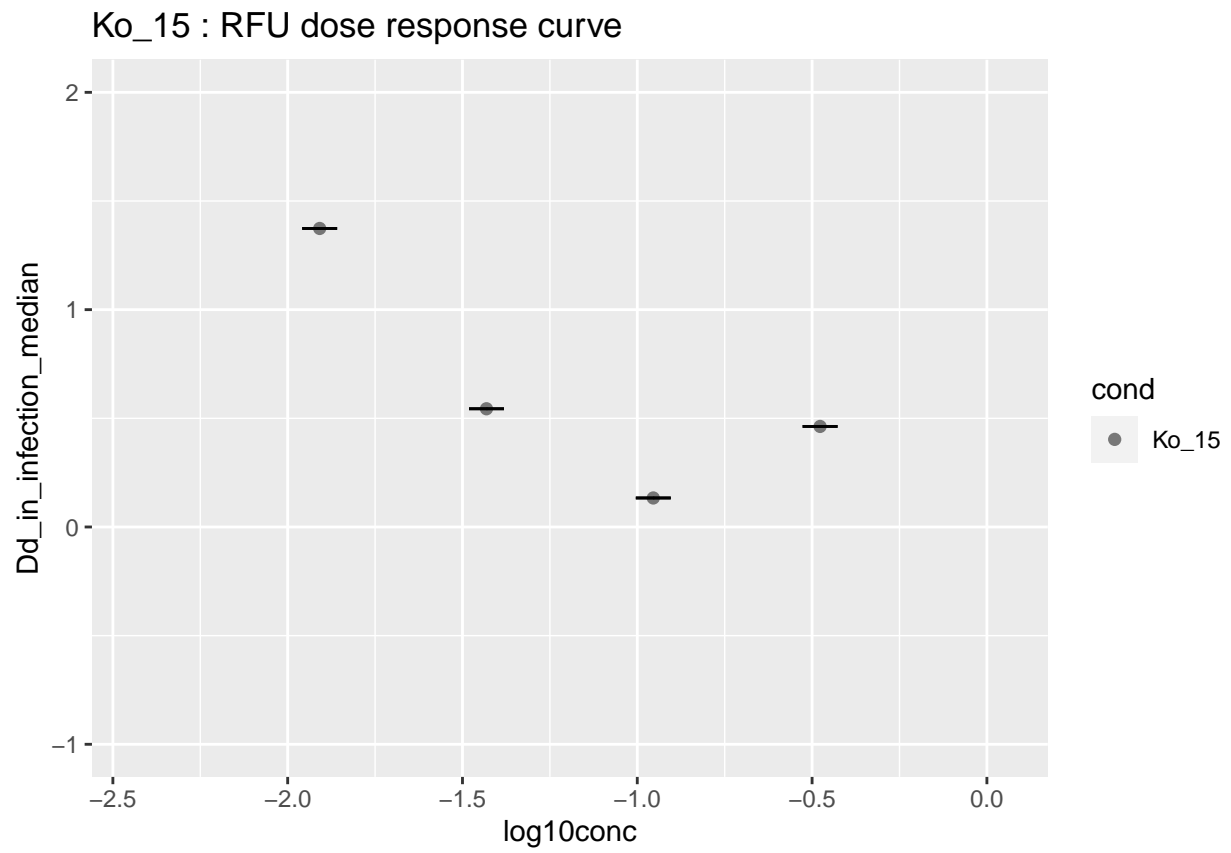


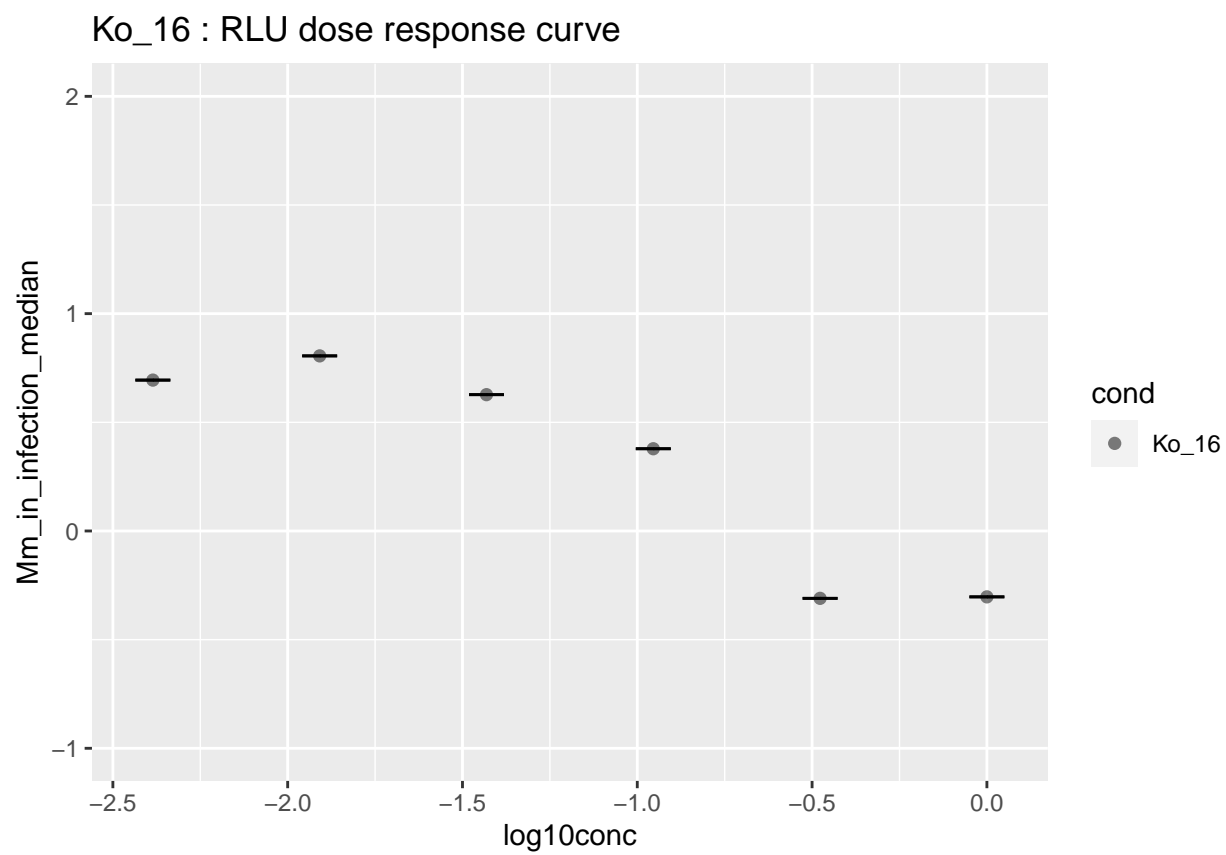


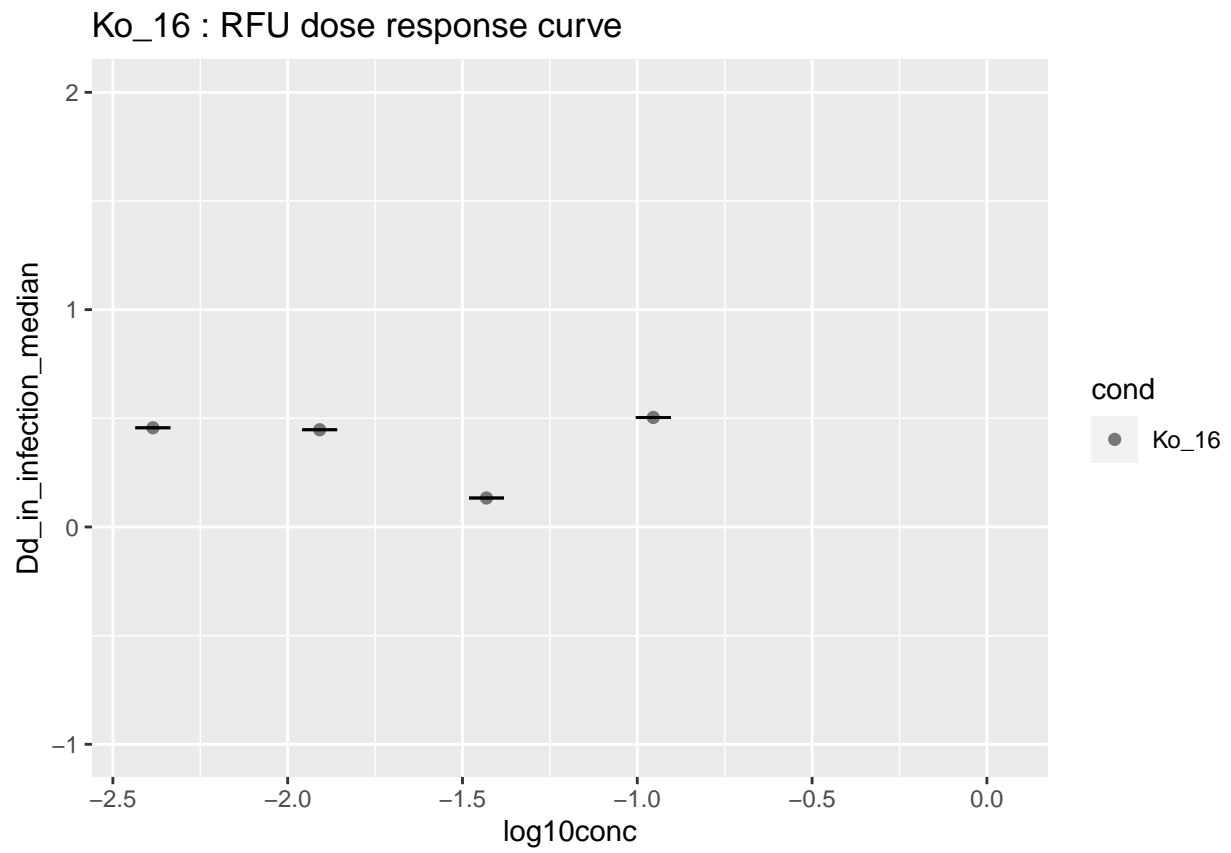


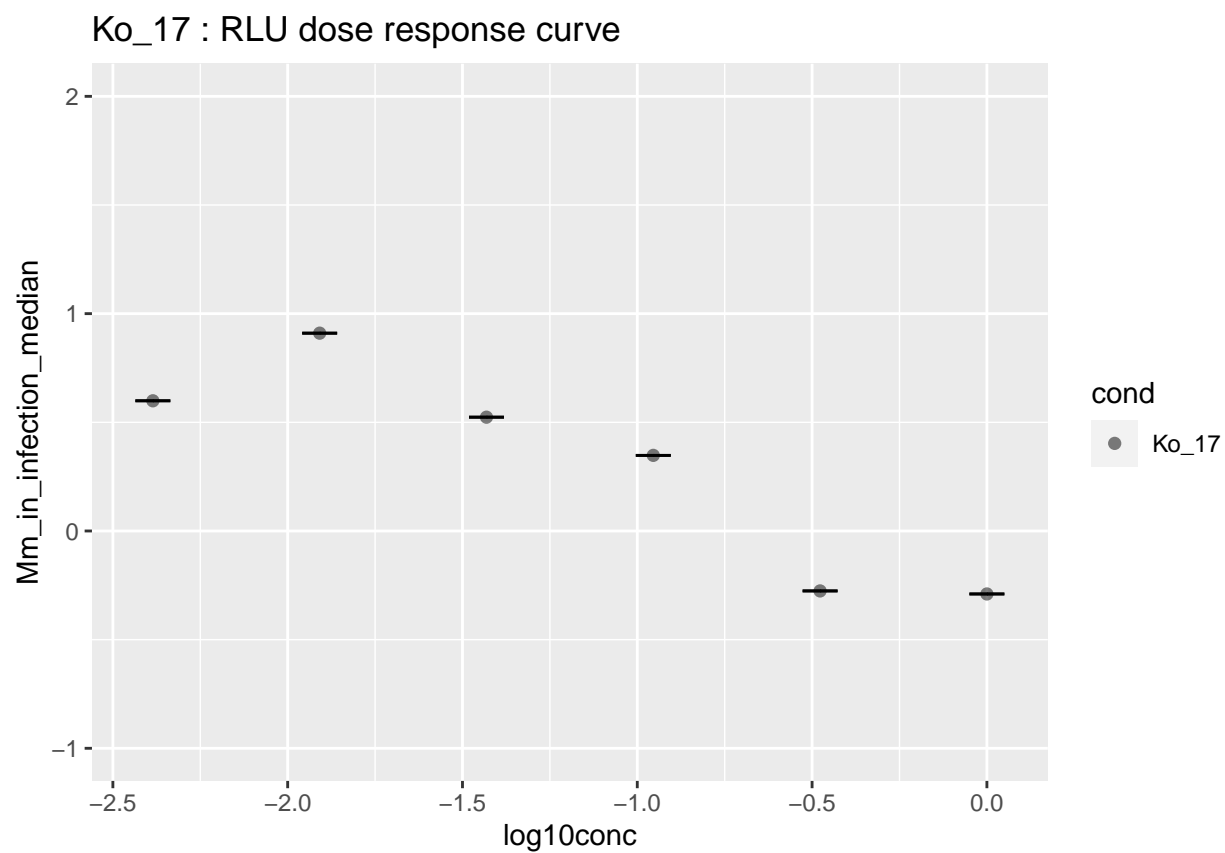


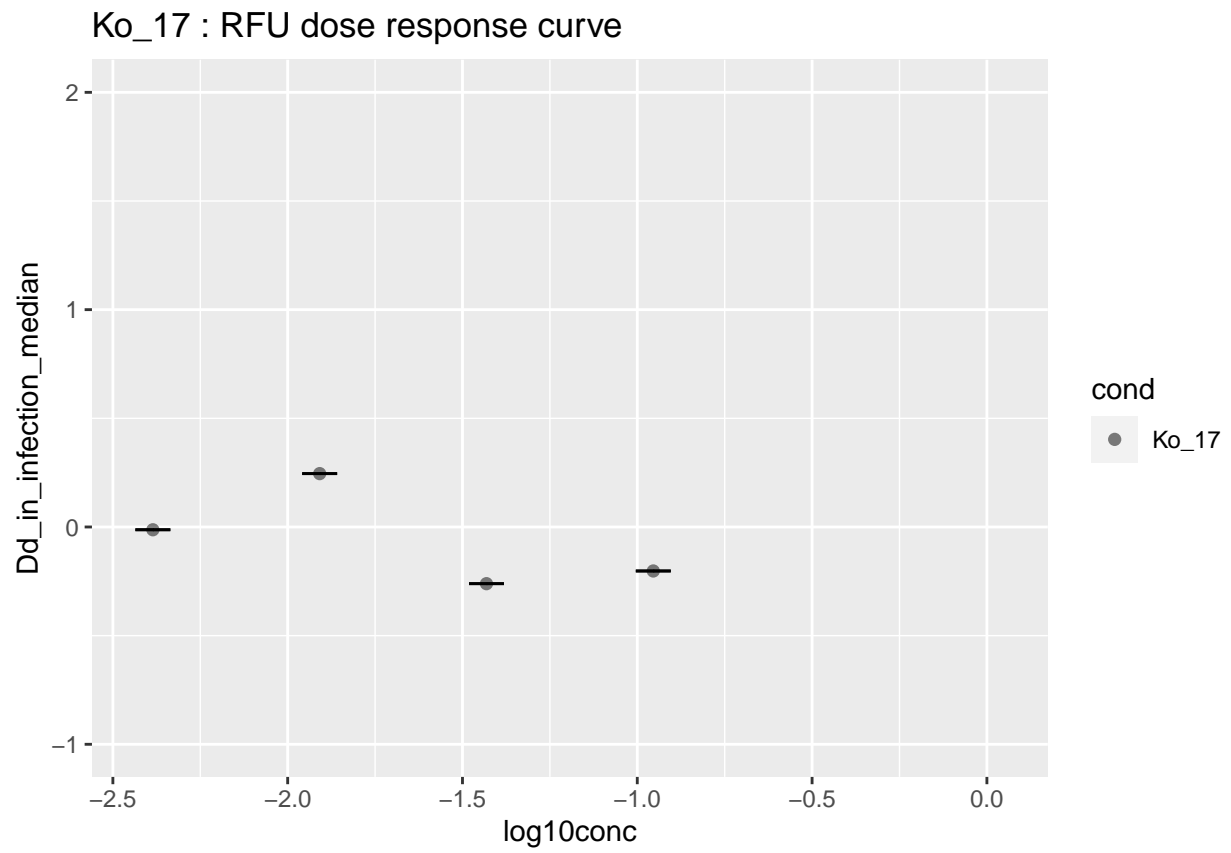


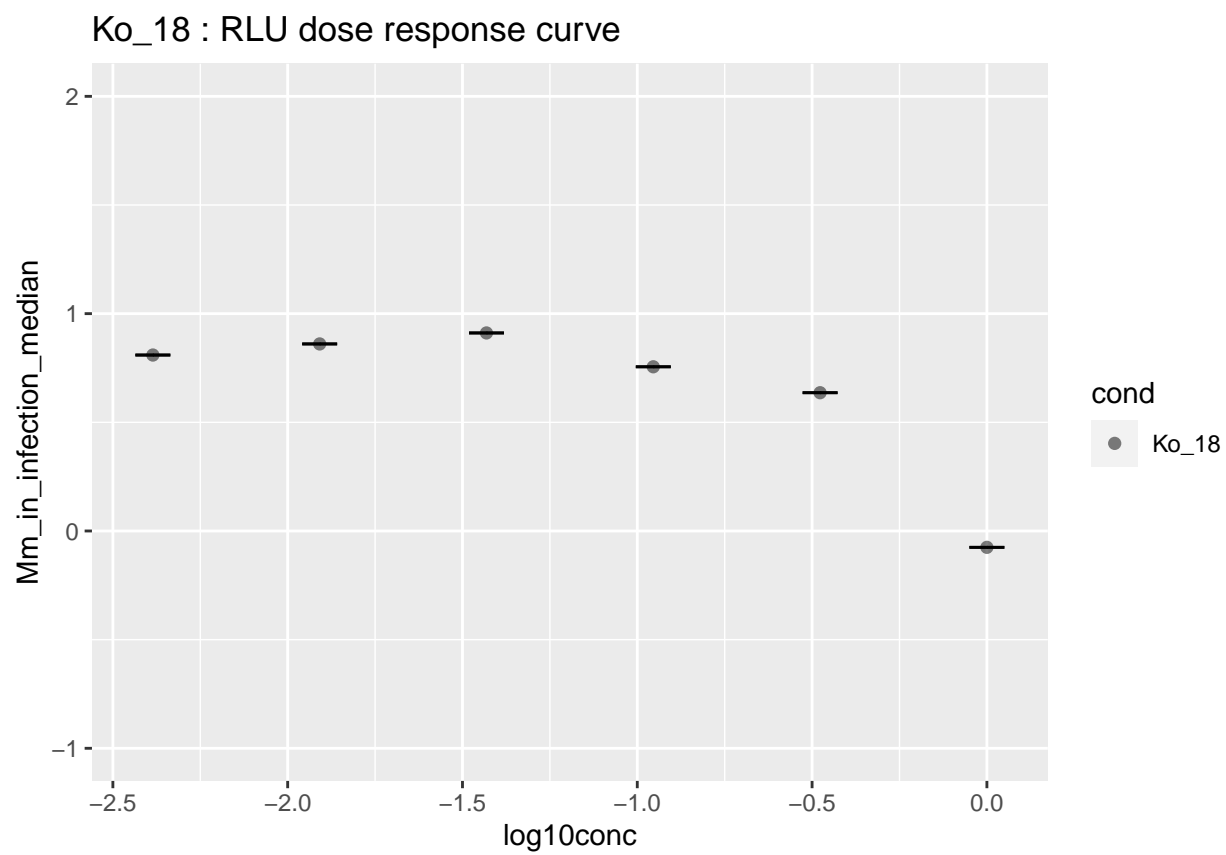


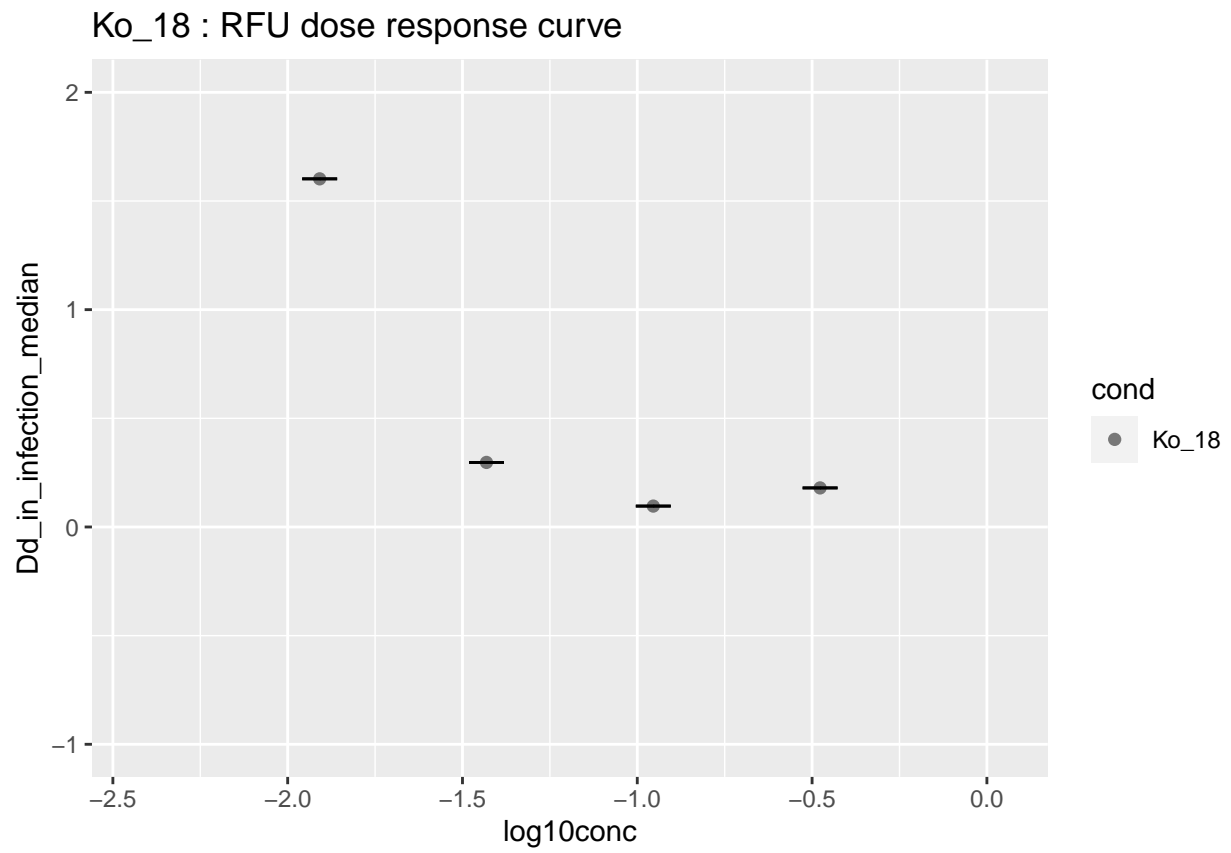


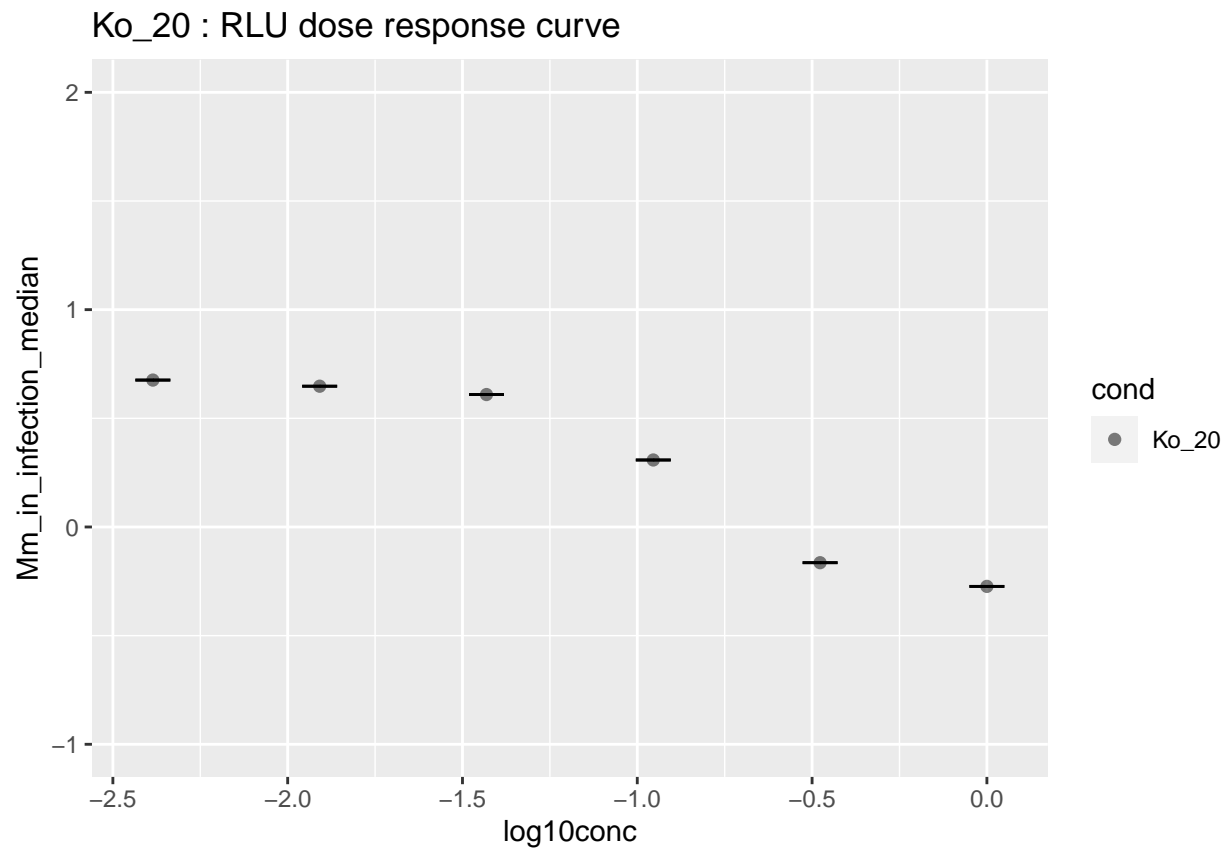


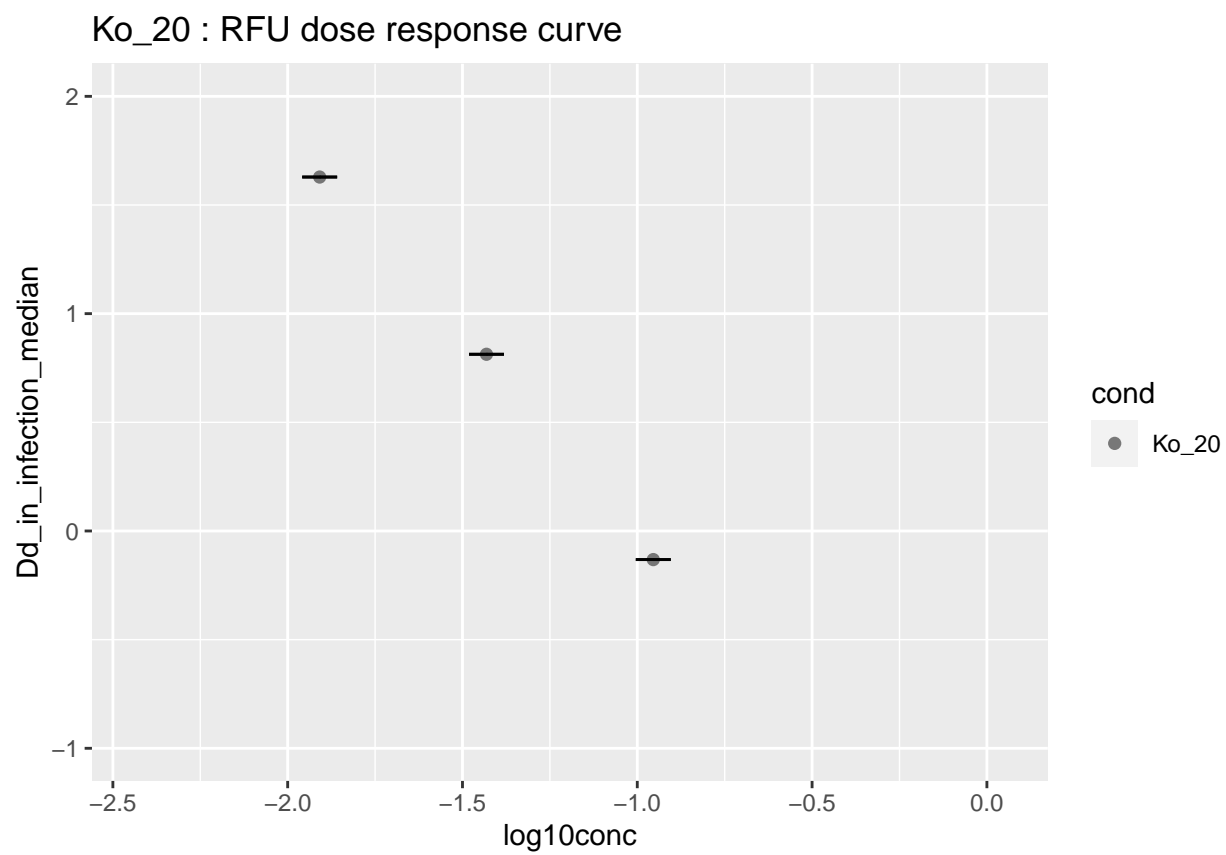


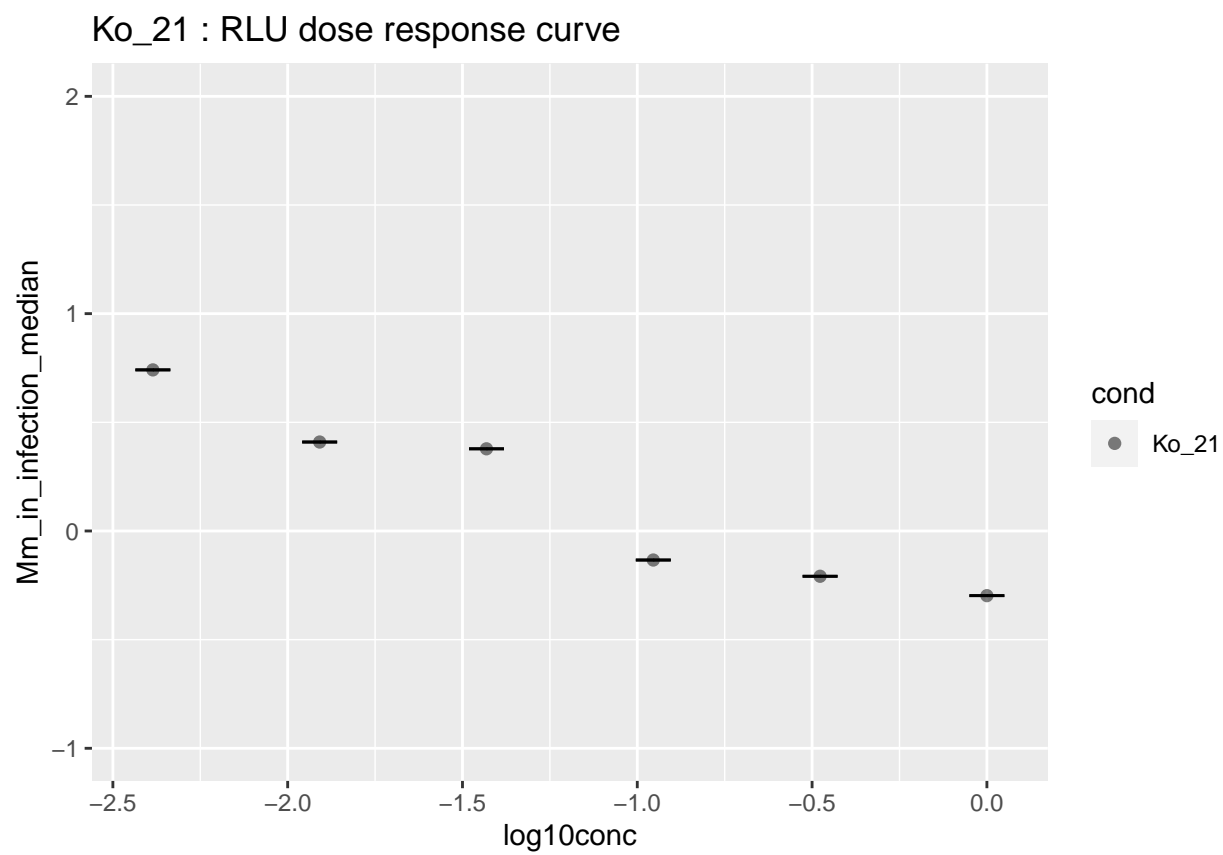


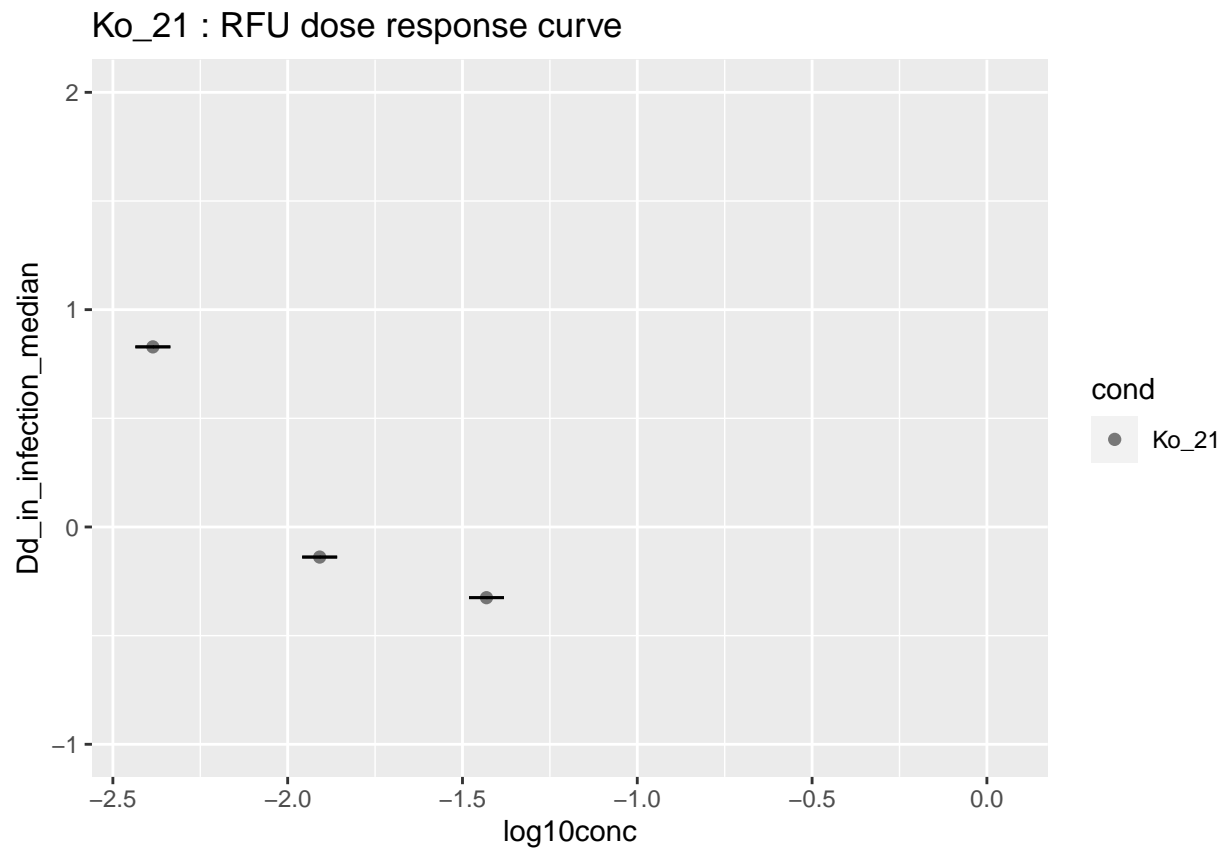


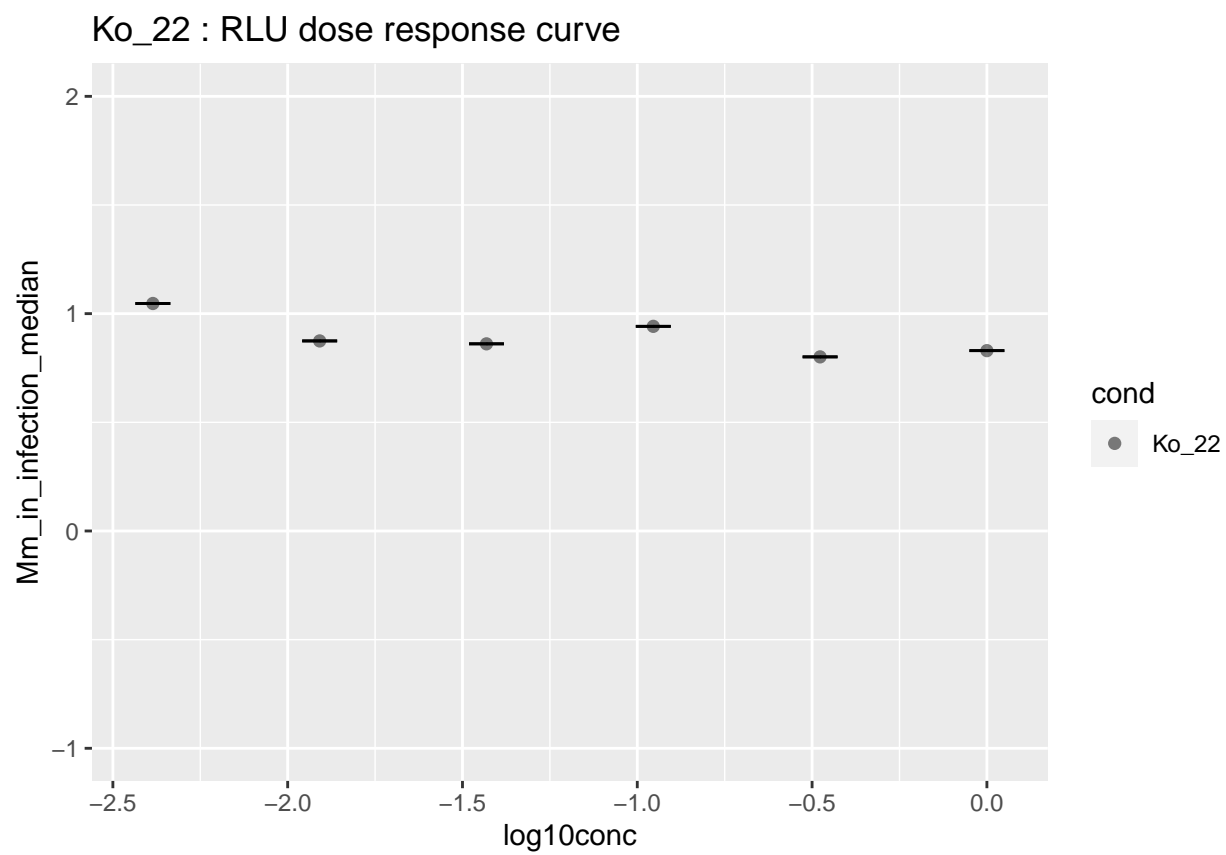


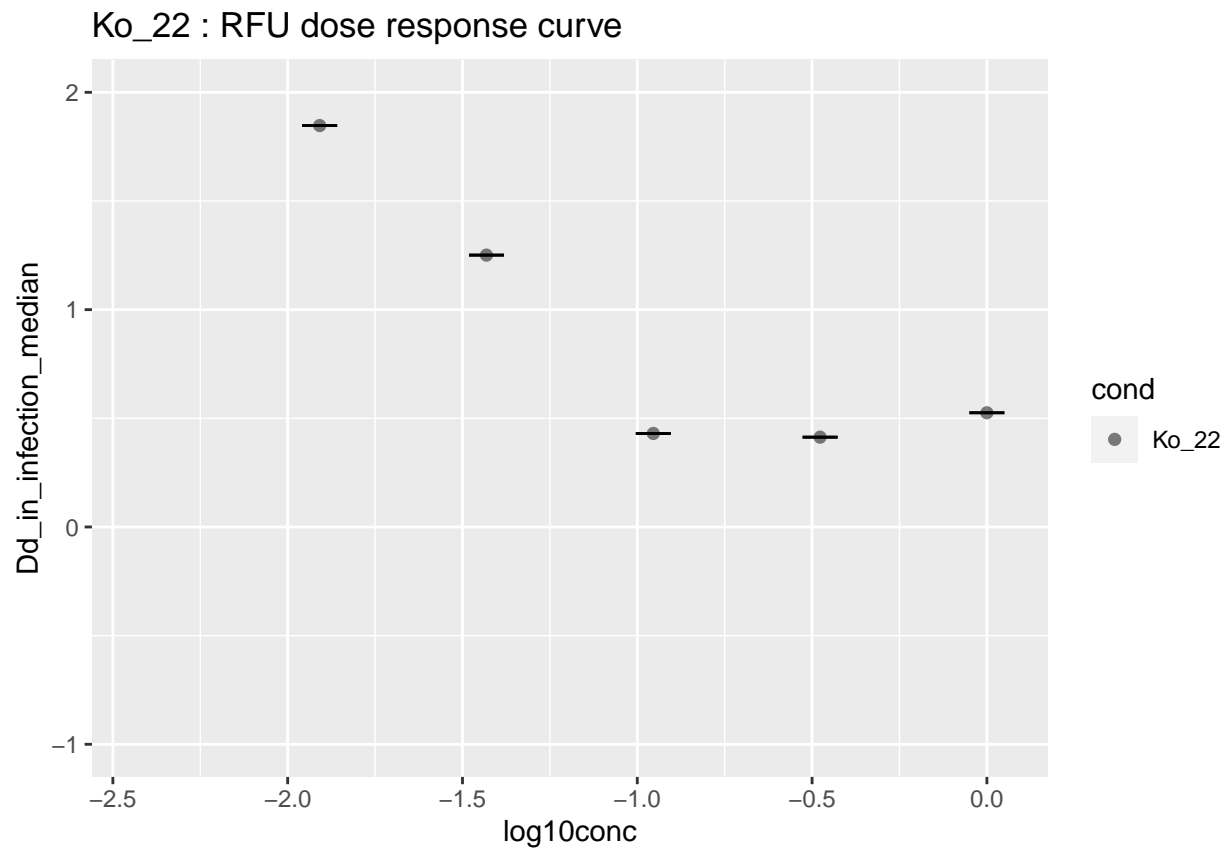


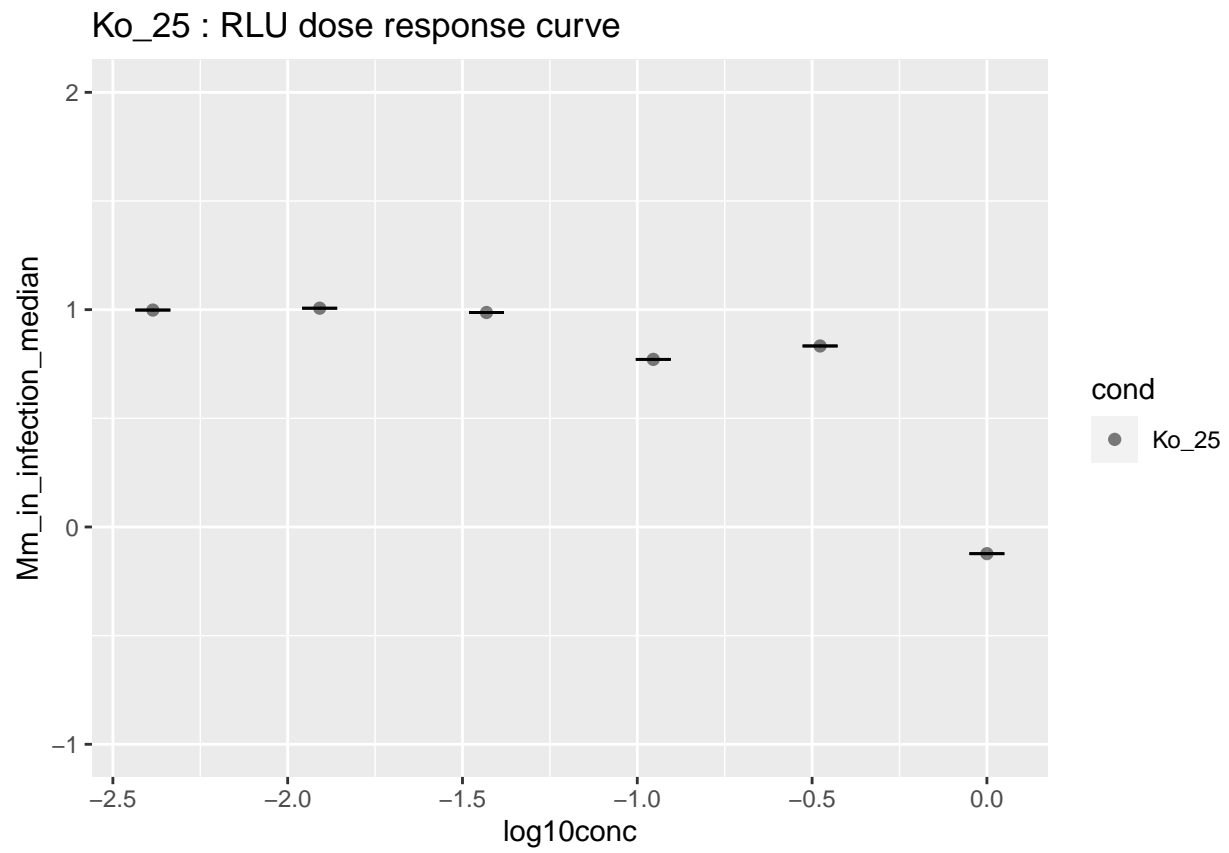


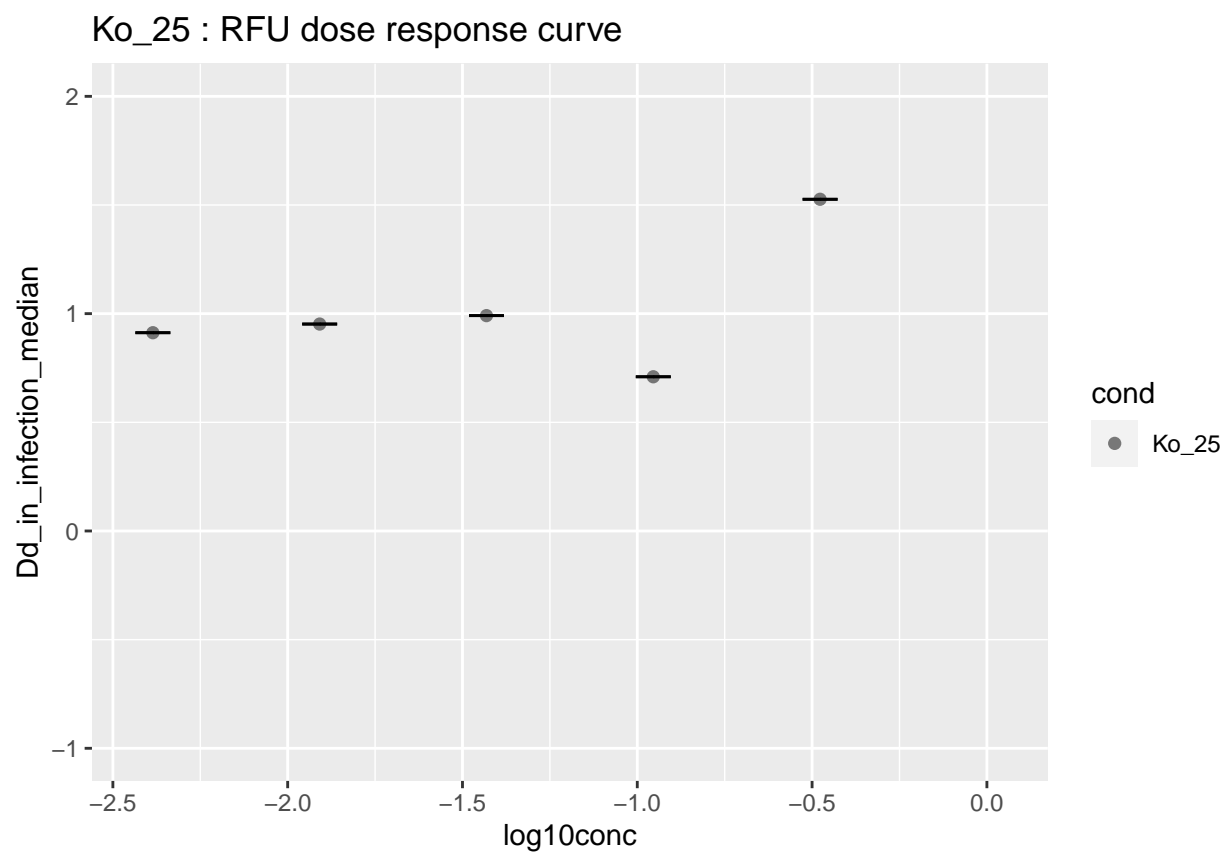


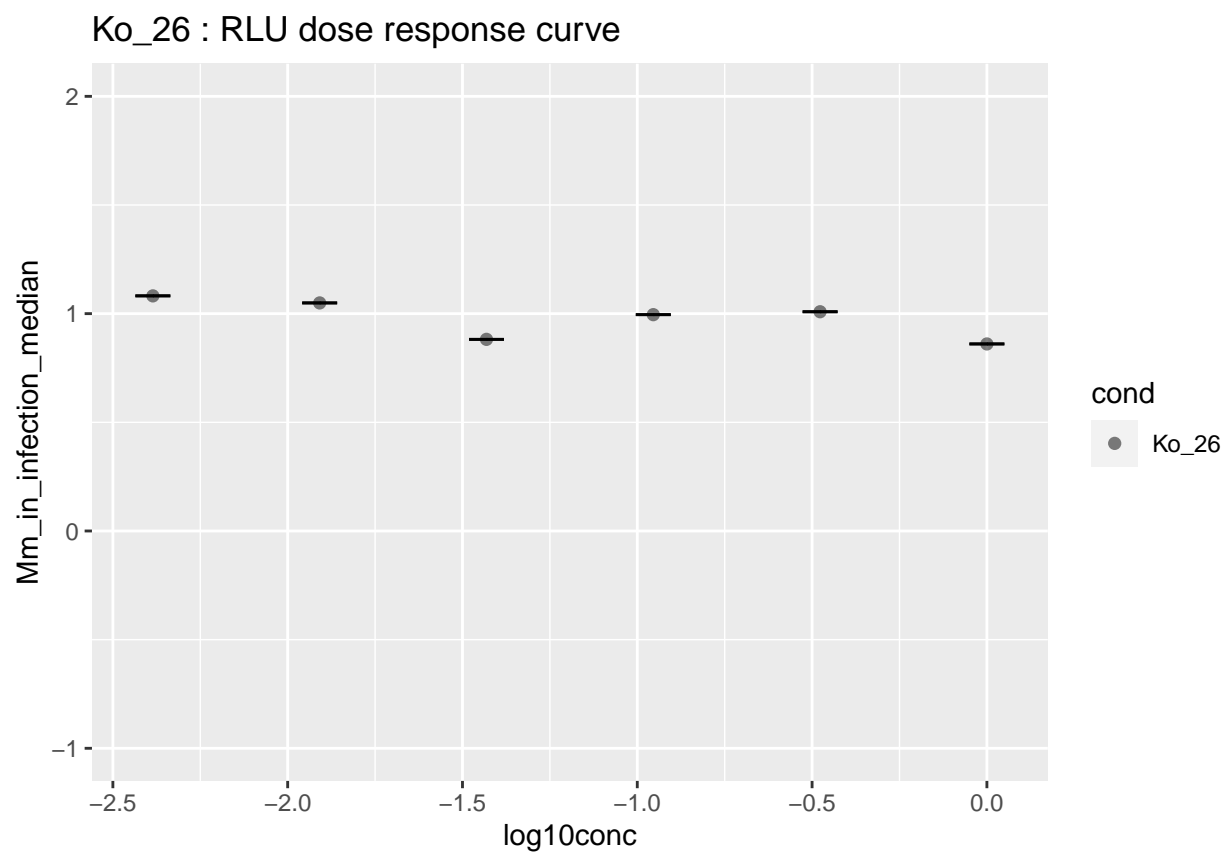


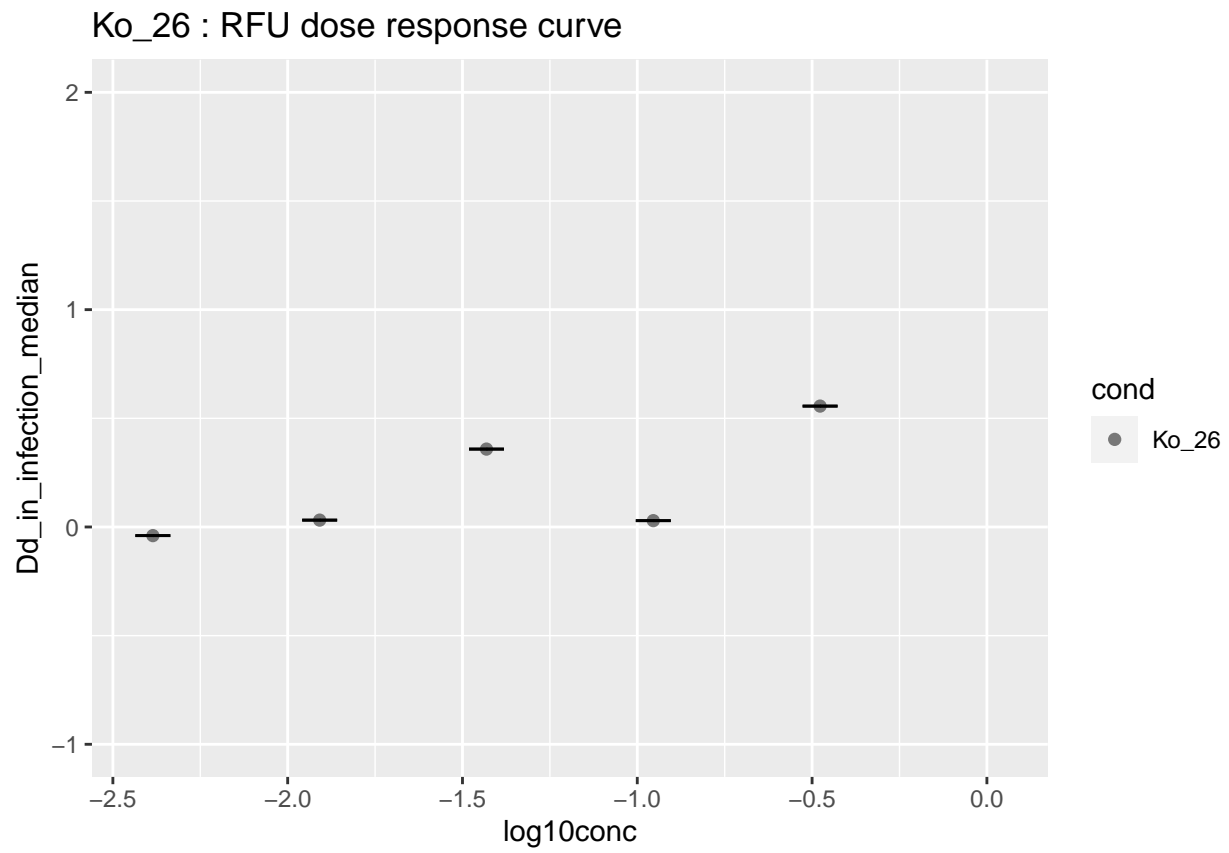


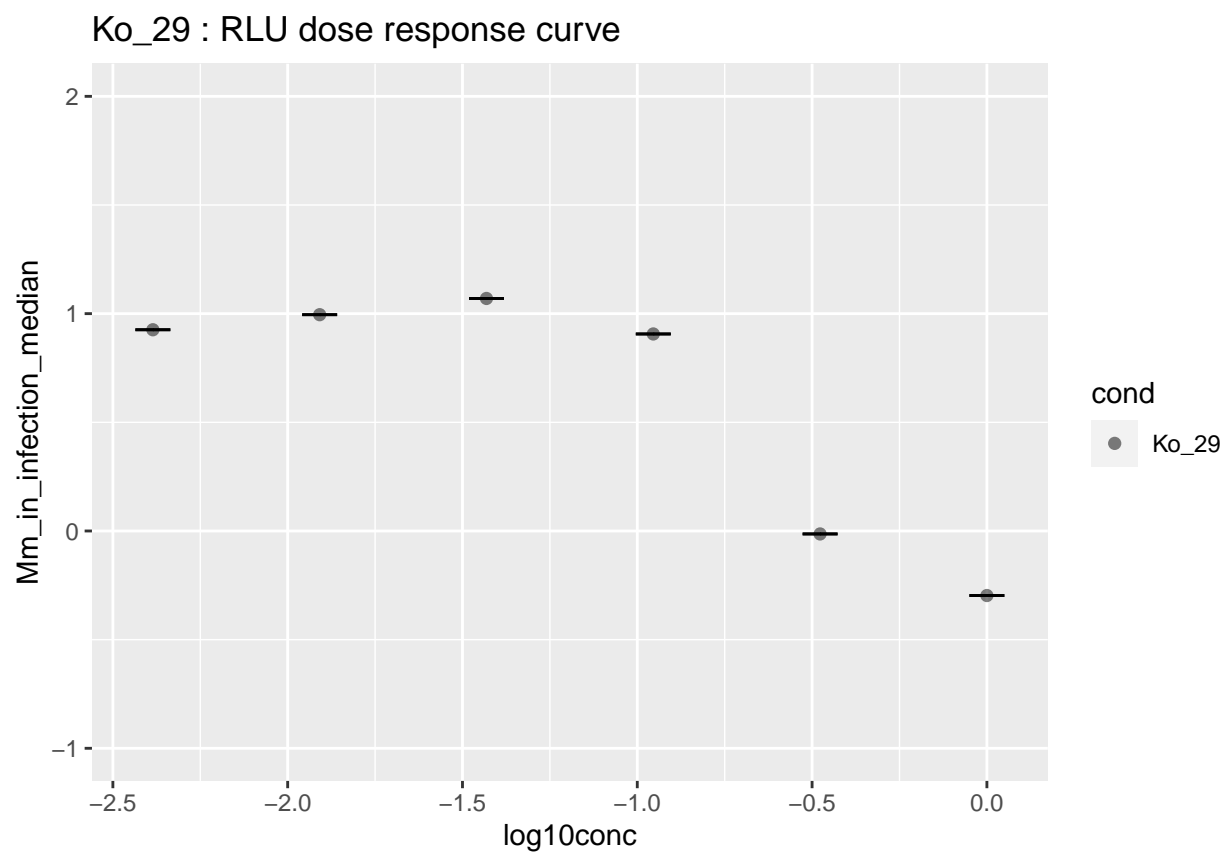


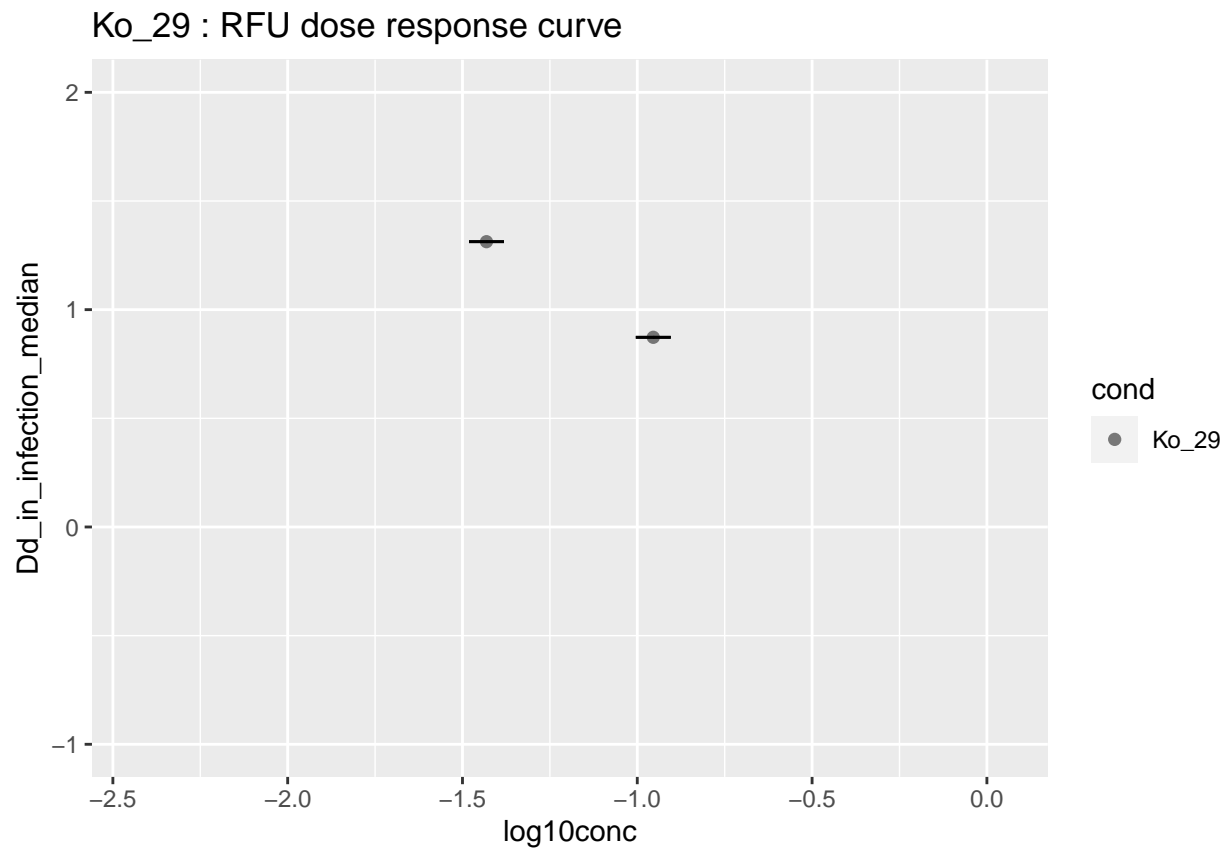


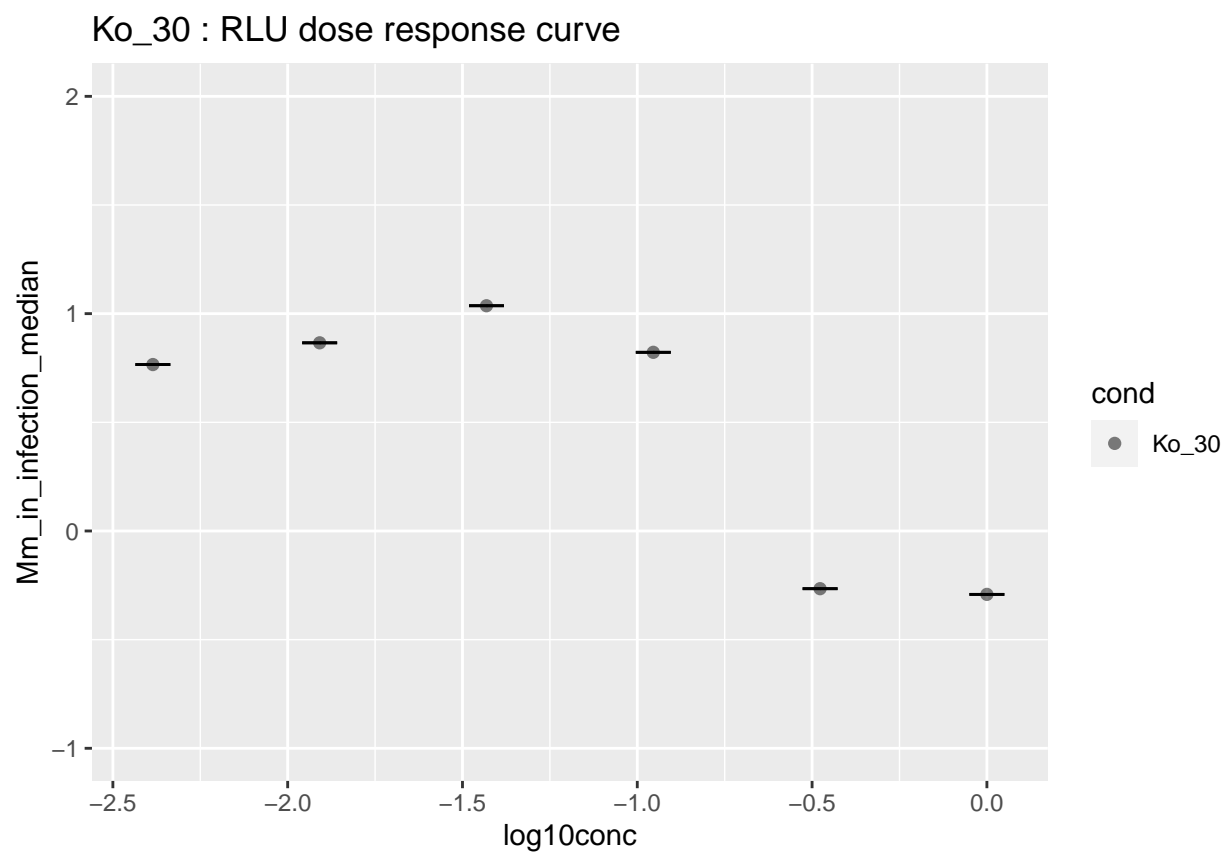


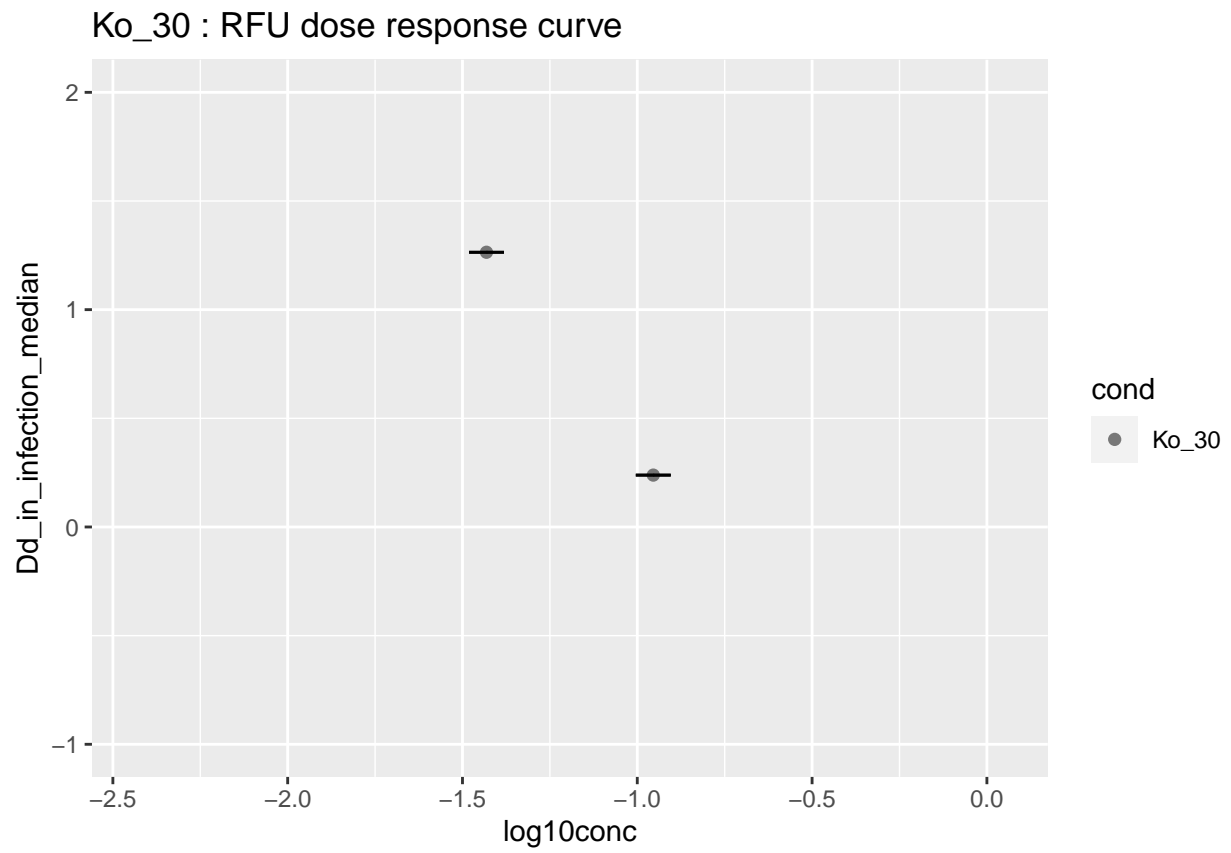


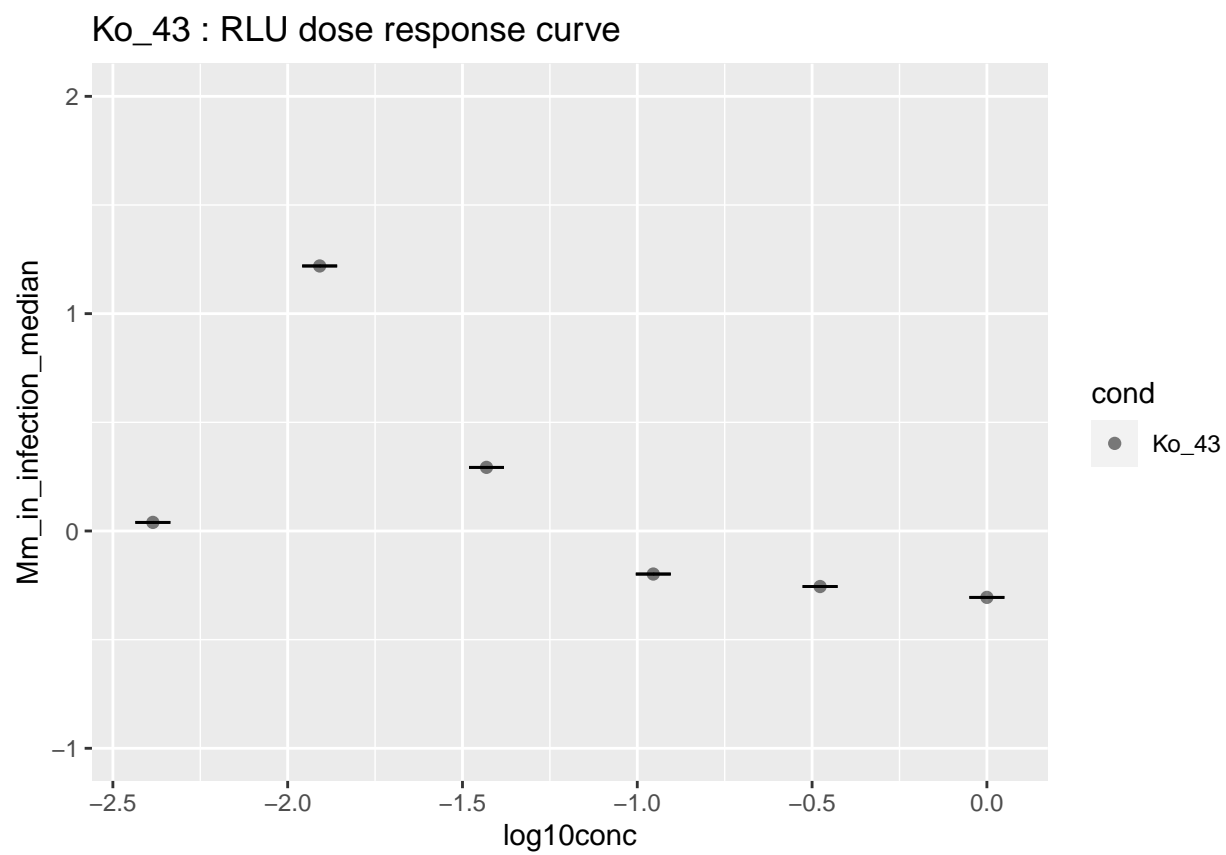


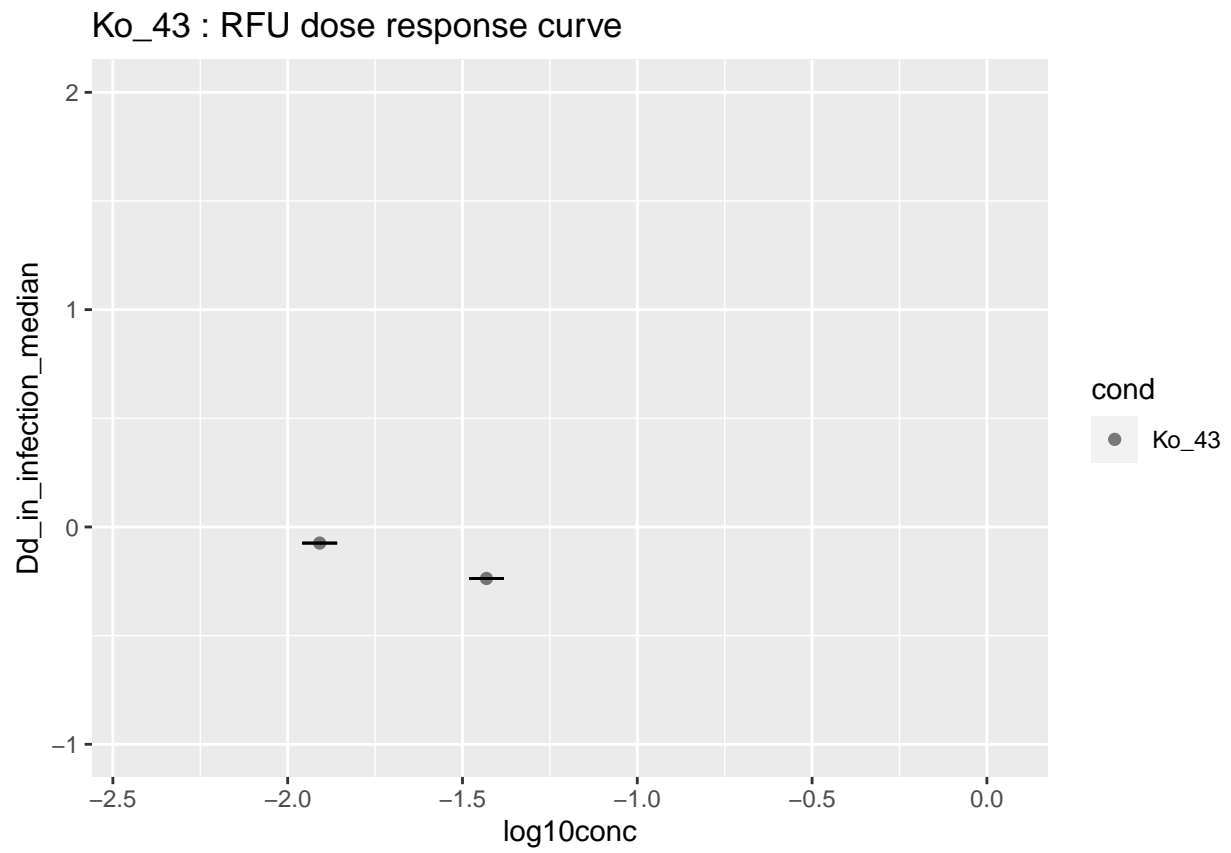


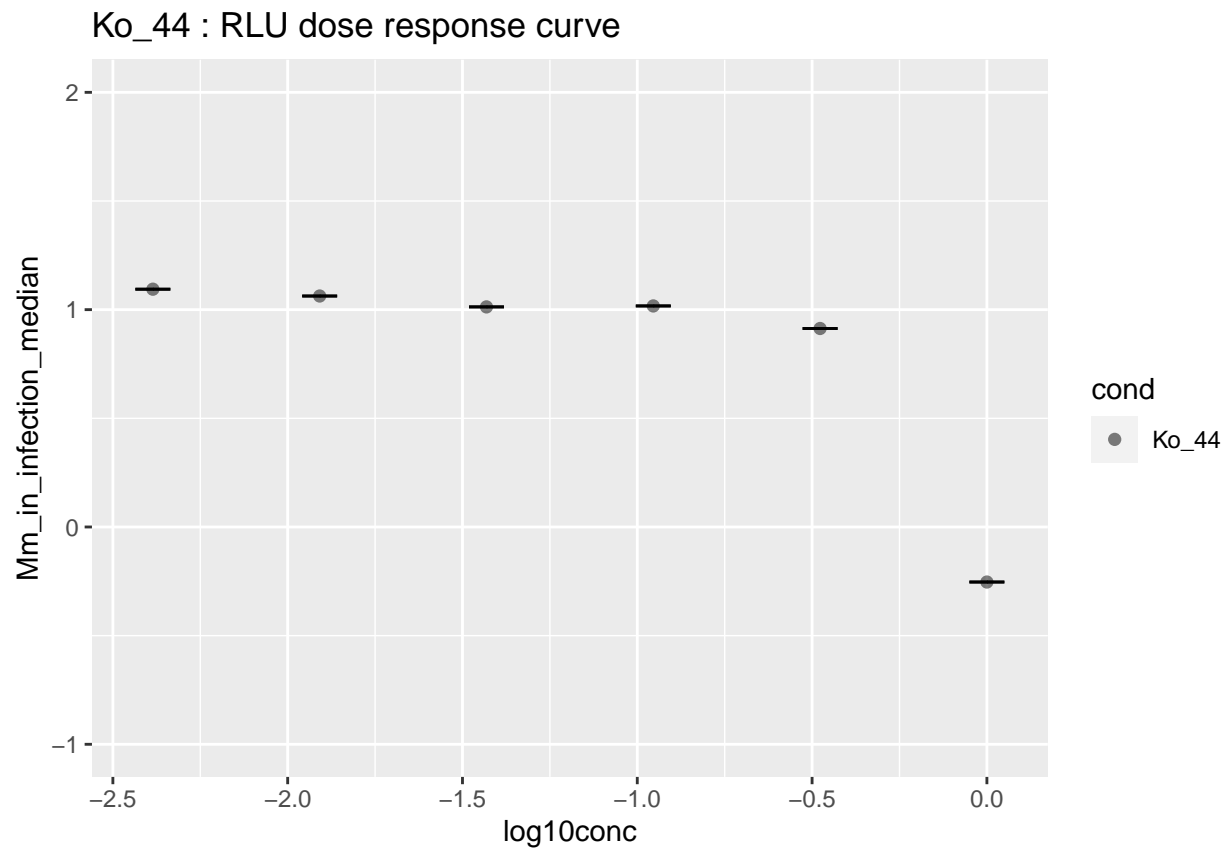


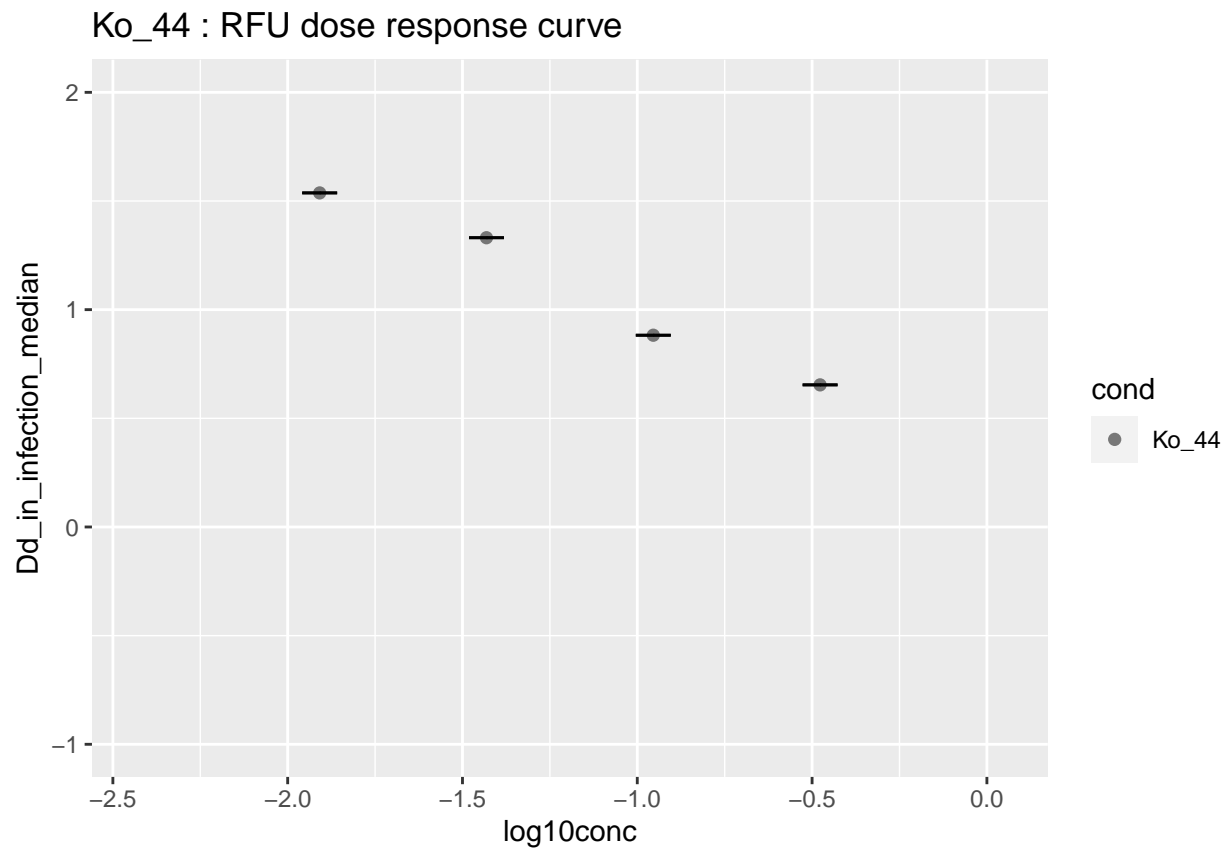


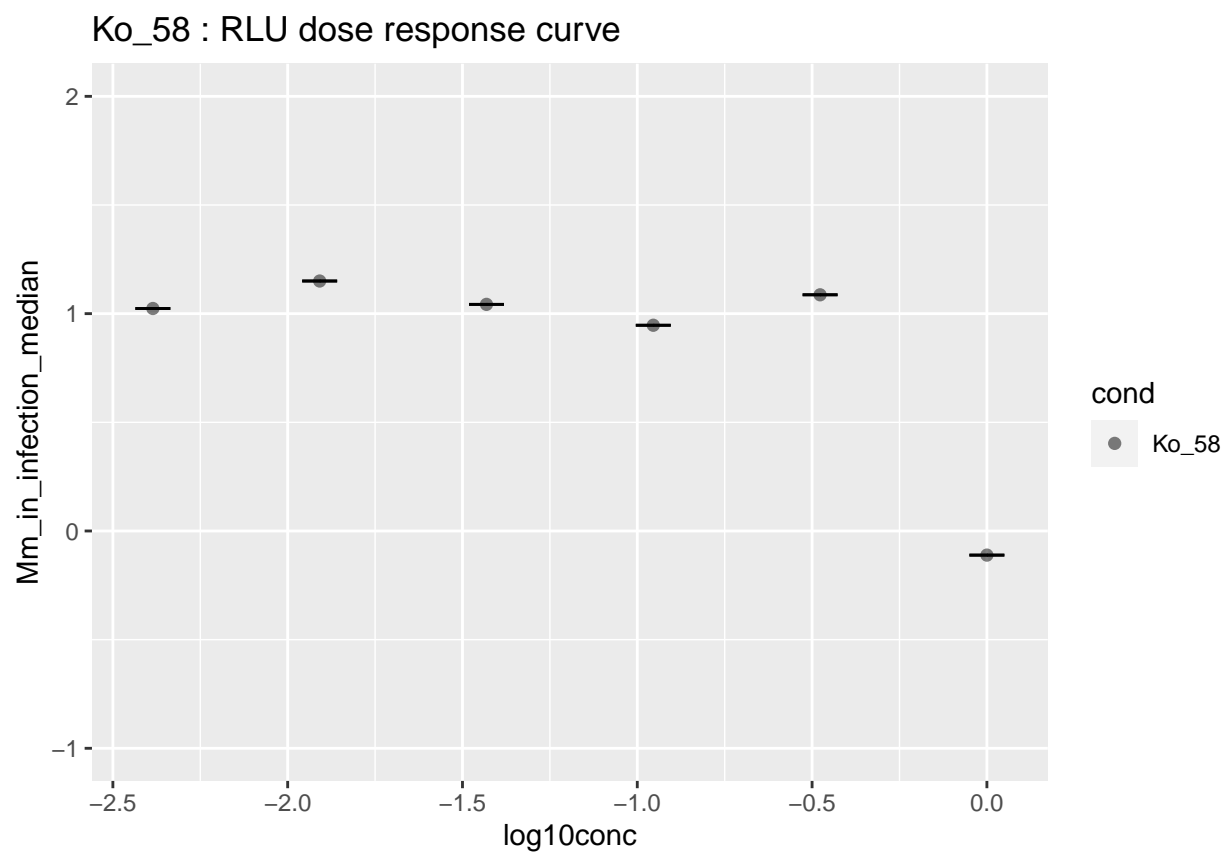


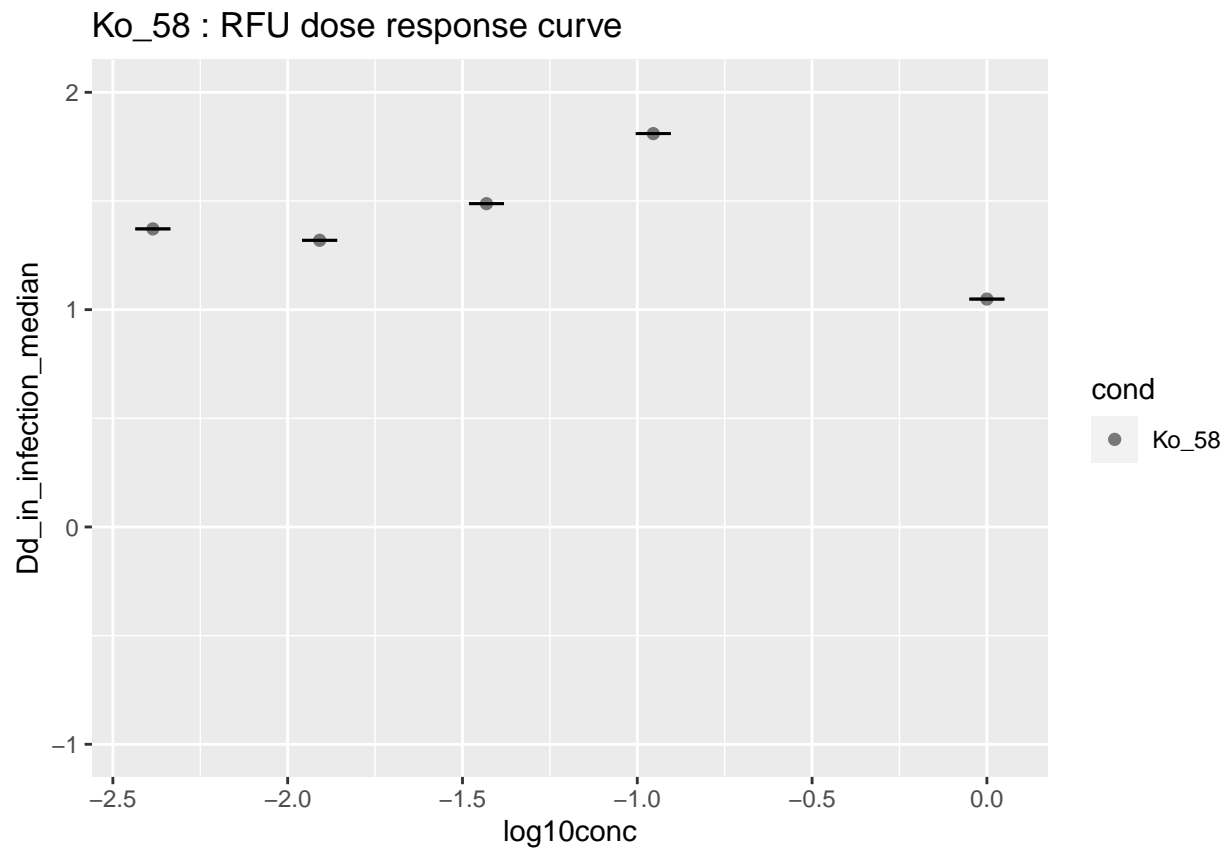


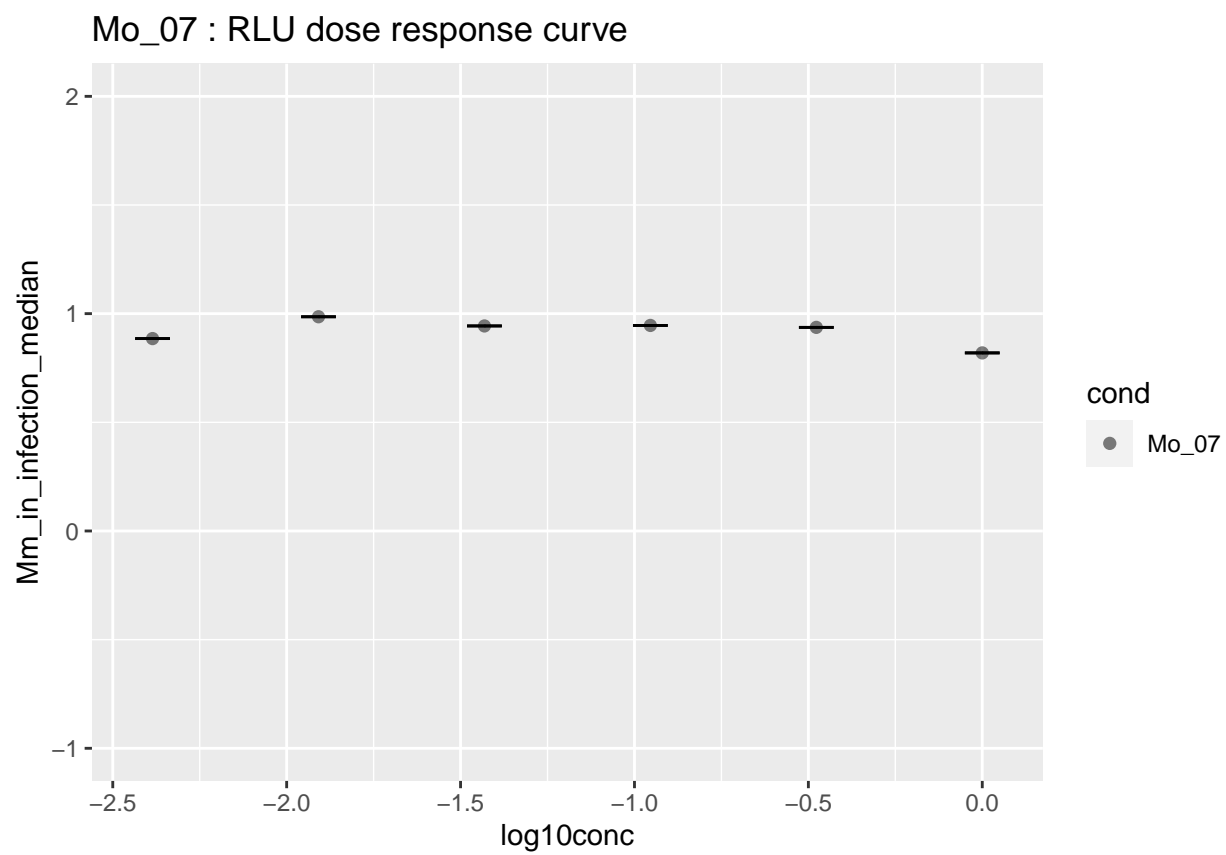


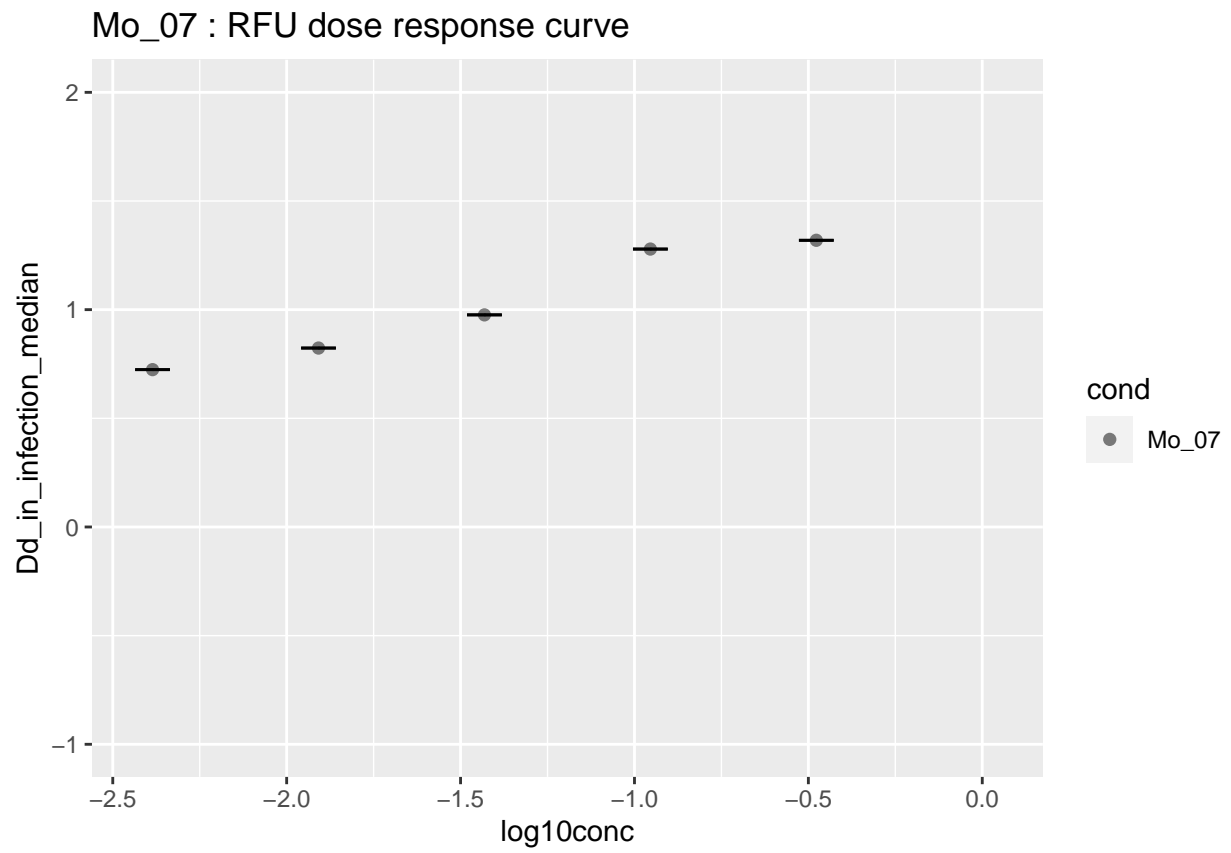


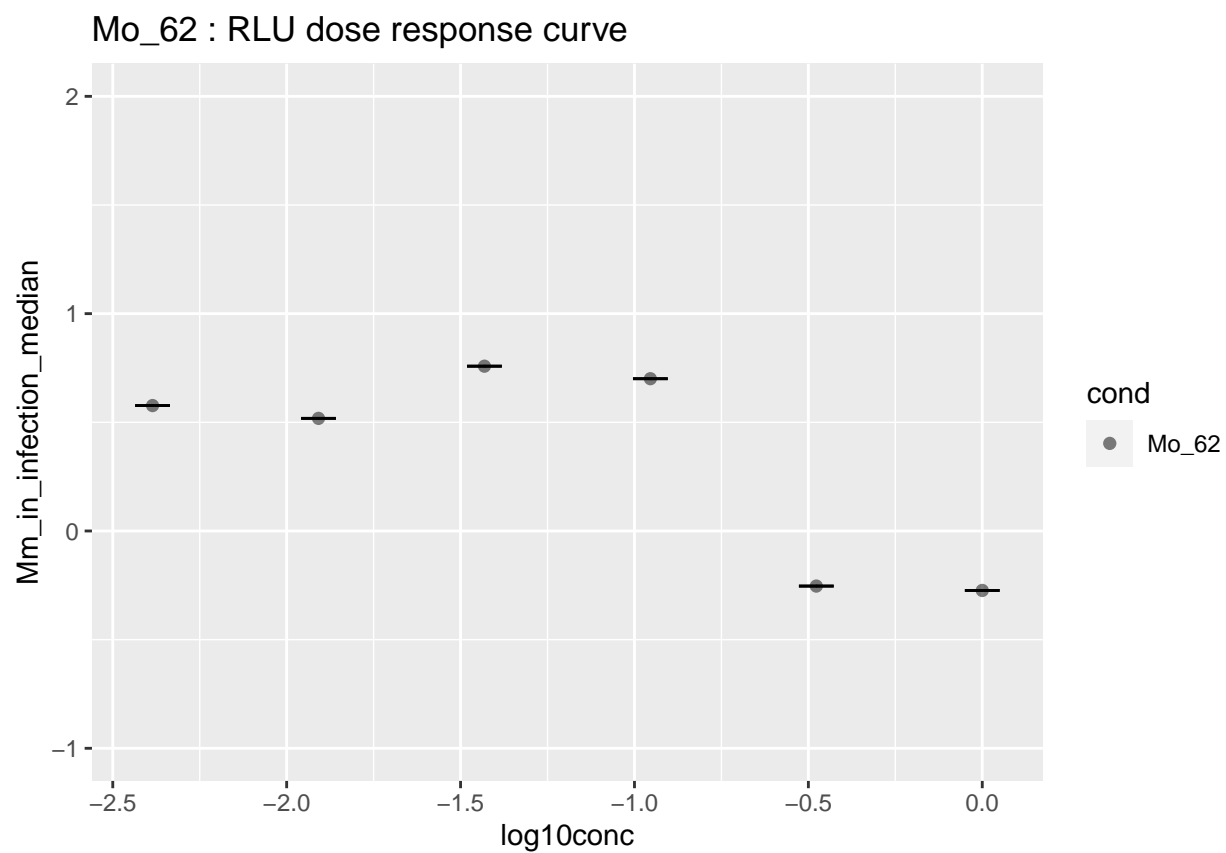


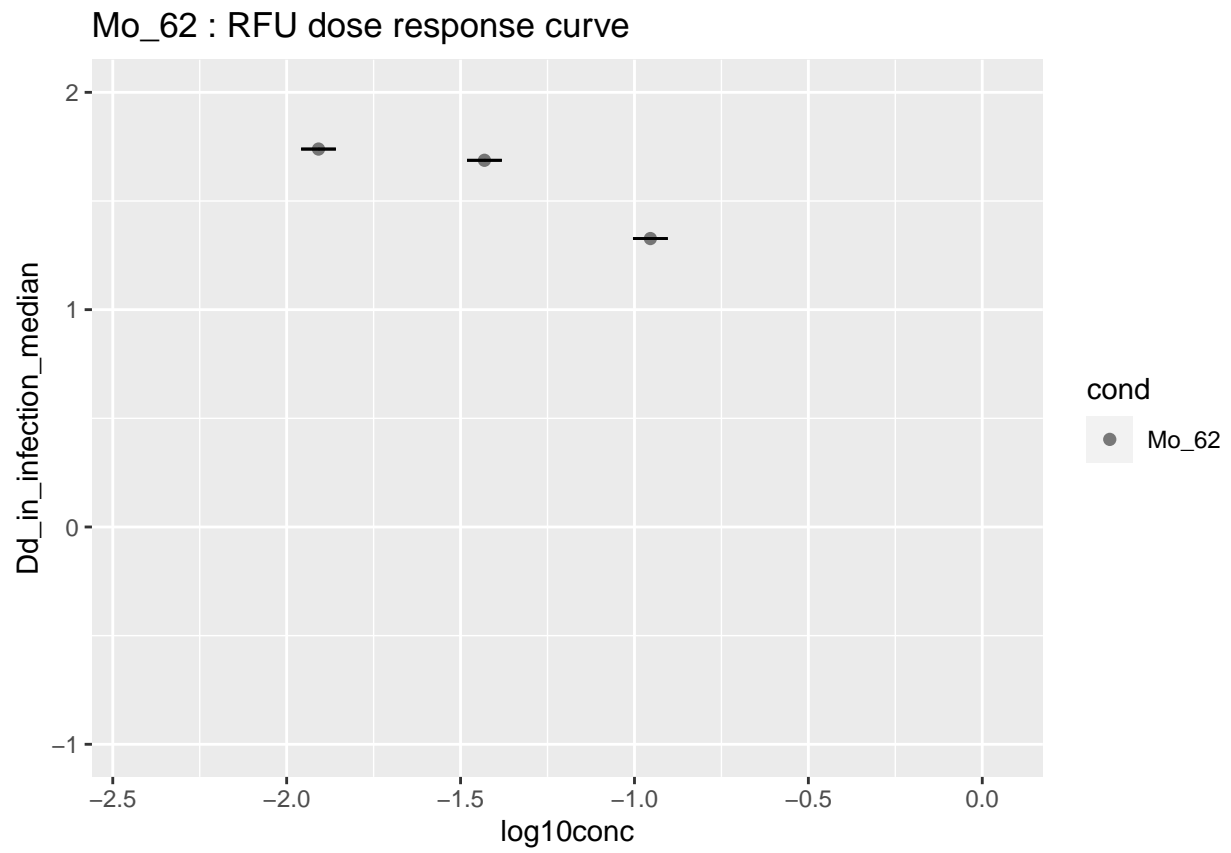


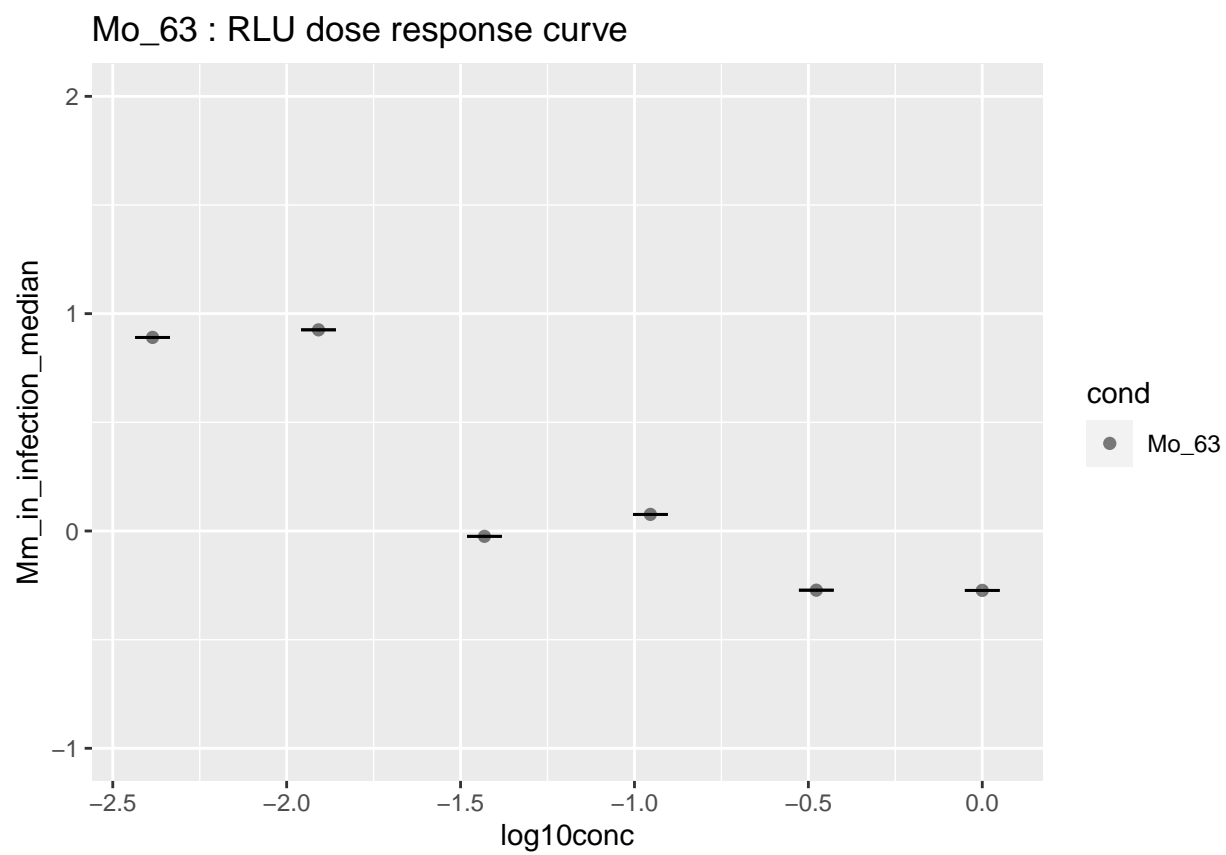


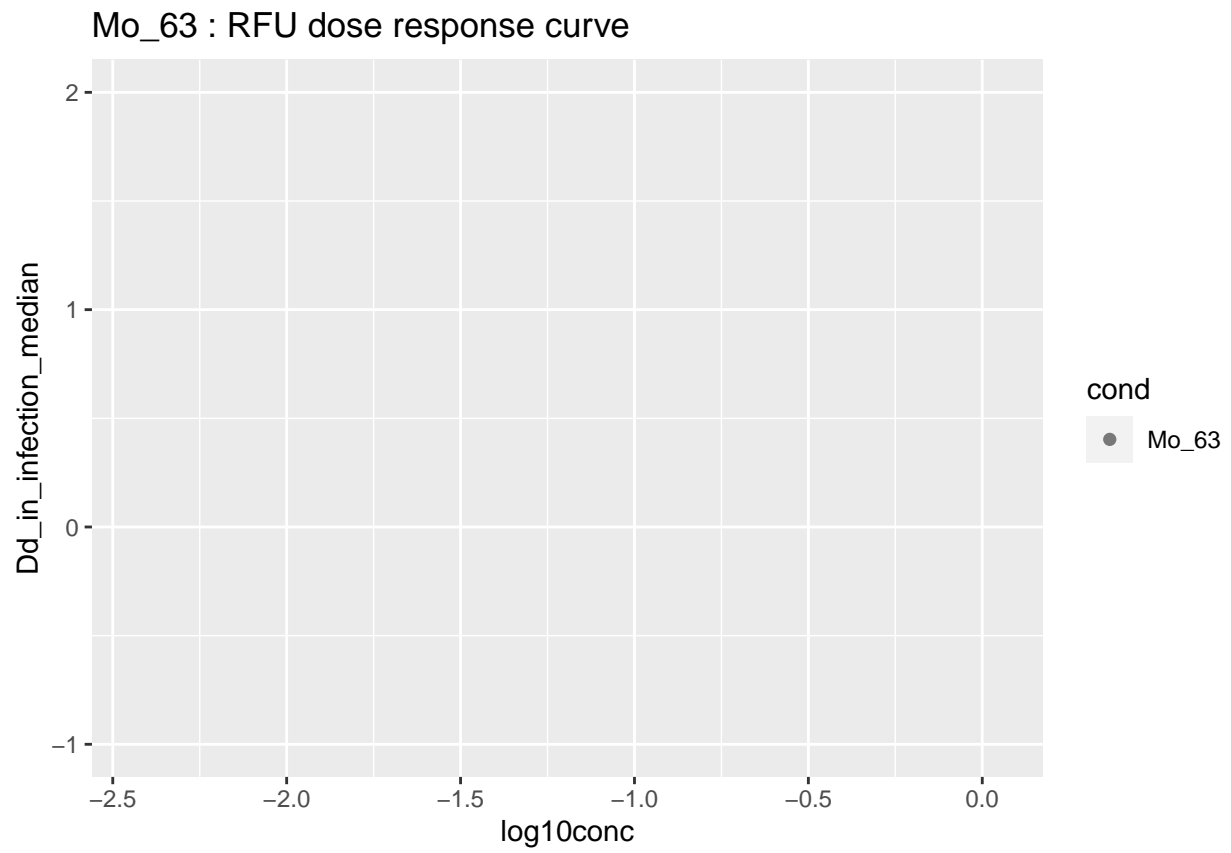


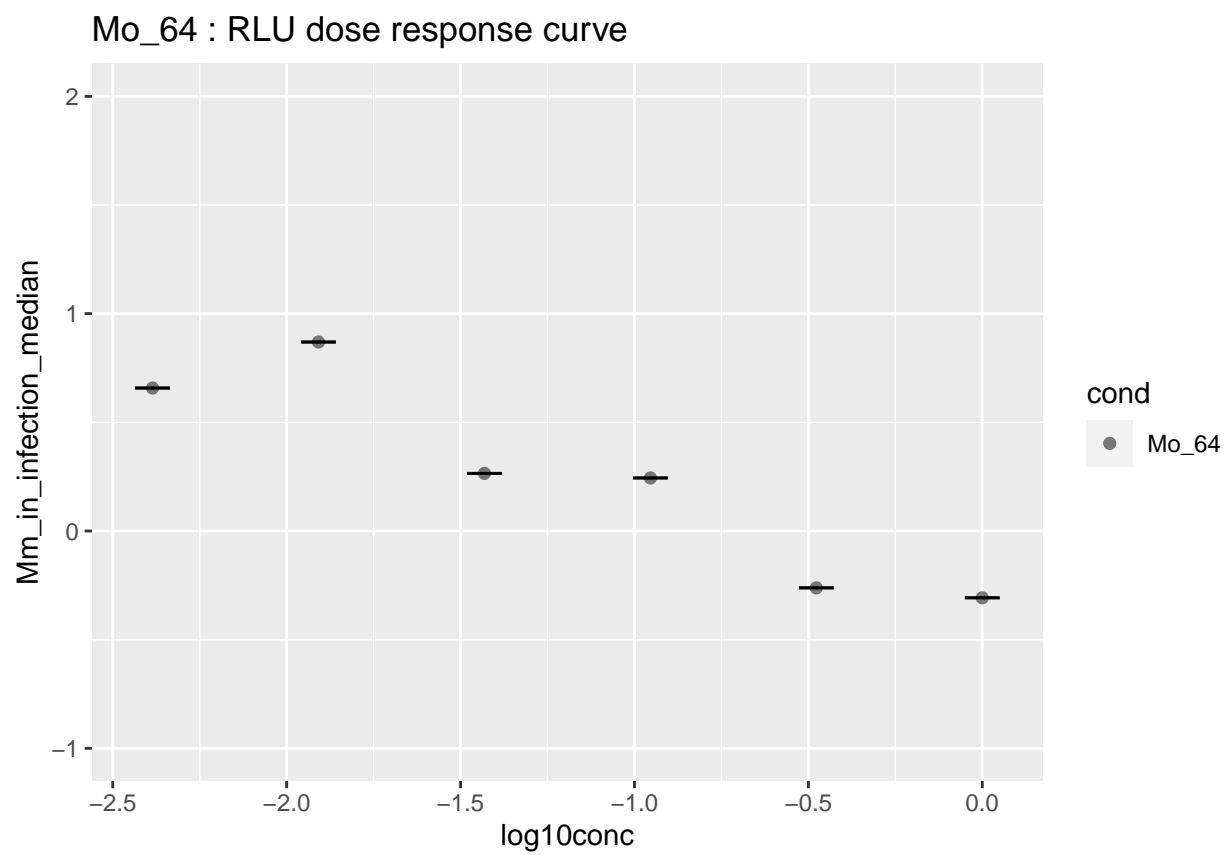


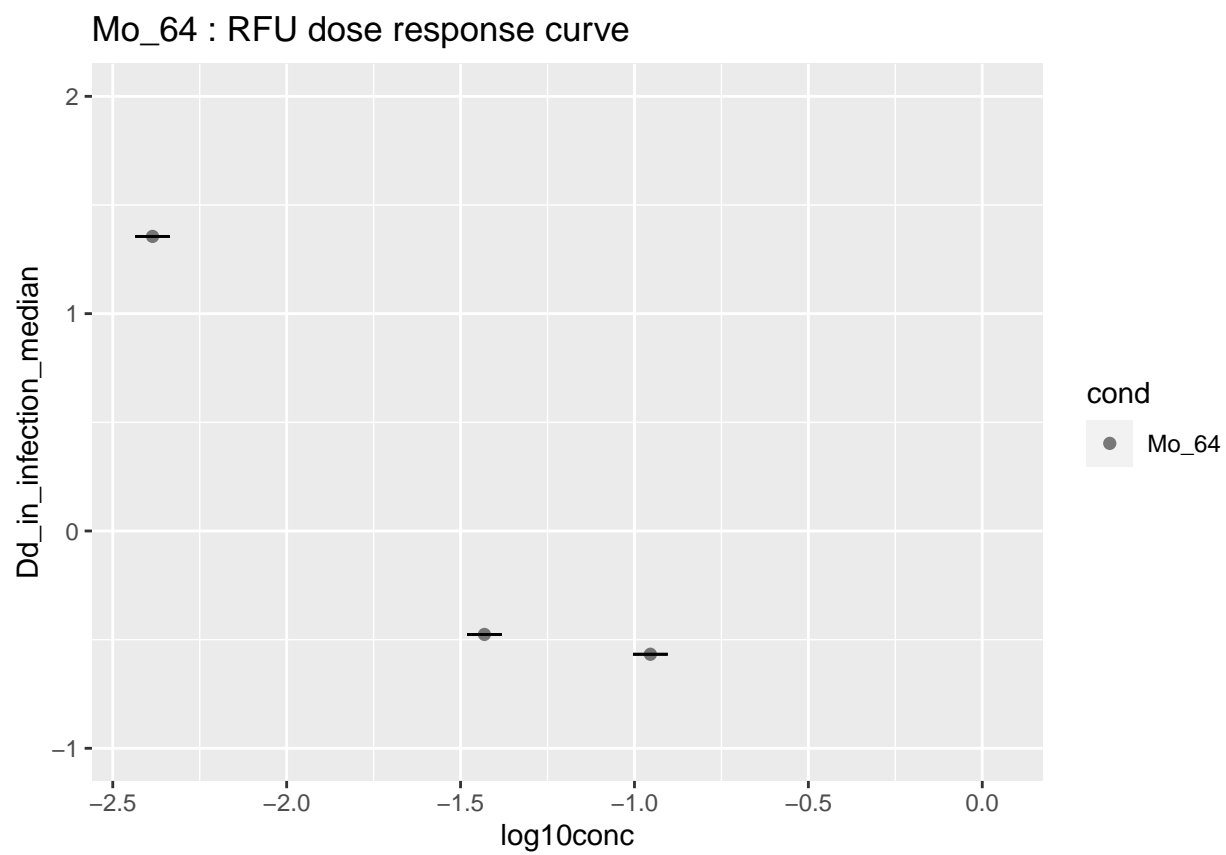


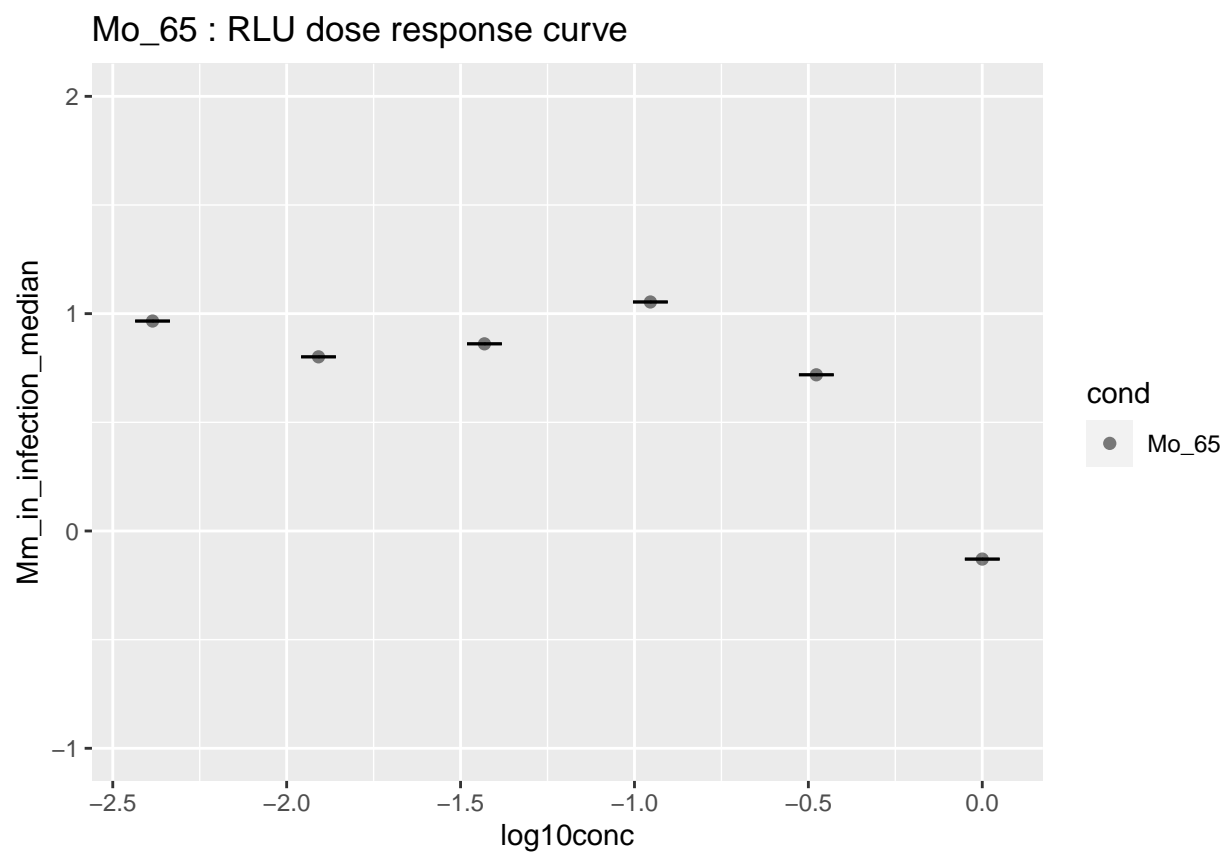


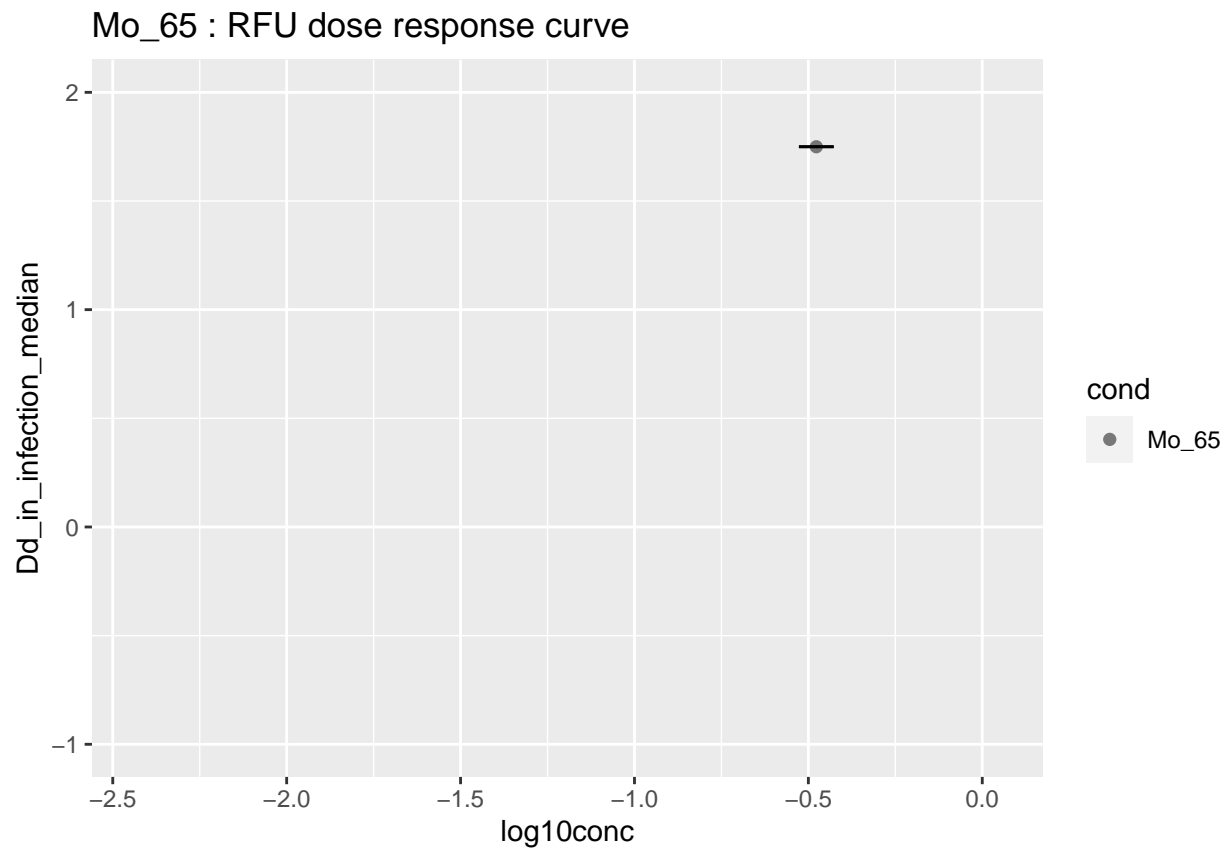


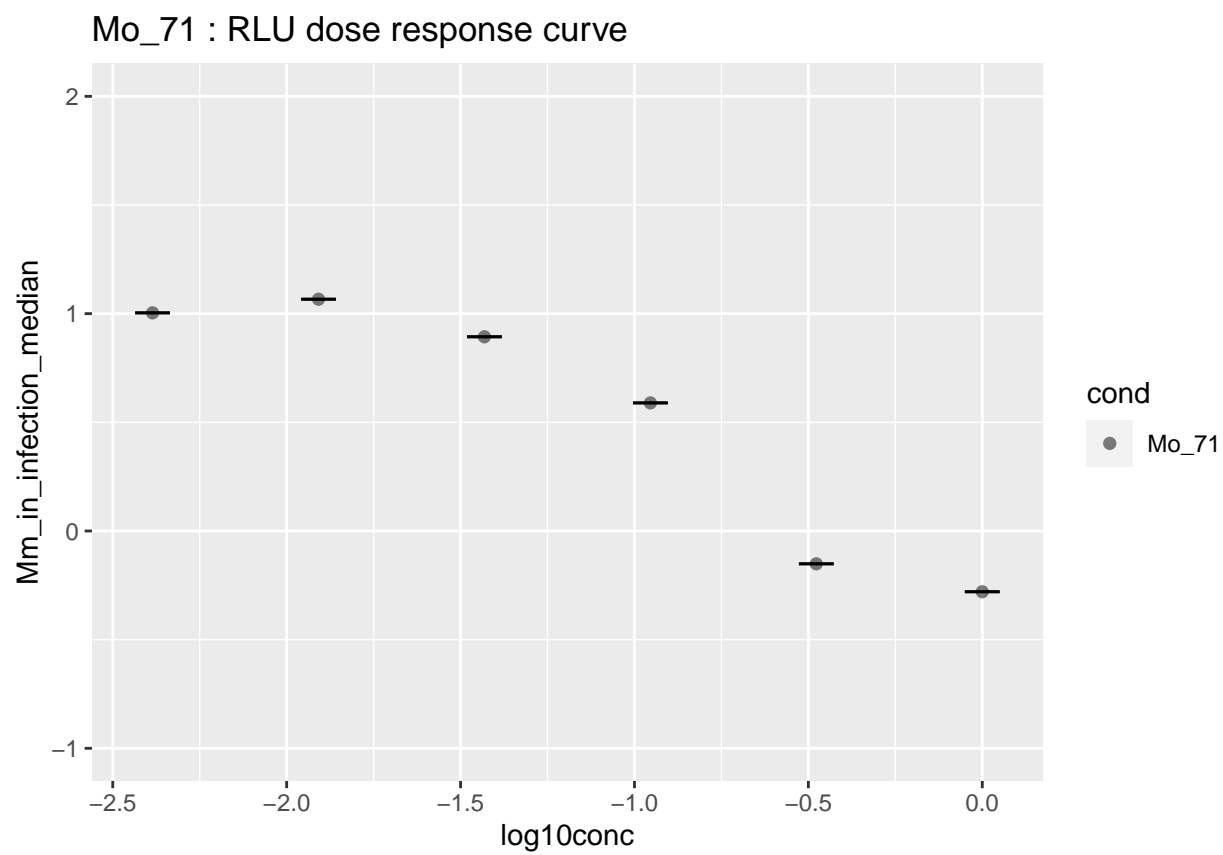


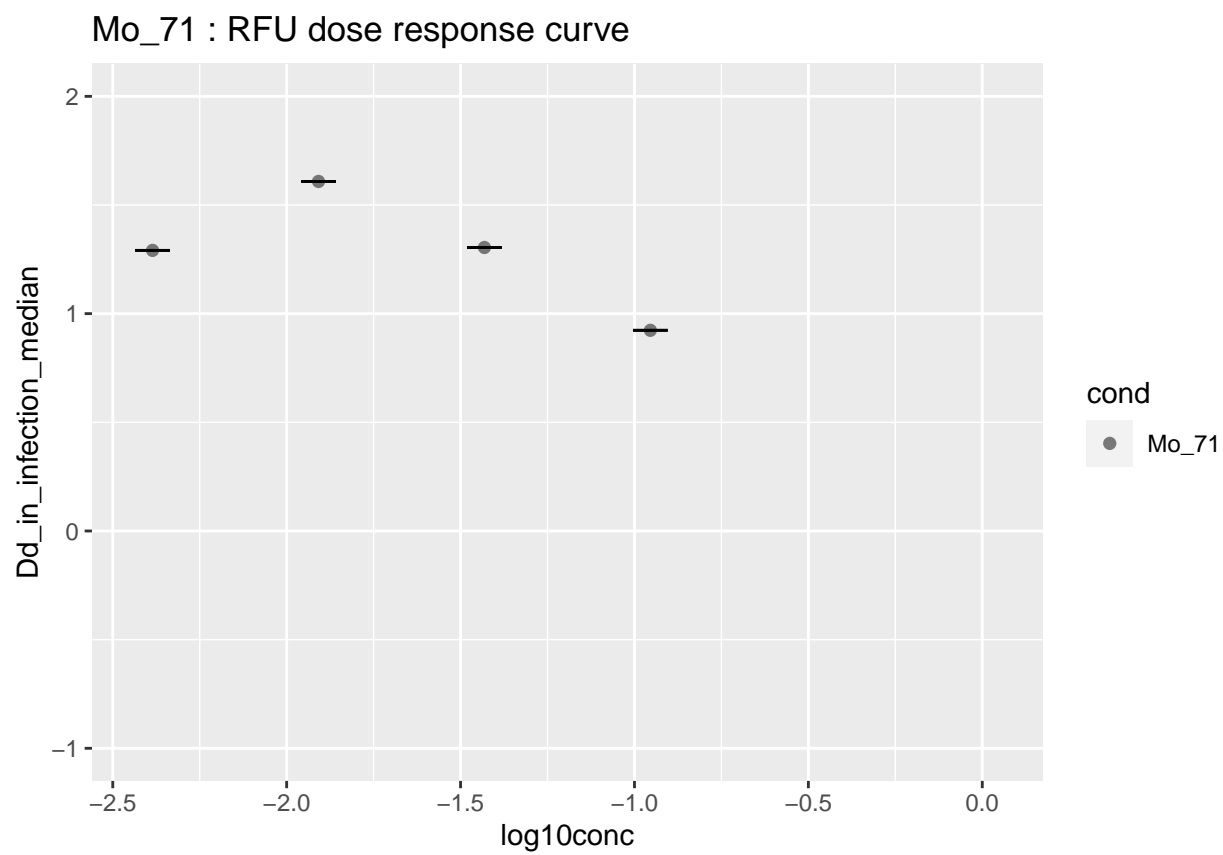


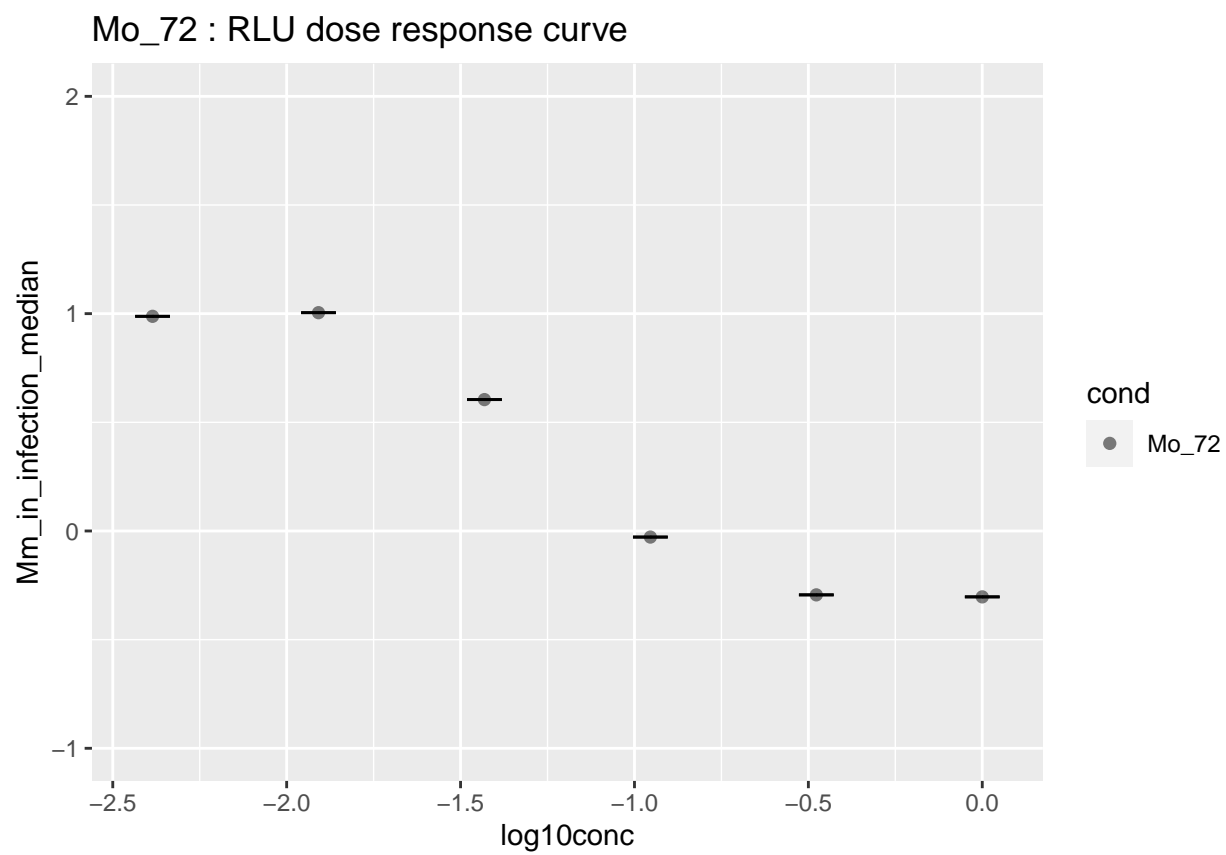


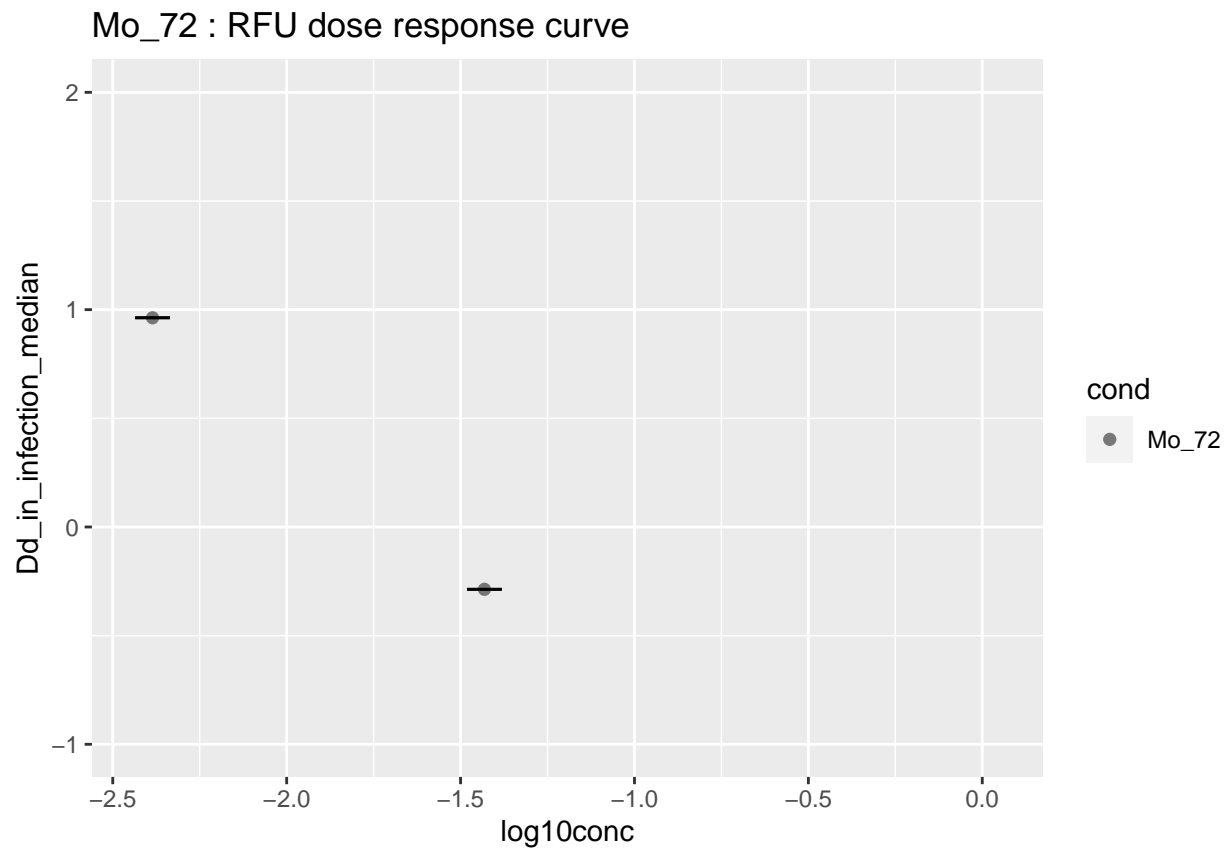


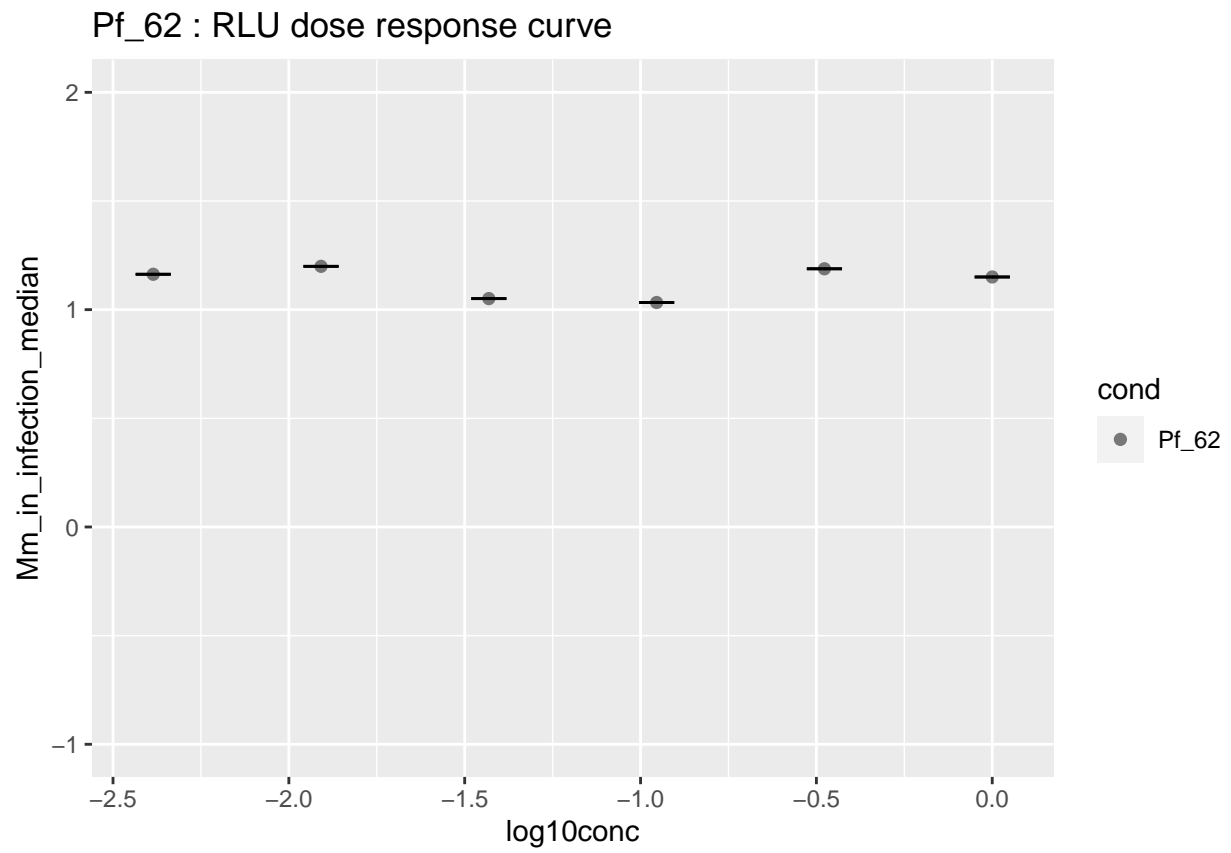


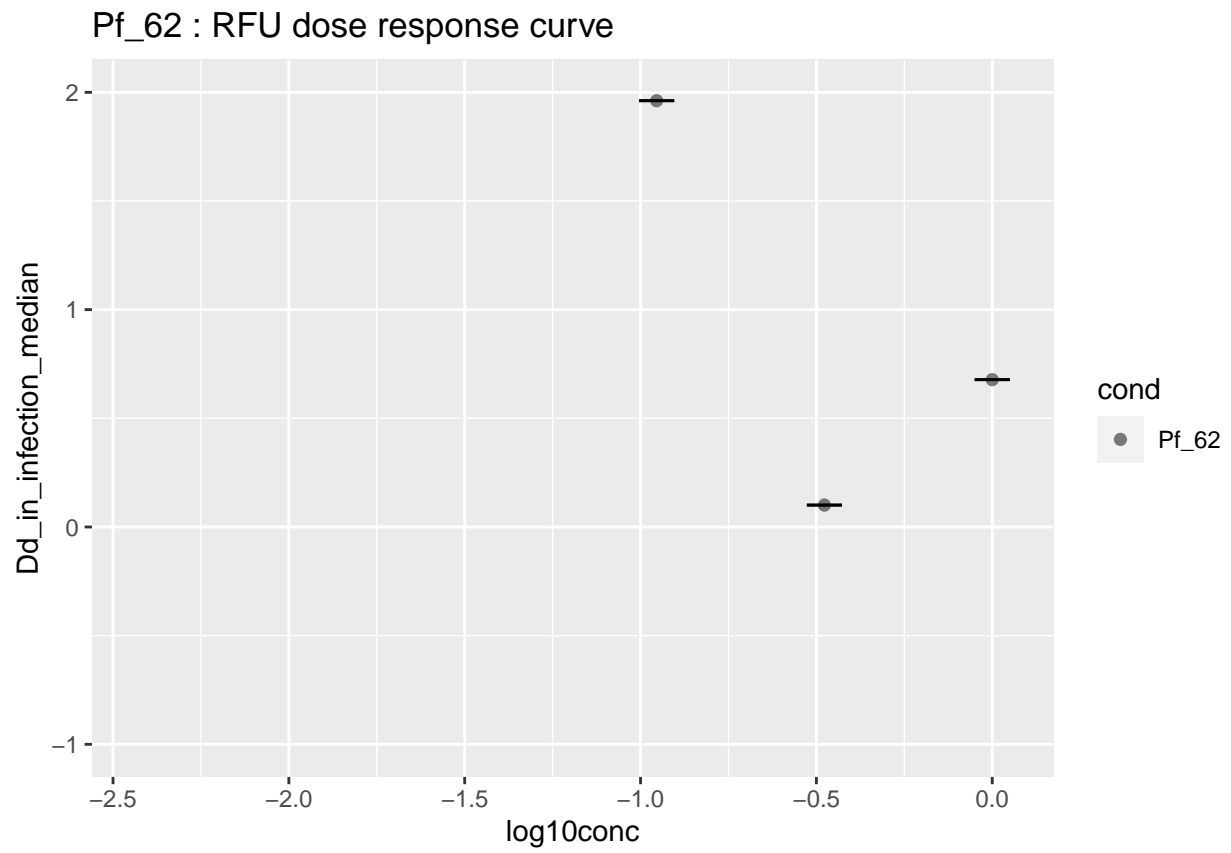


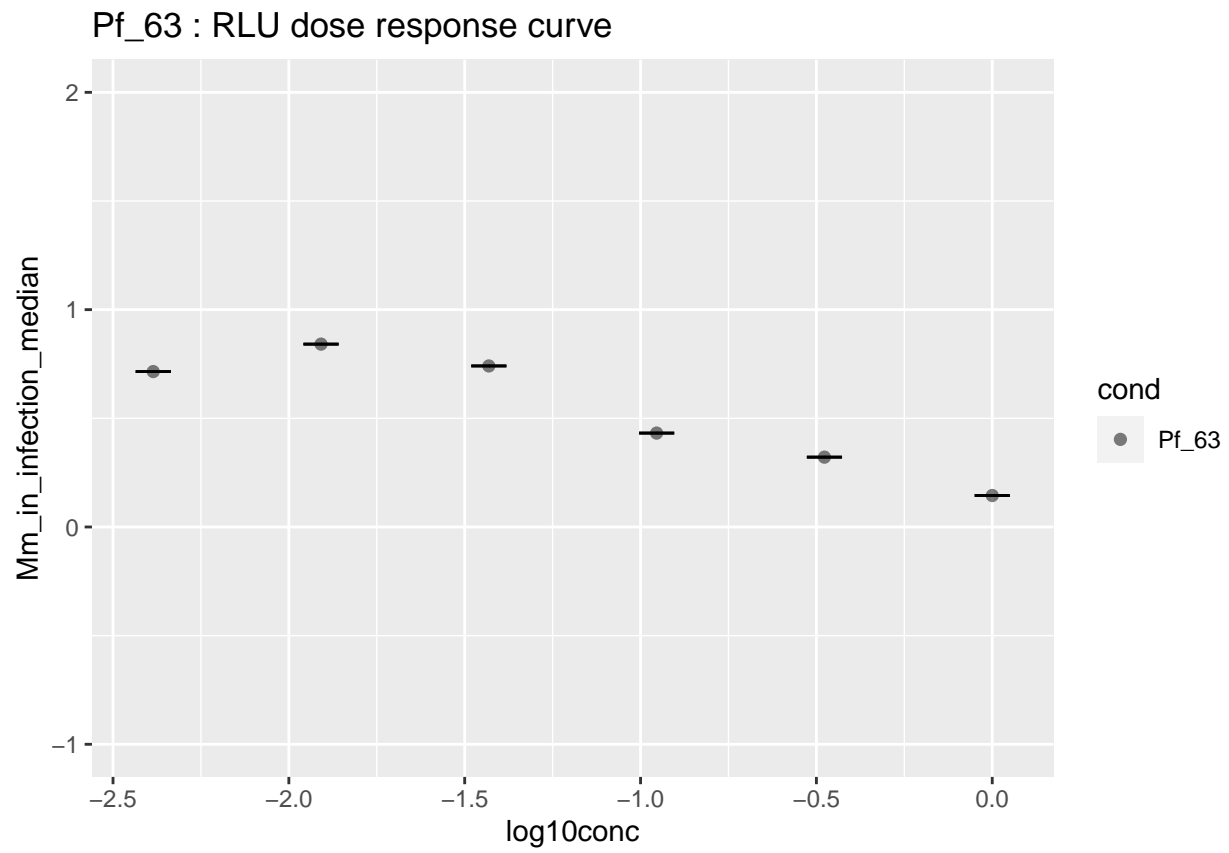


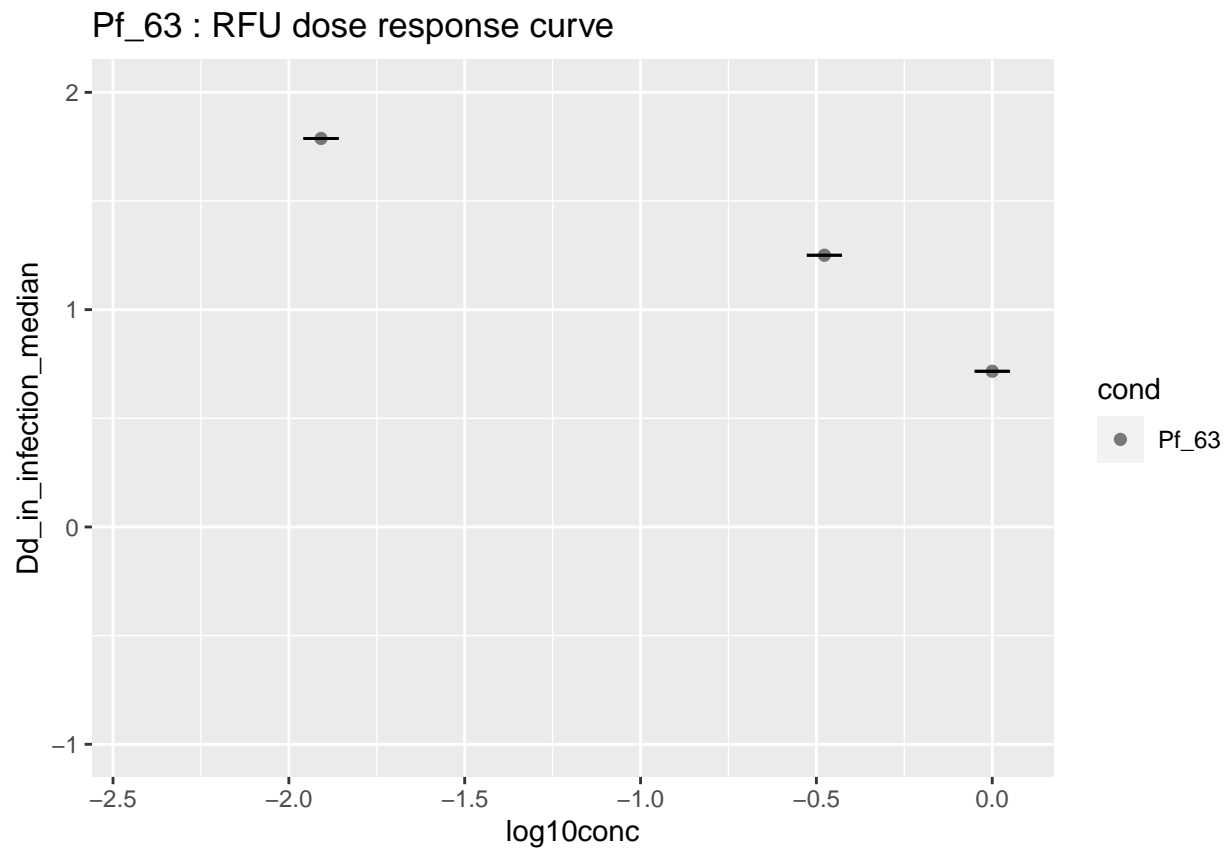


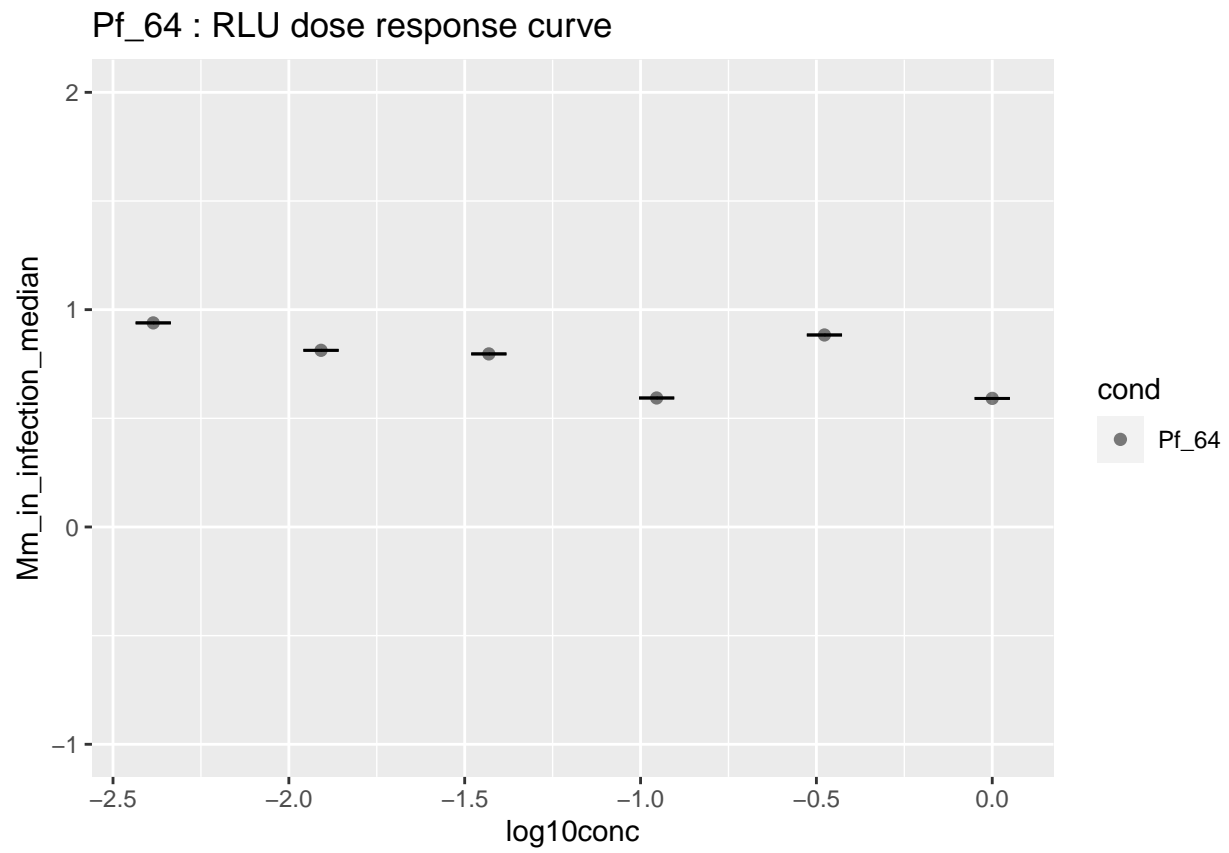


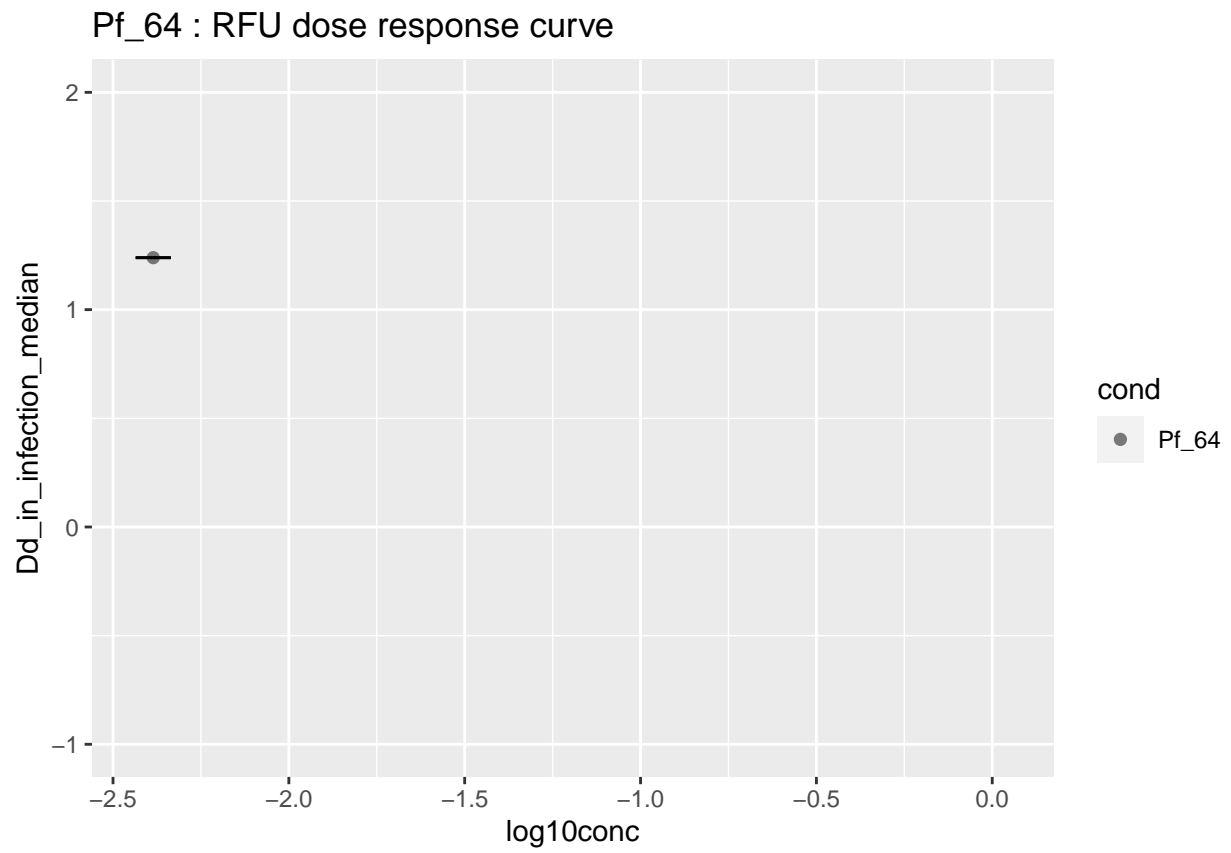


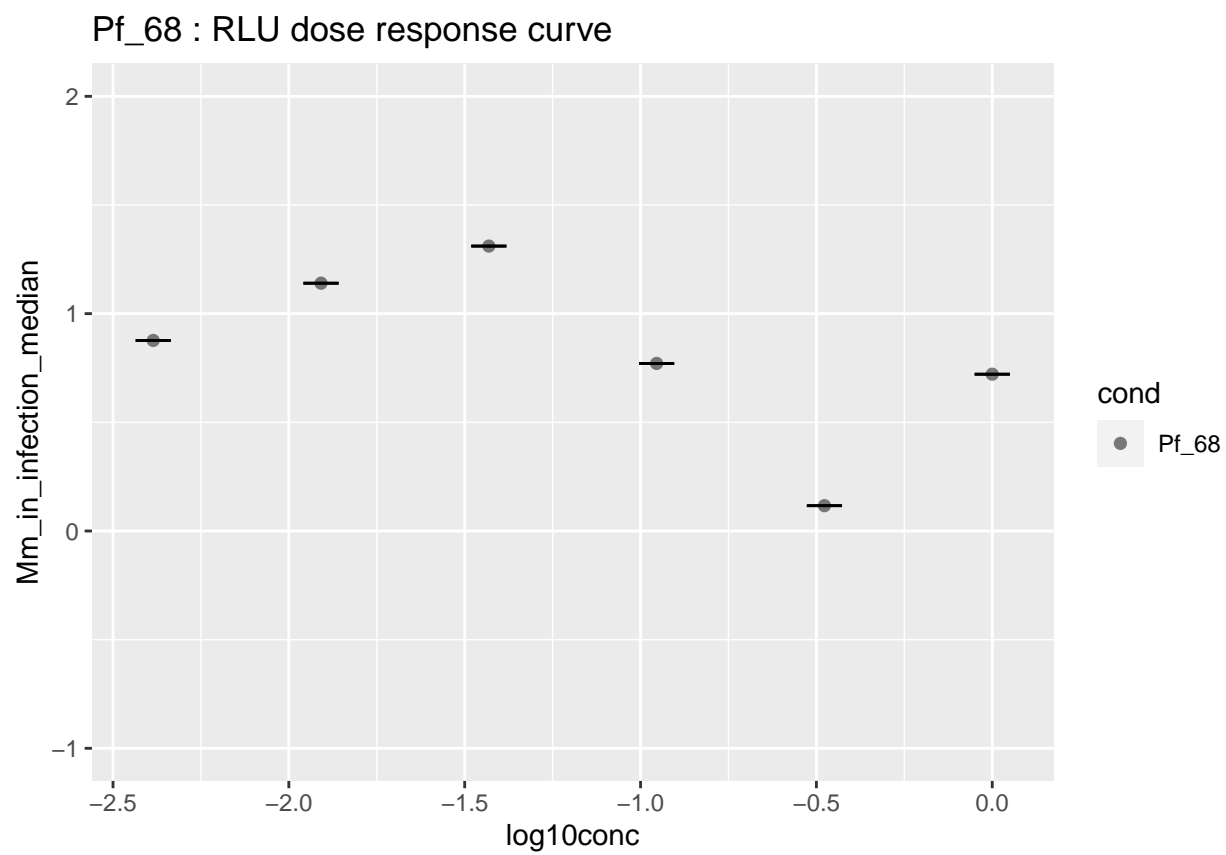


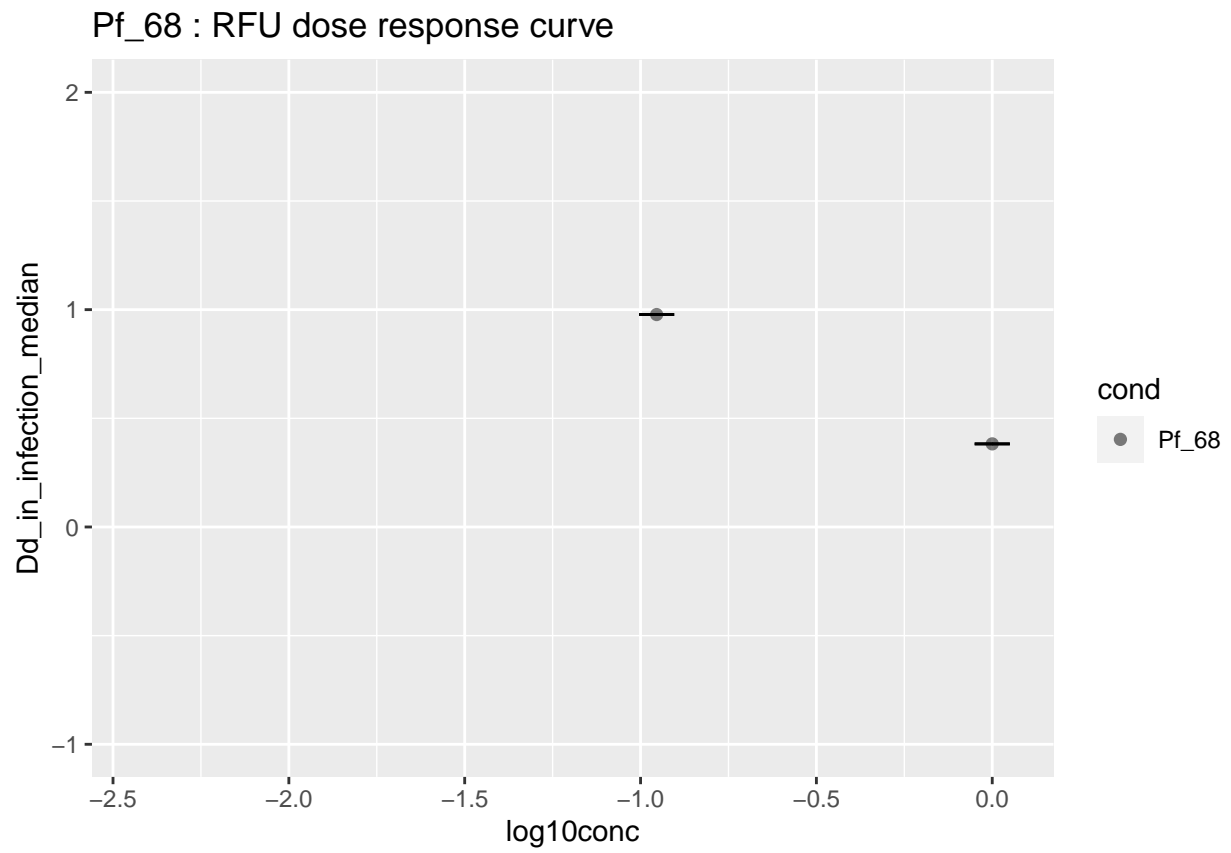


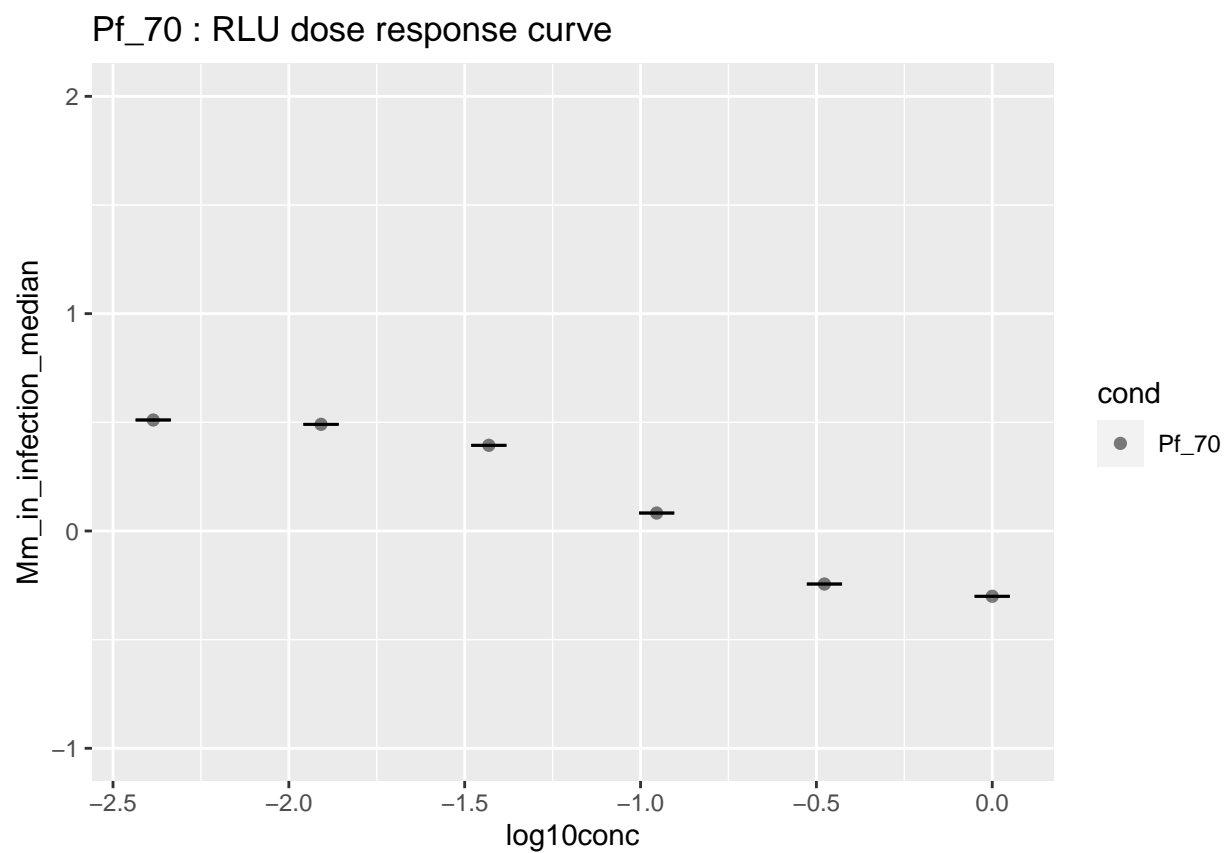


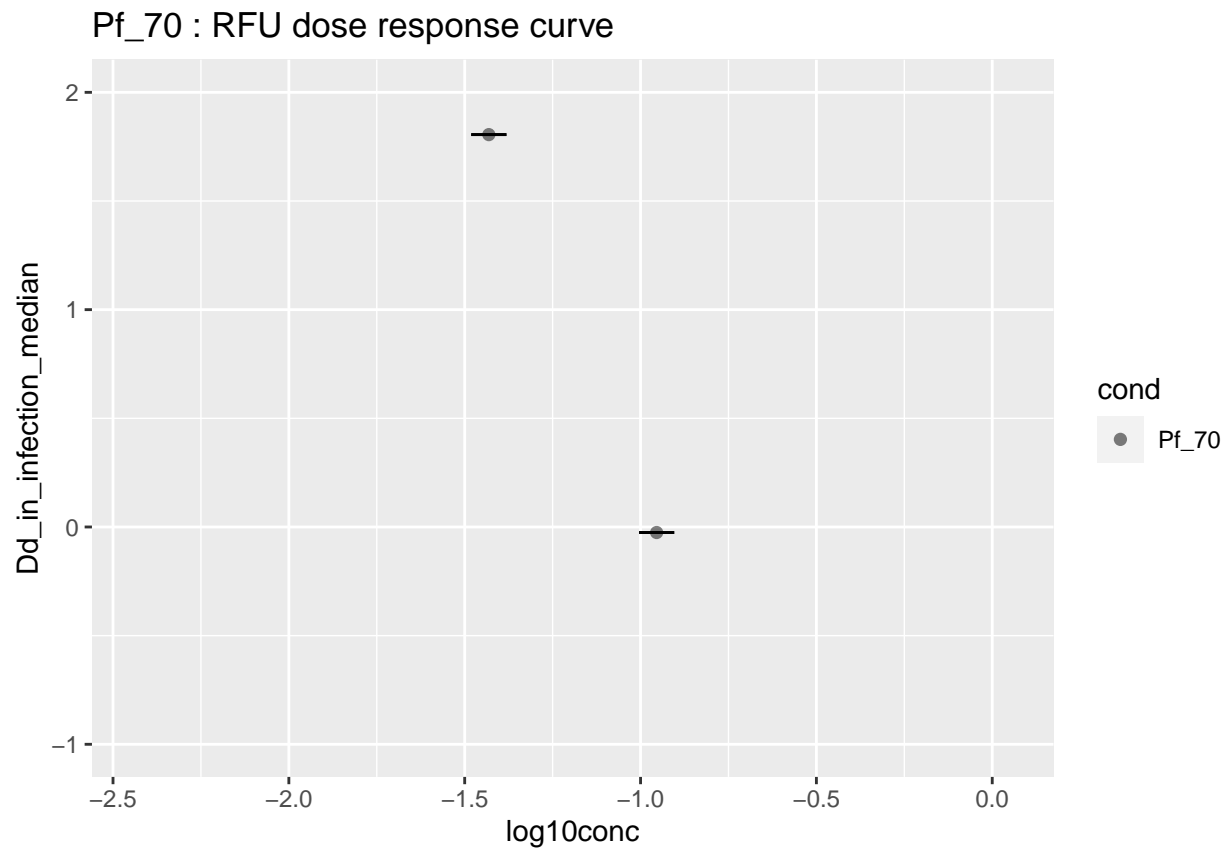


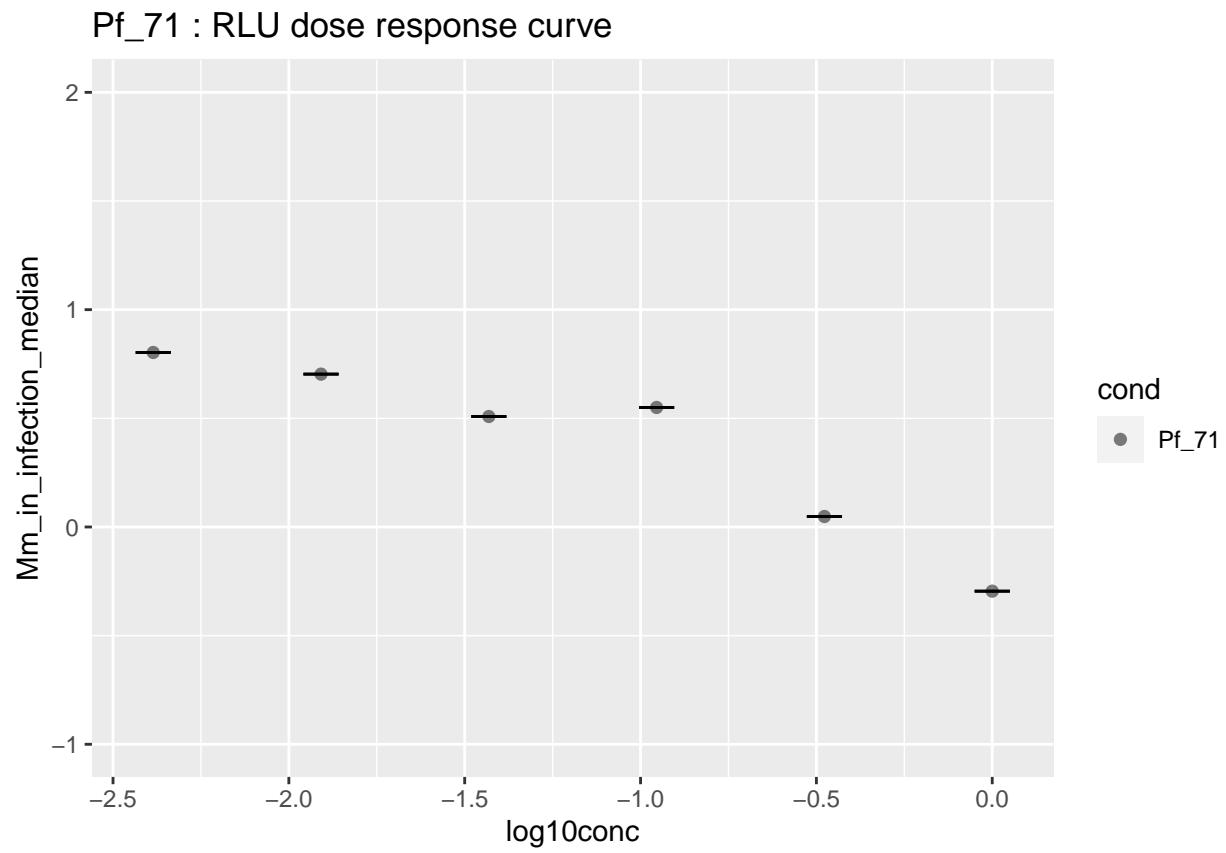


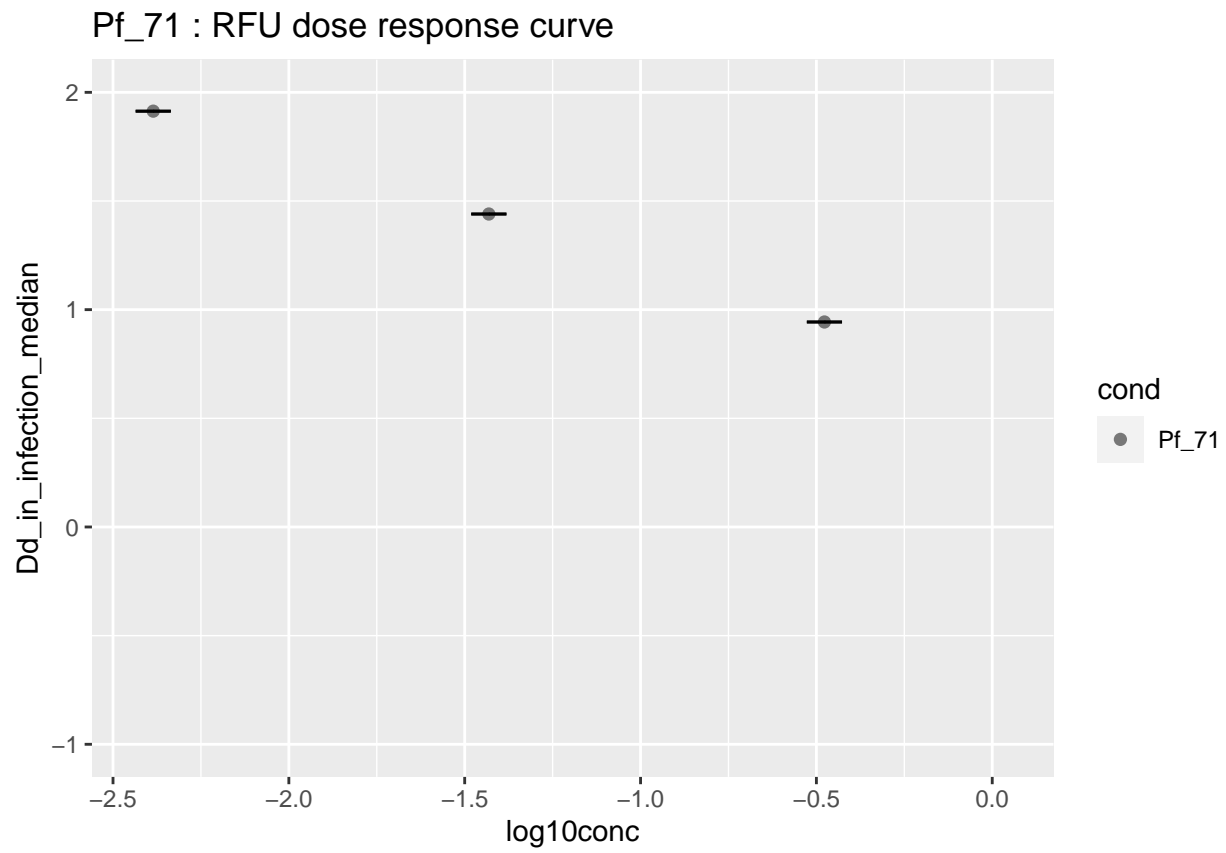


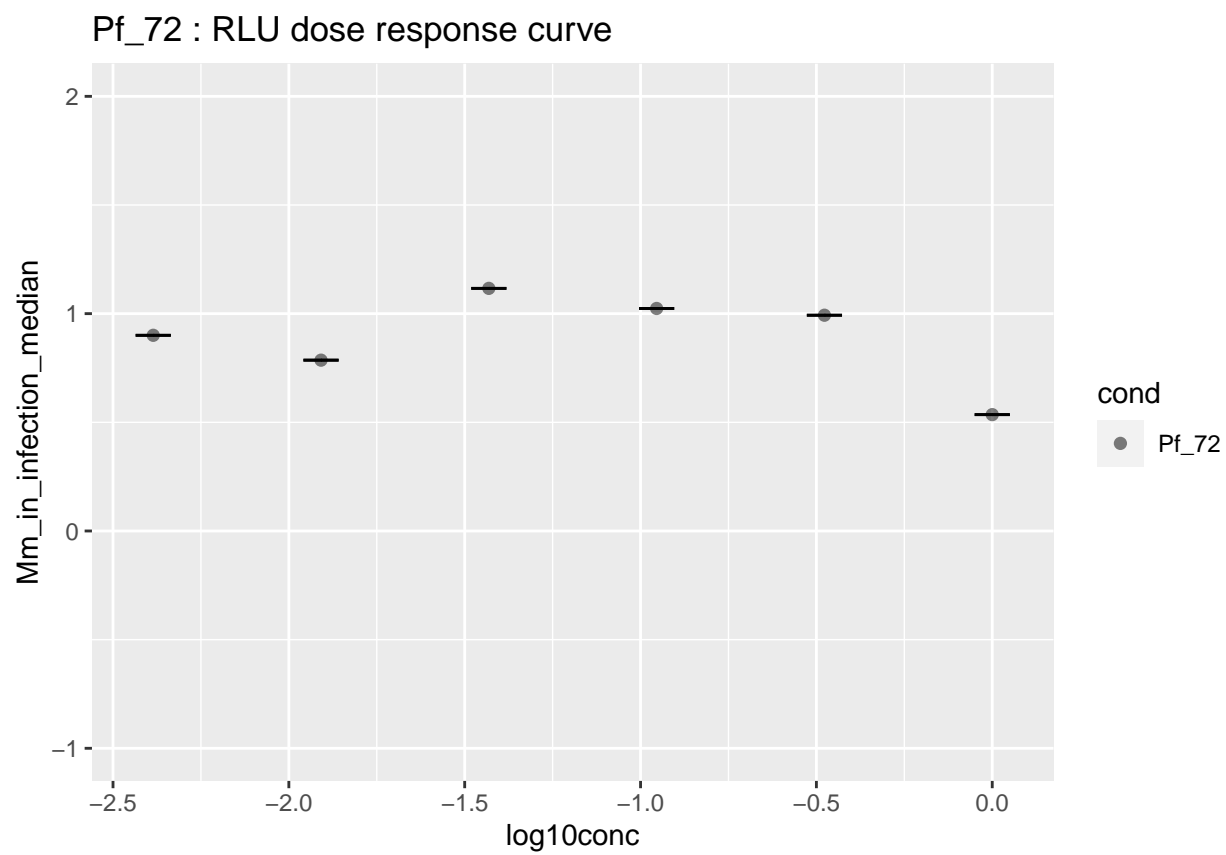


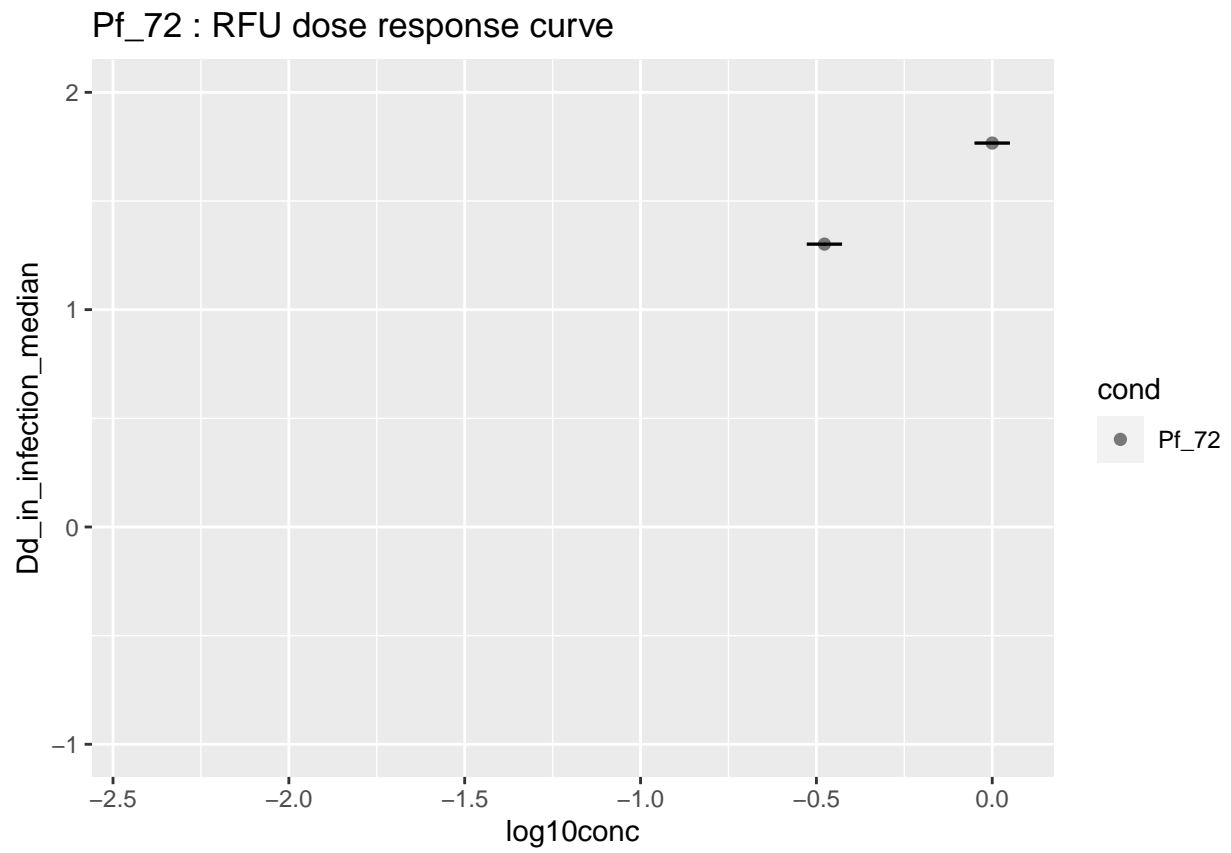


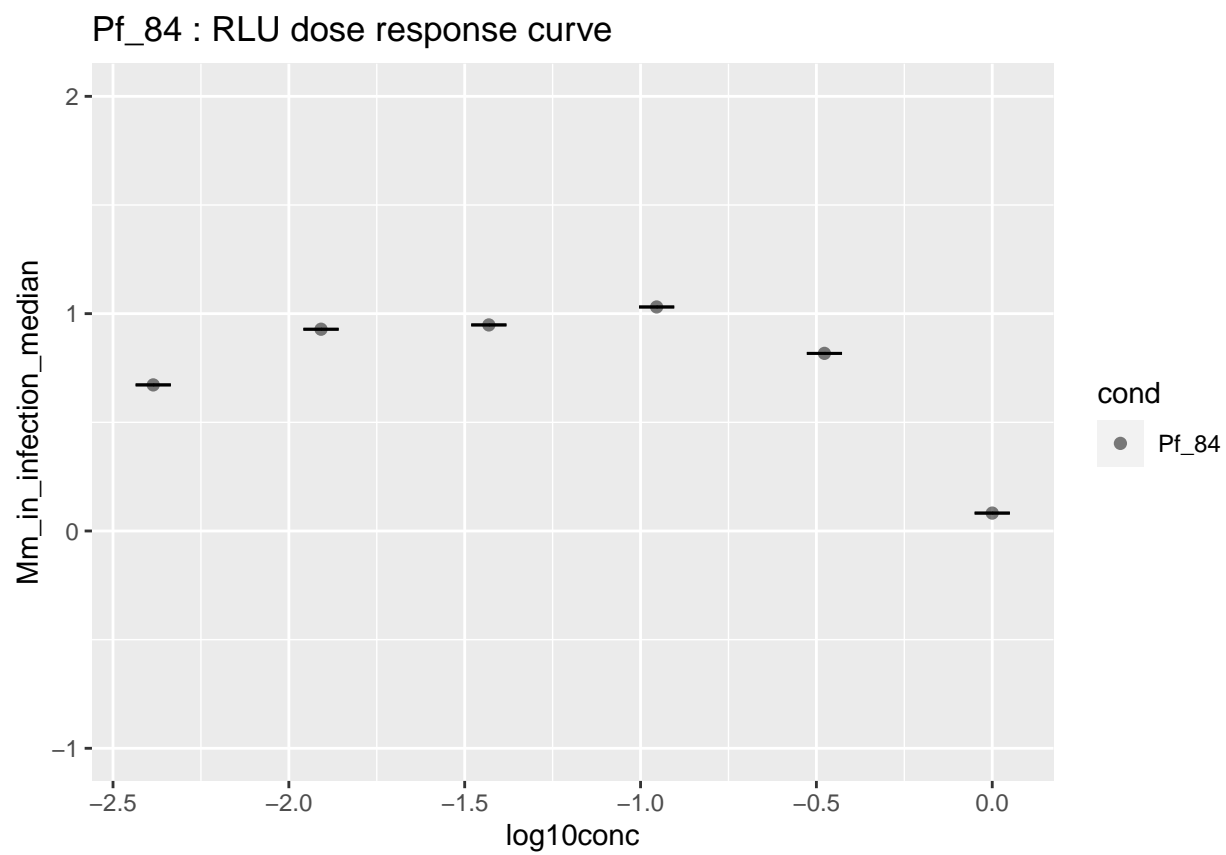


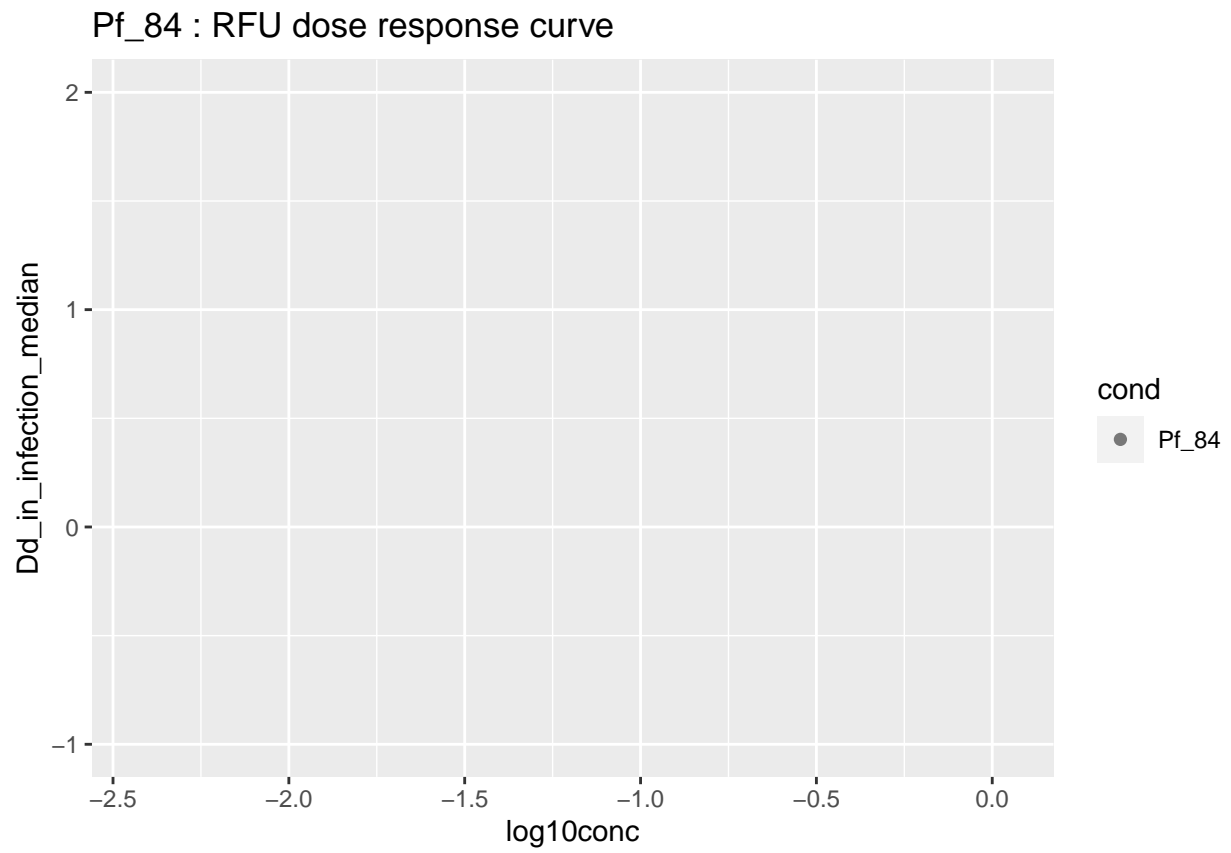




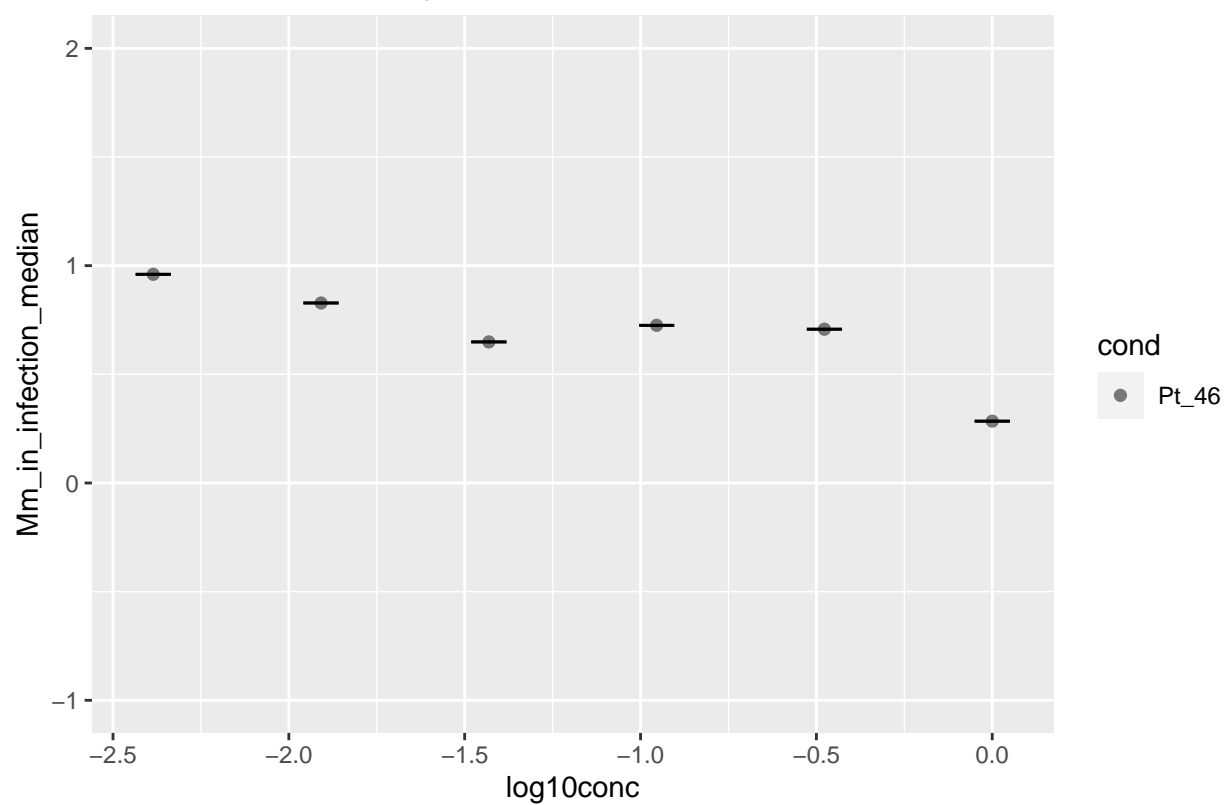


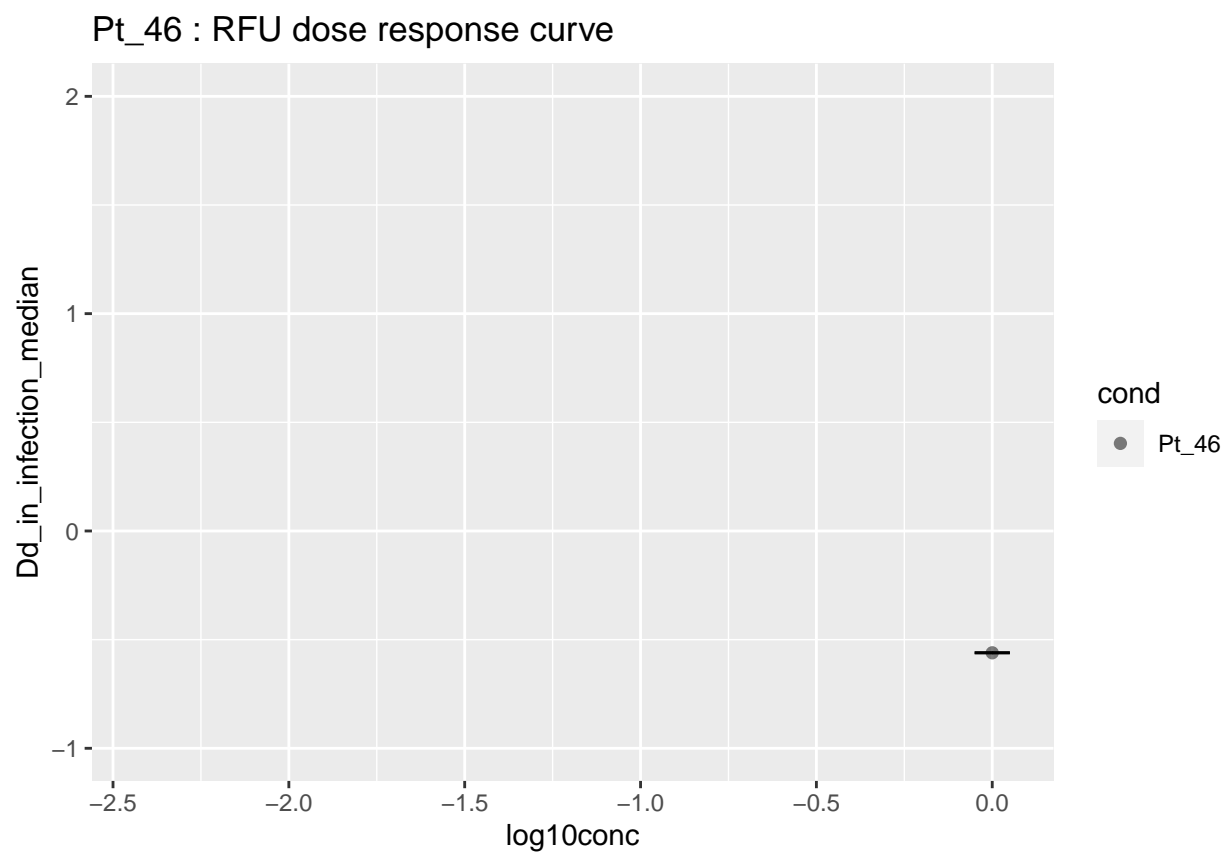


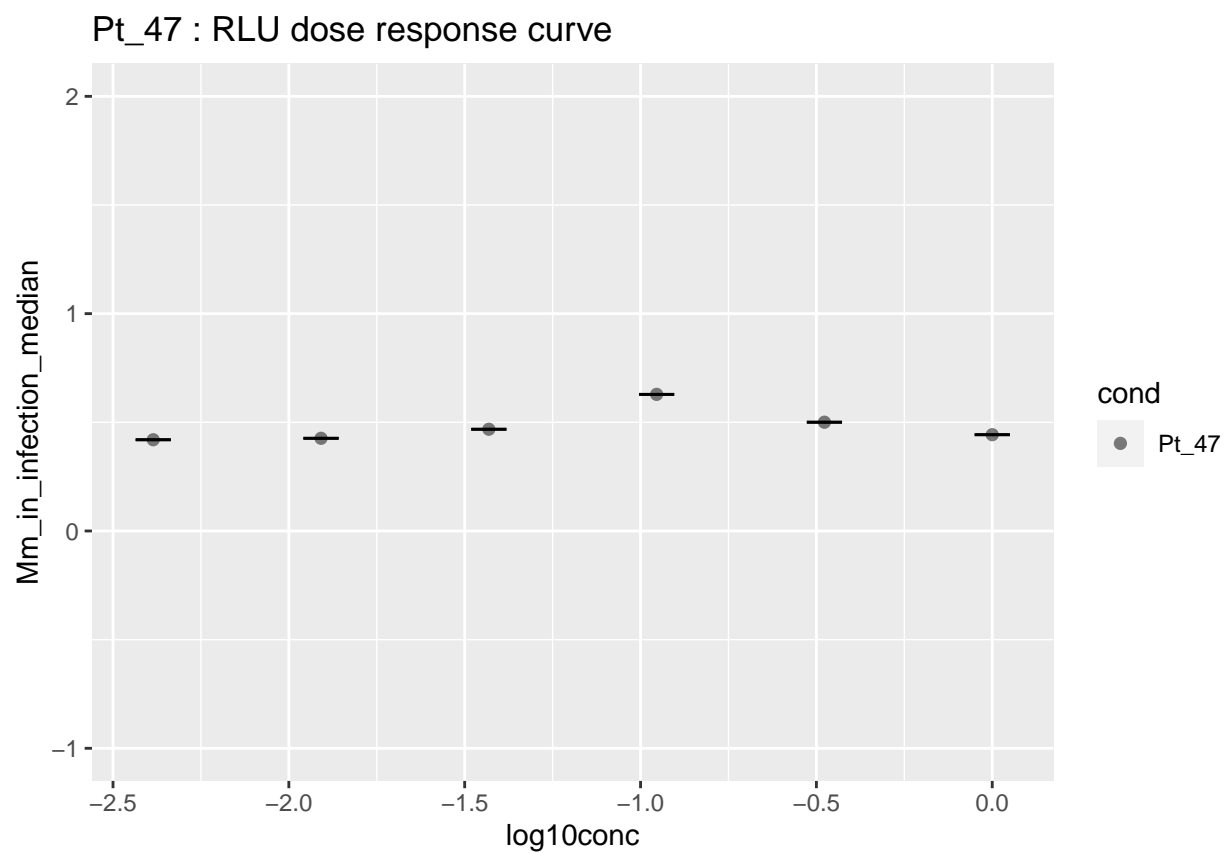


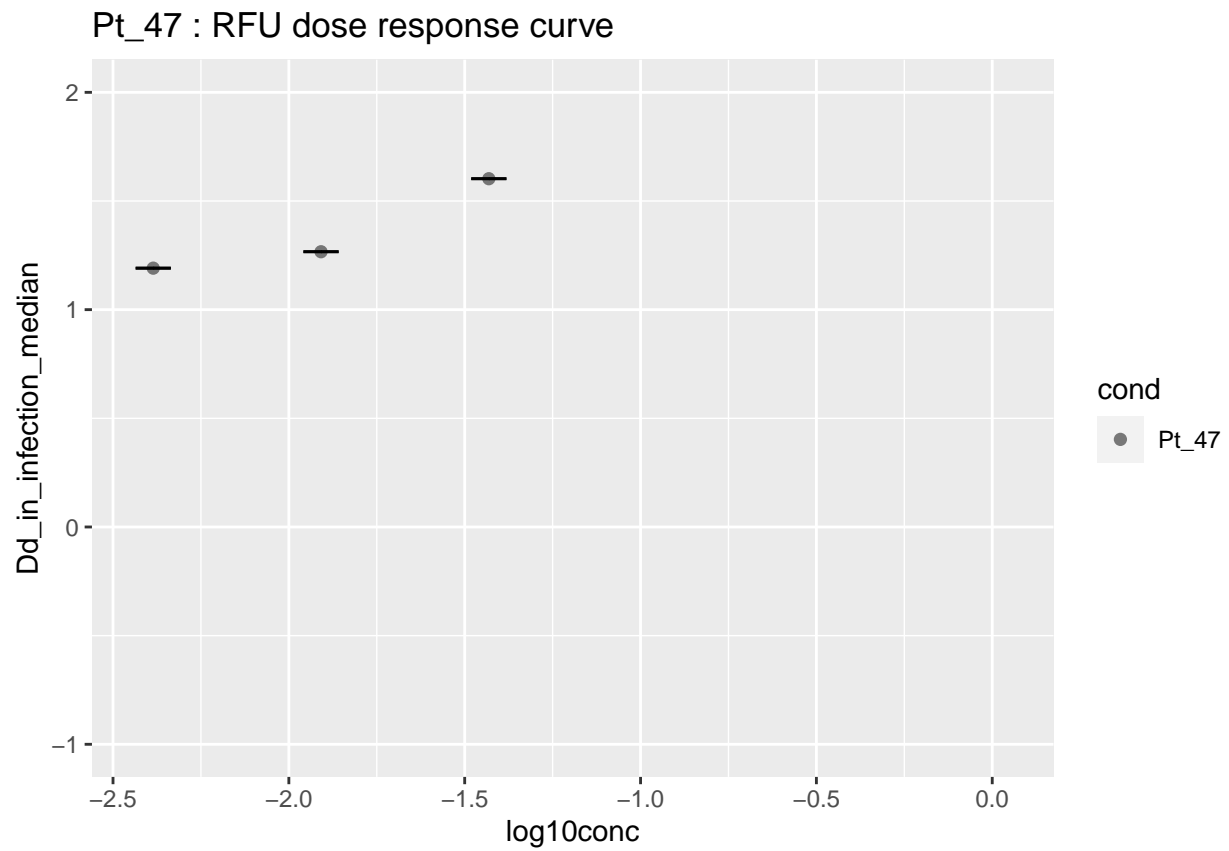


Pt_46 : RLU dose response curve

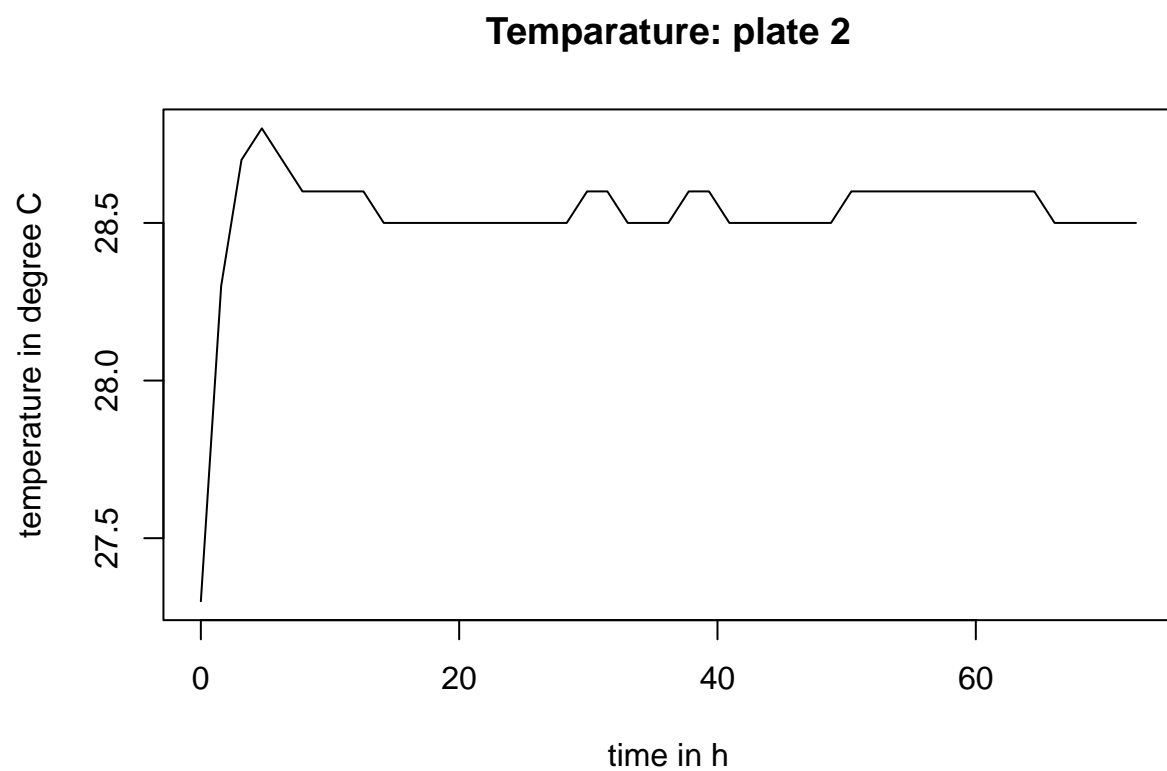




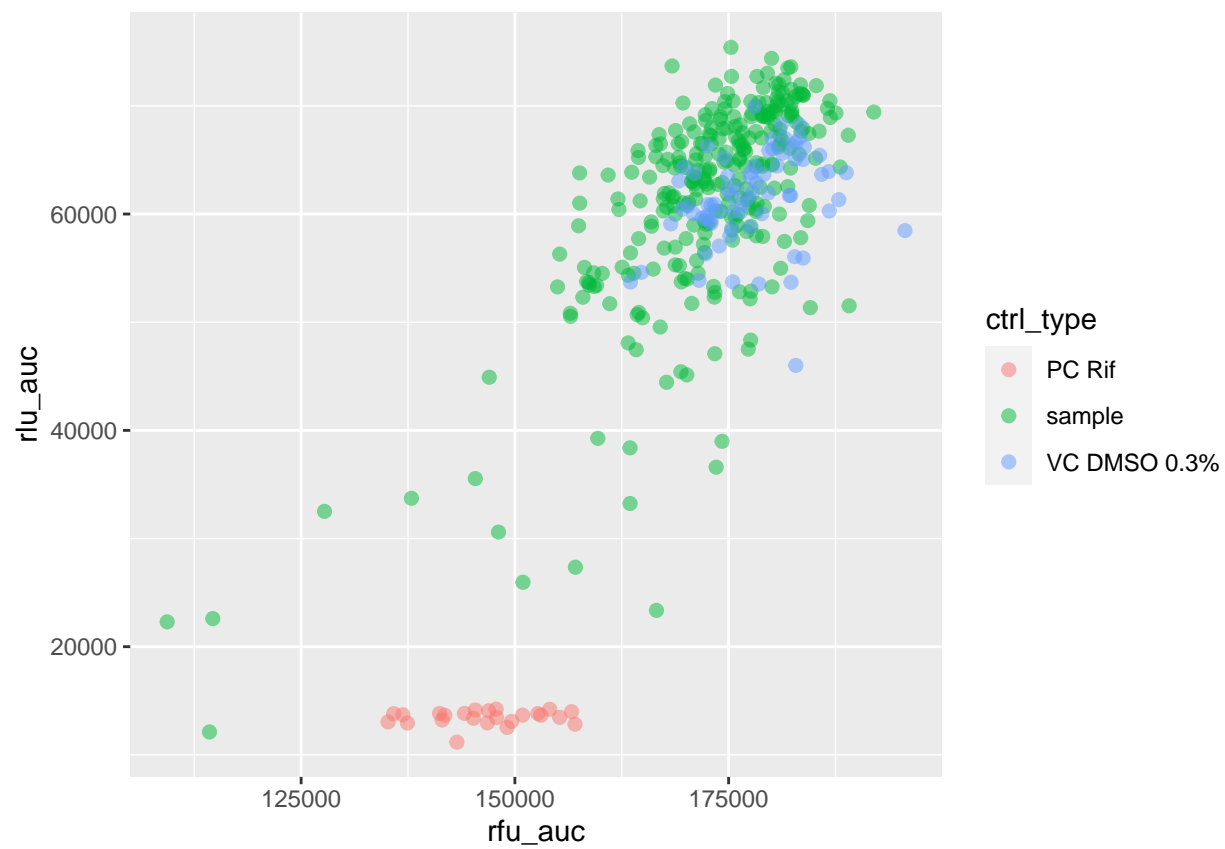


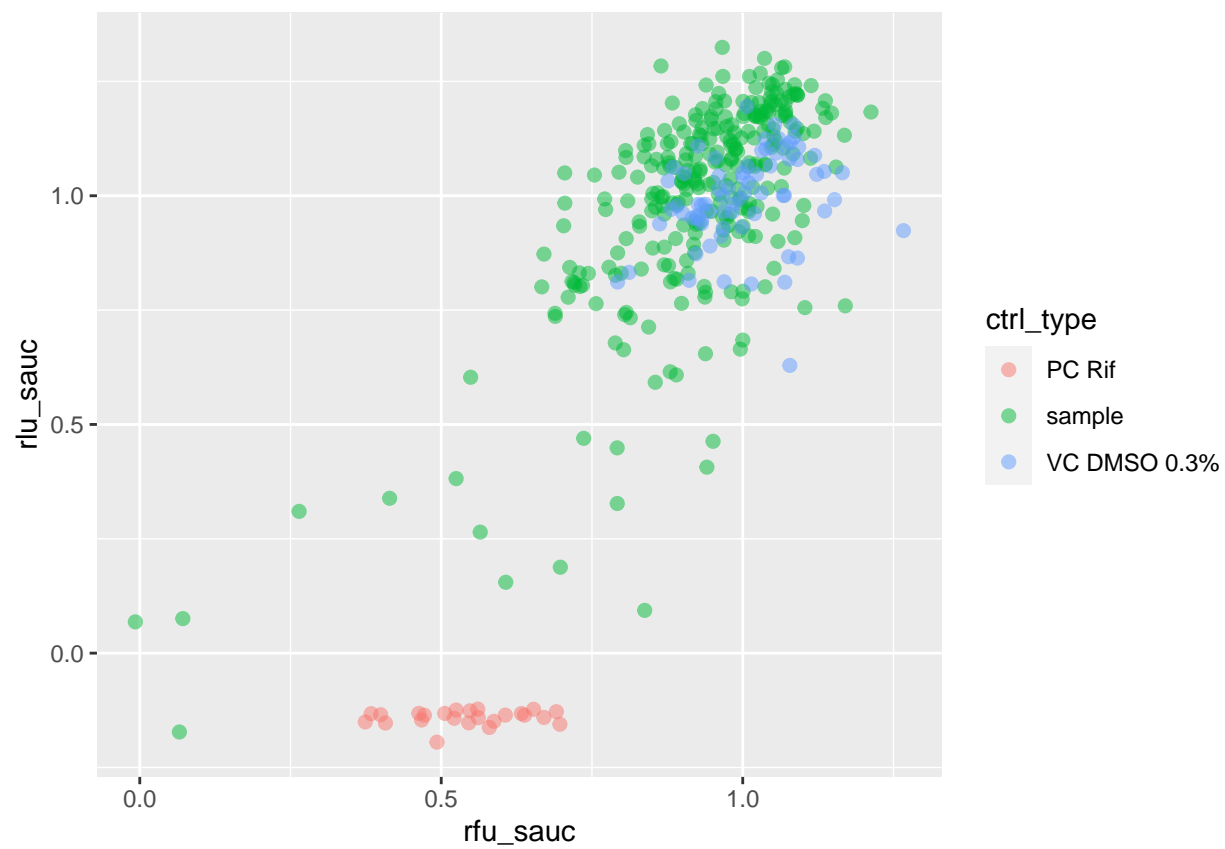


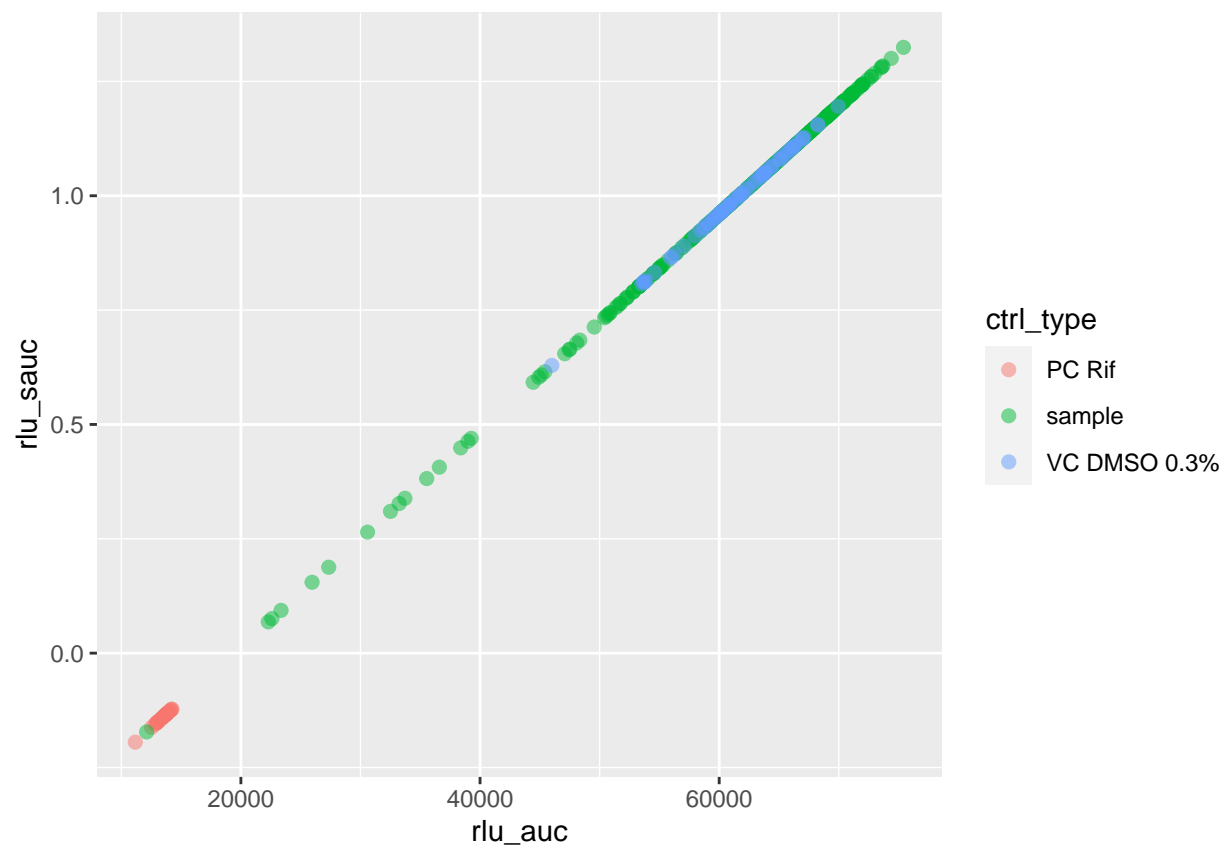
```
## [1] "analysis for plate03.xlsx"
```

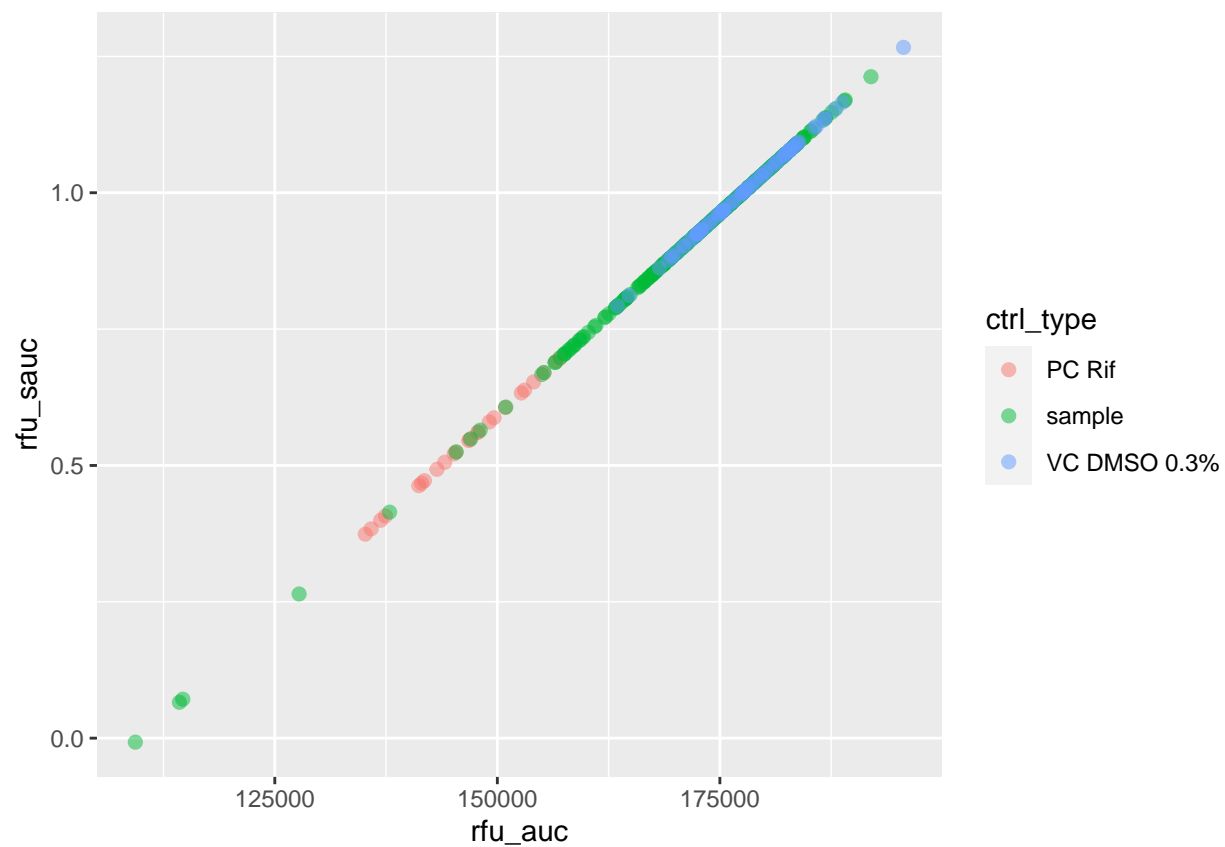


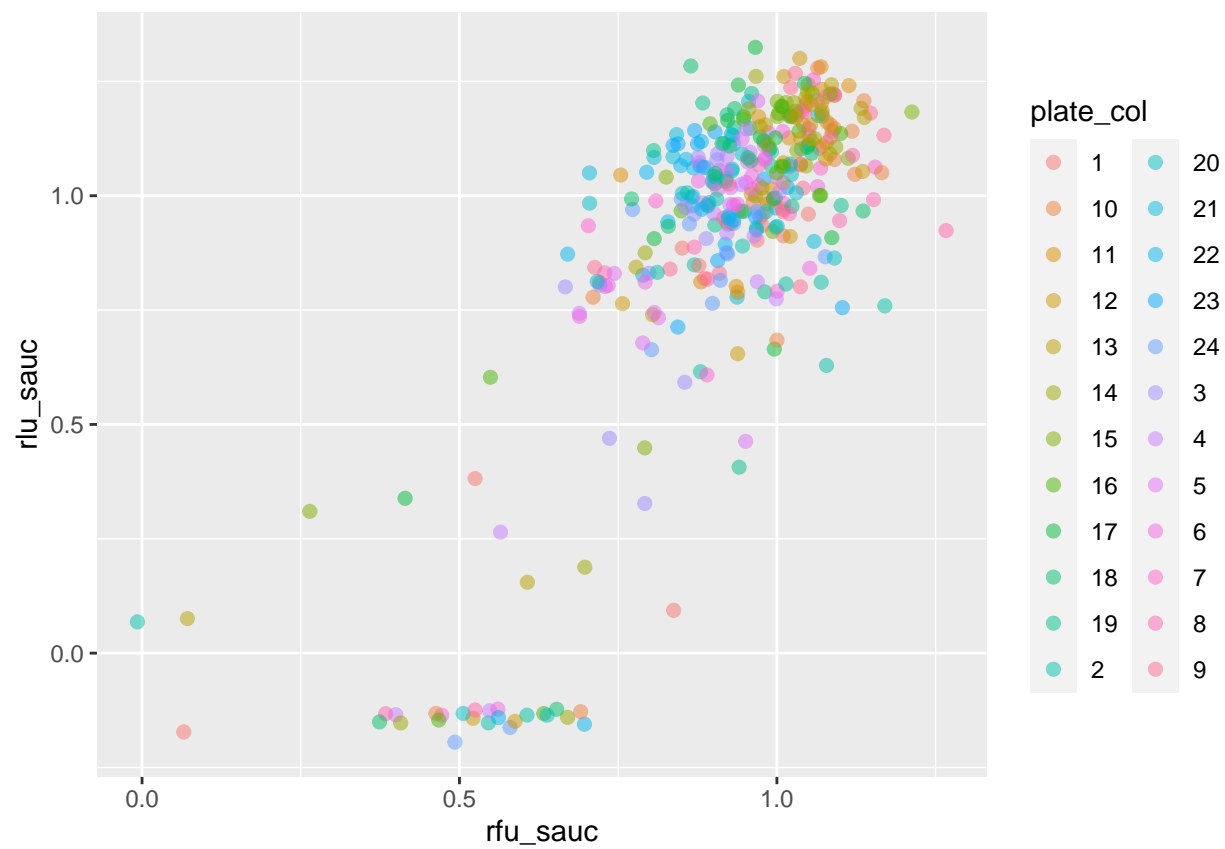
```
## [1] "VC DMSO 0.3% Robust z'-factor of rlu for plate plate03.xlsx, biorep 1 : "  
## [1] 0.73
```

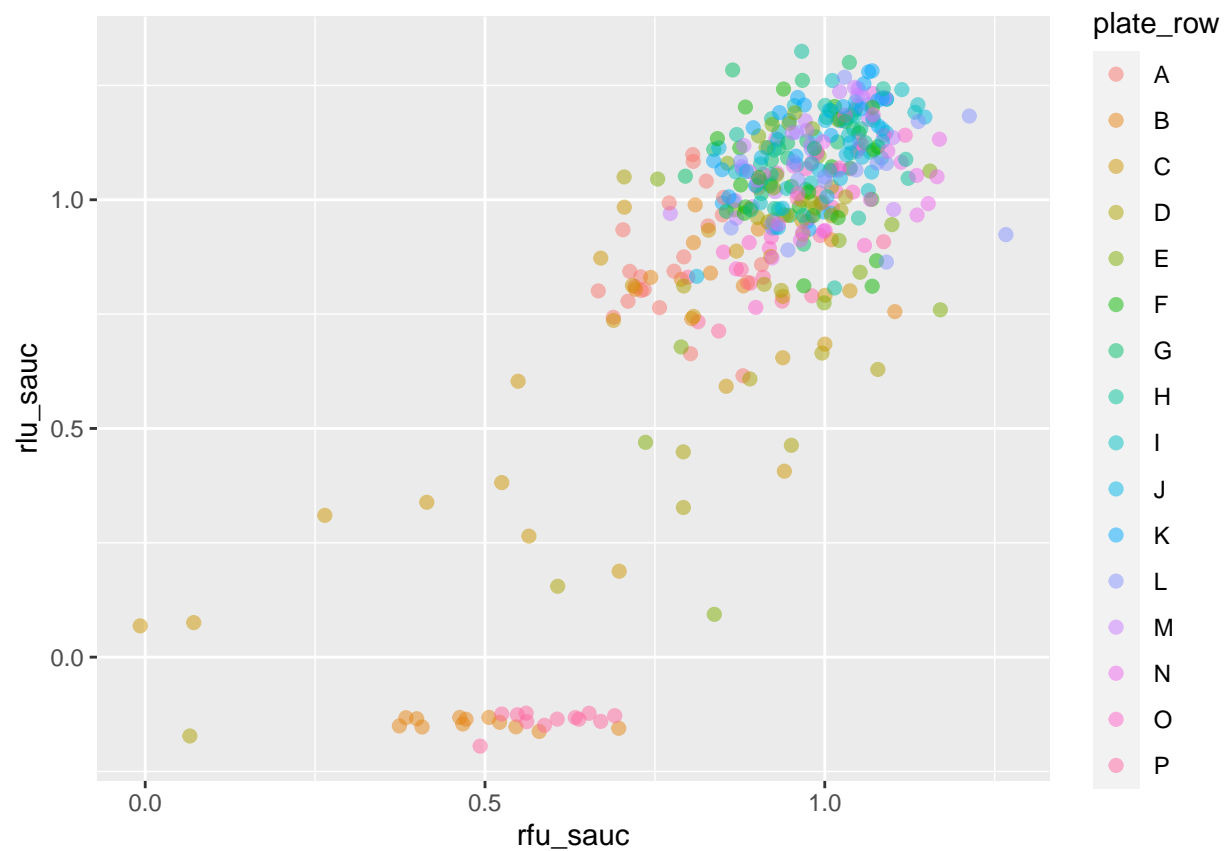





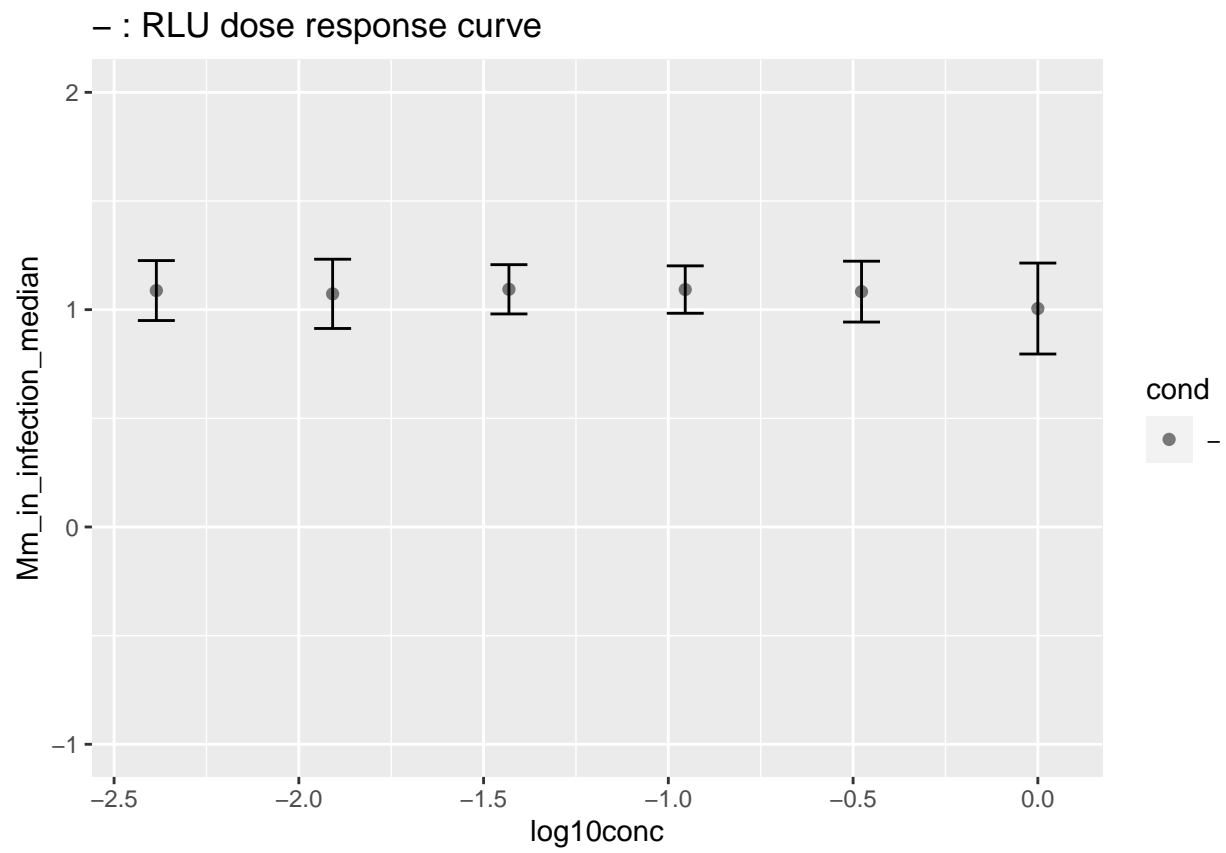


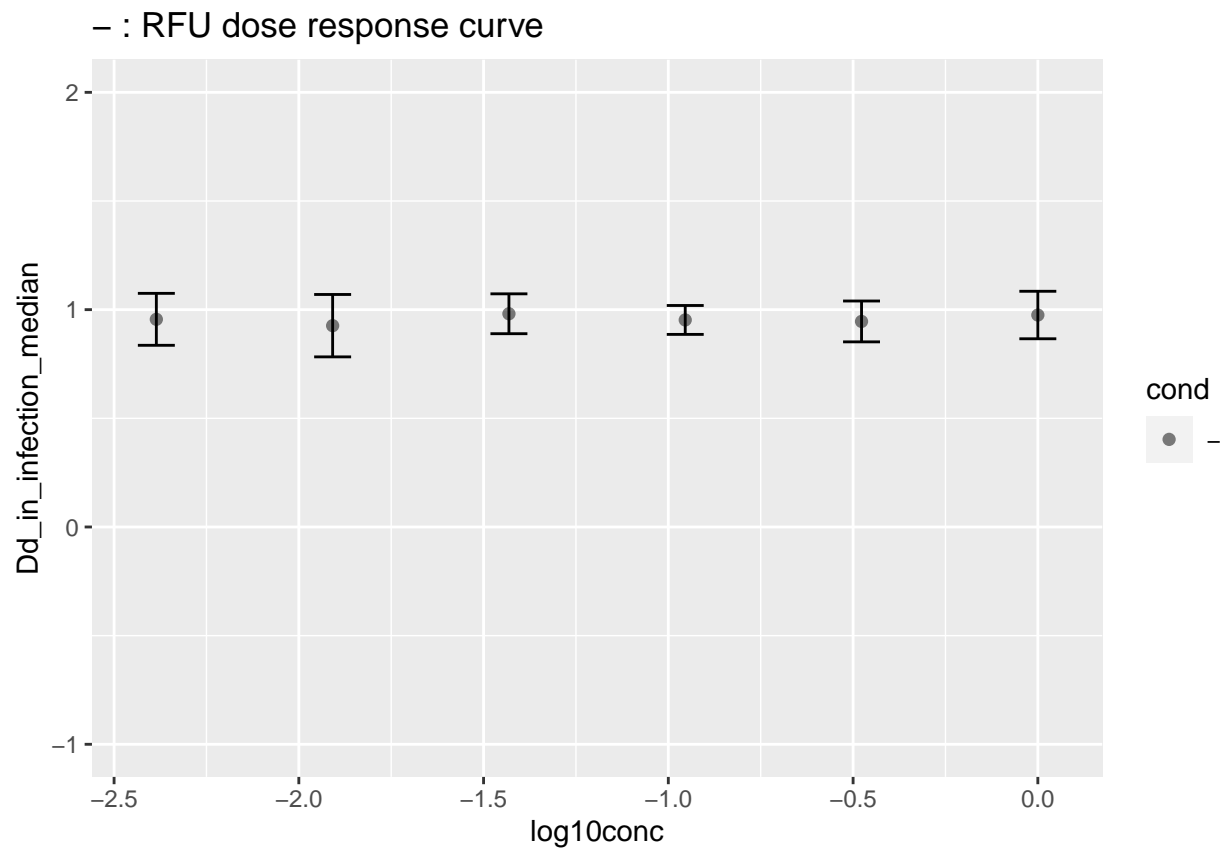


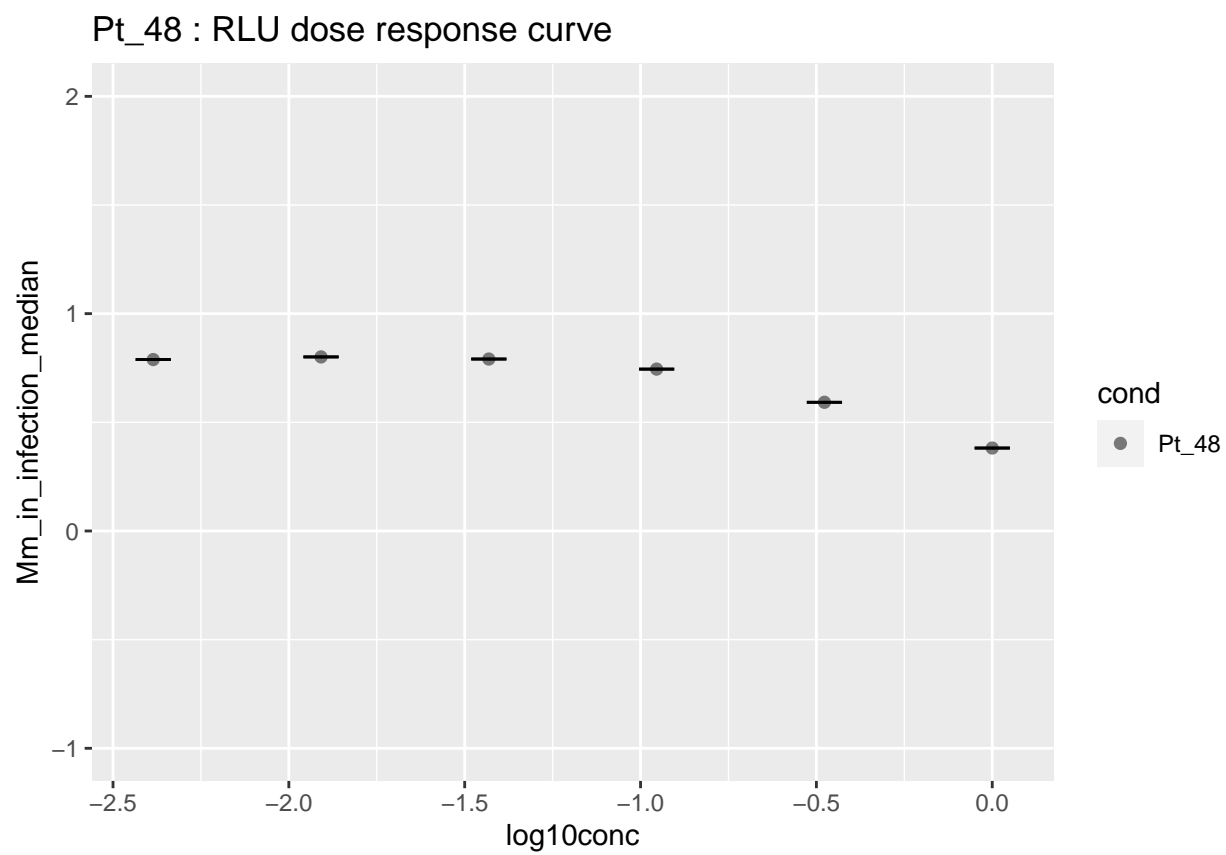


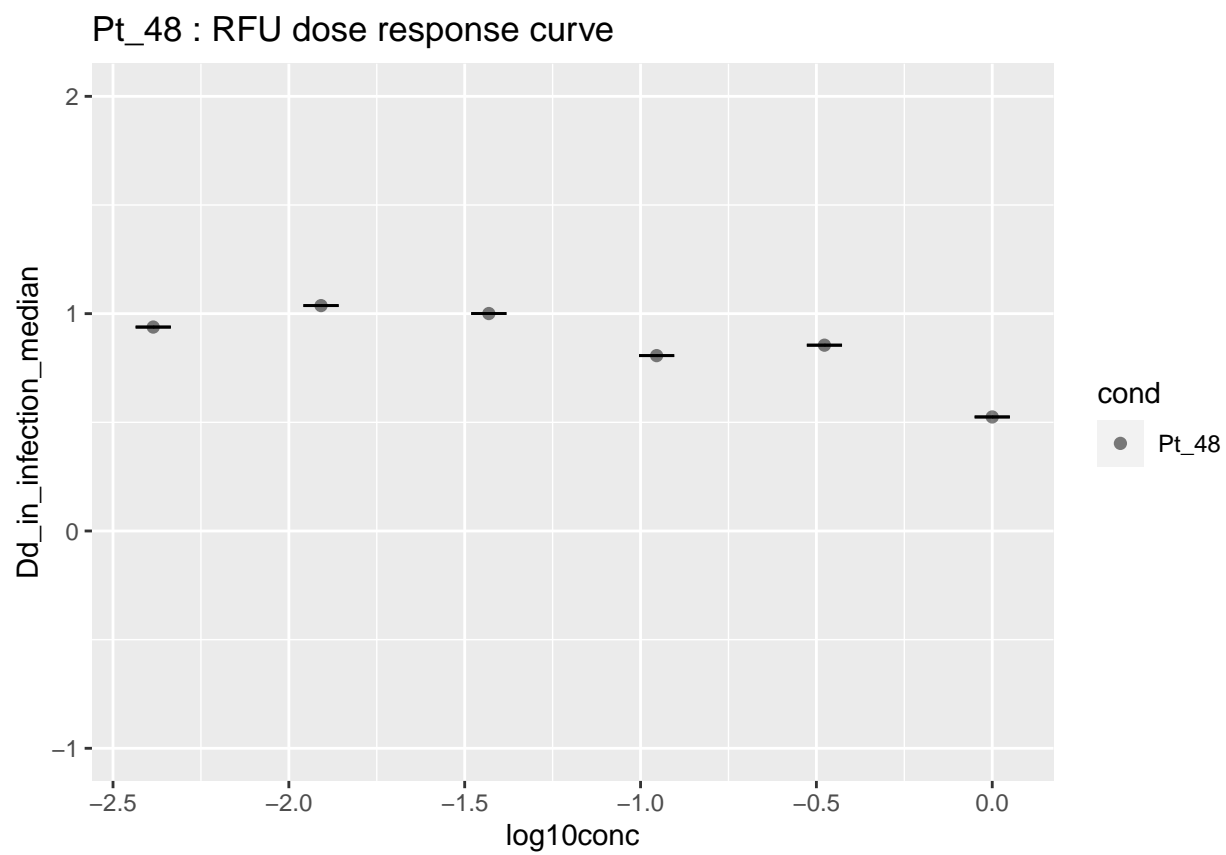


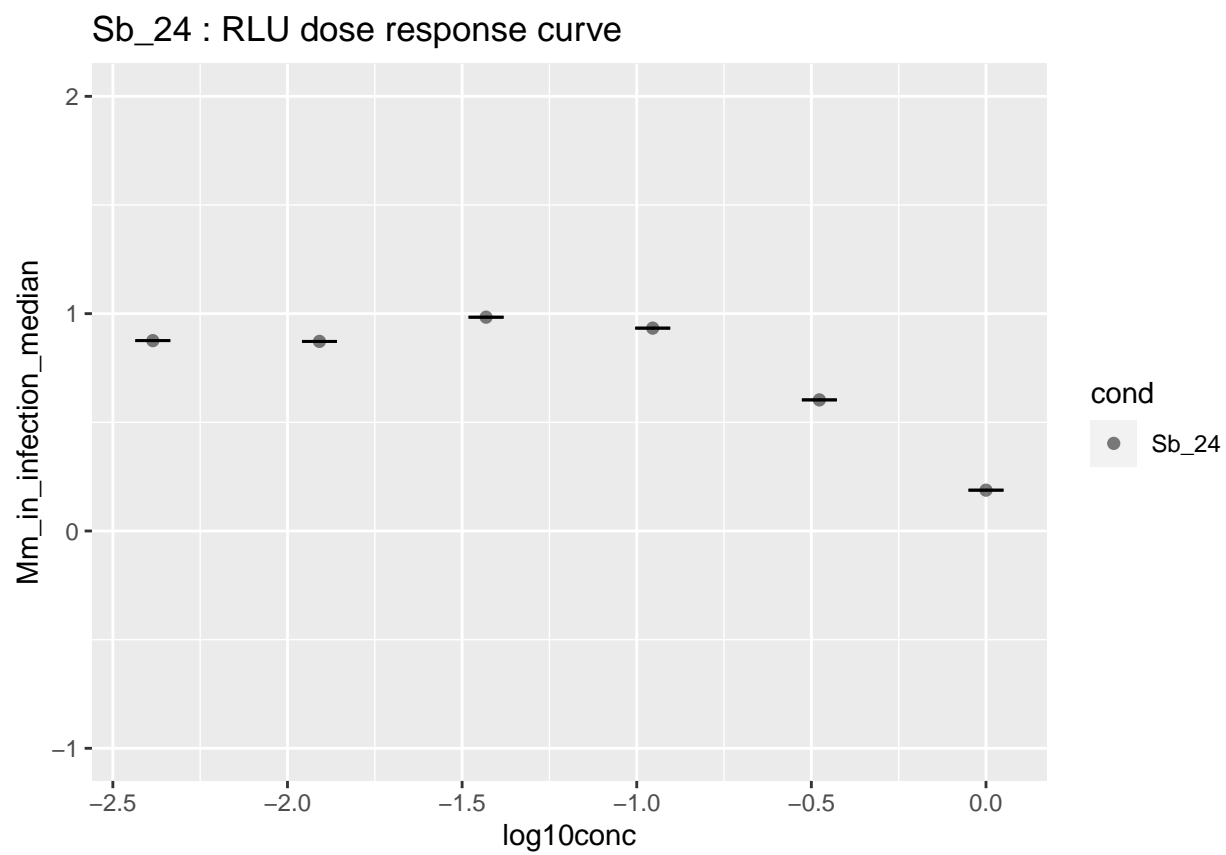
```
## [1] "Dose response curves over all bioreps within this plate"
```

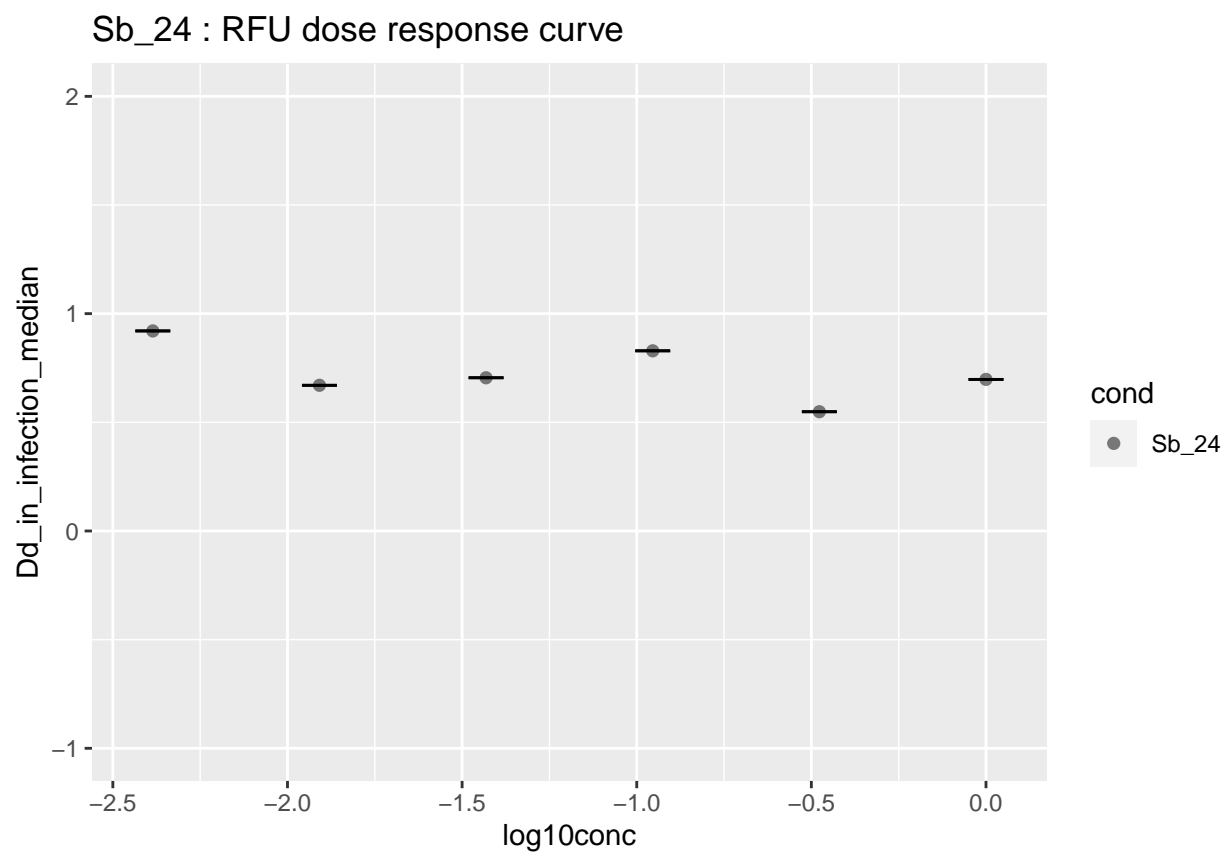


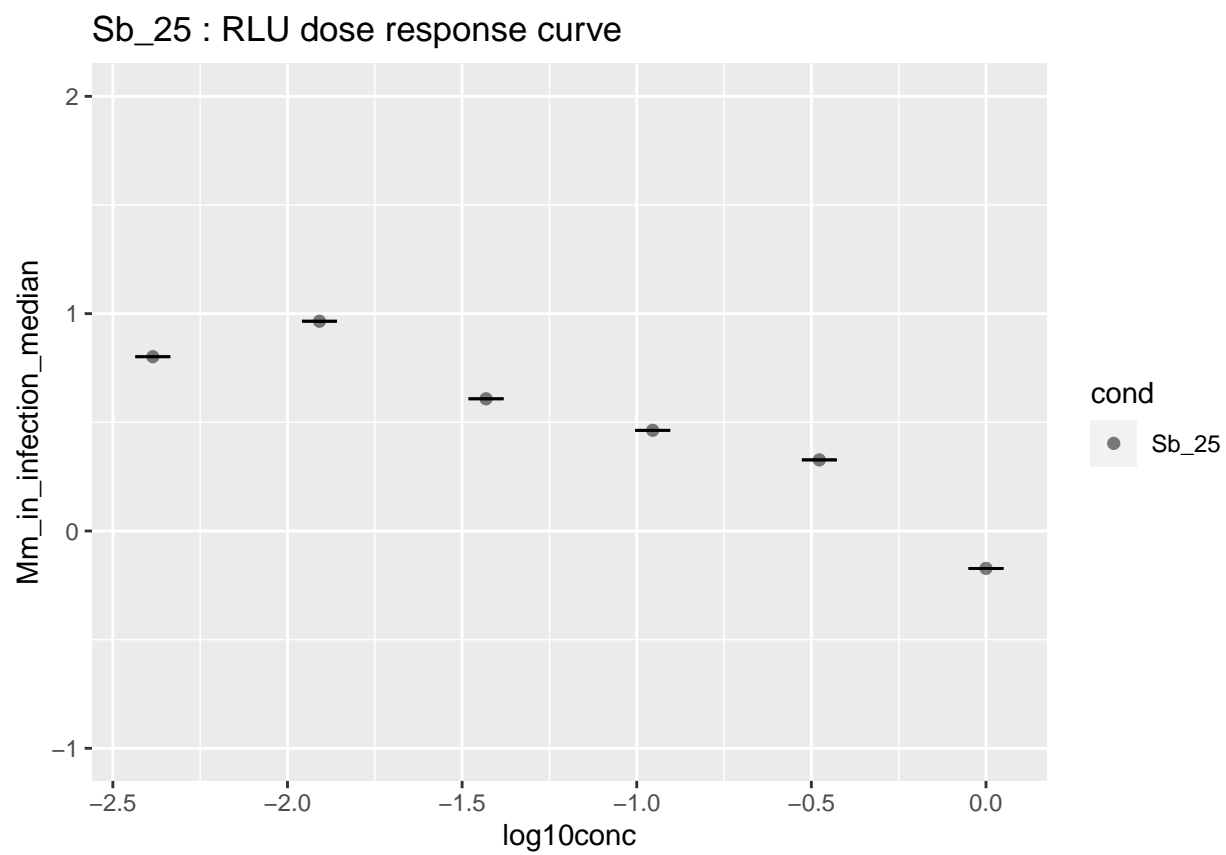


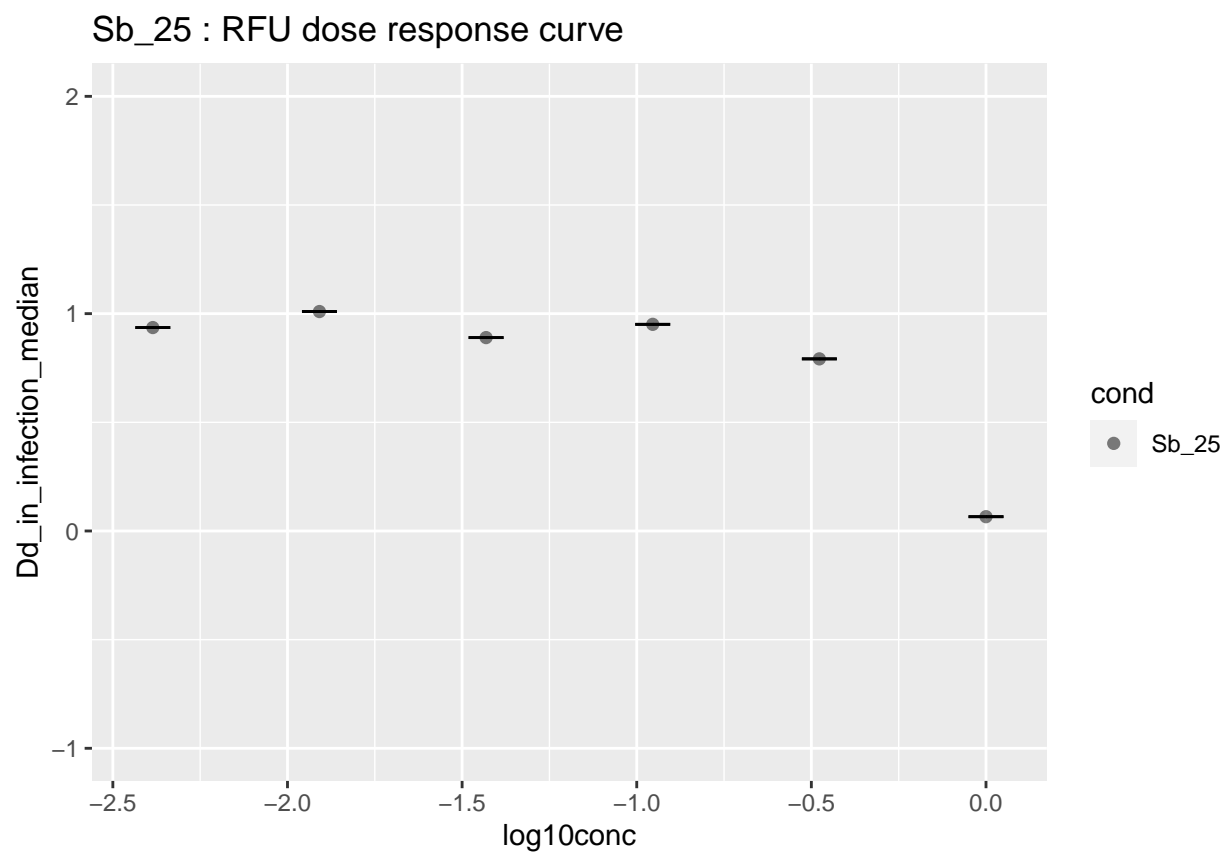


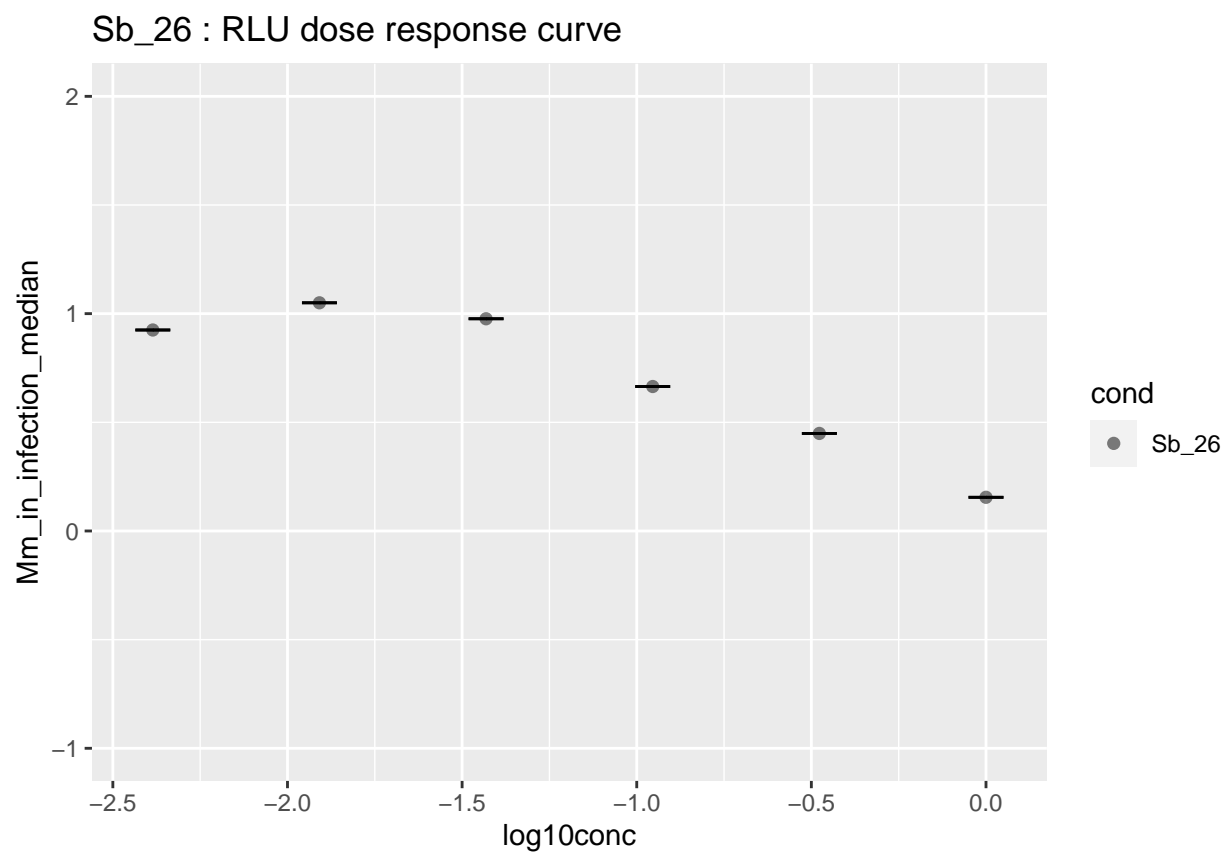


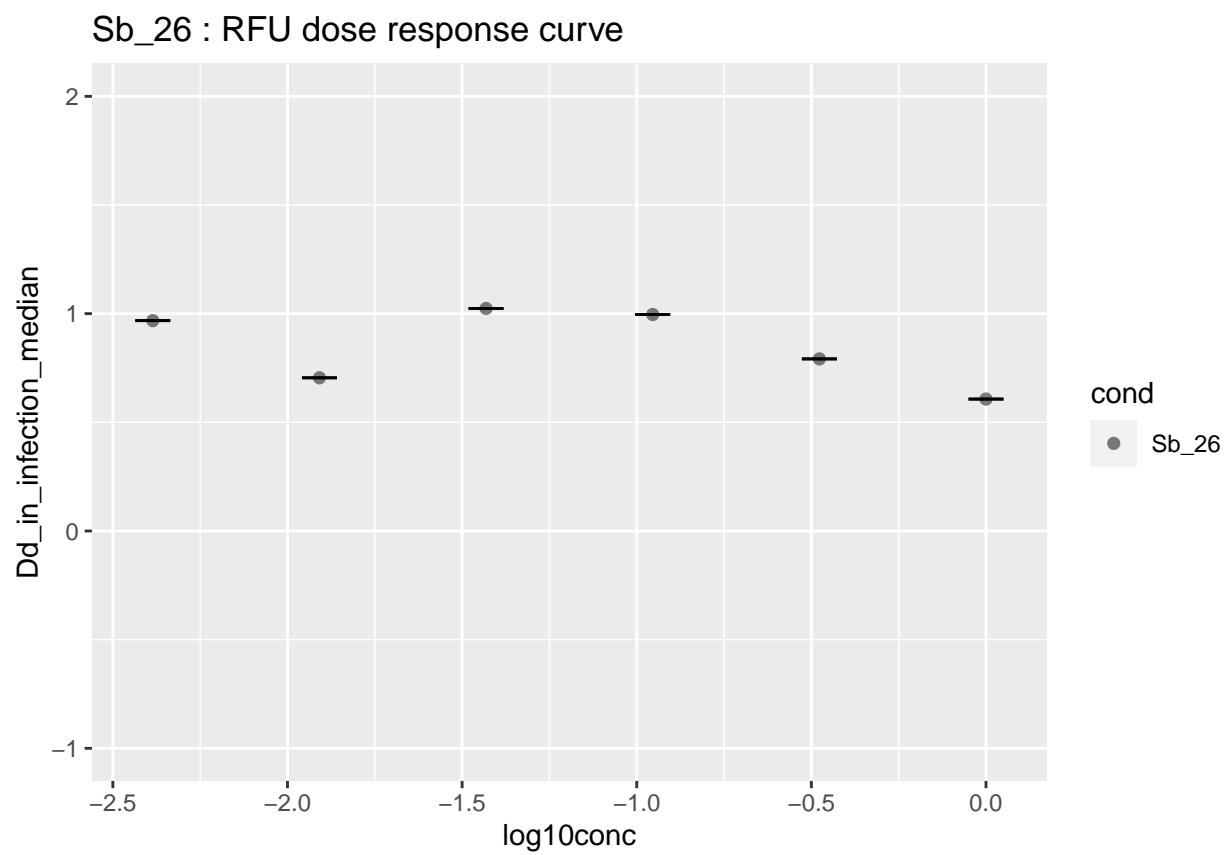


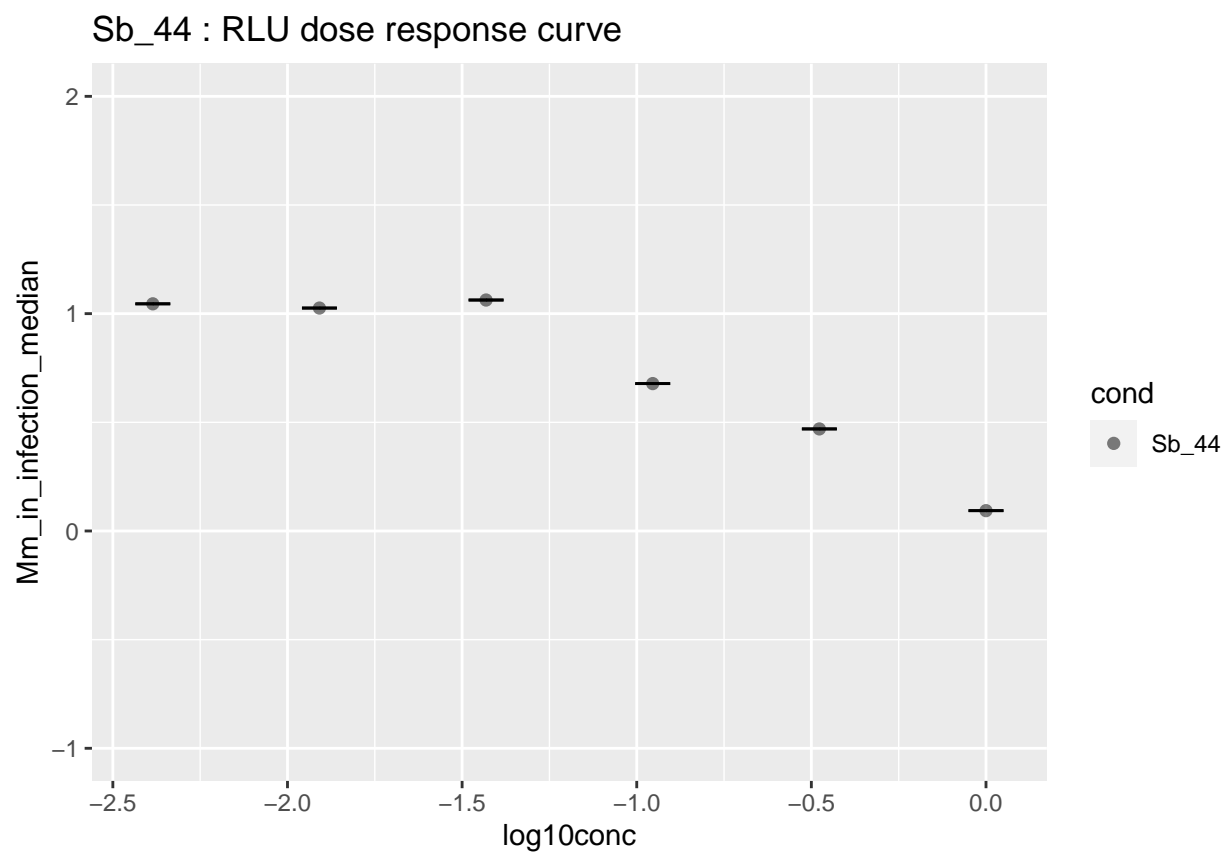


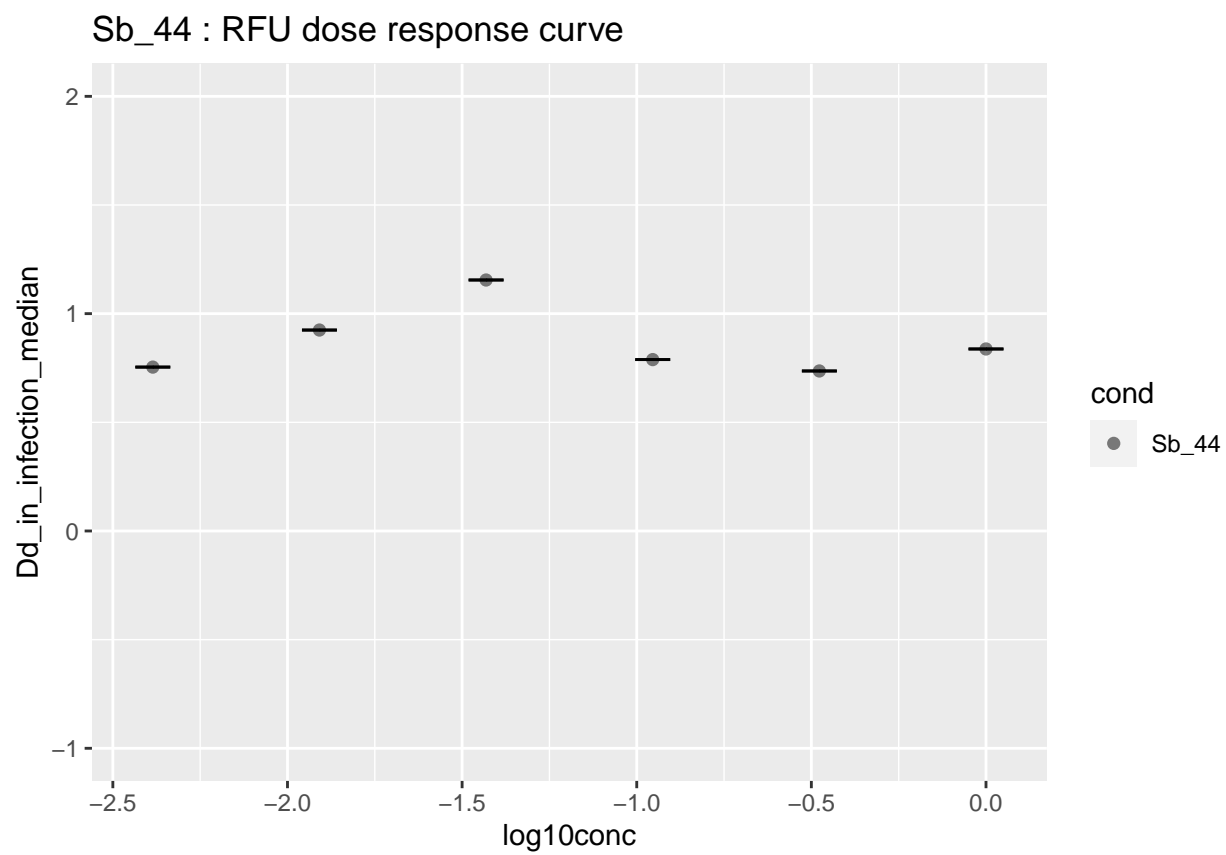


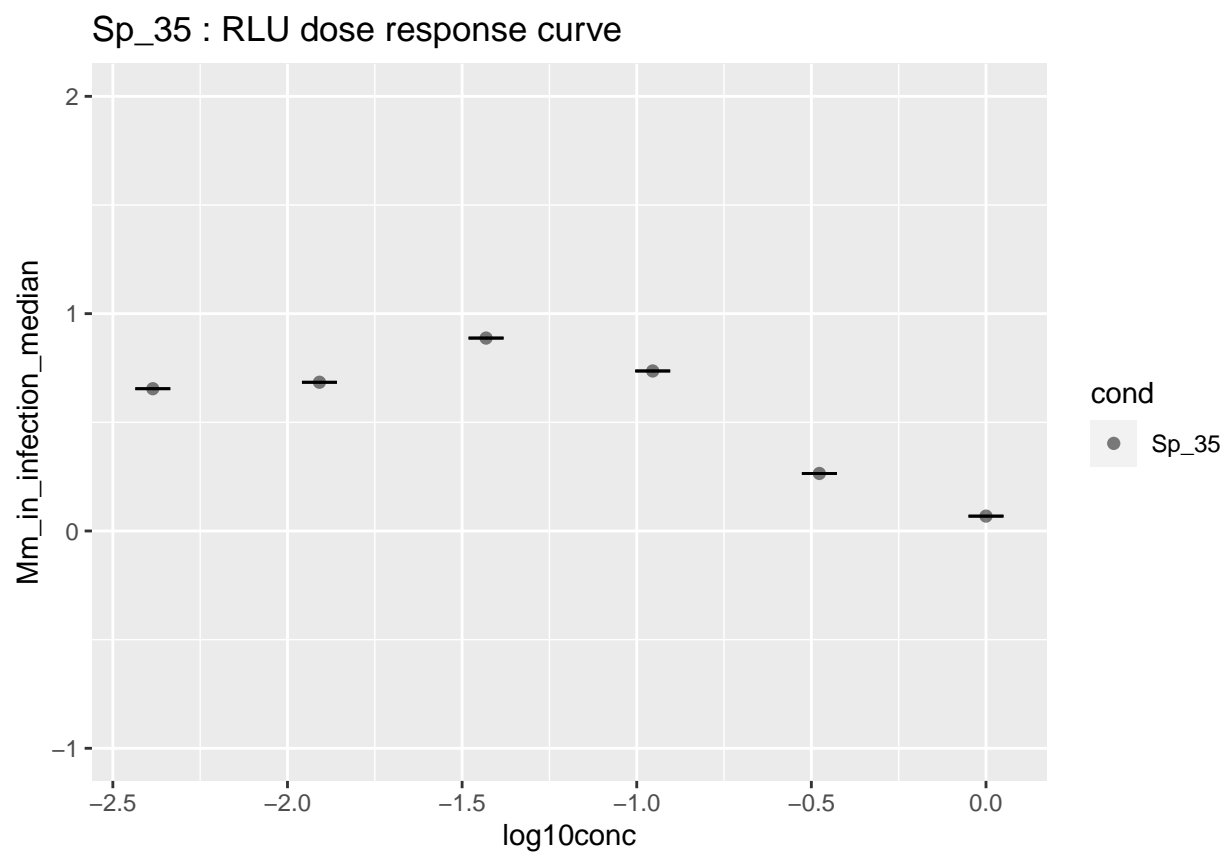


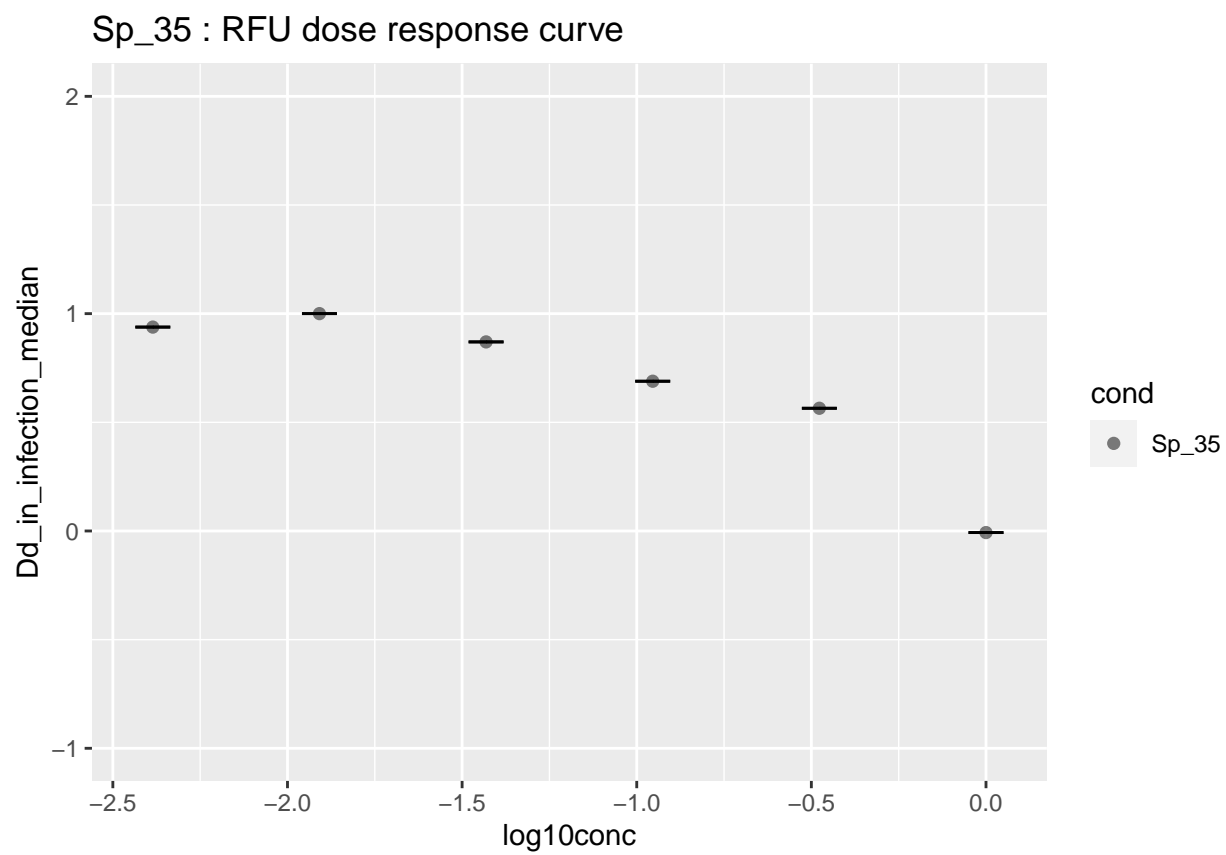


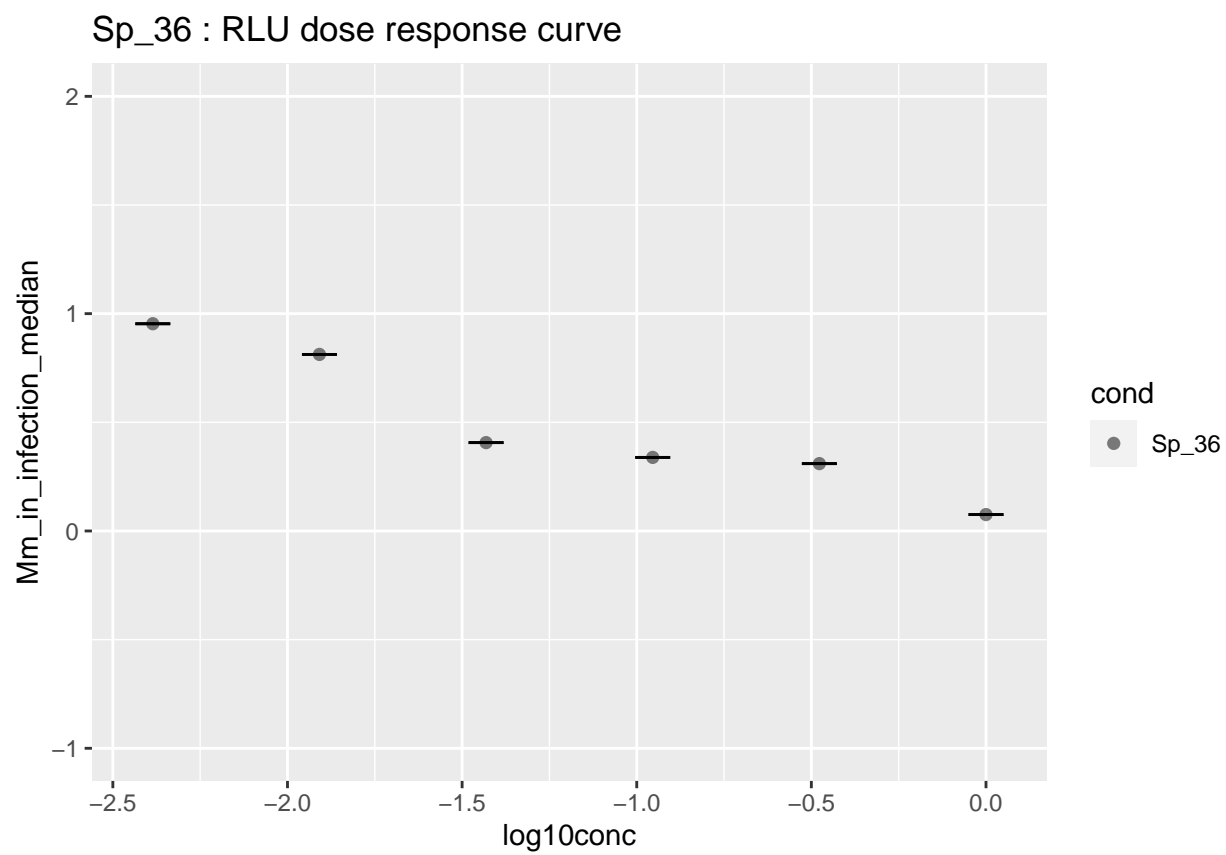


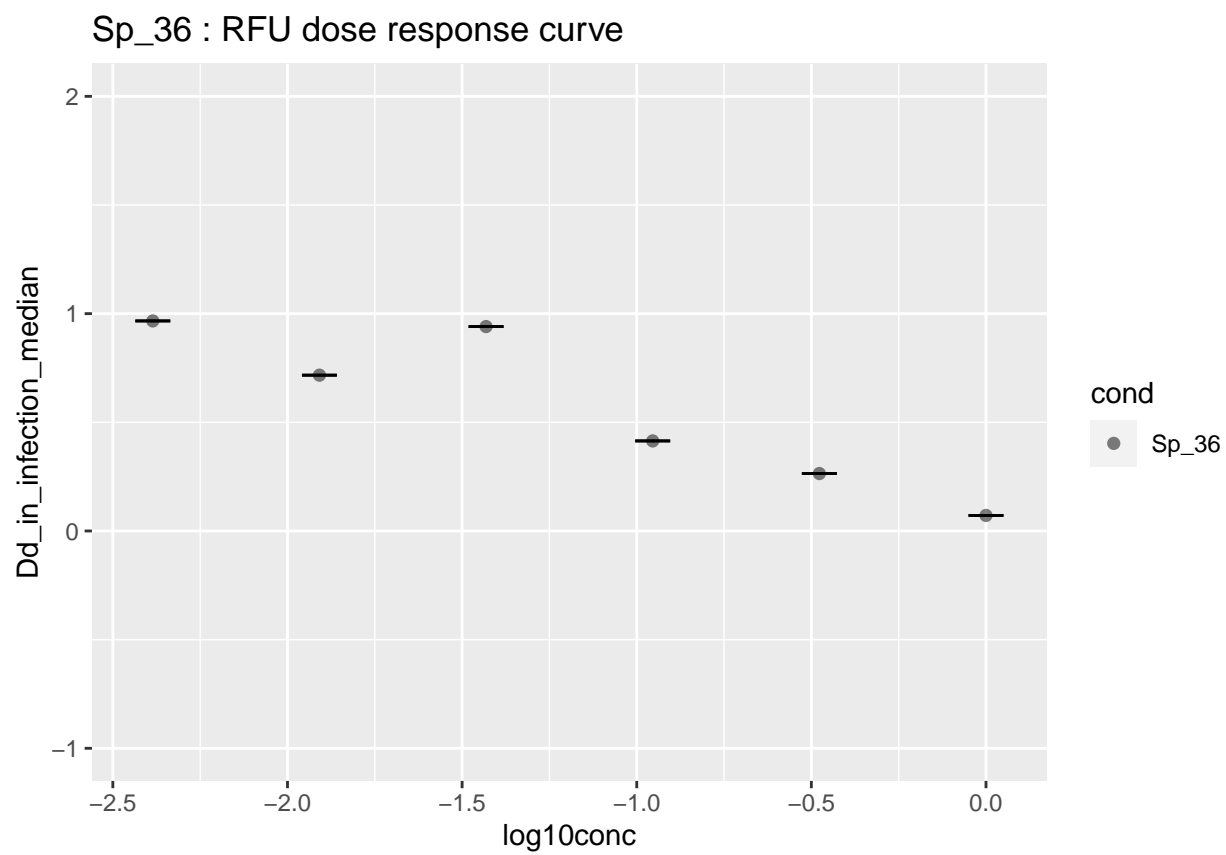


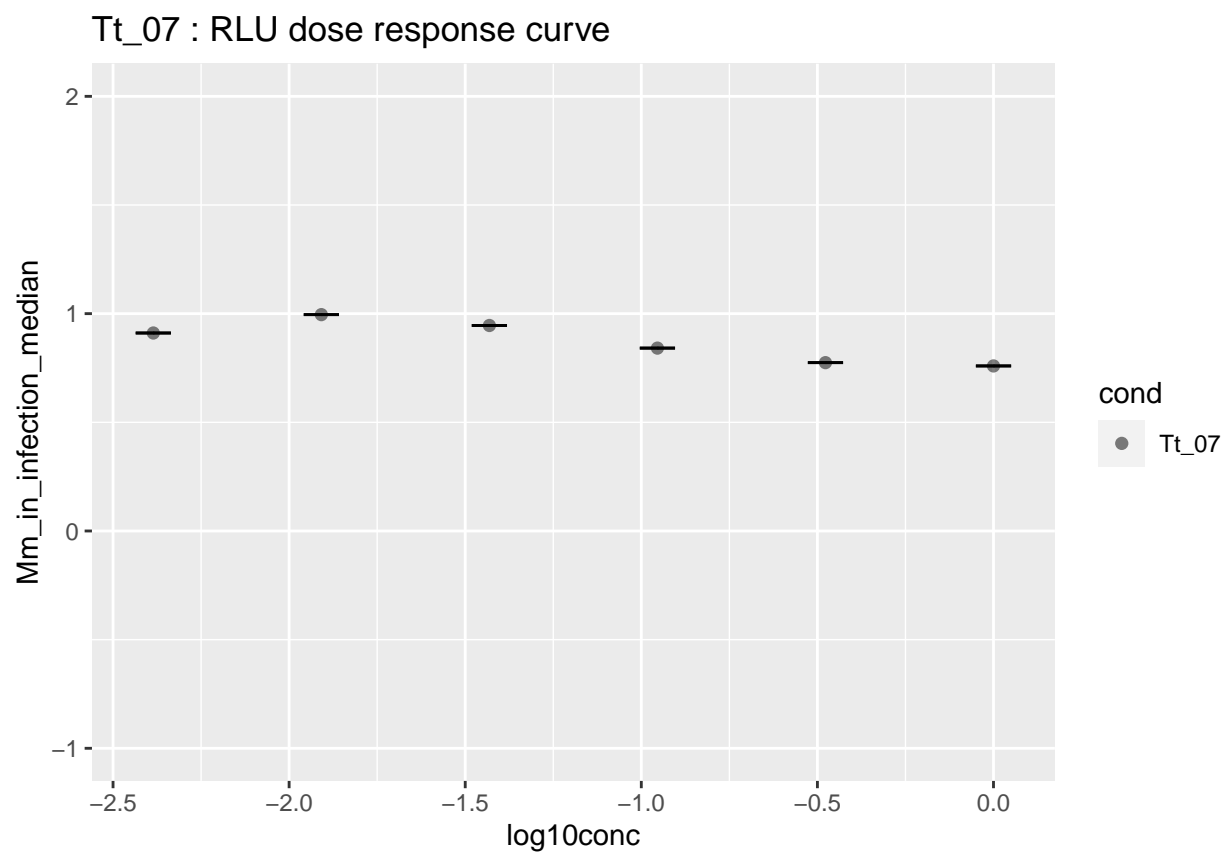


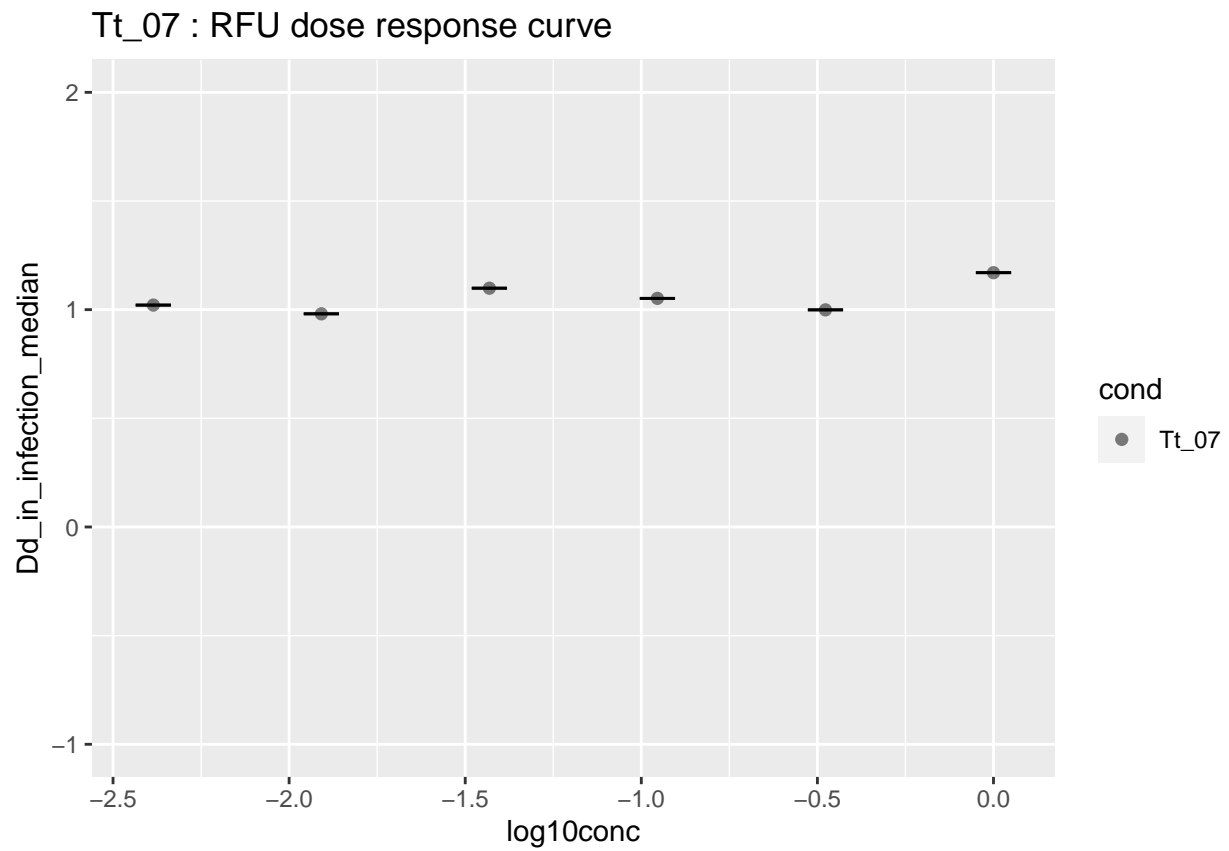






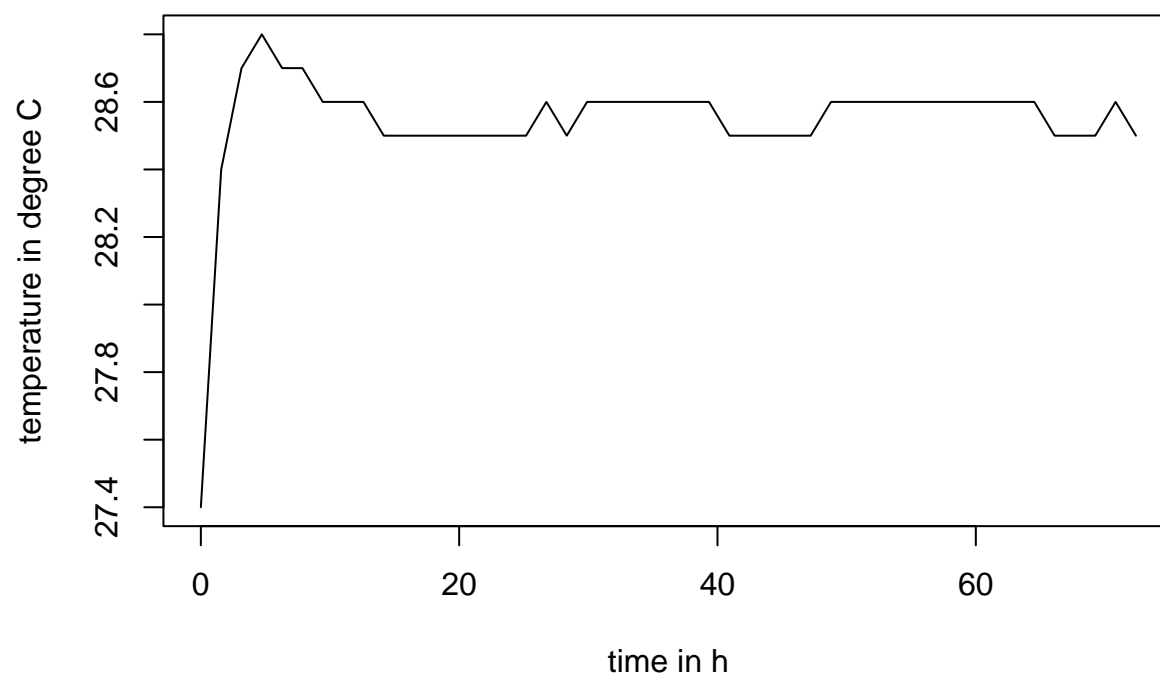




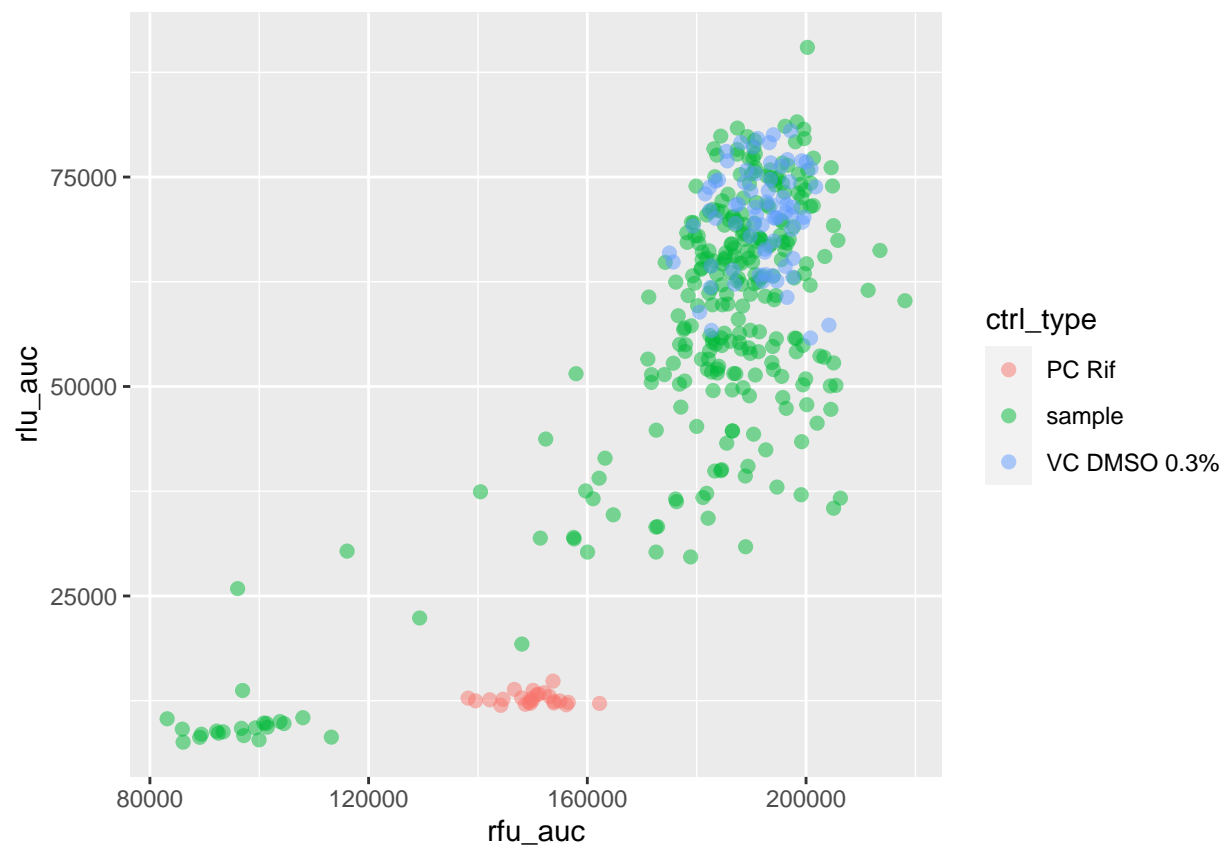


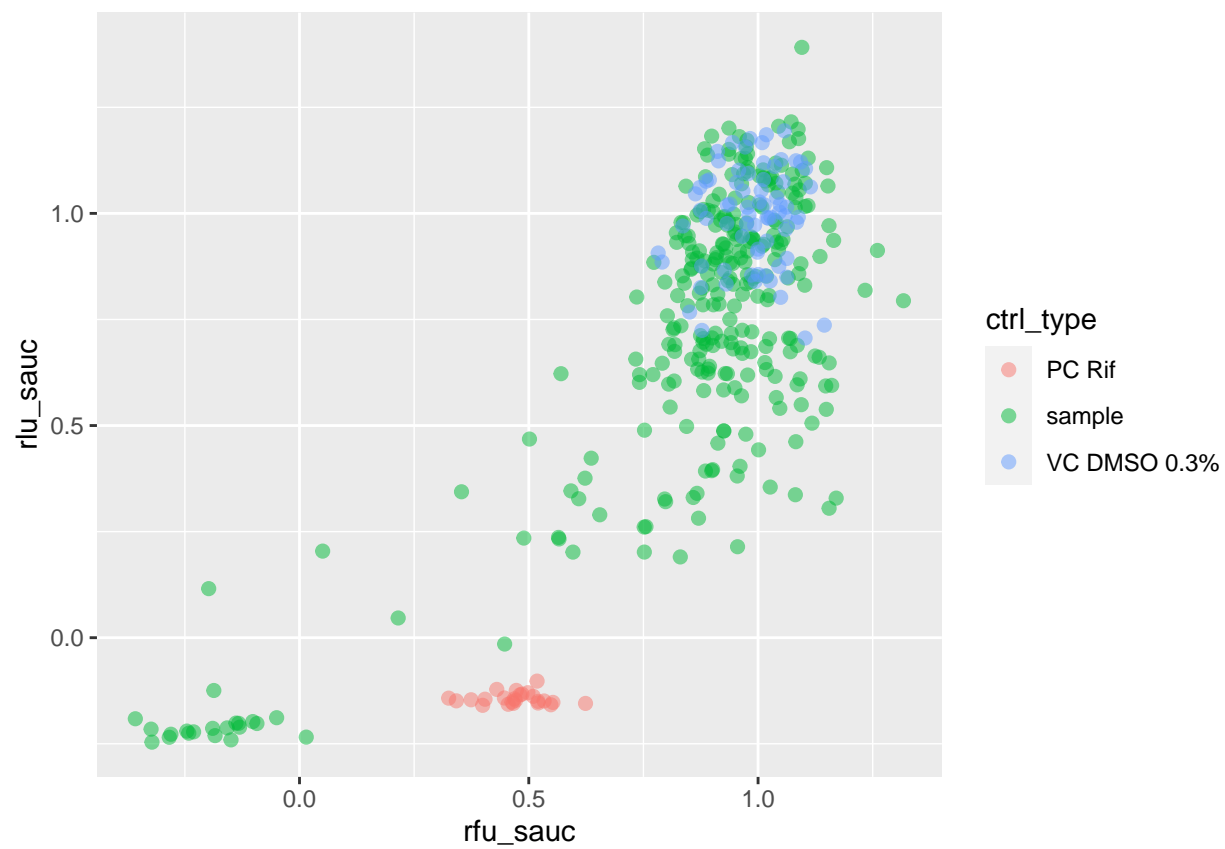
```
## [1] "analysis for plate04.xlsx"
```

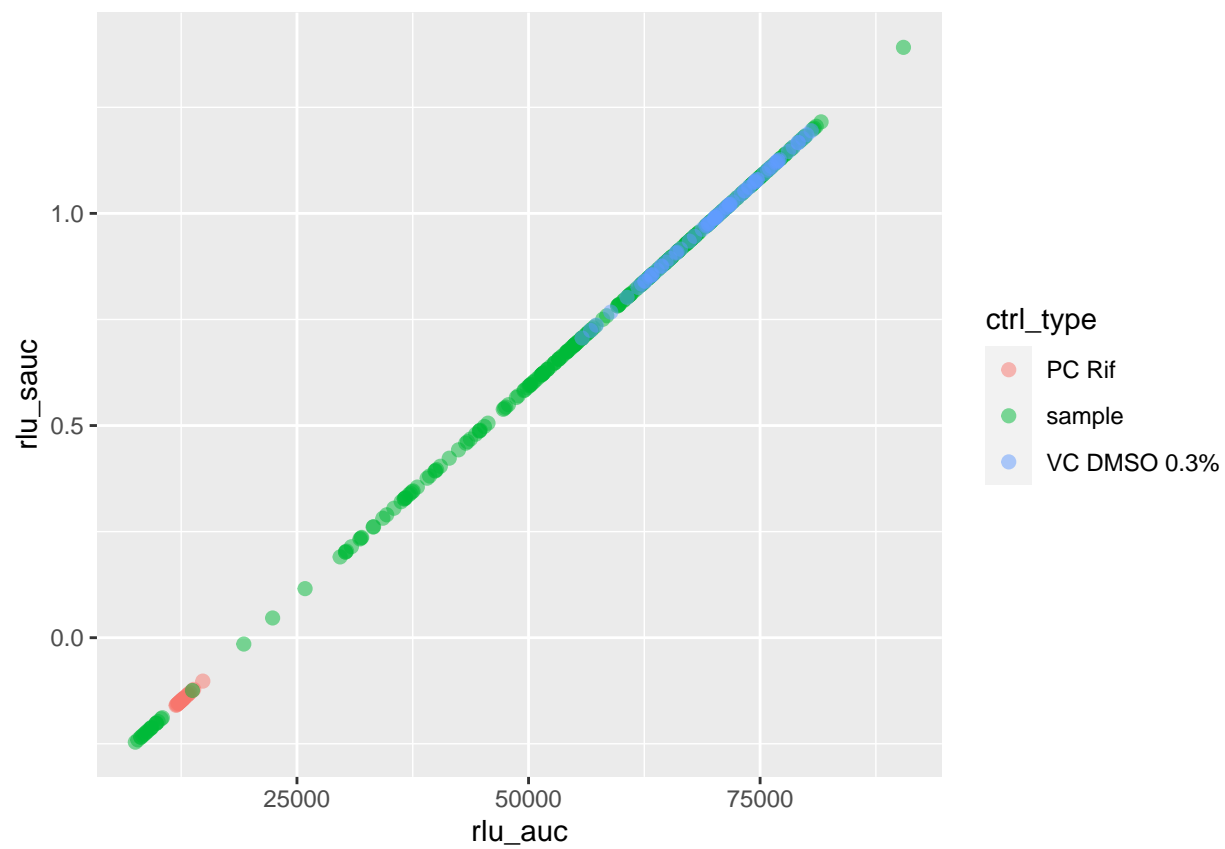

Temperature: plate 3

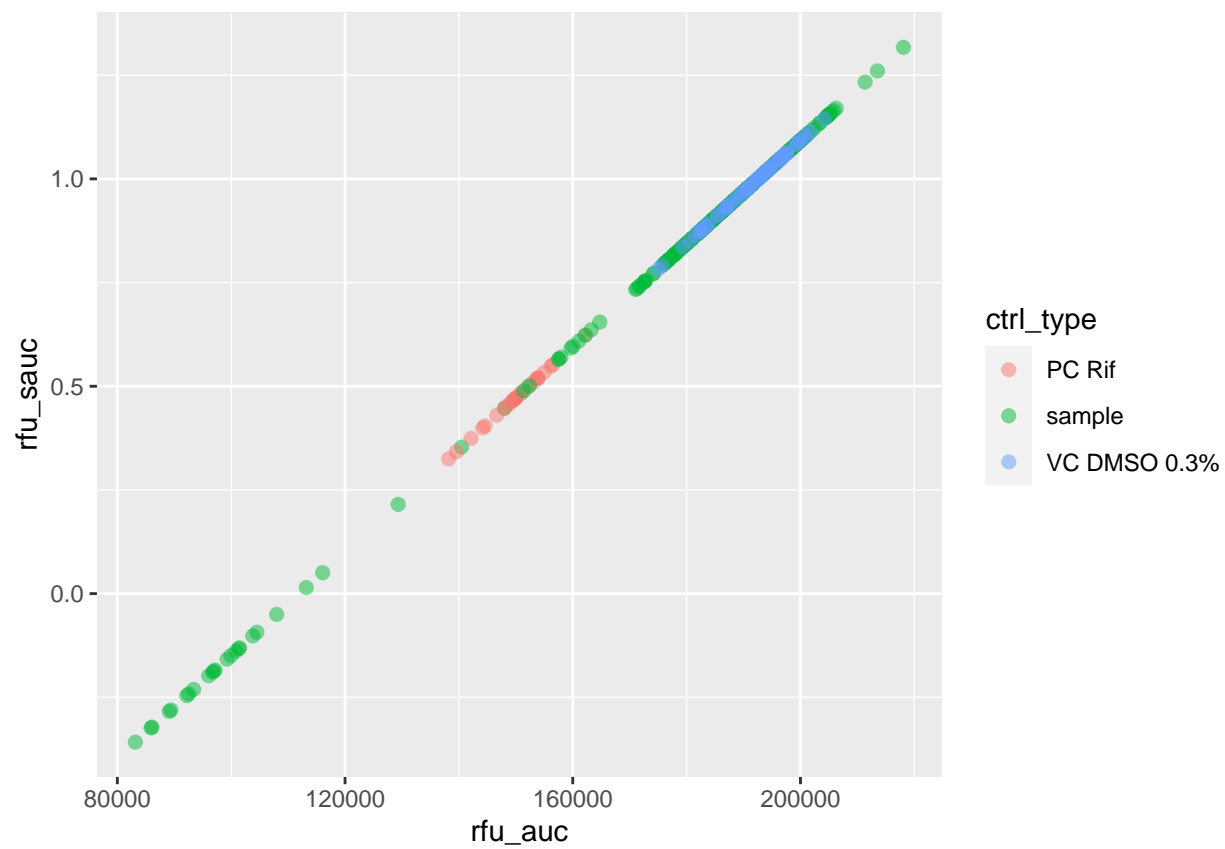


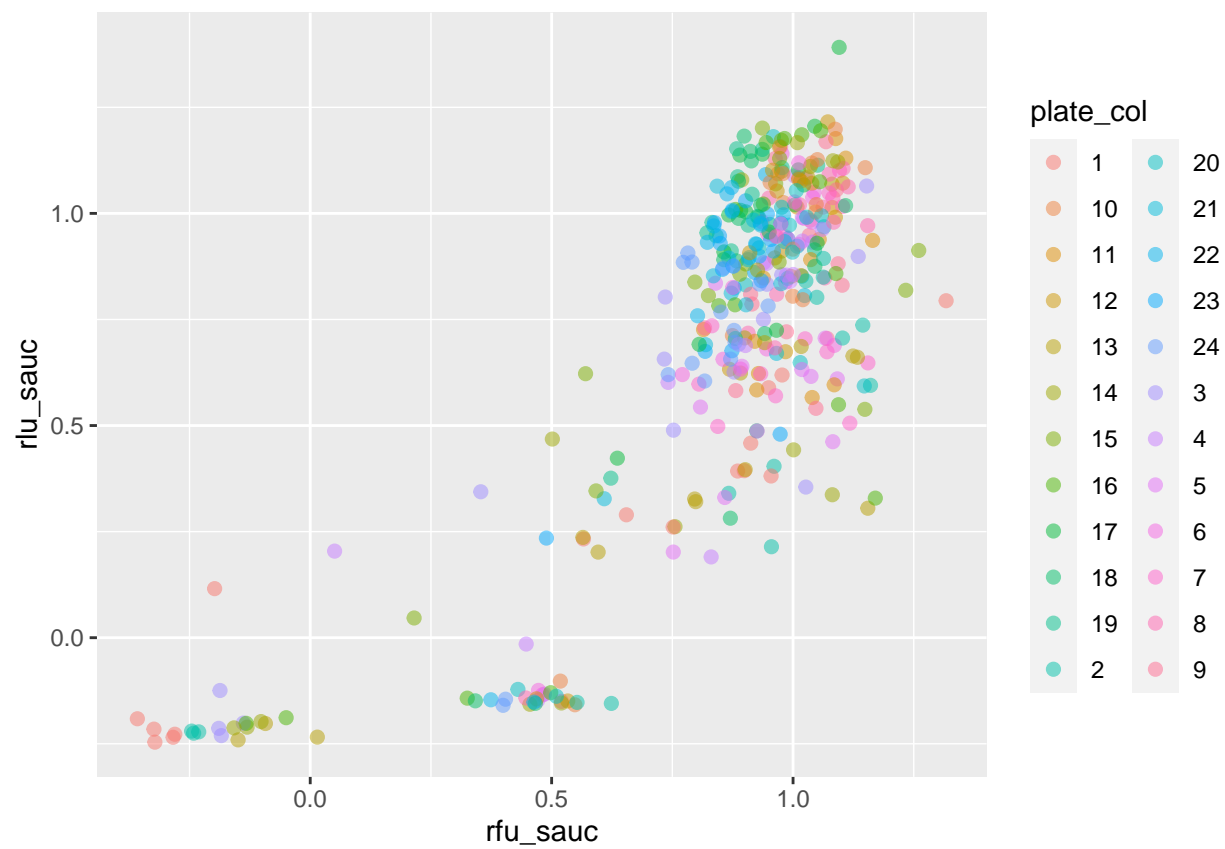
```
## [1] "VC DMSO 0.3% Robust z'-factor of rlu for plate plate04.xlsx, biorep 2 : "  
## [1] 0.63
```

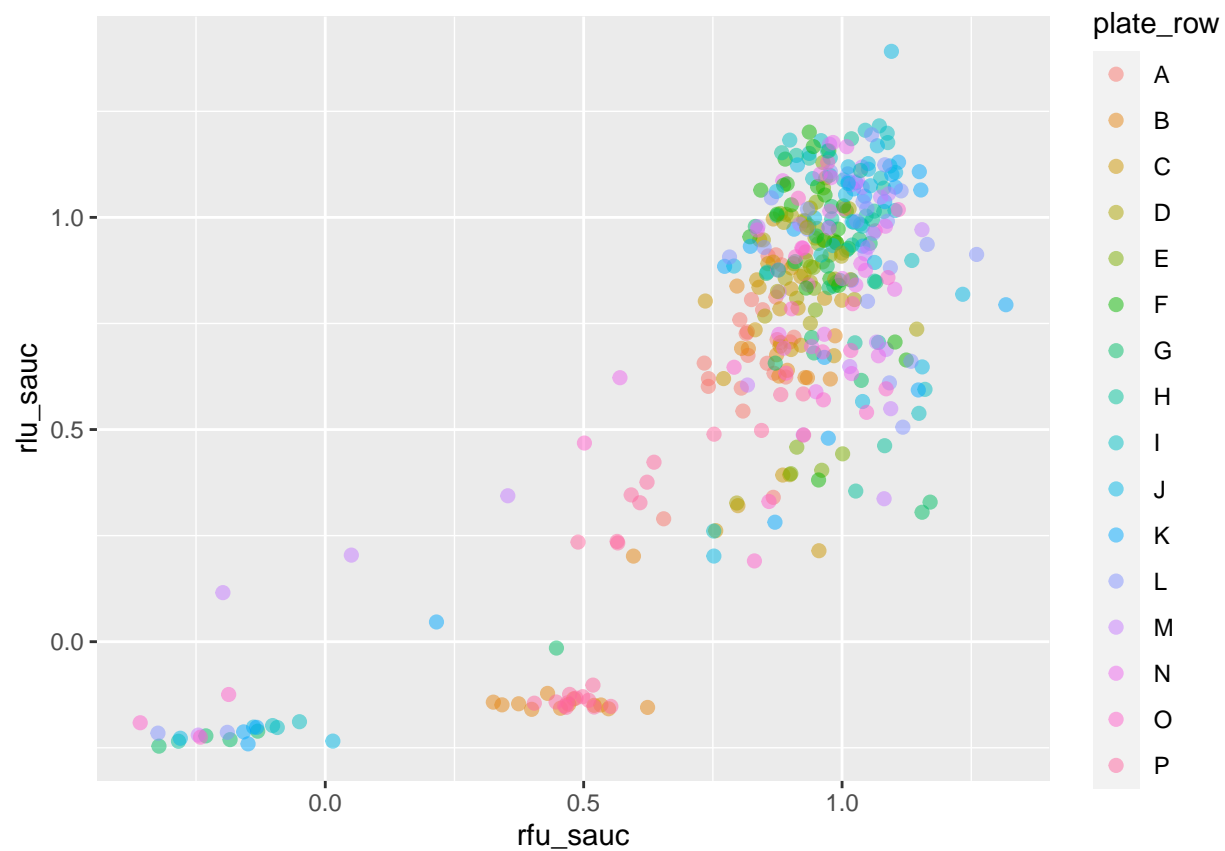




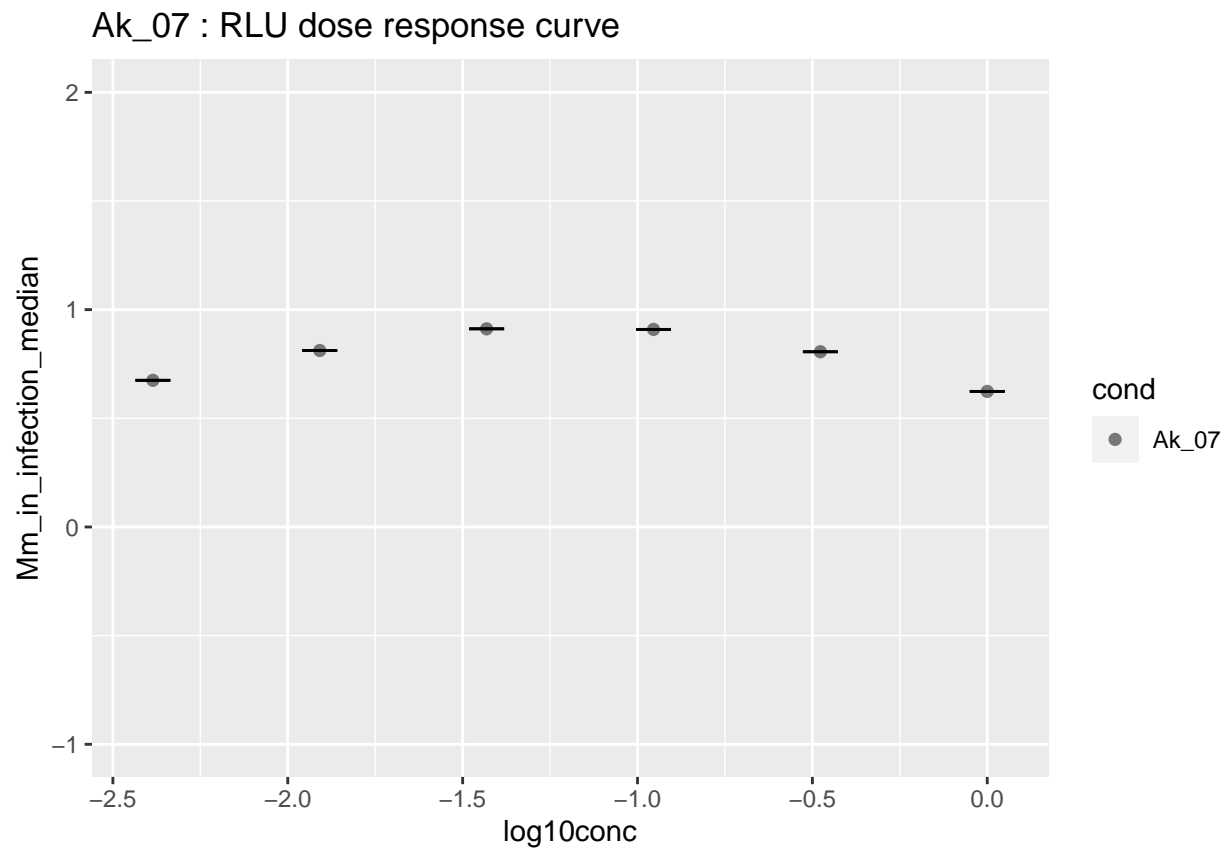


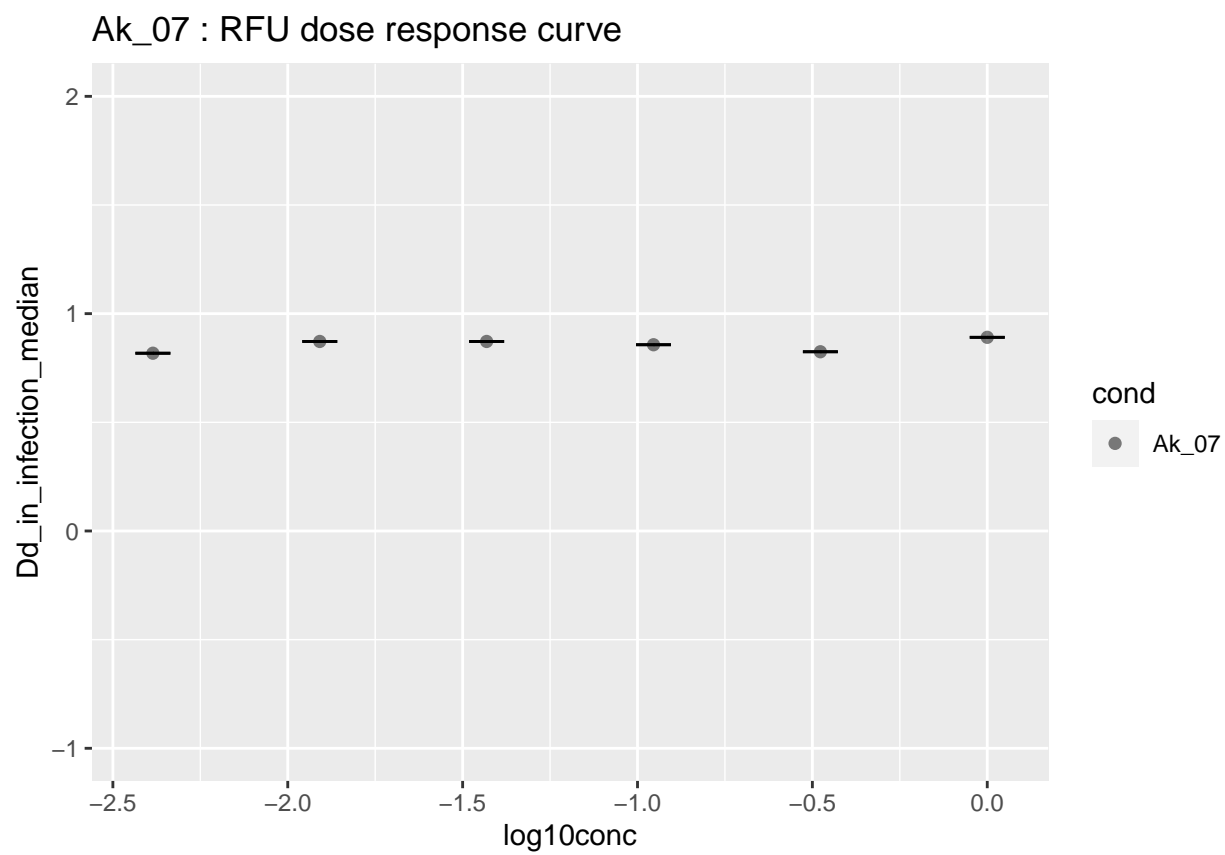


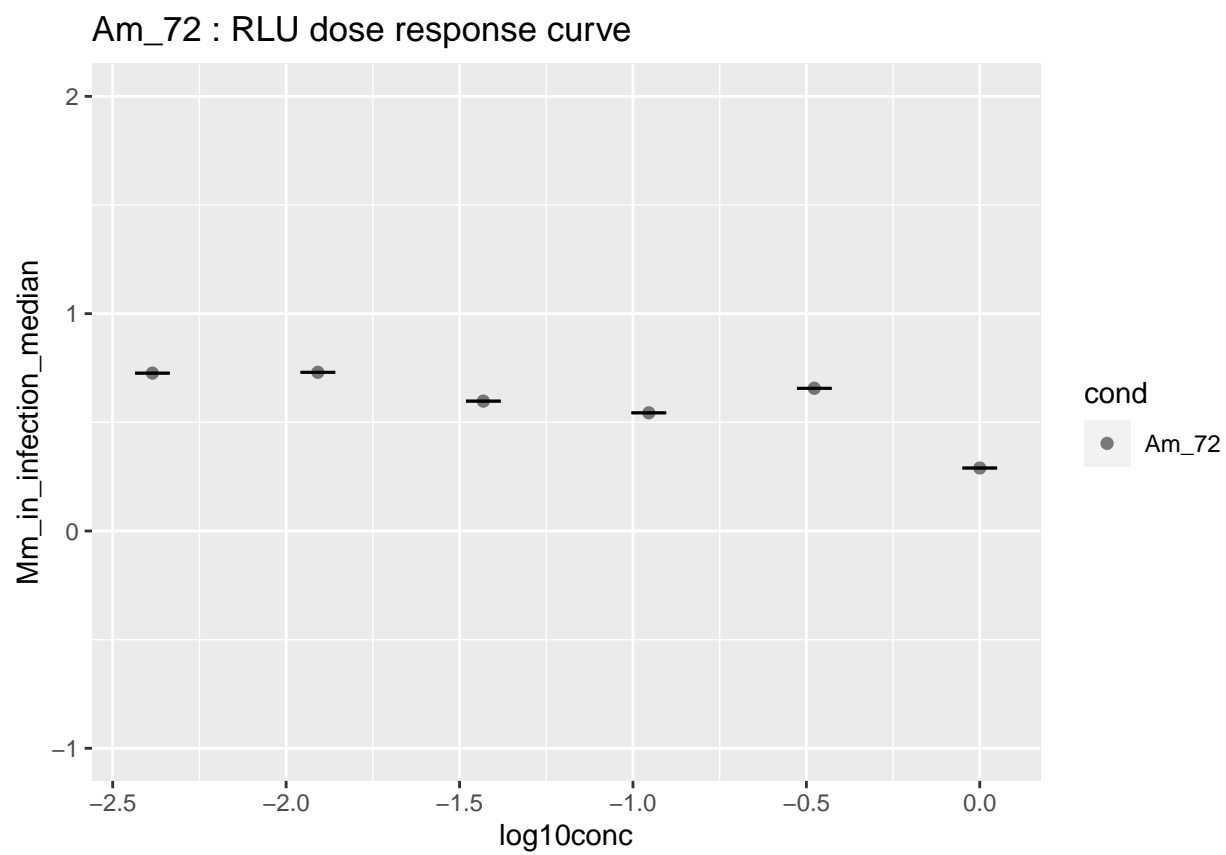


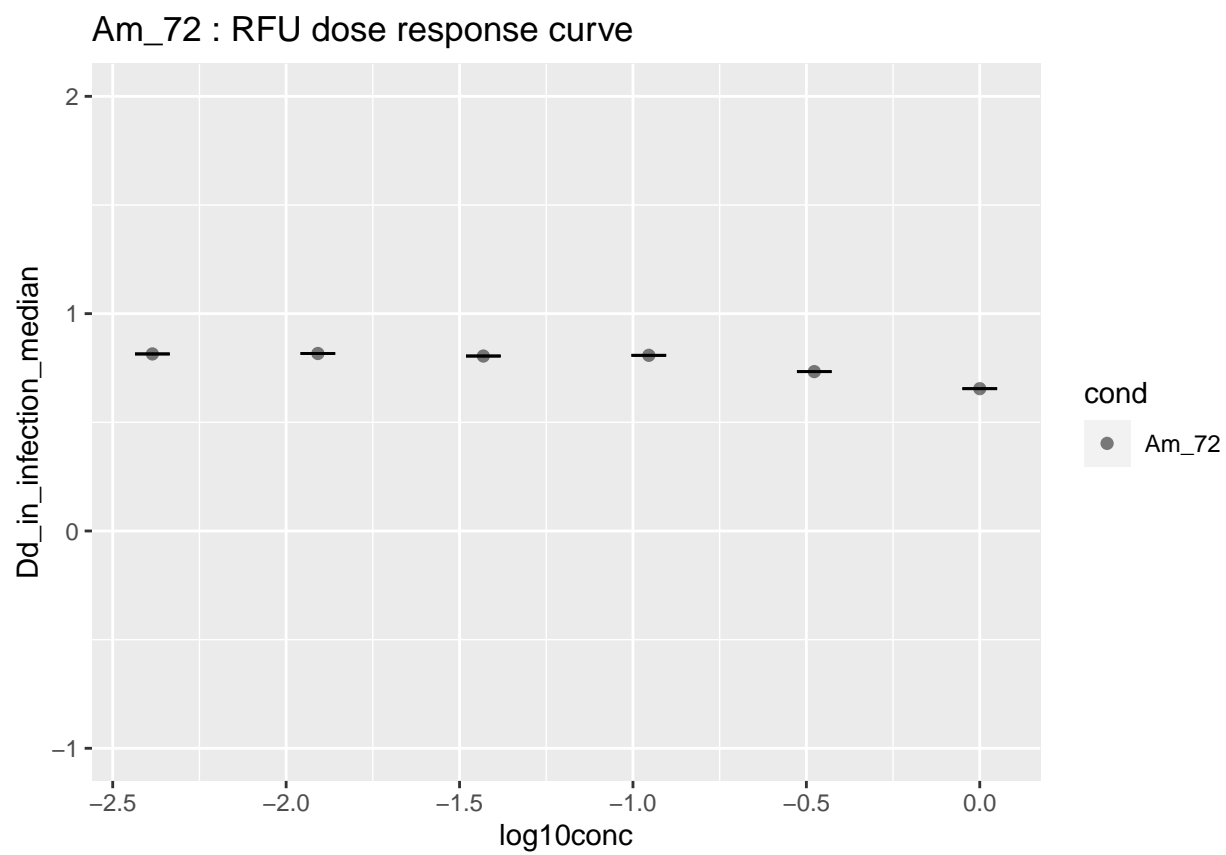


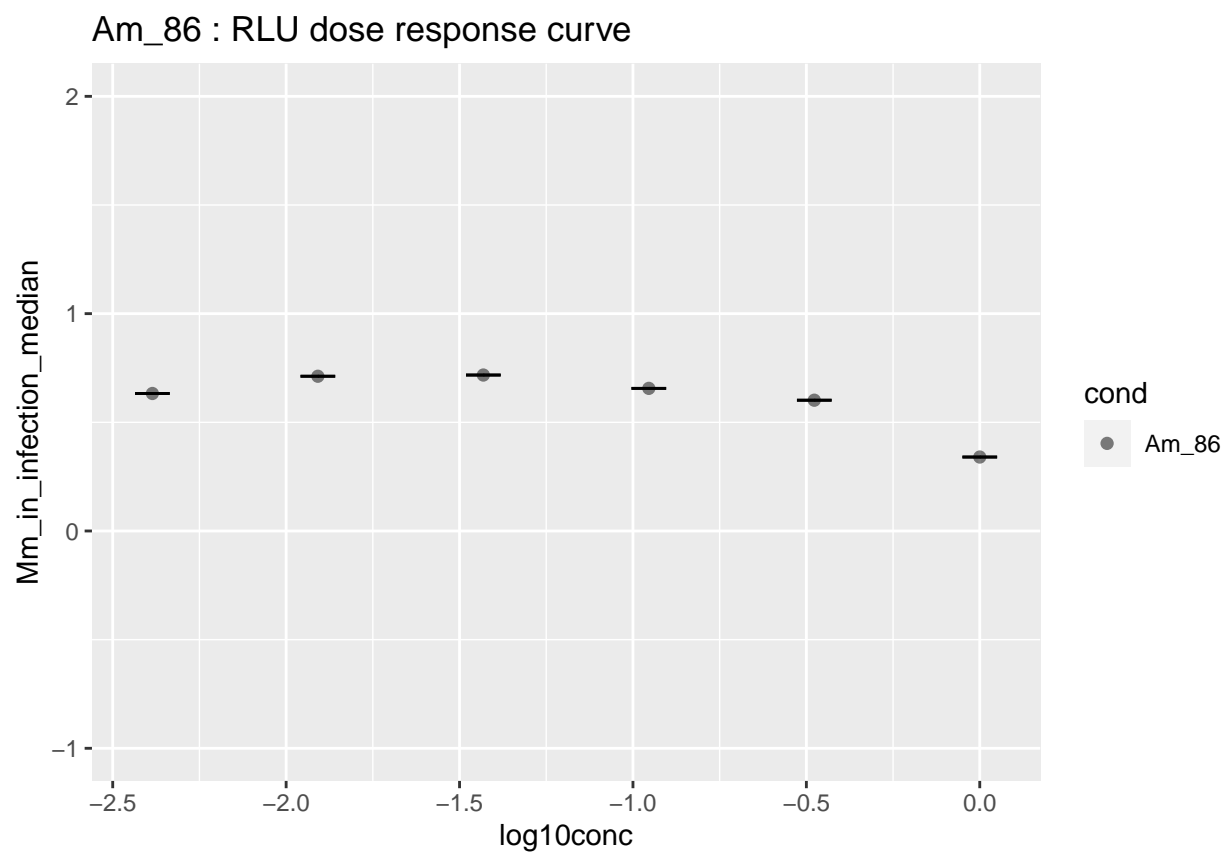
```
## [1] "Dose response curves over all bioreps within this plate"
```

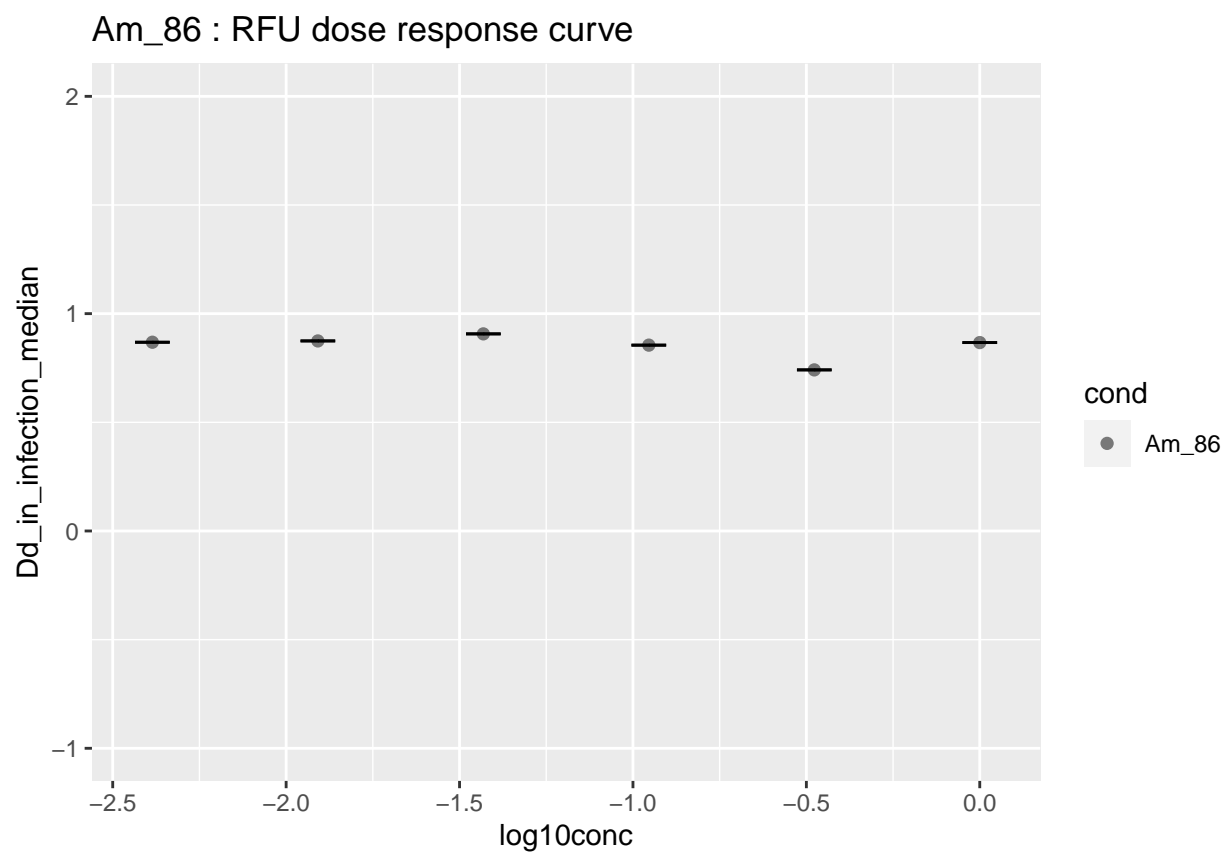


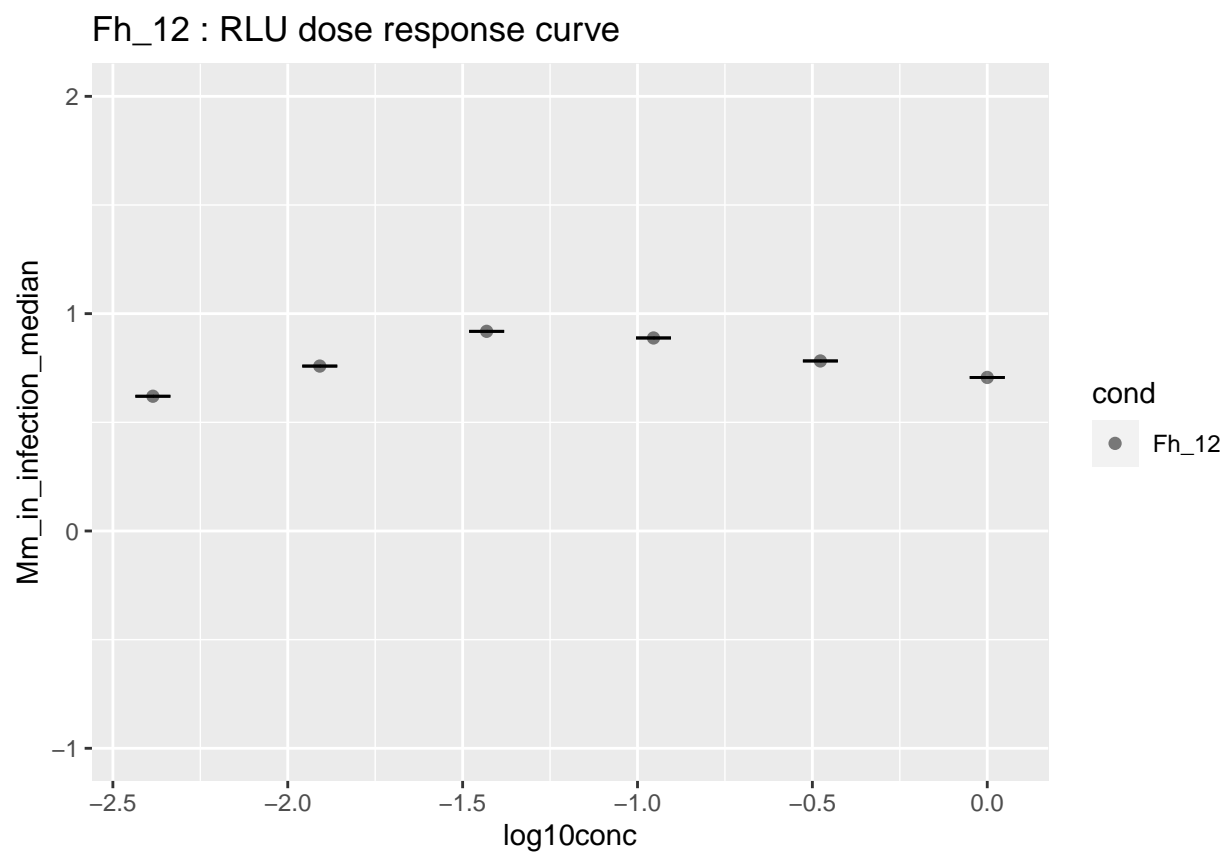


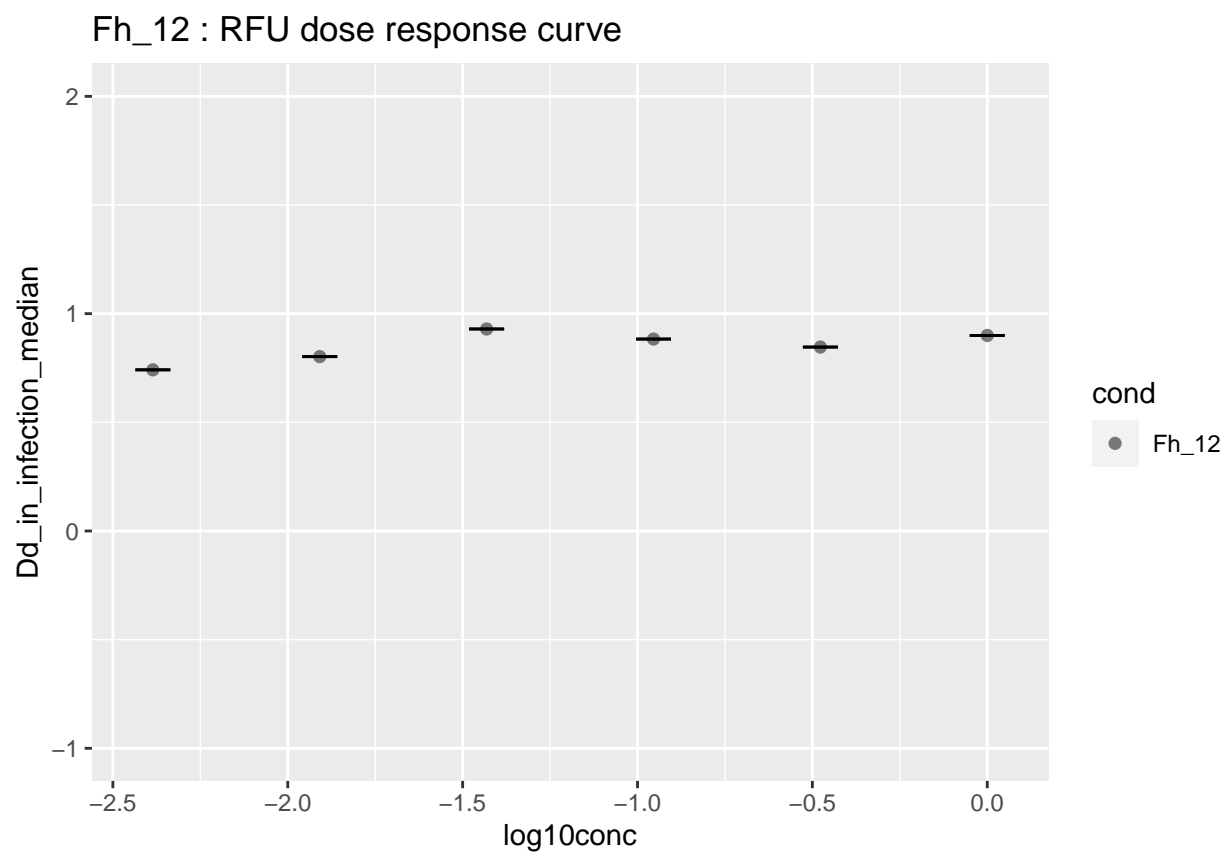


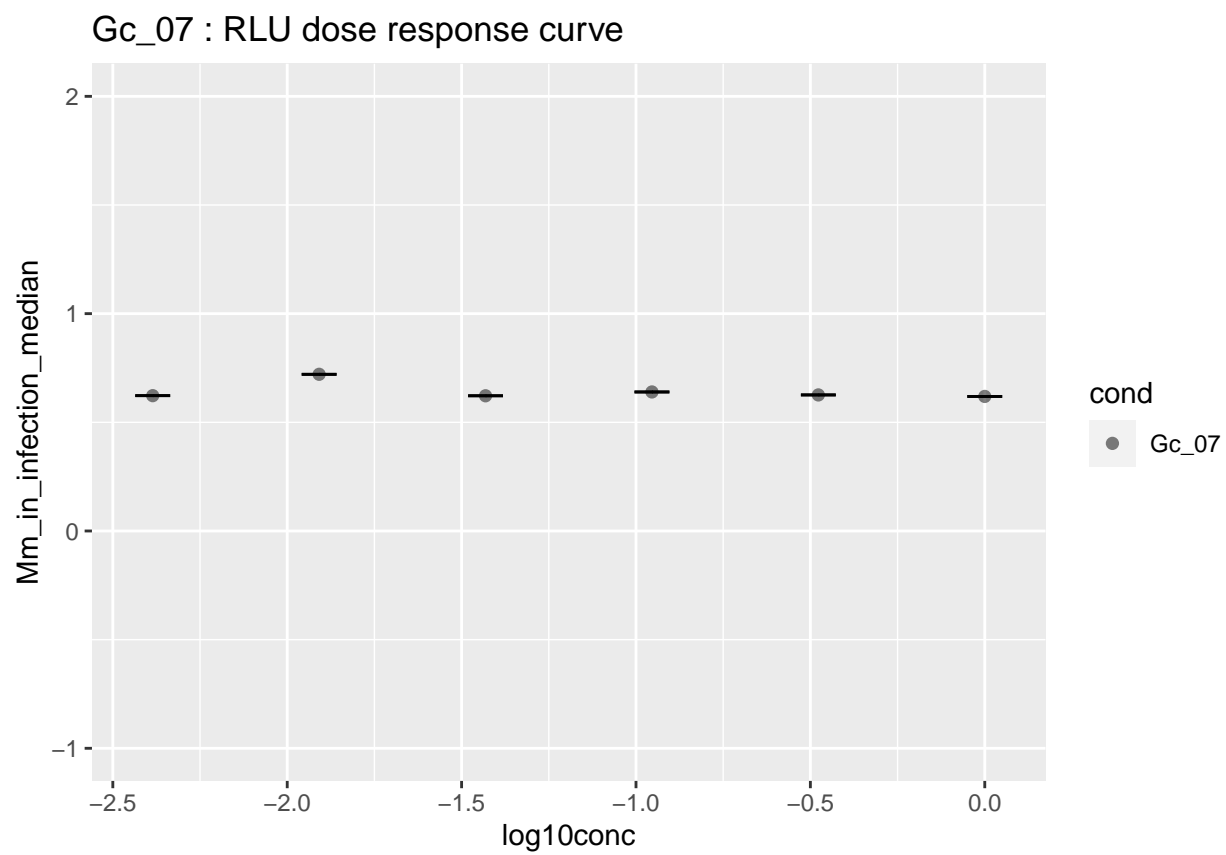


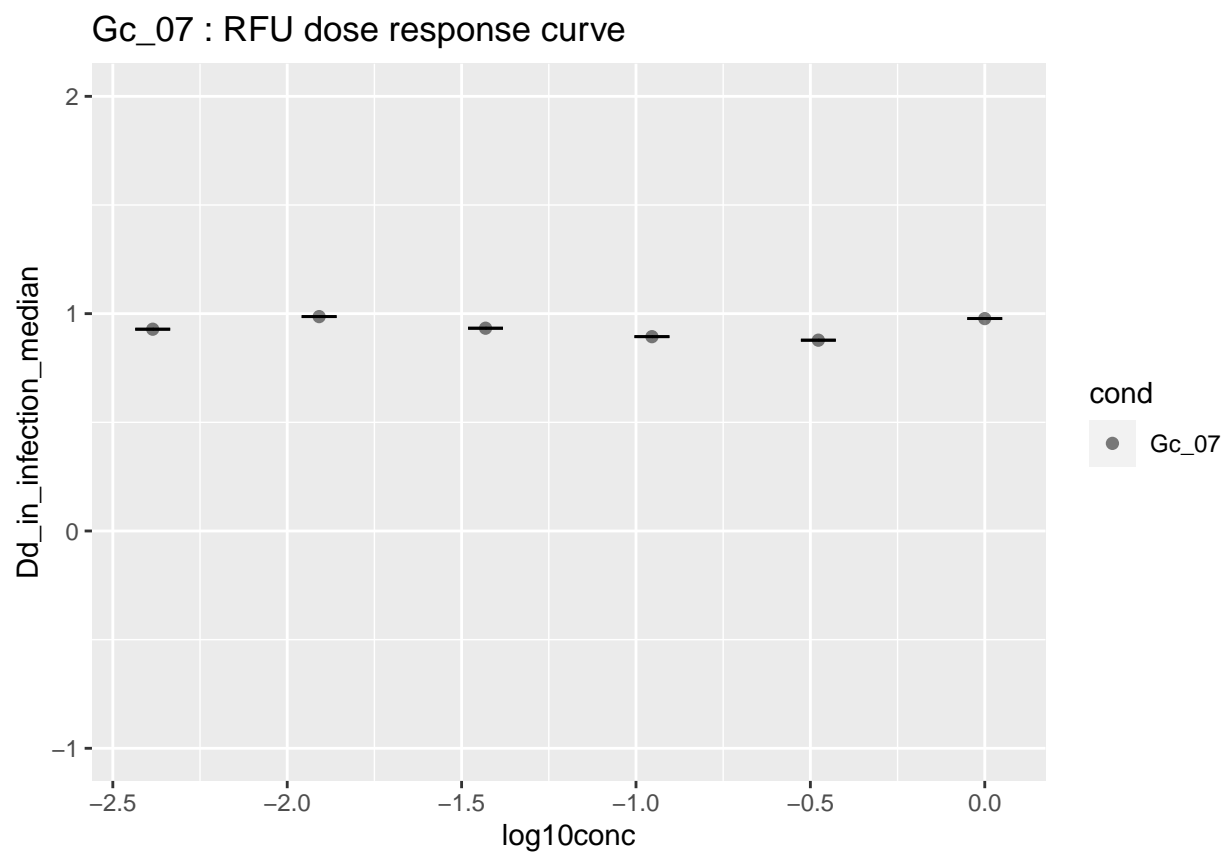


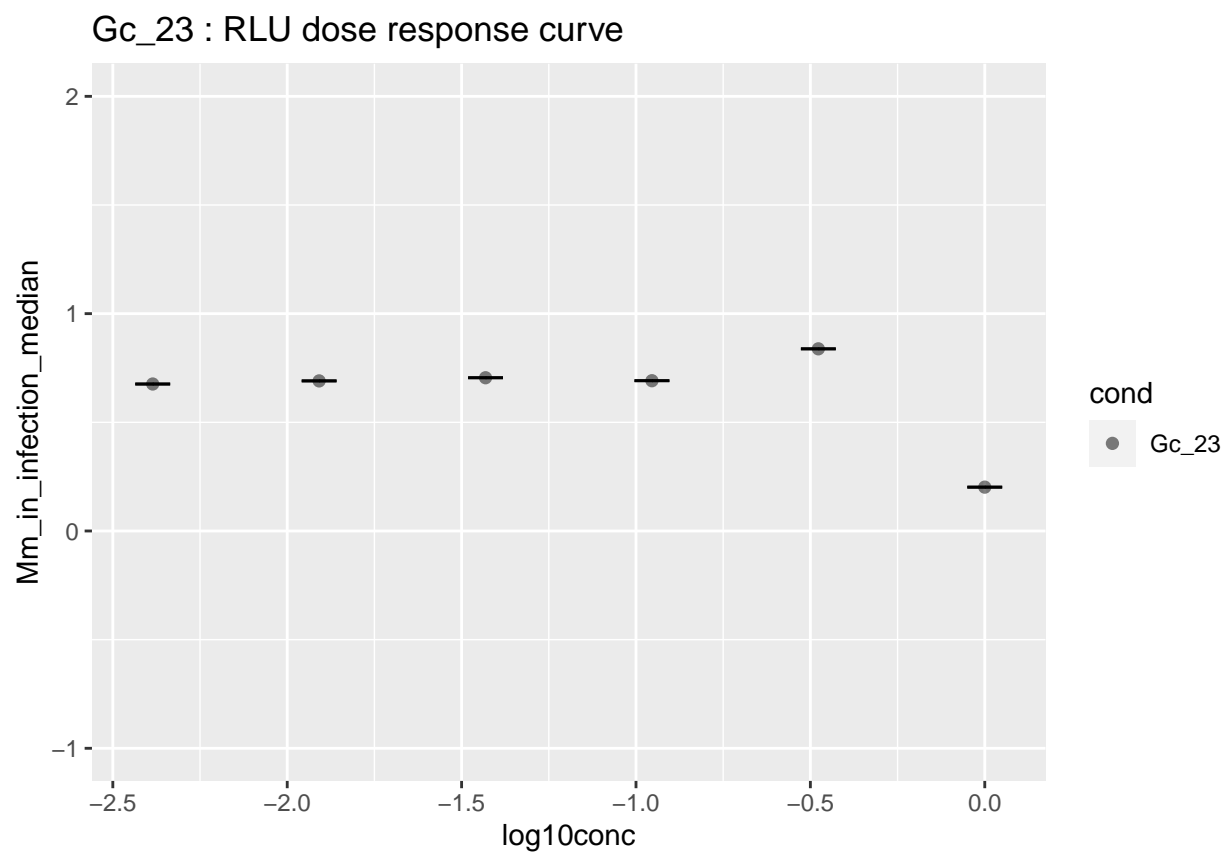


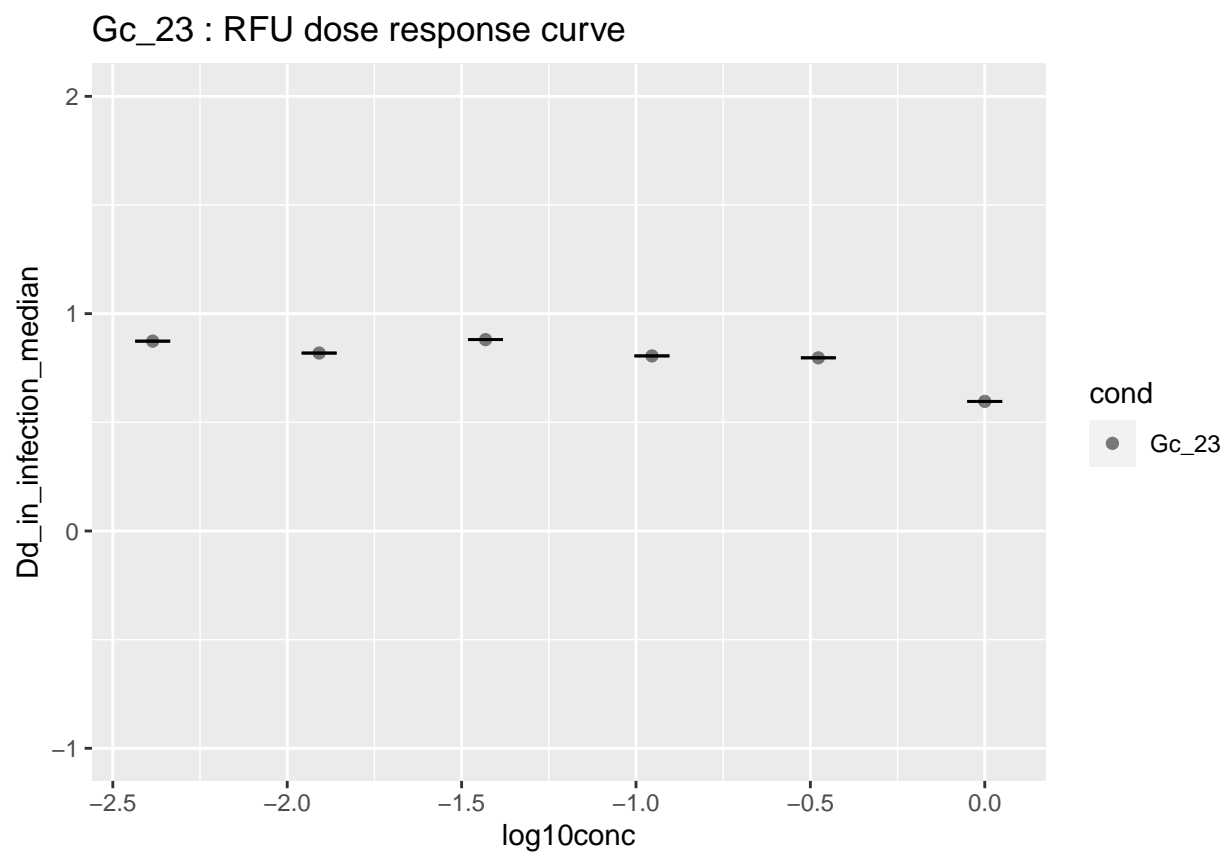


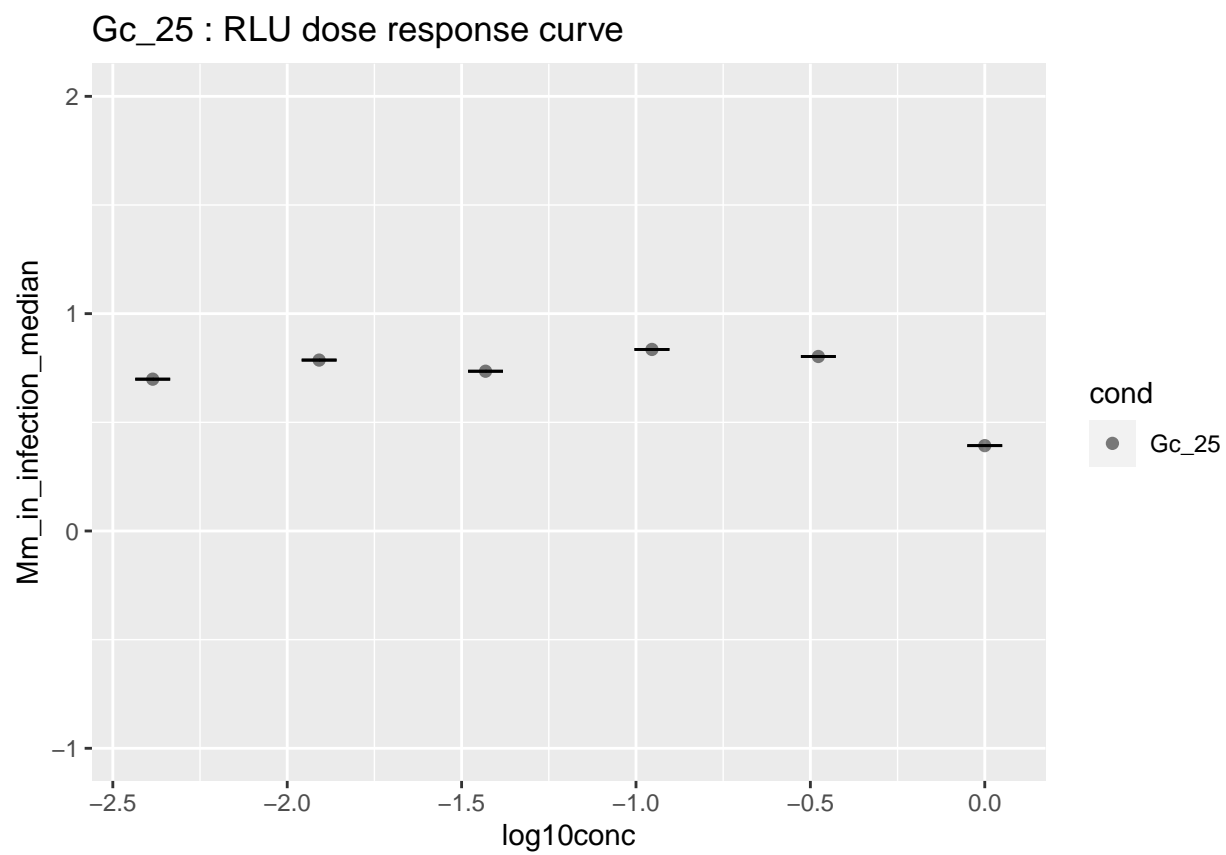


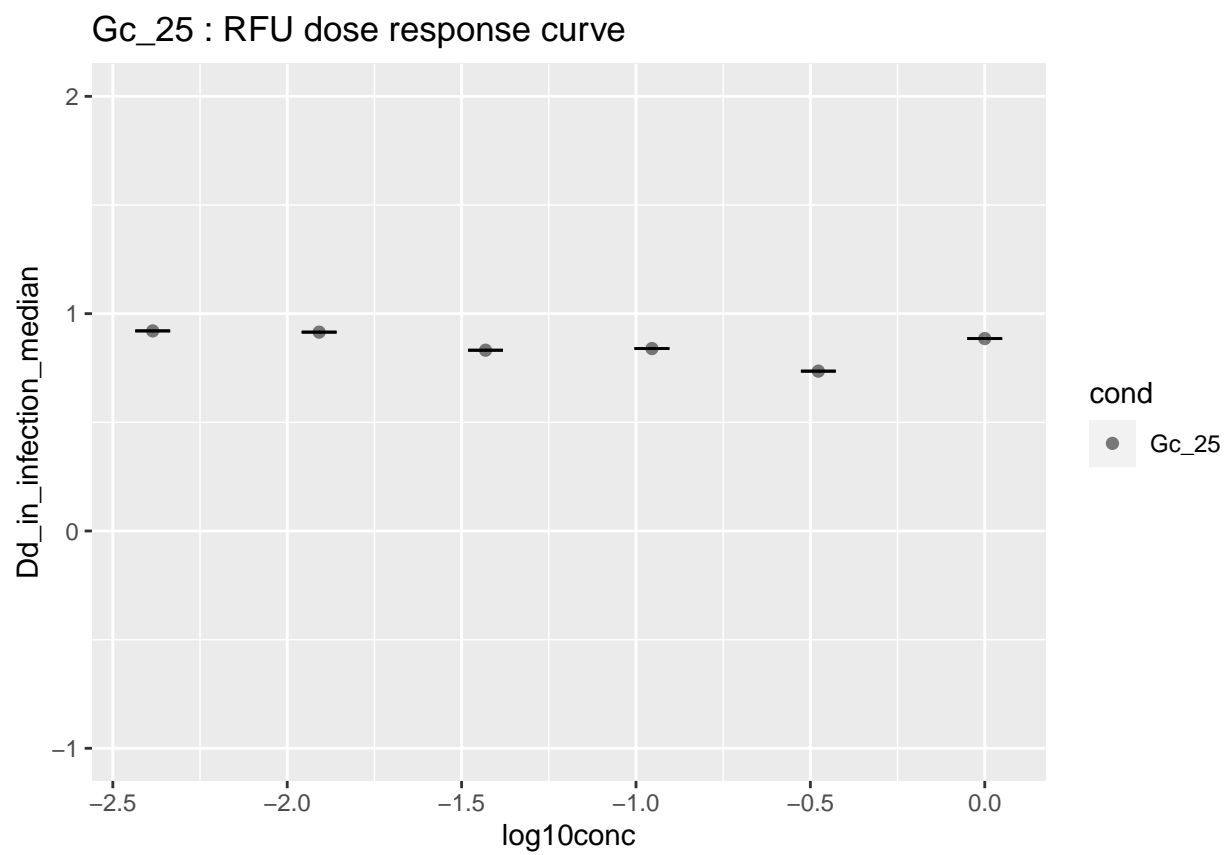


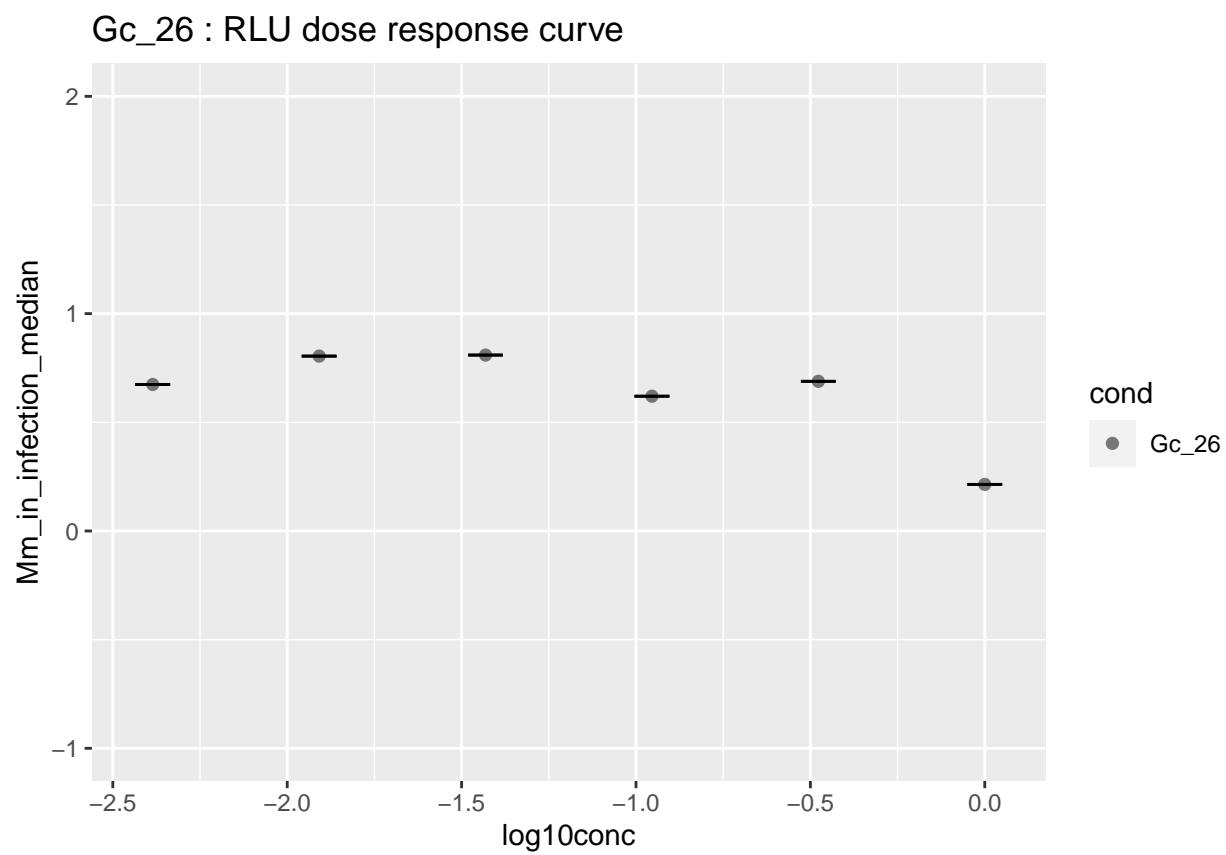


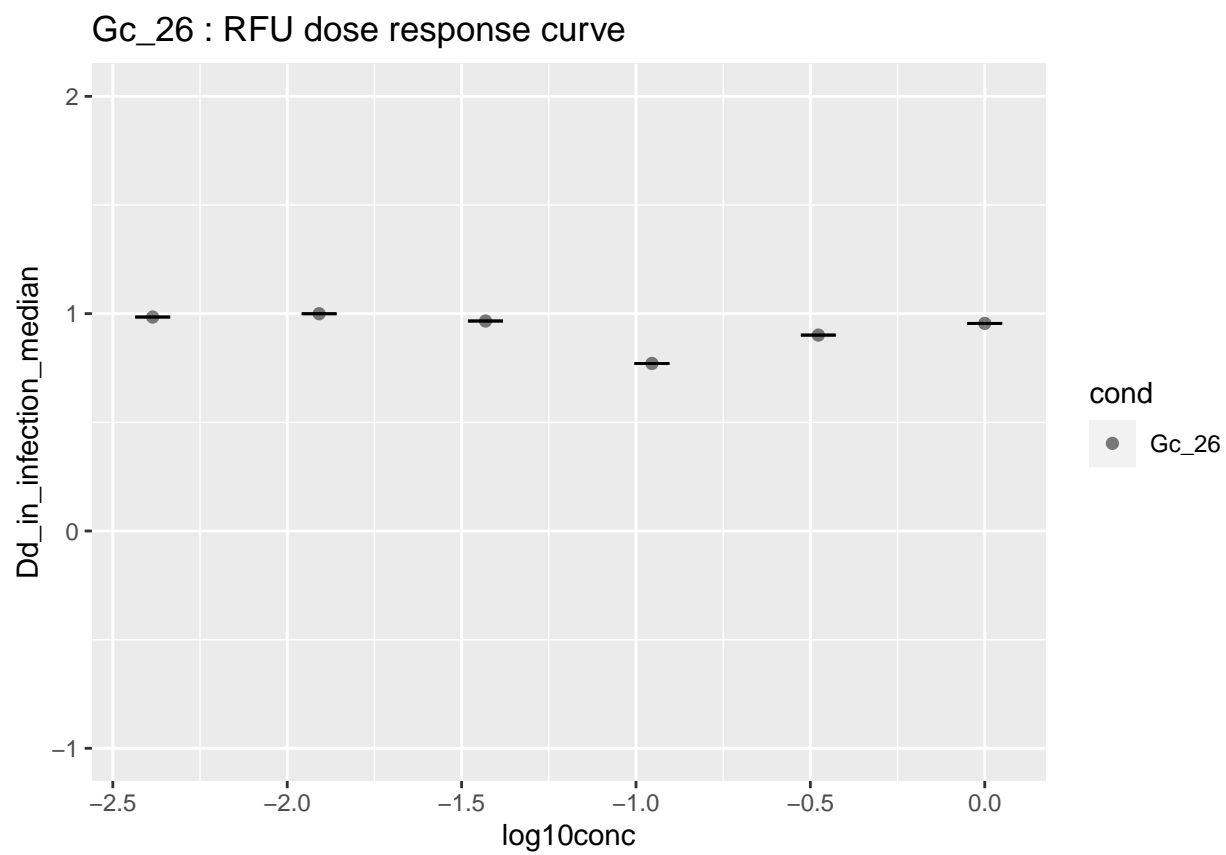


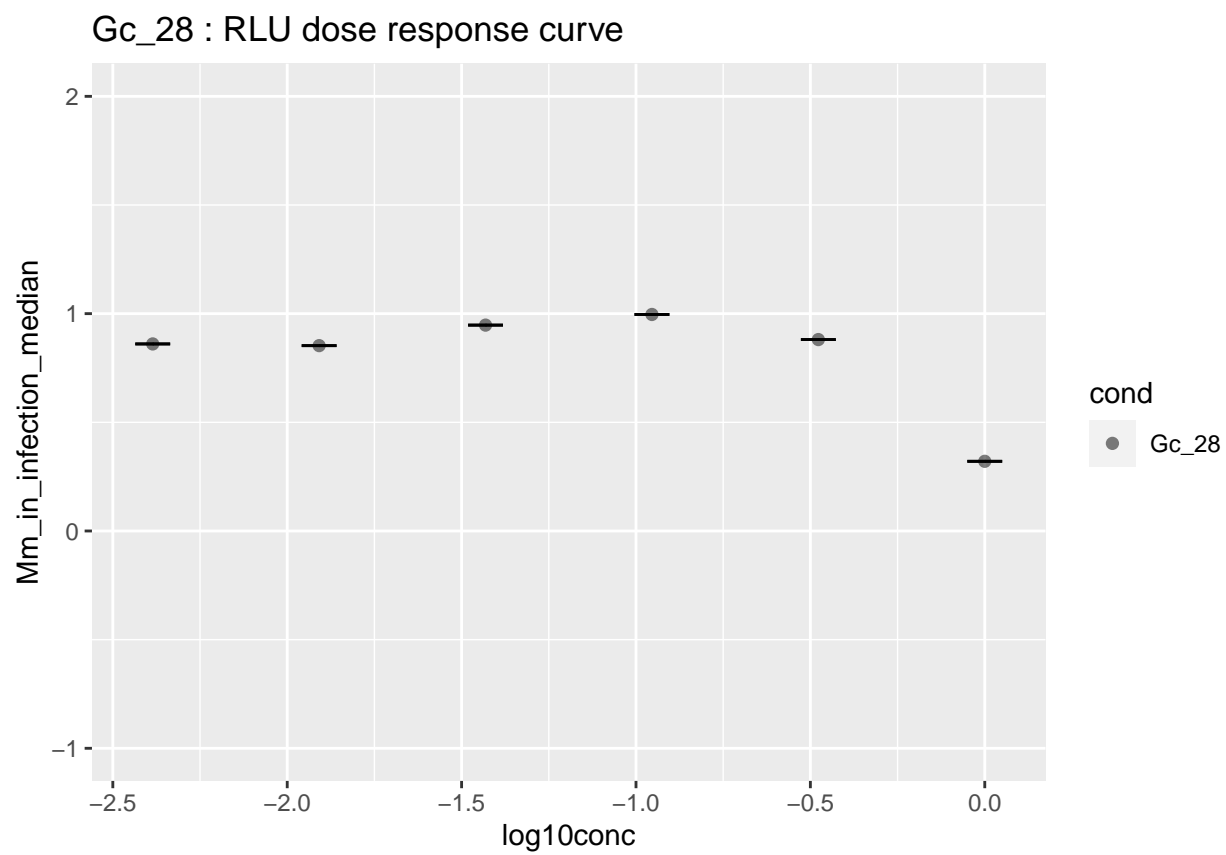


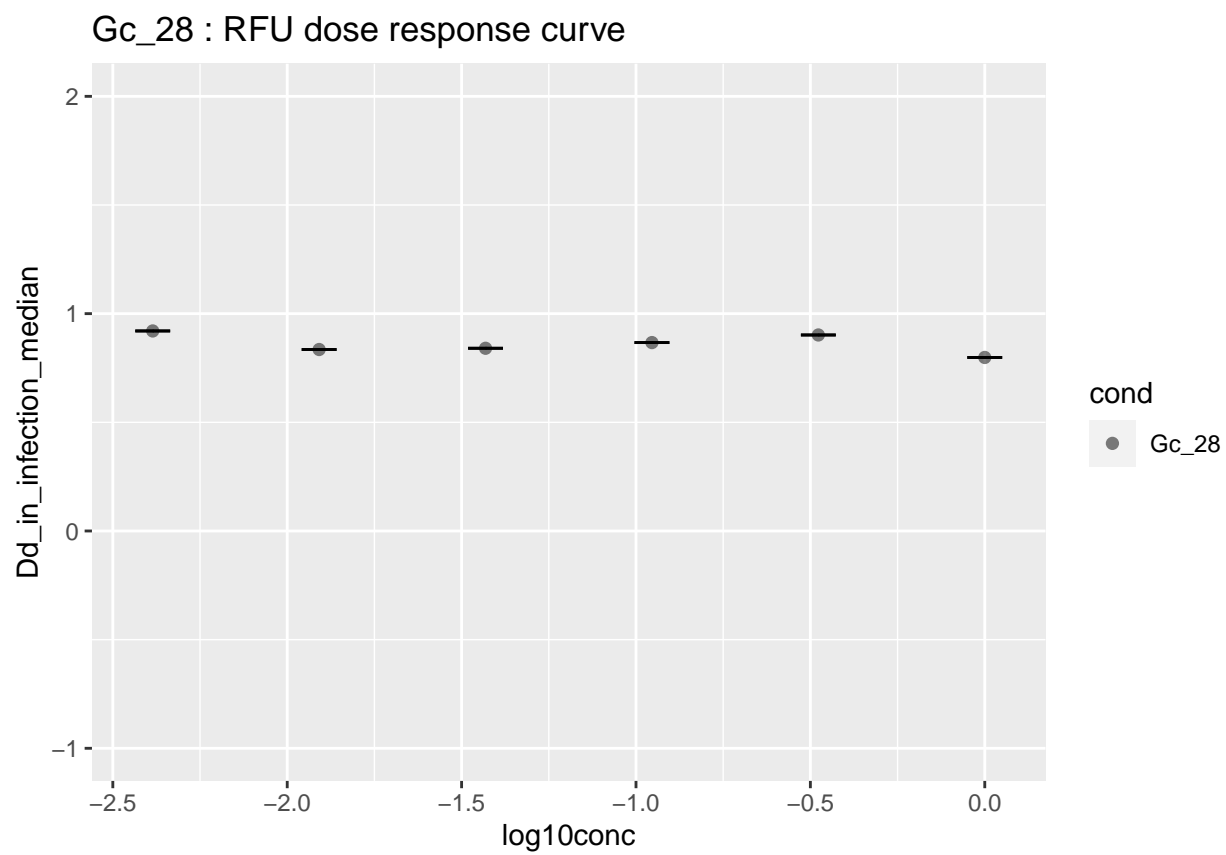


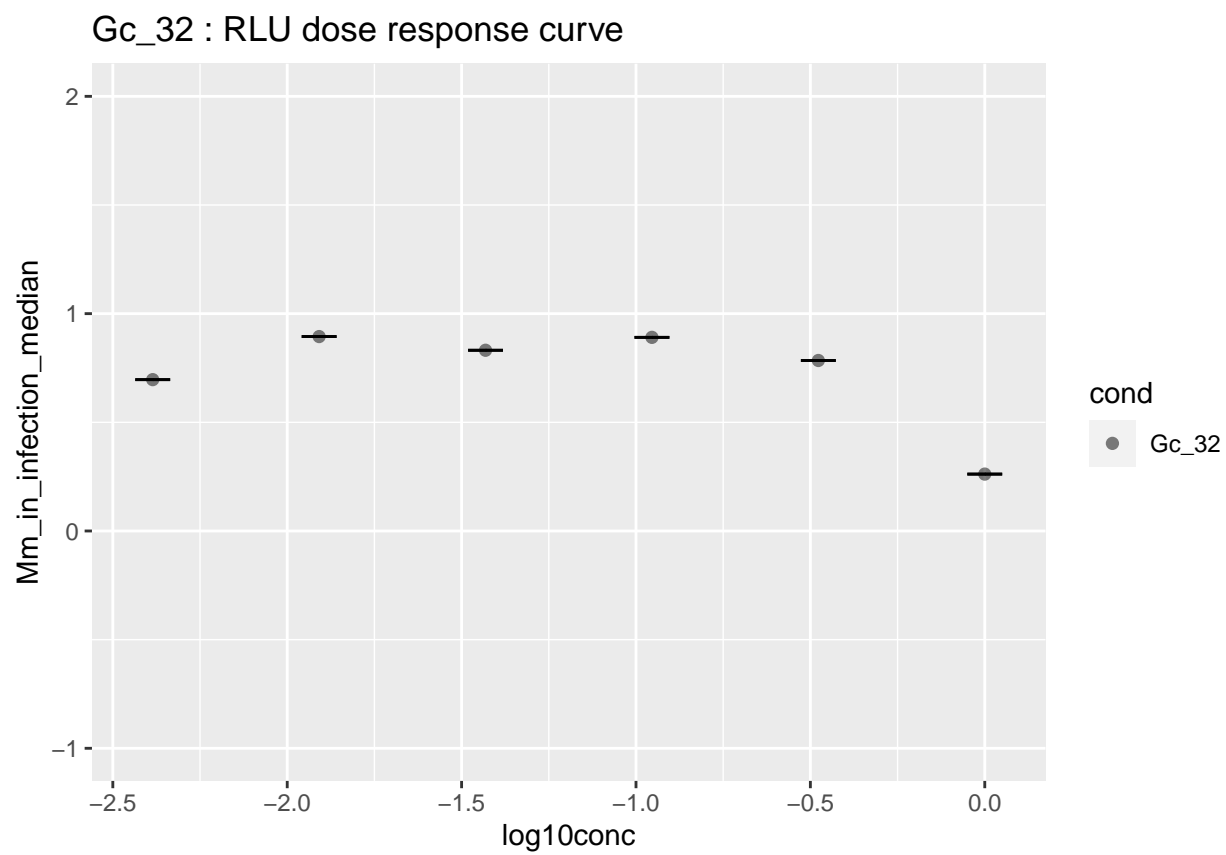


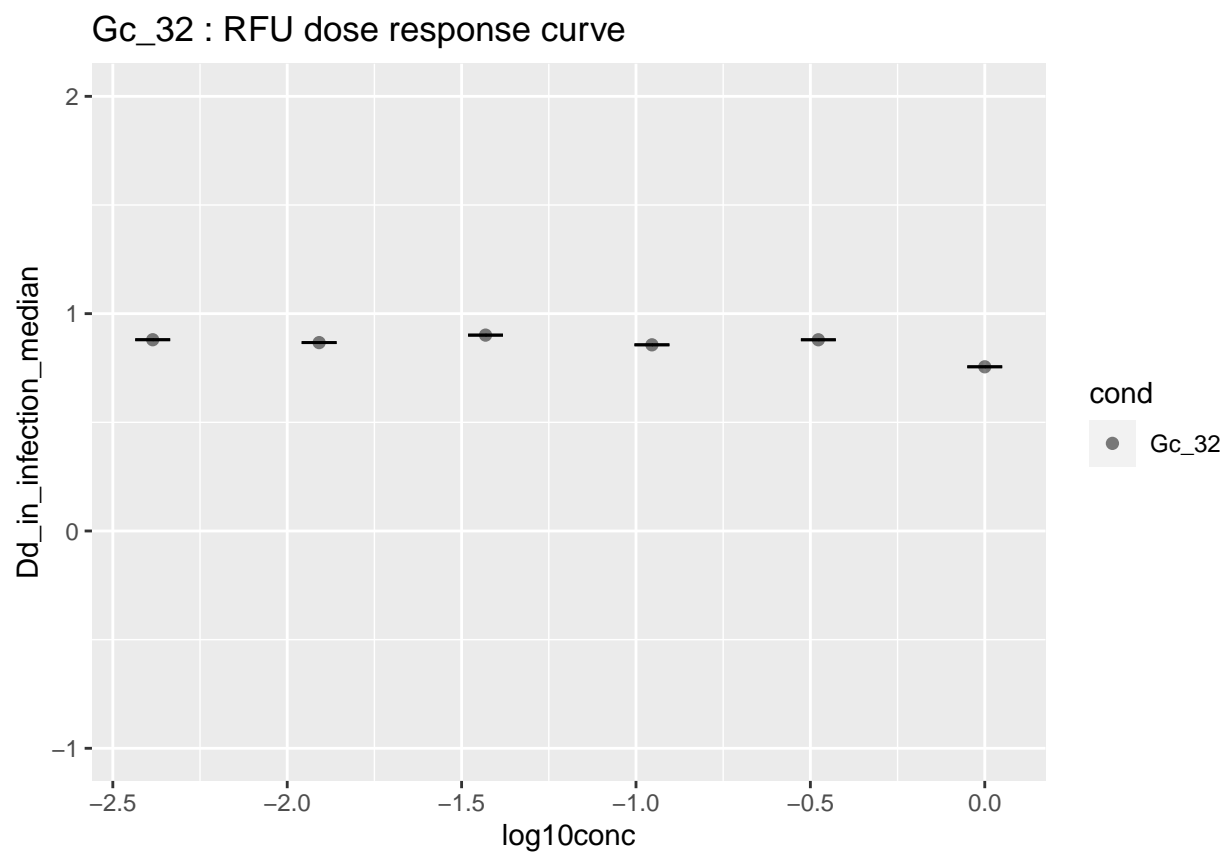


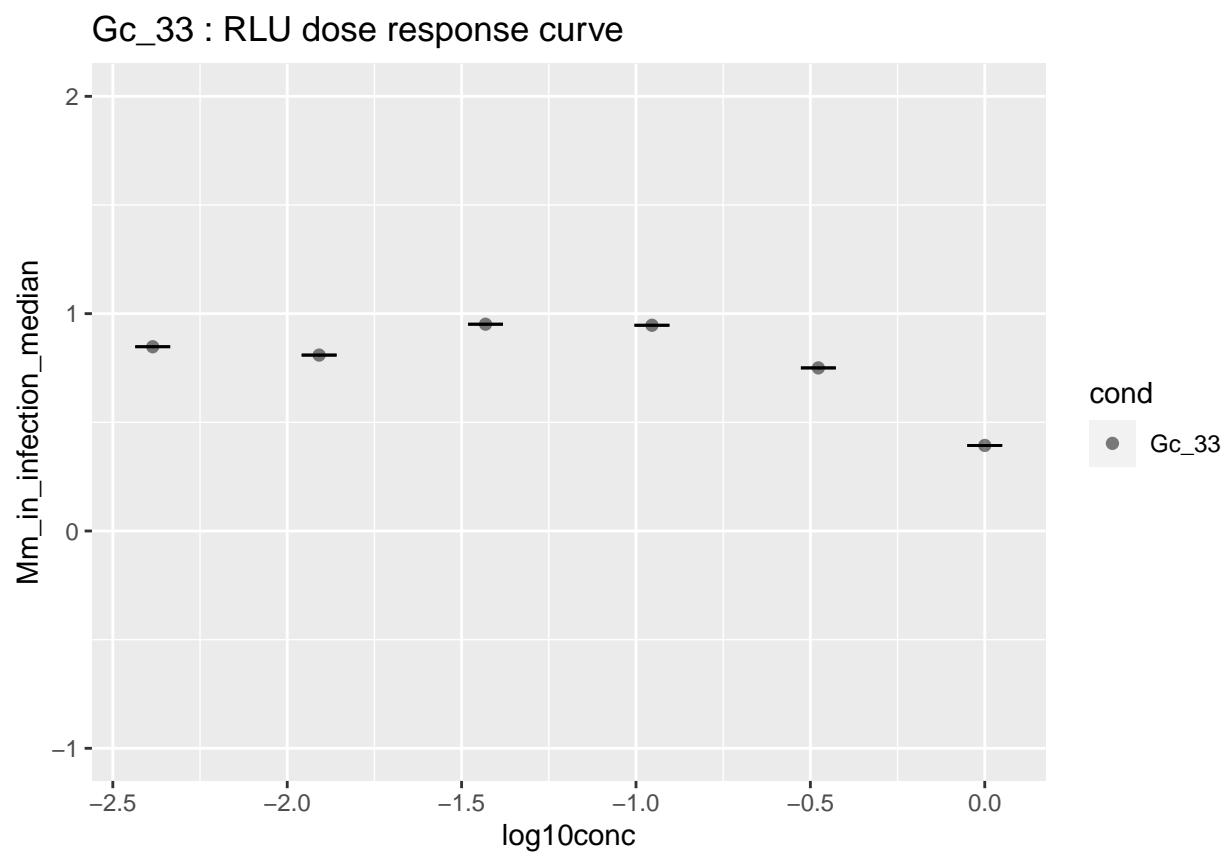


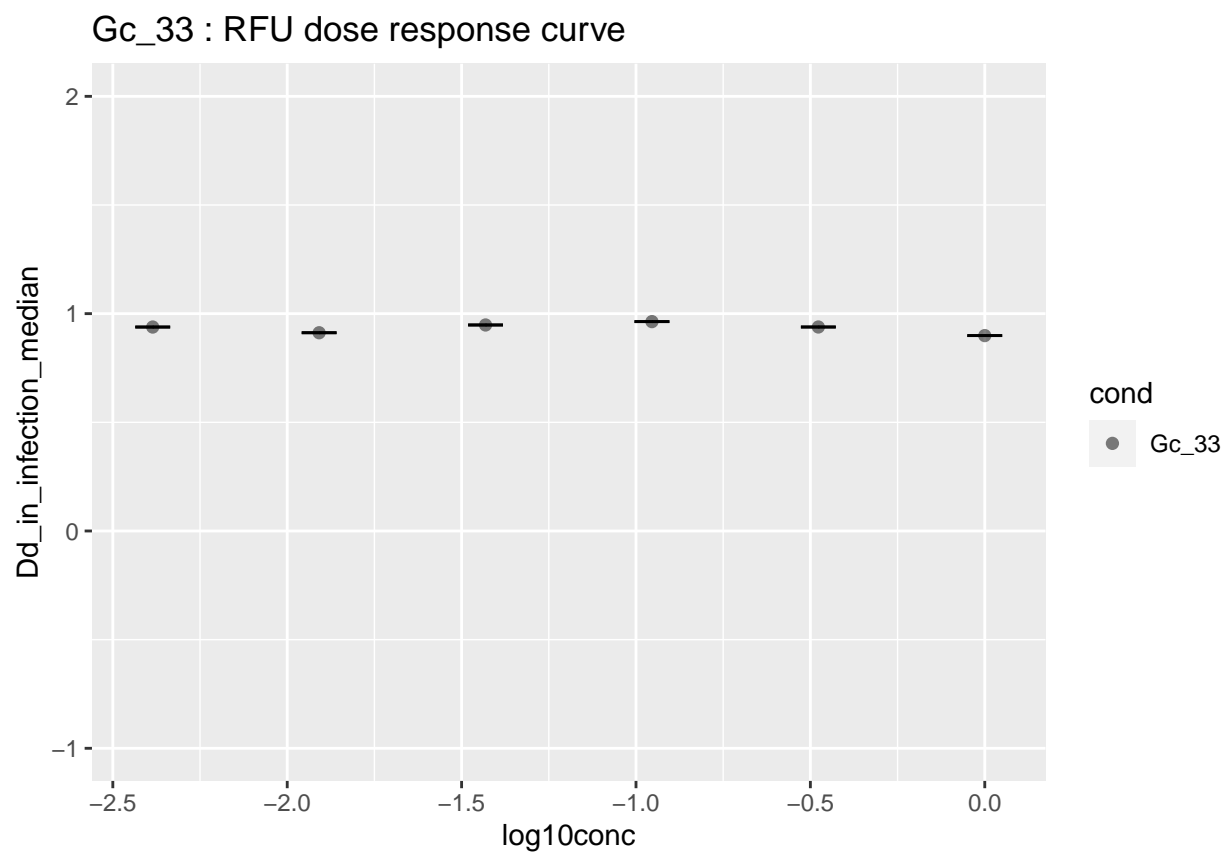


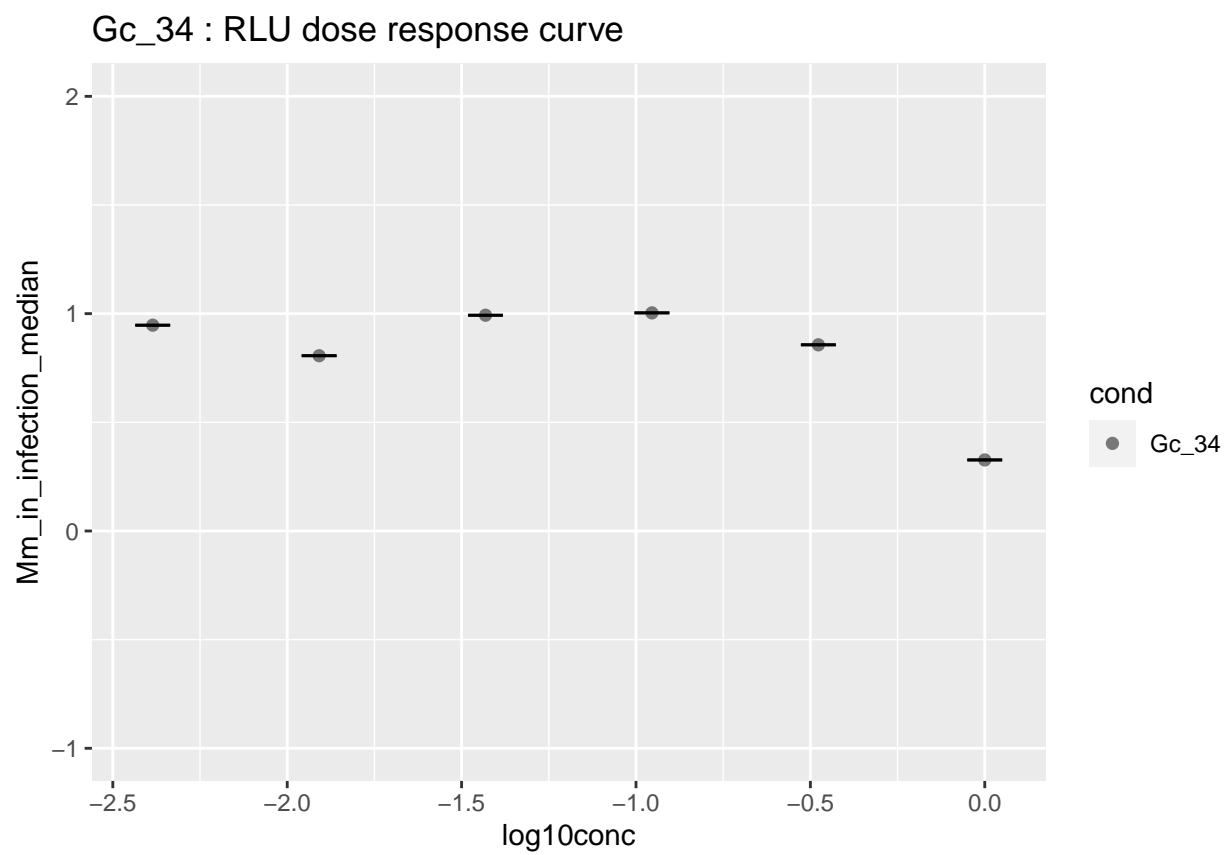


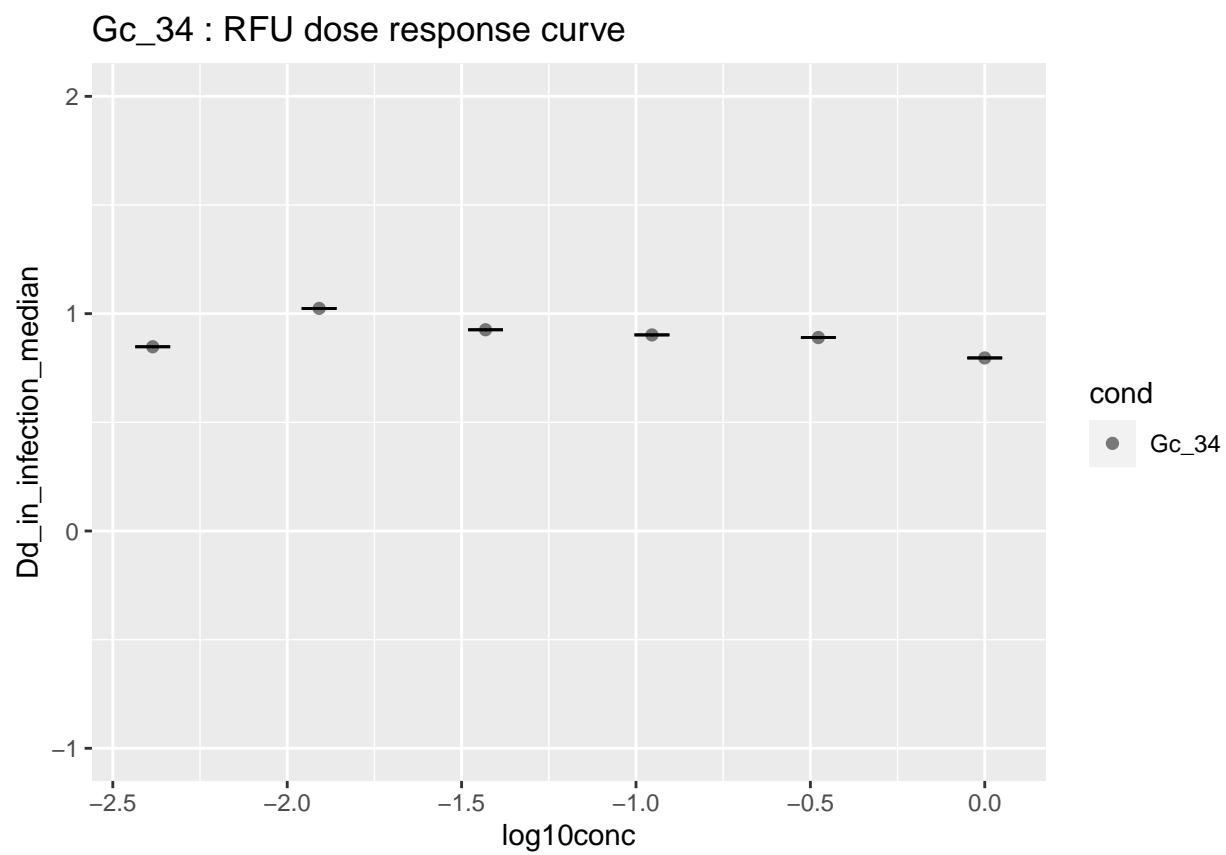


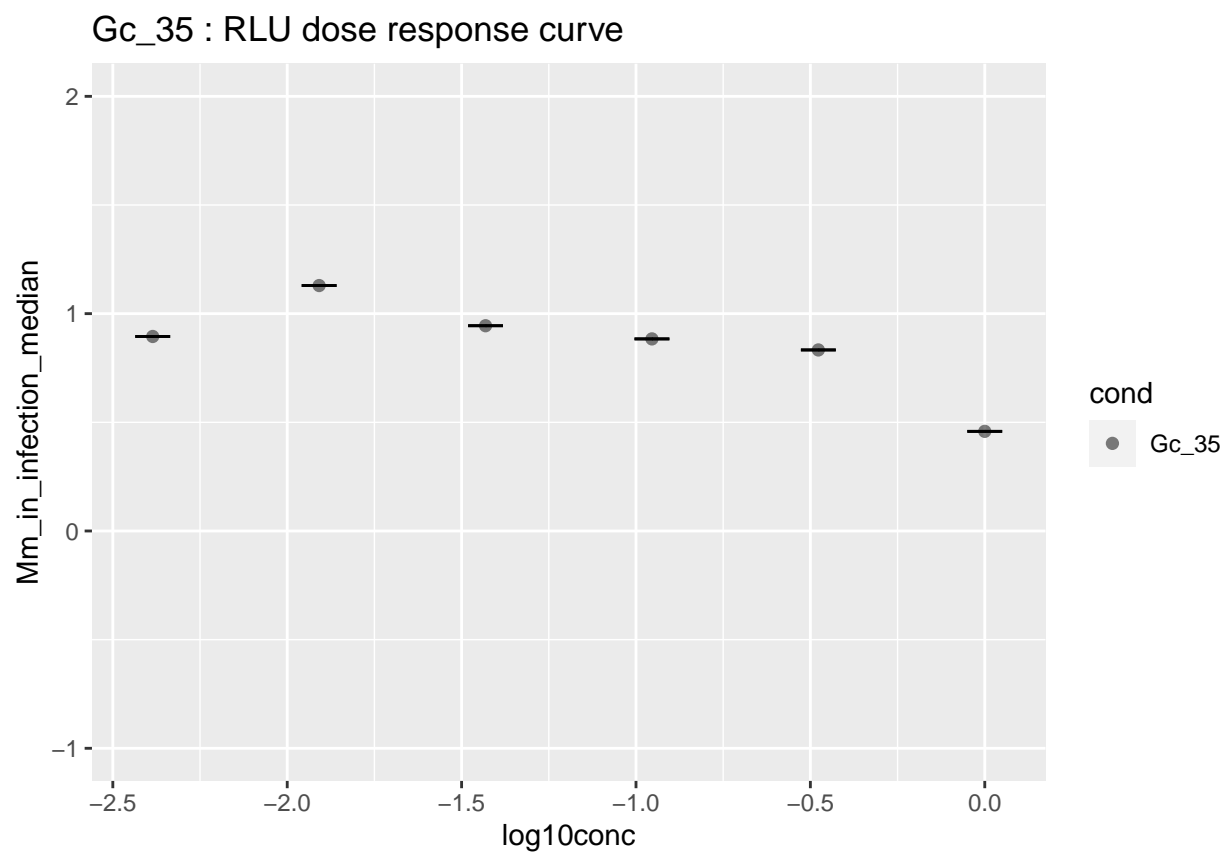


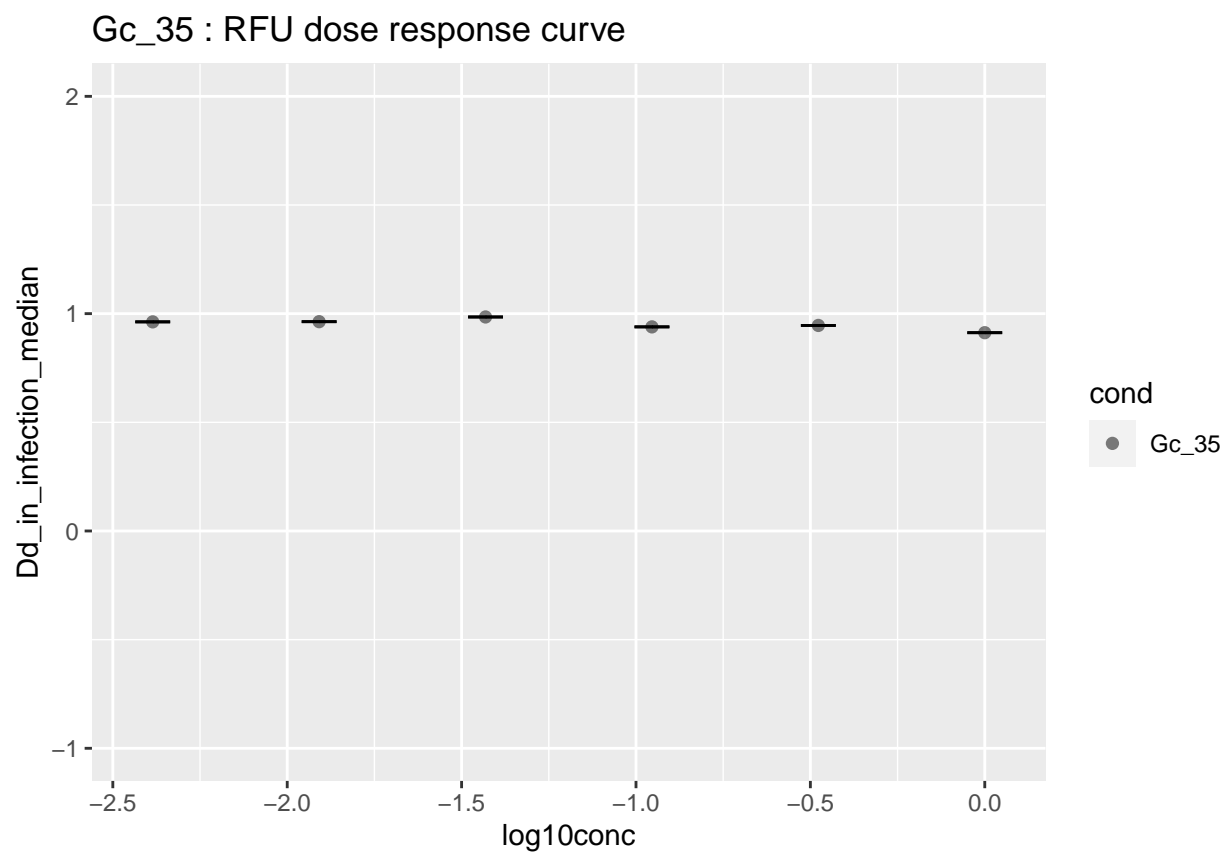


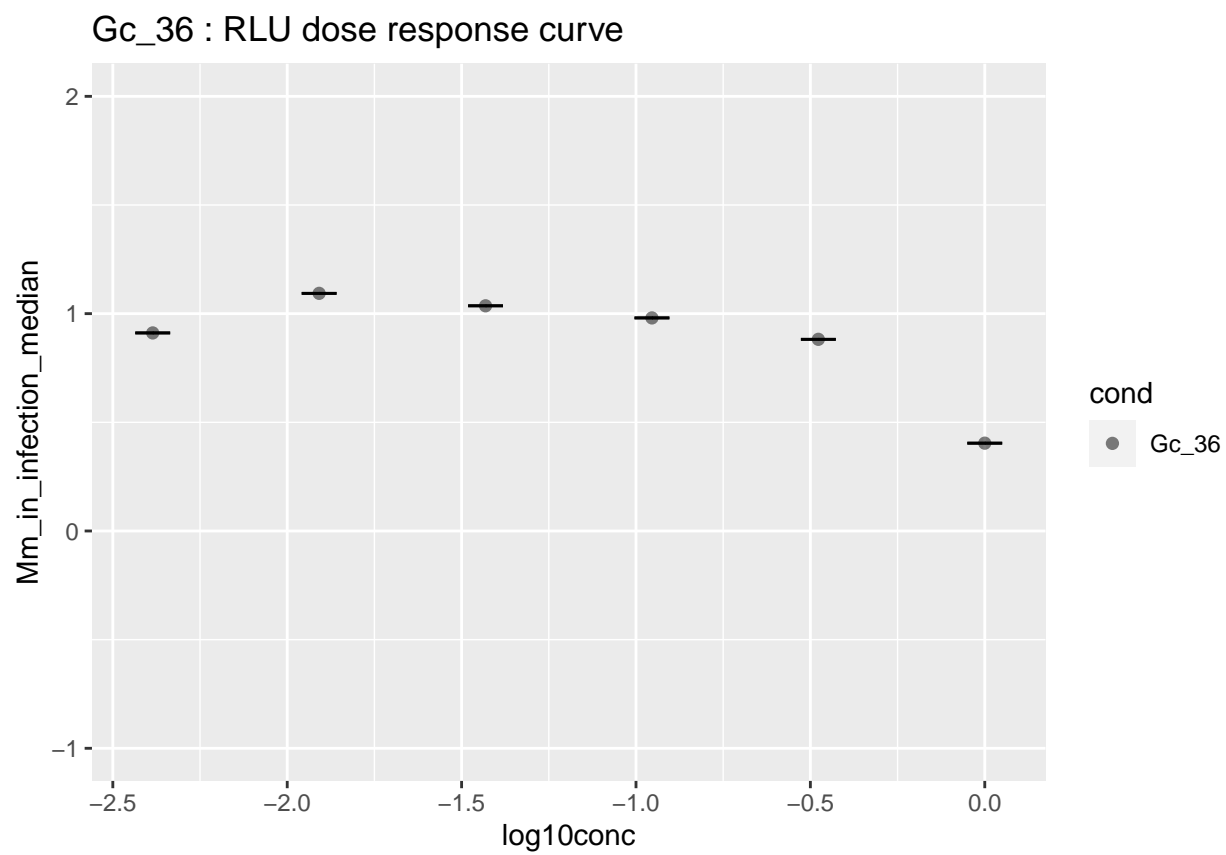


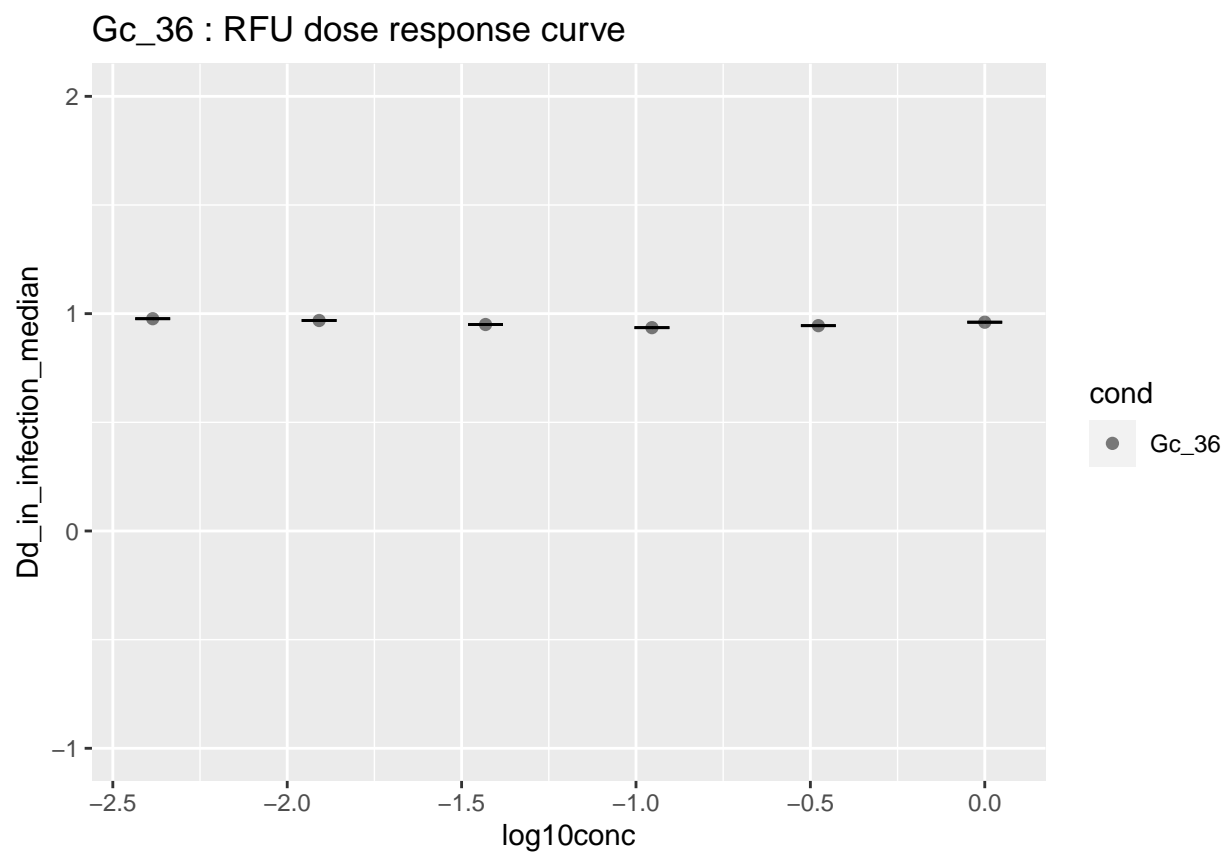


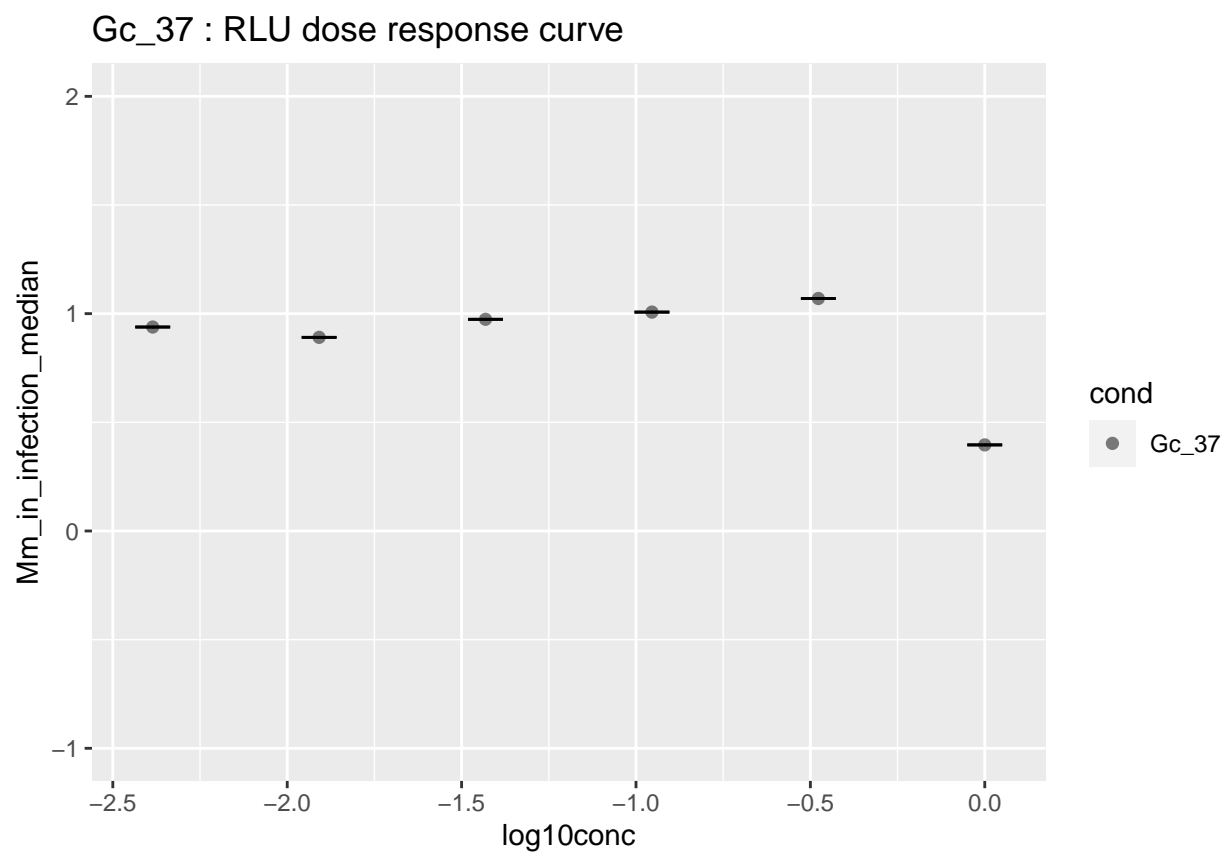


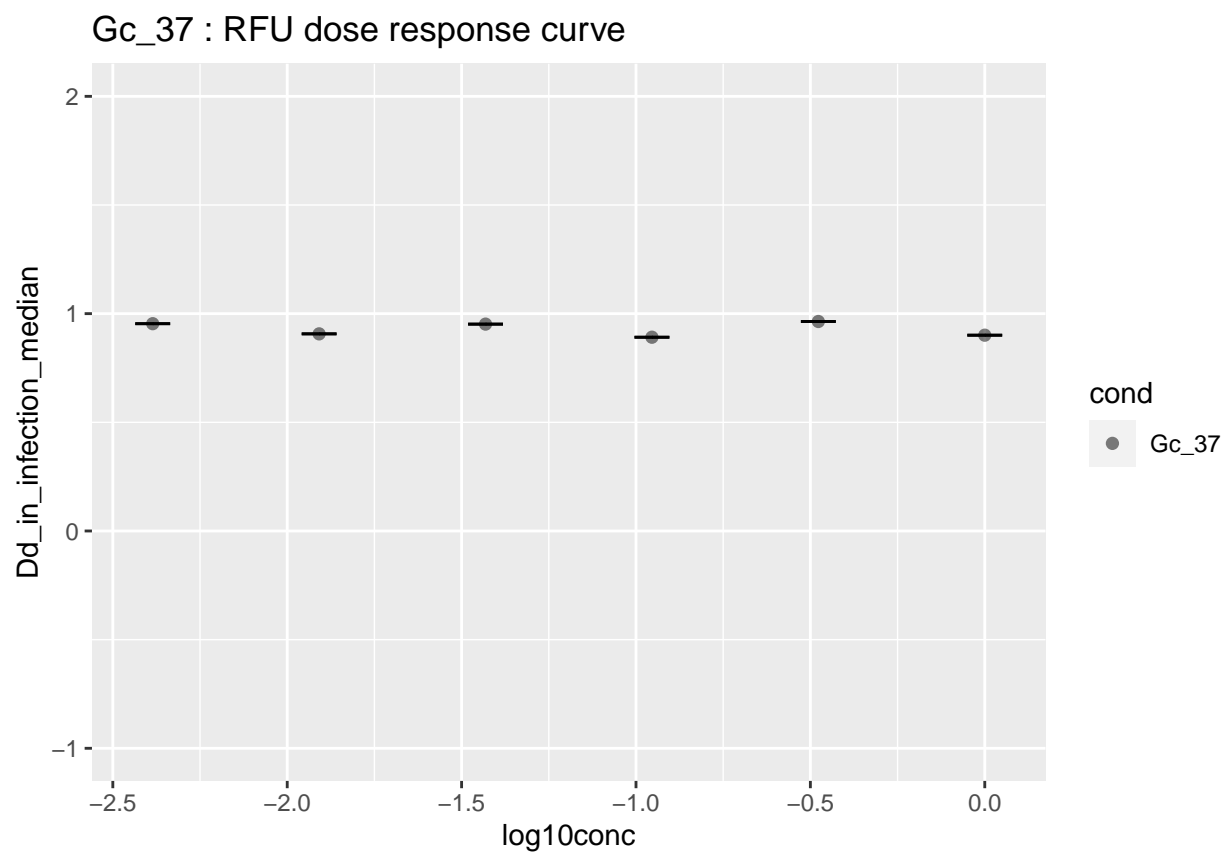


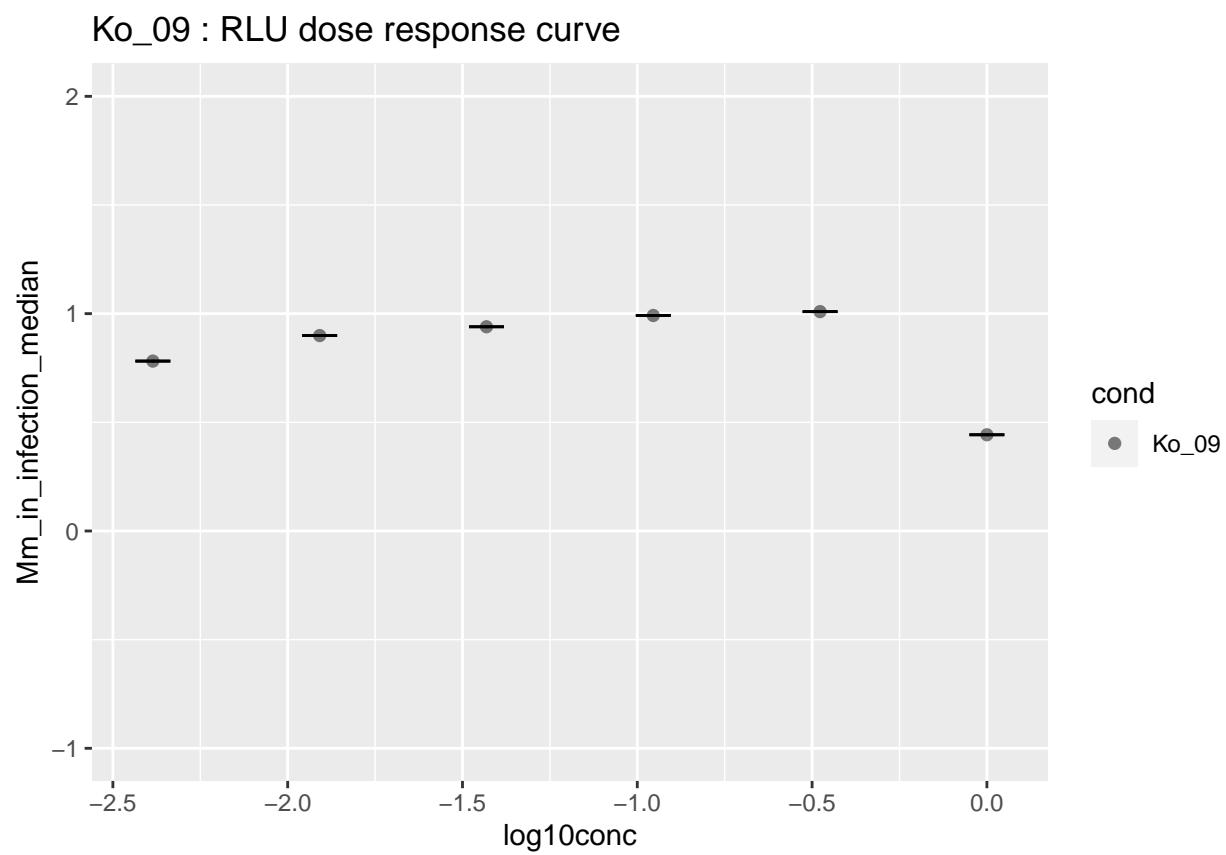


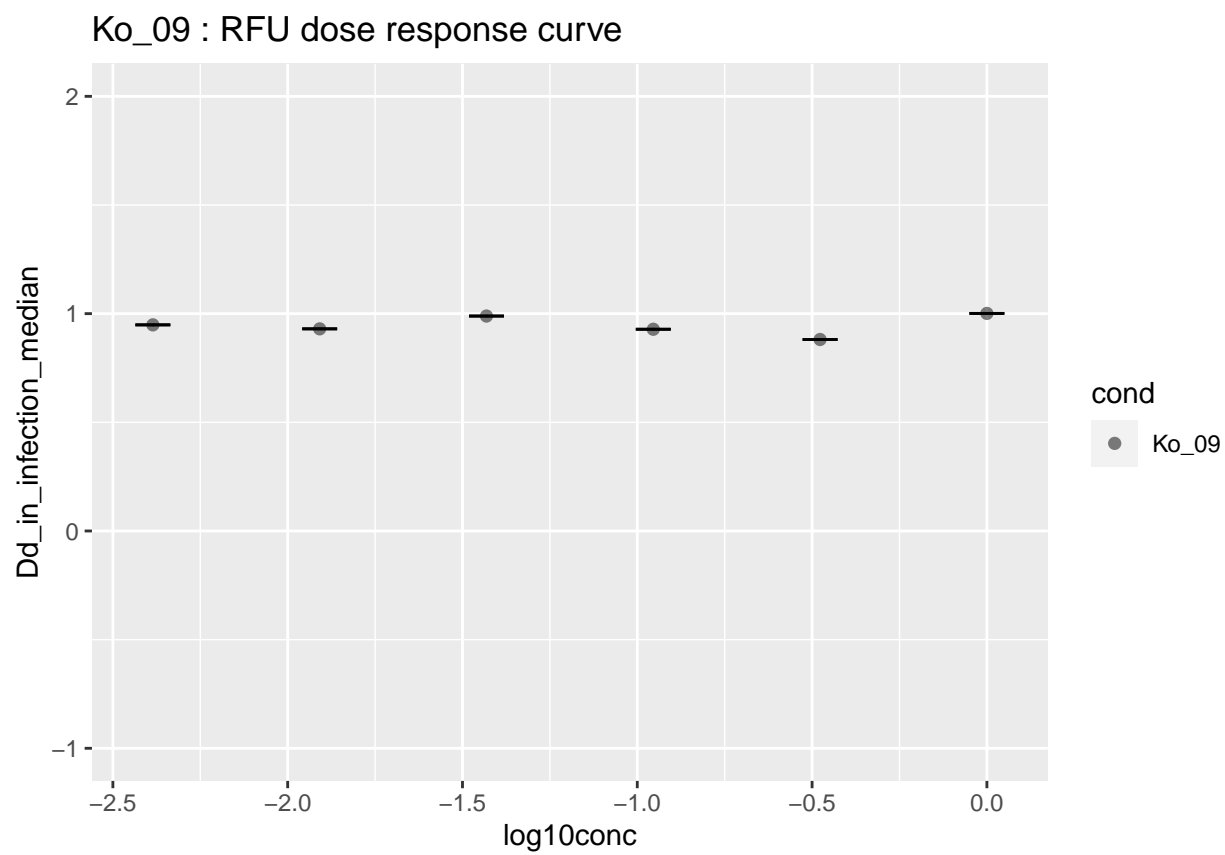


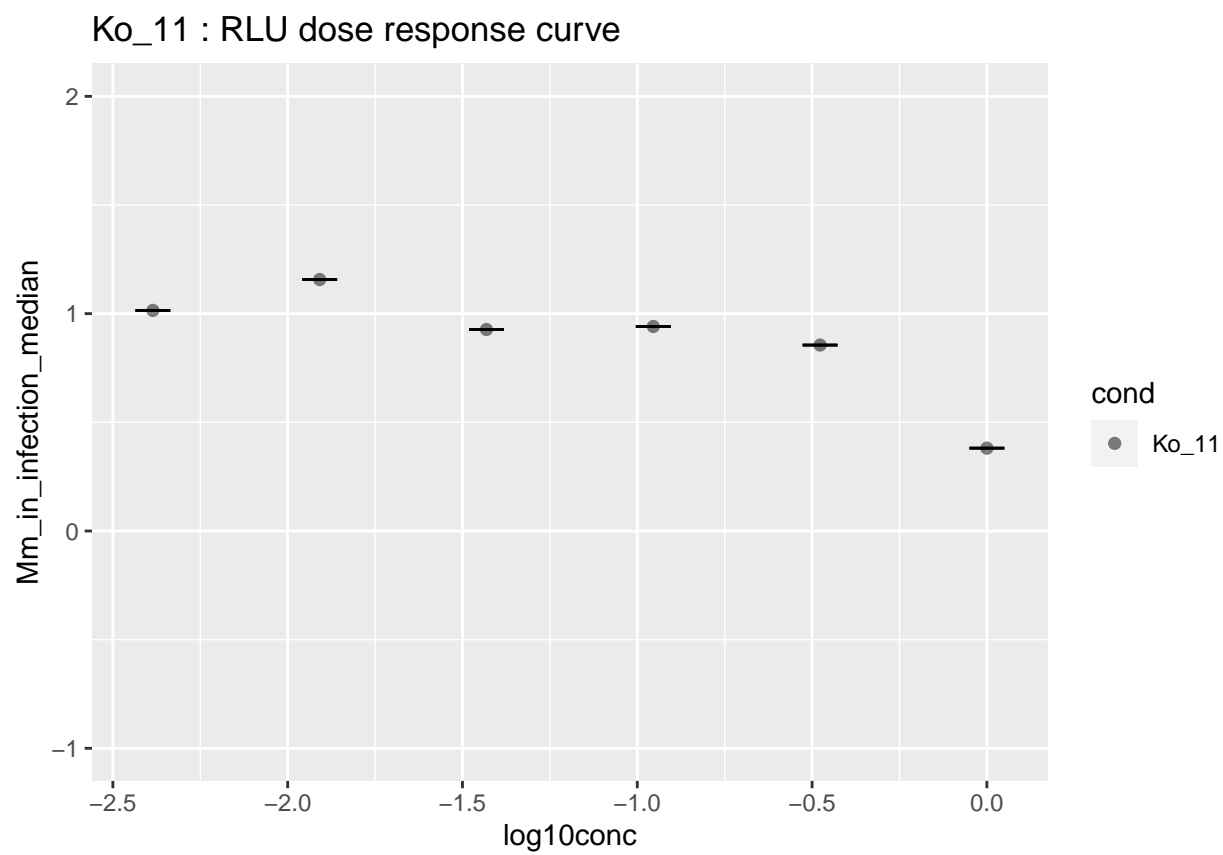


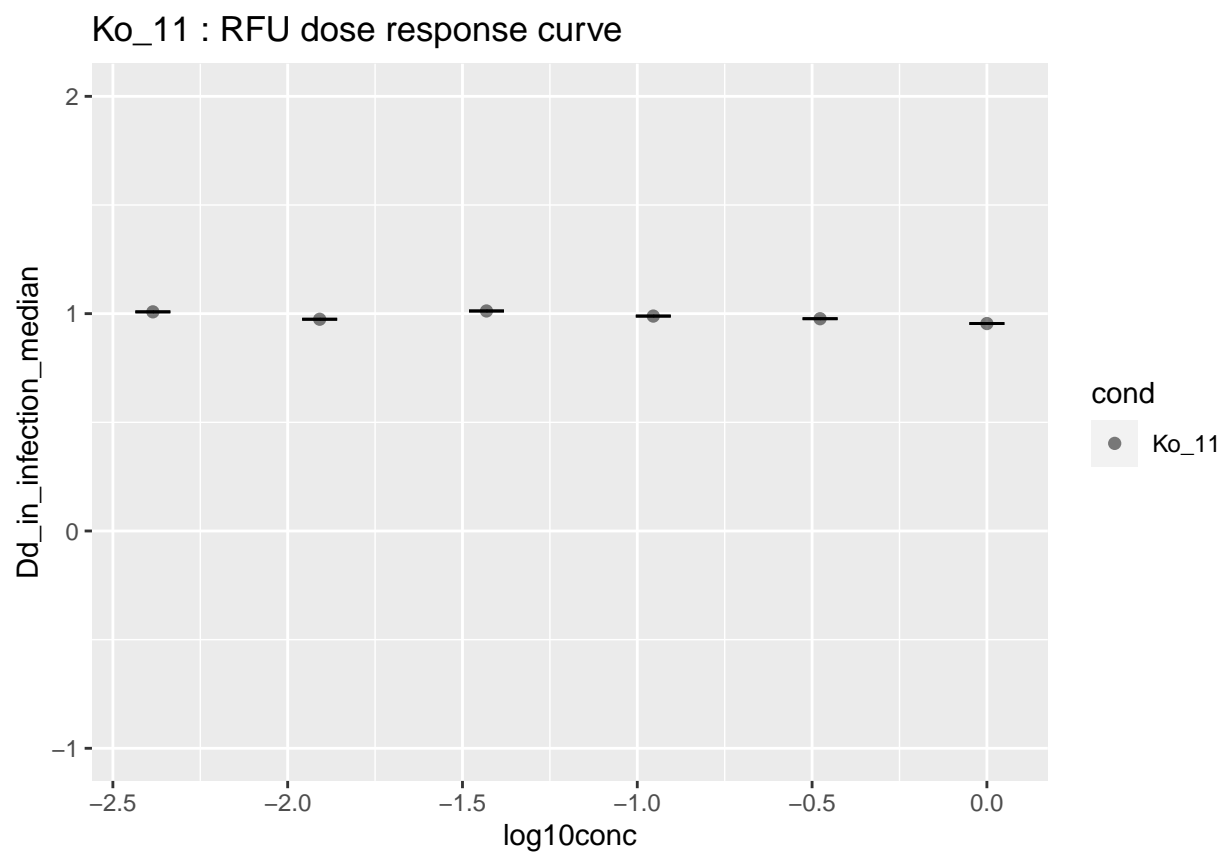


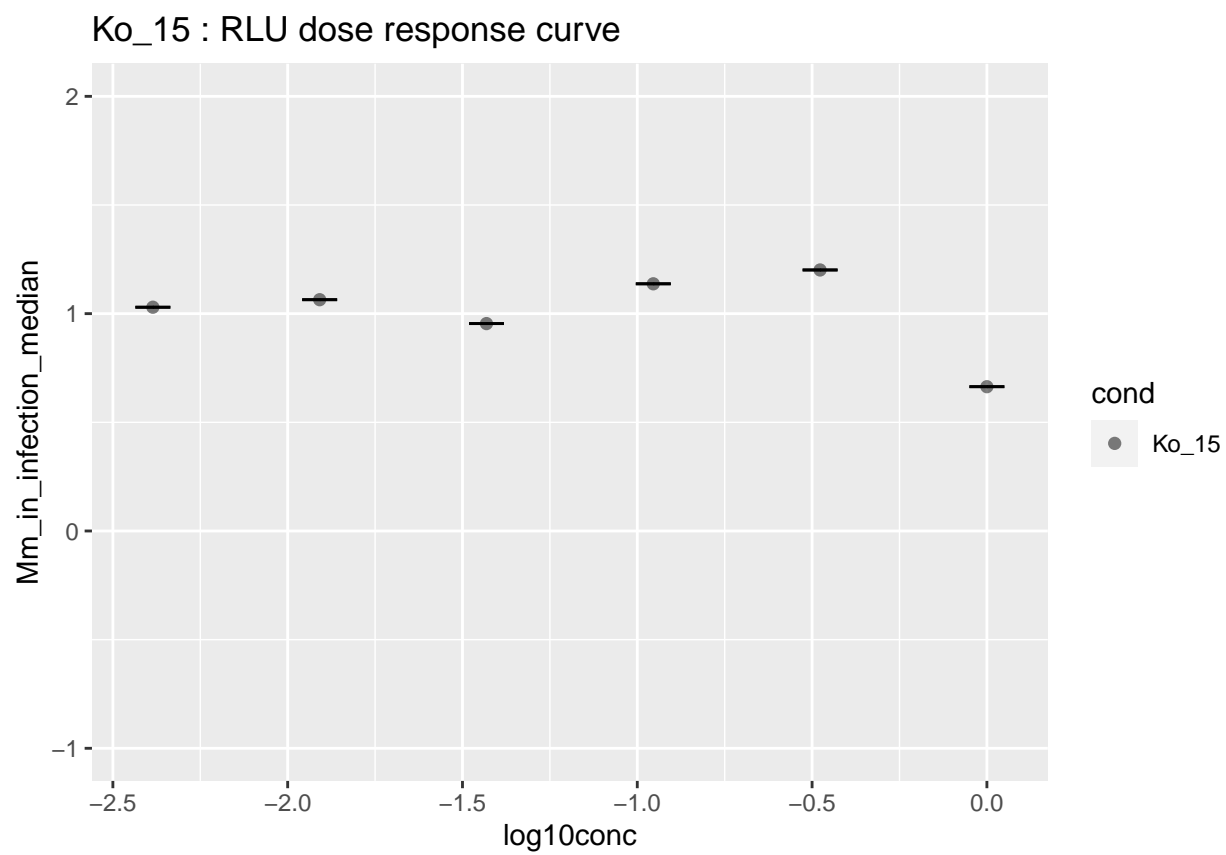


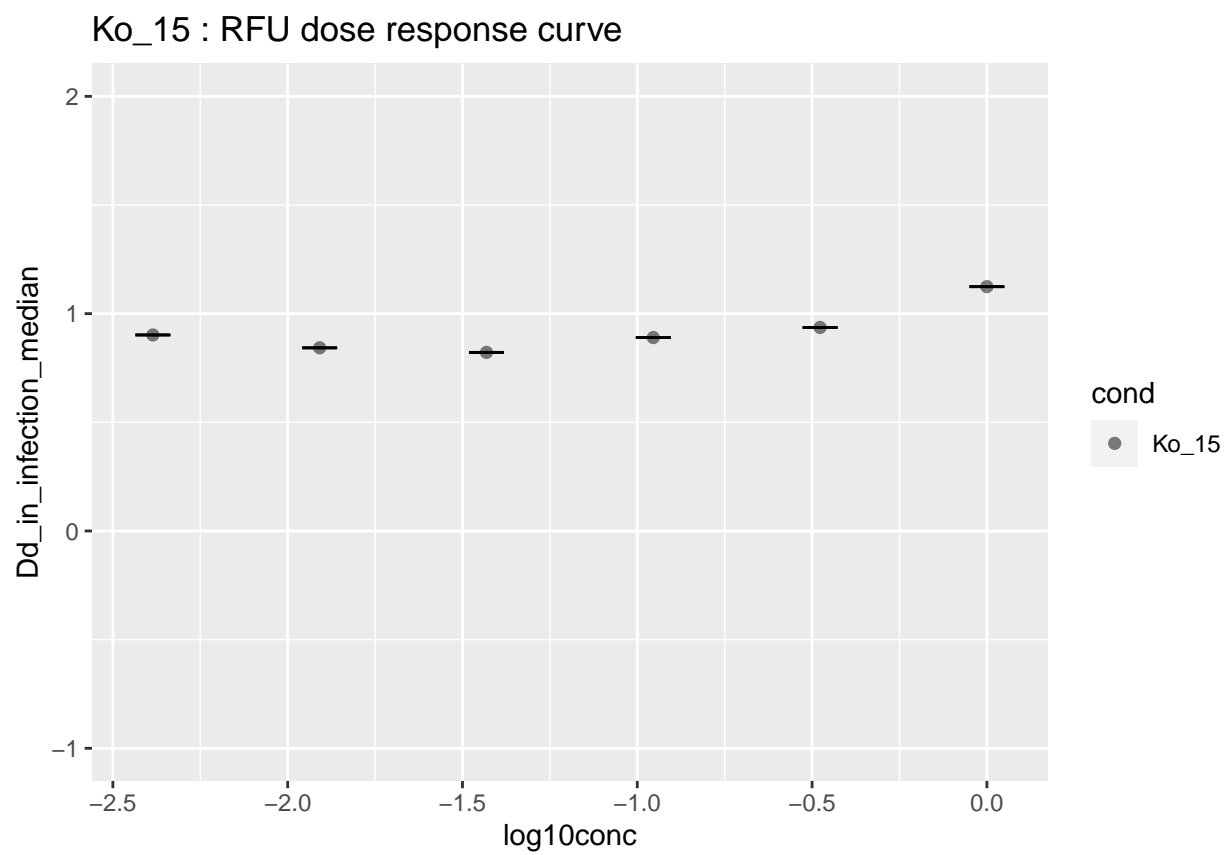


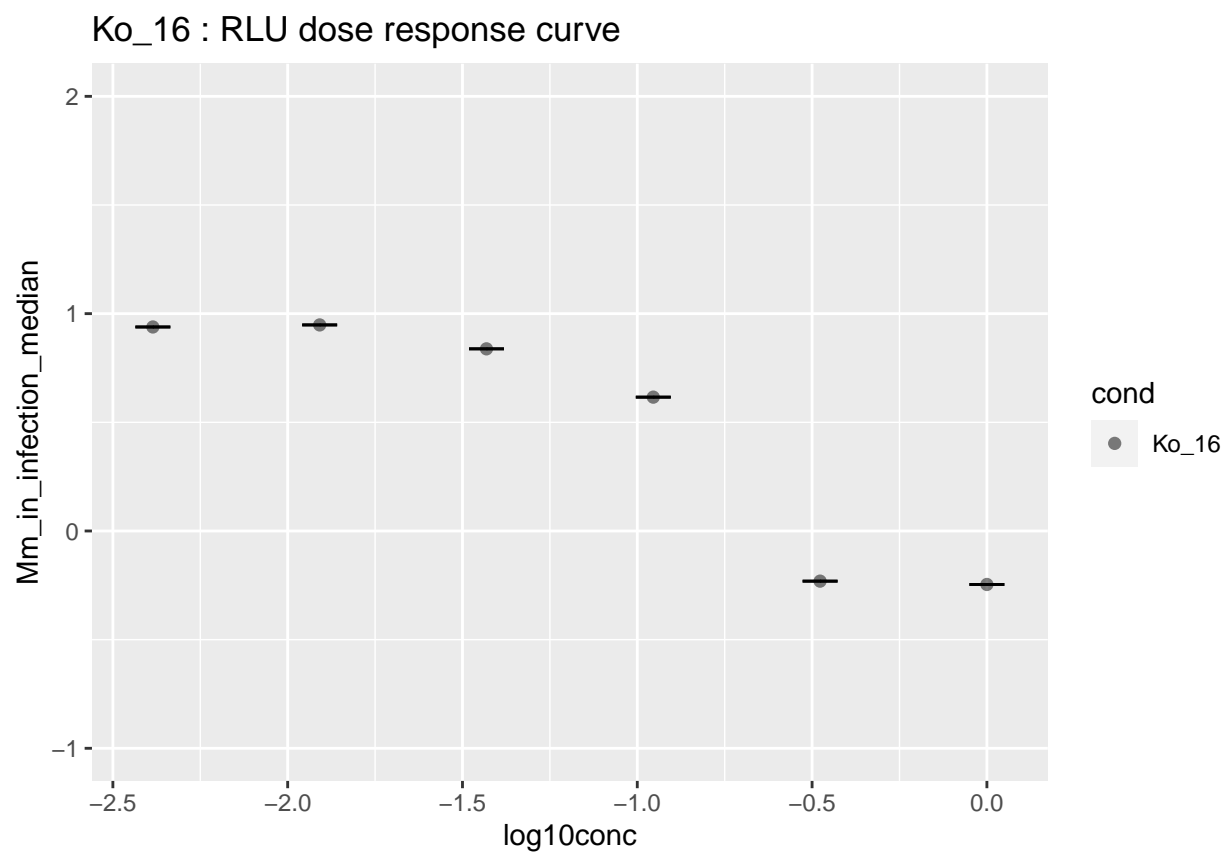


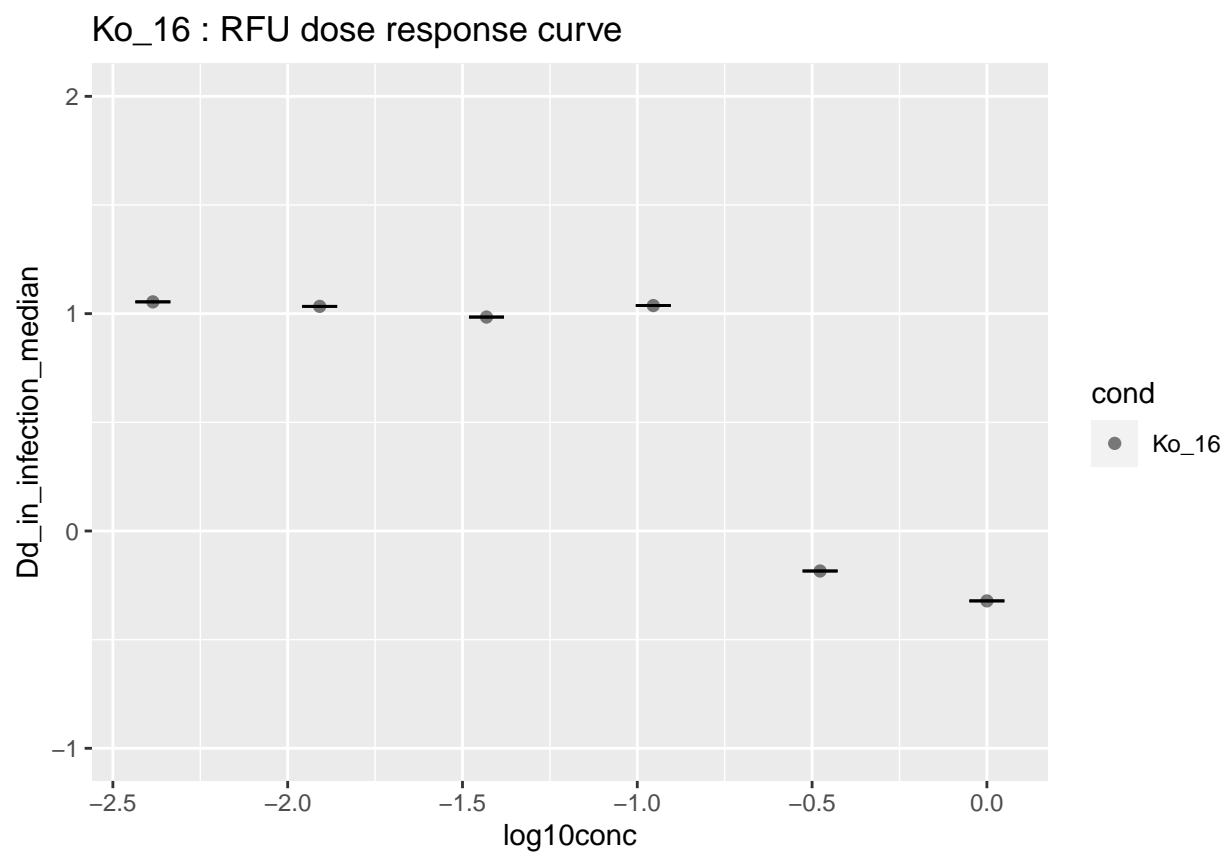


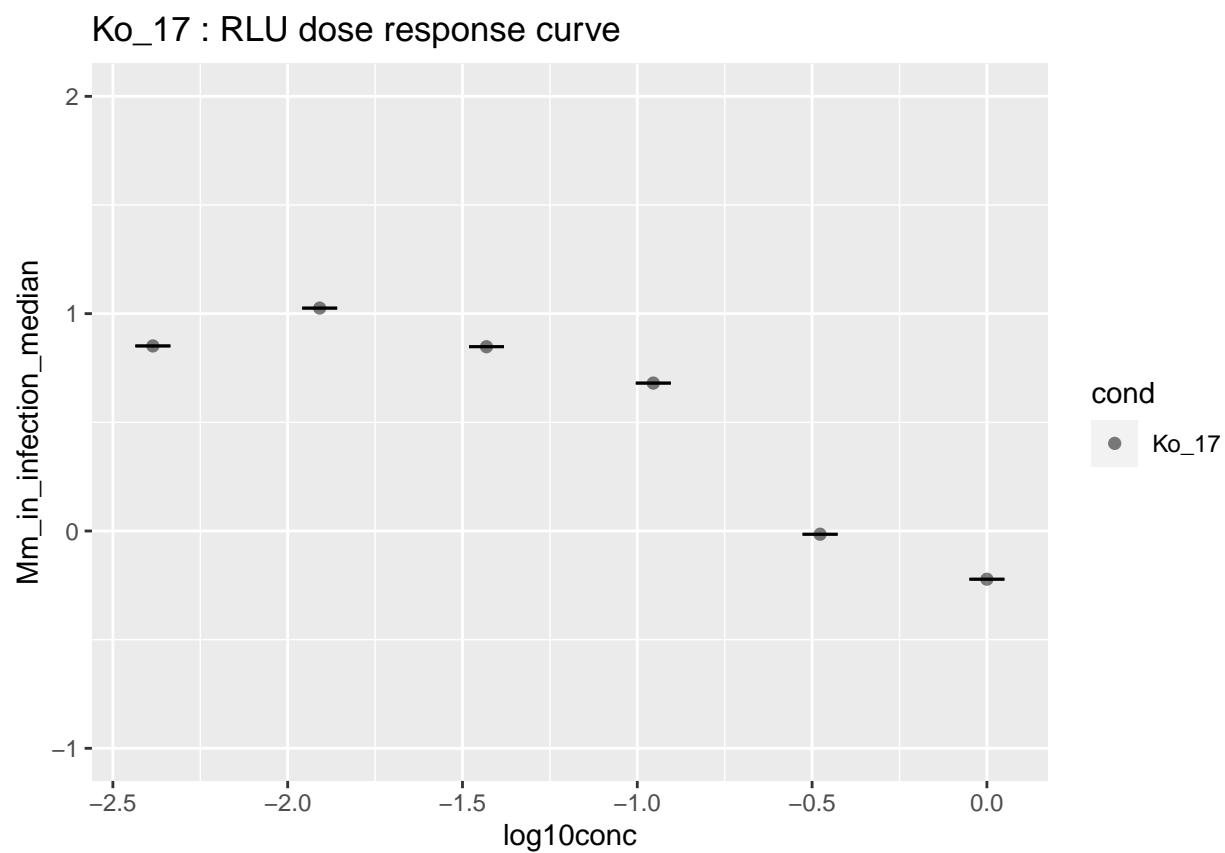


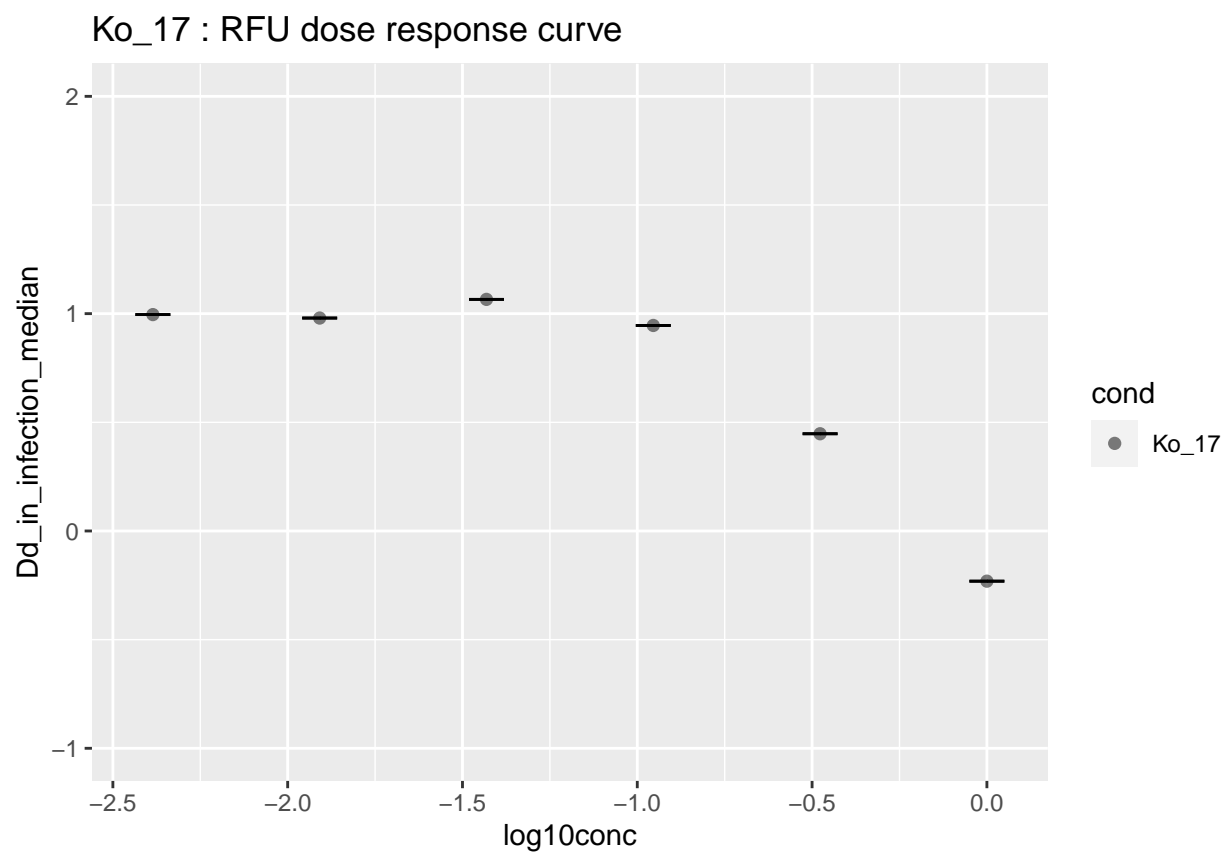


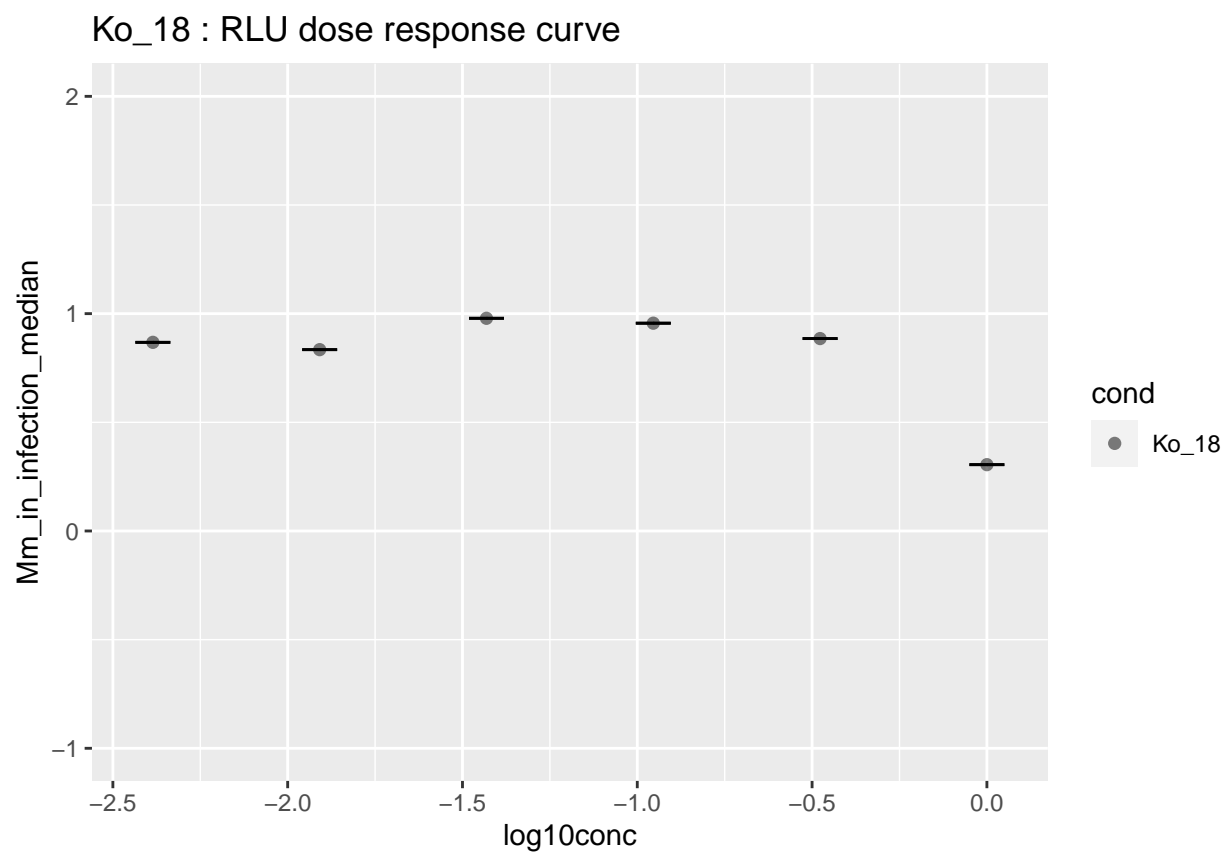


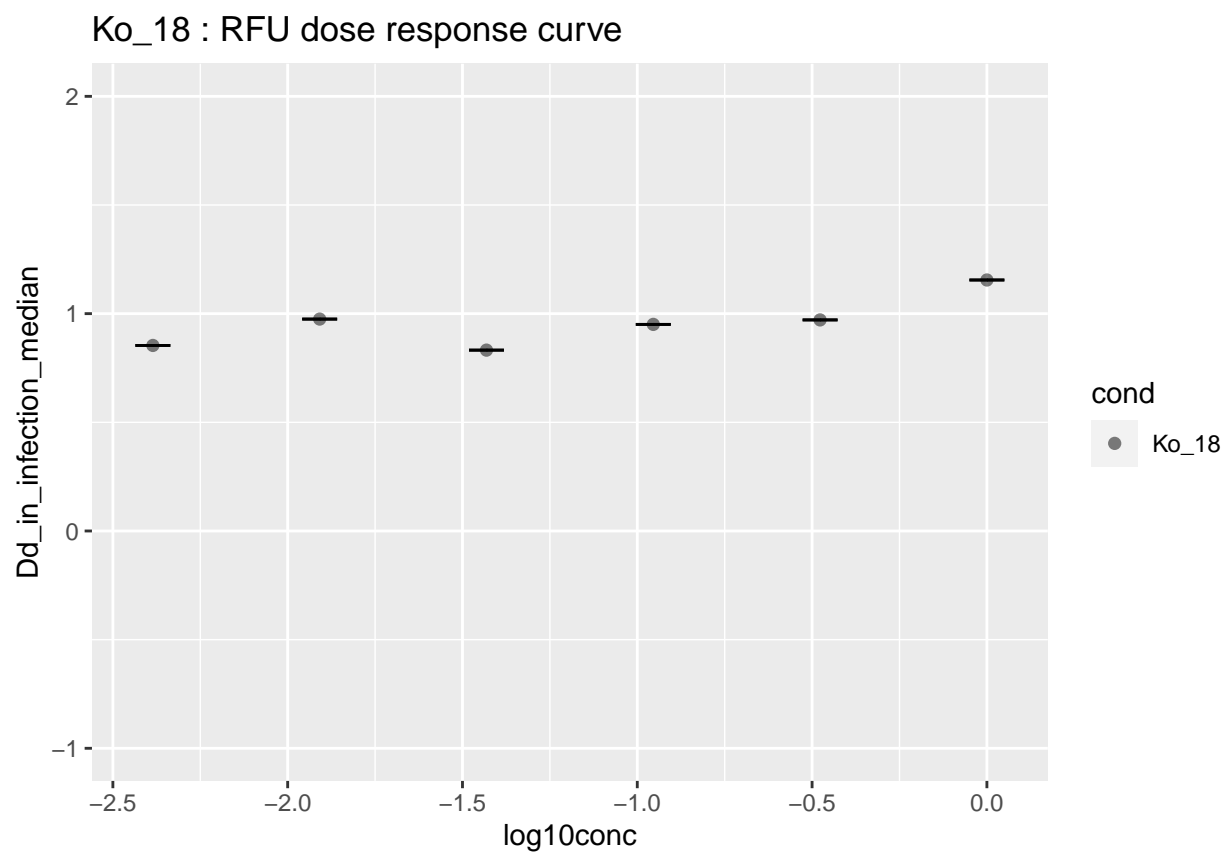


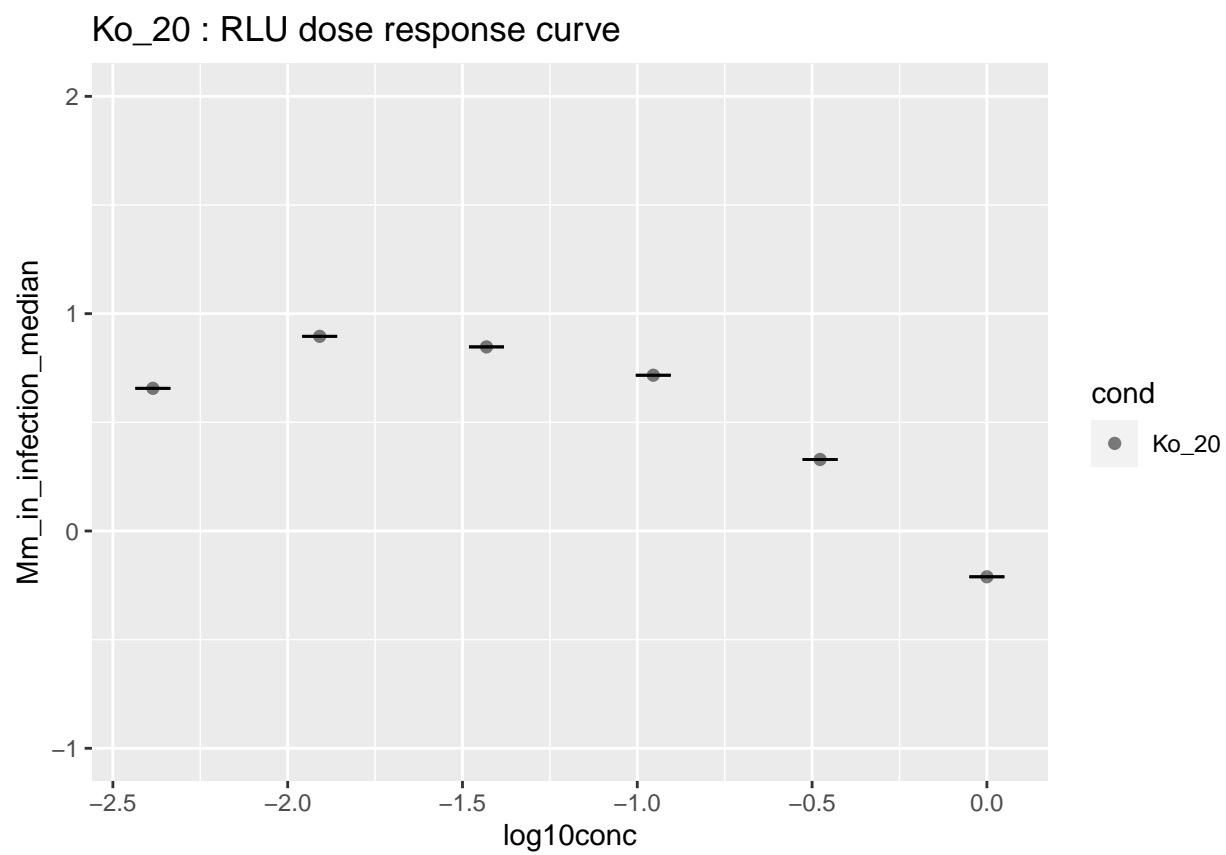


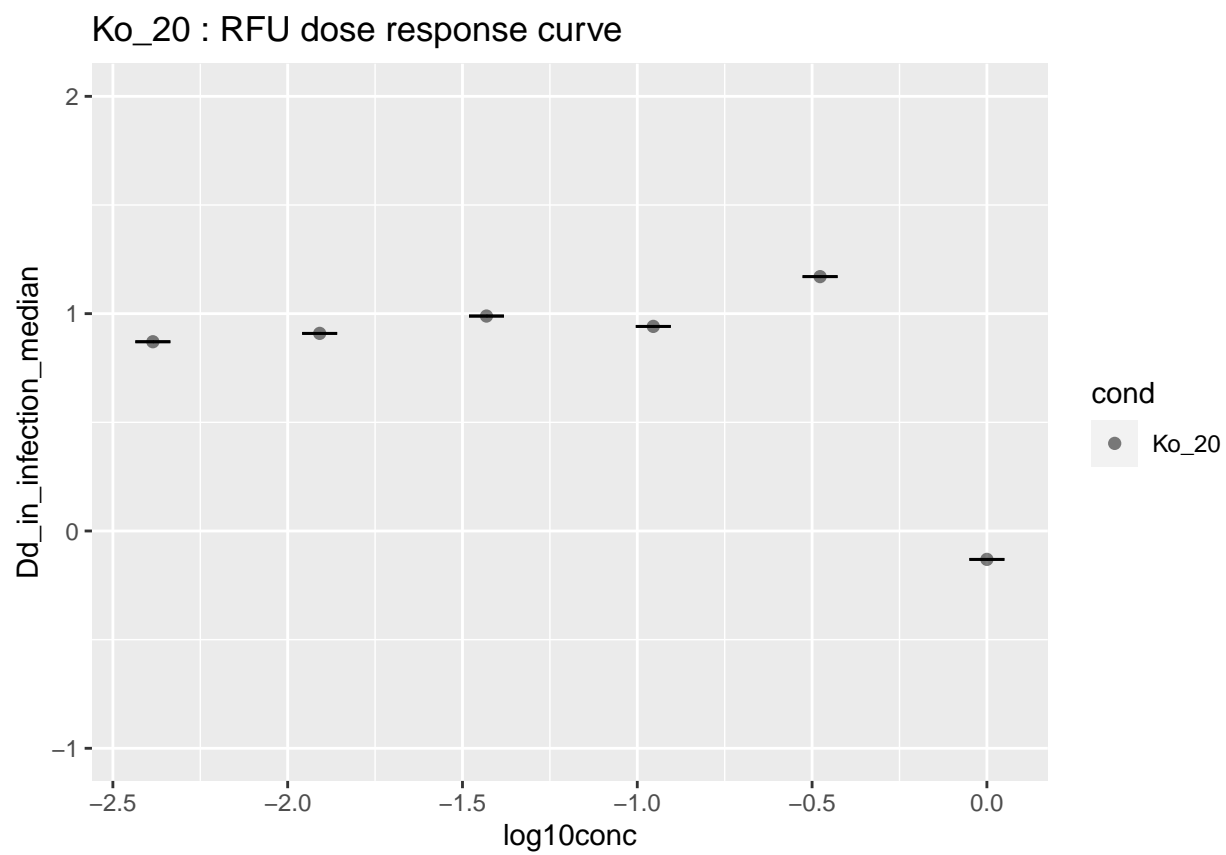


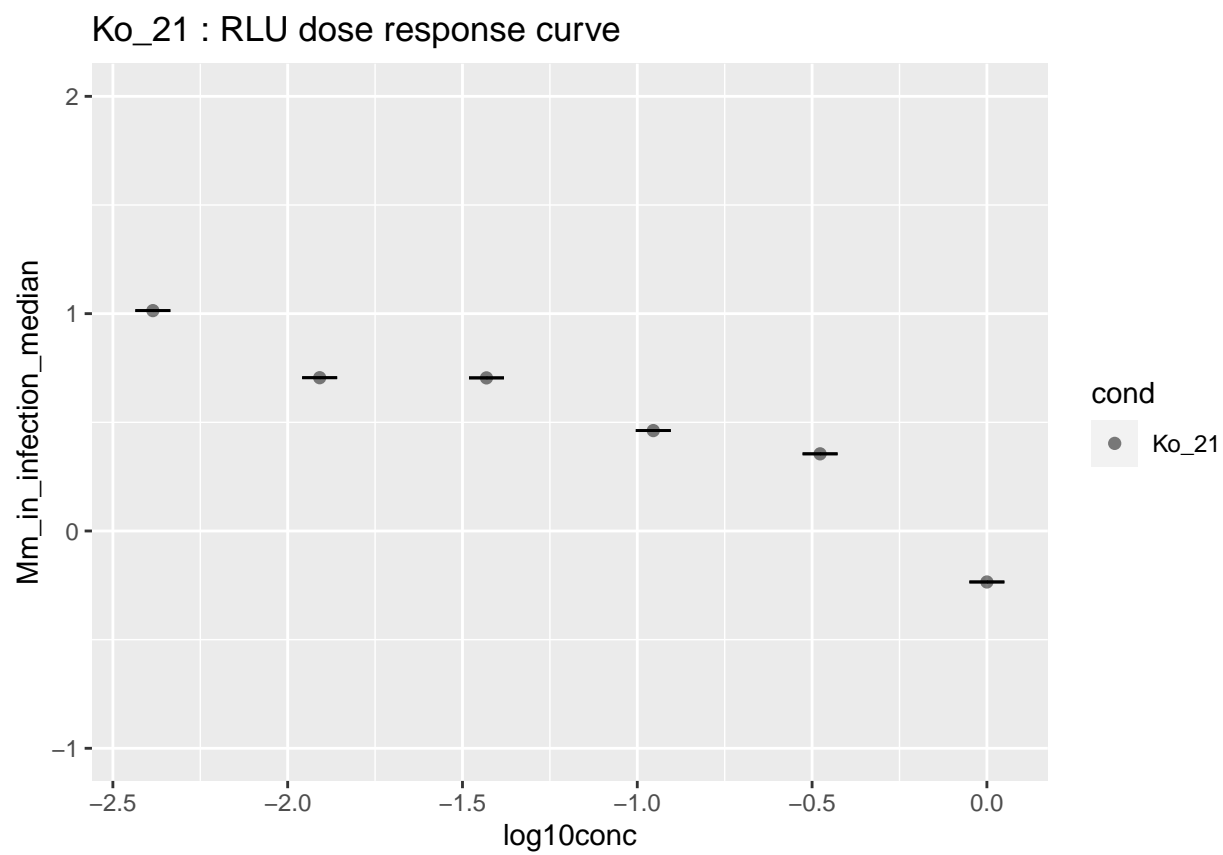


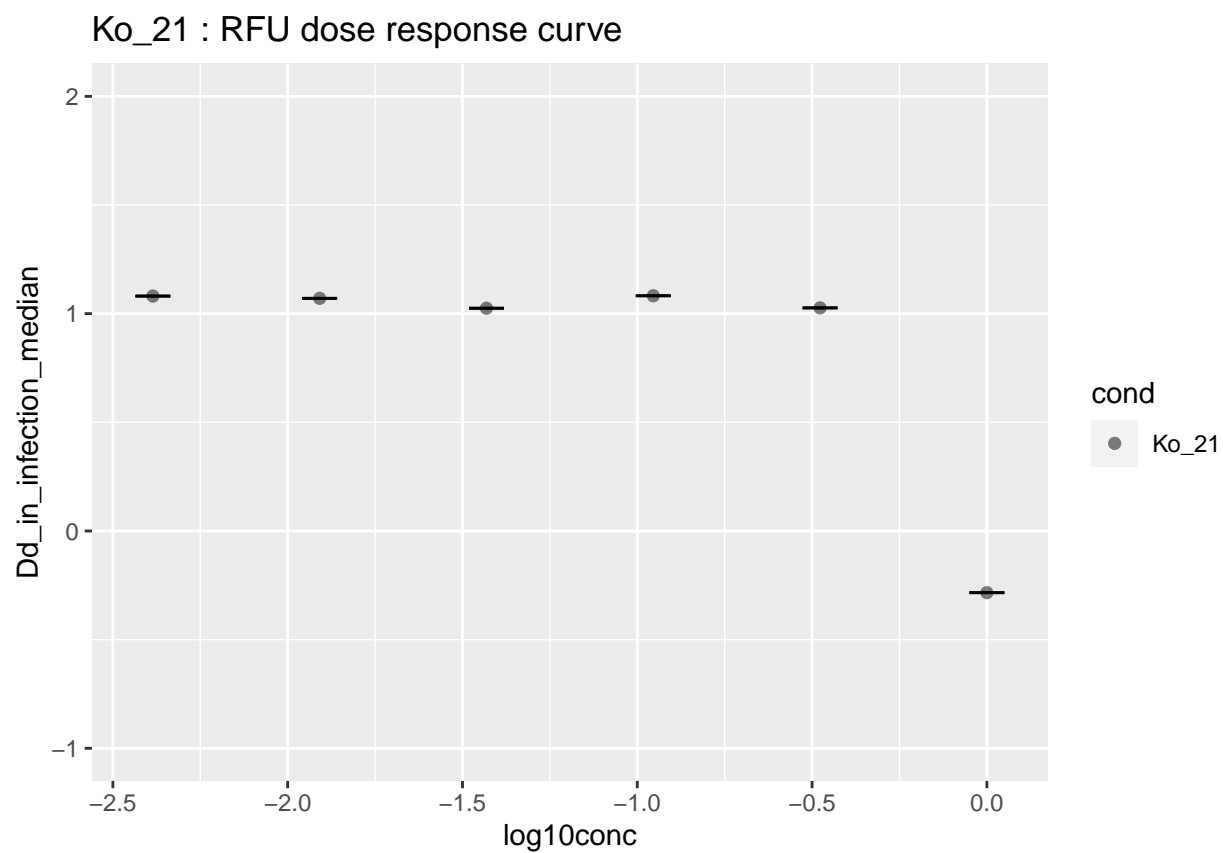


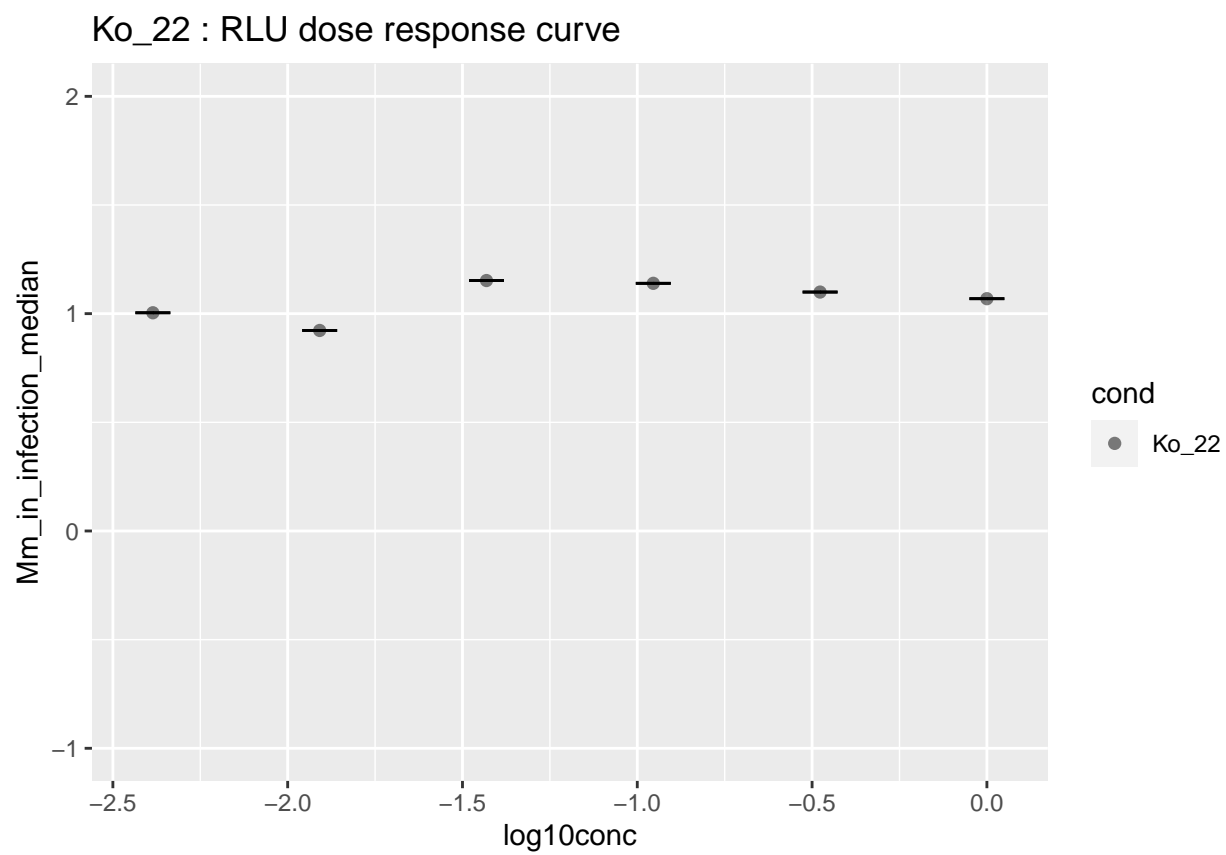


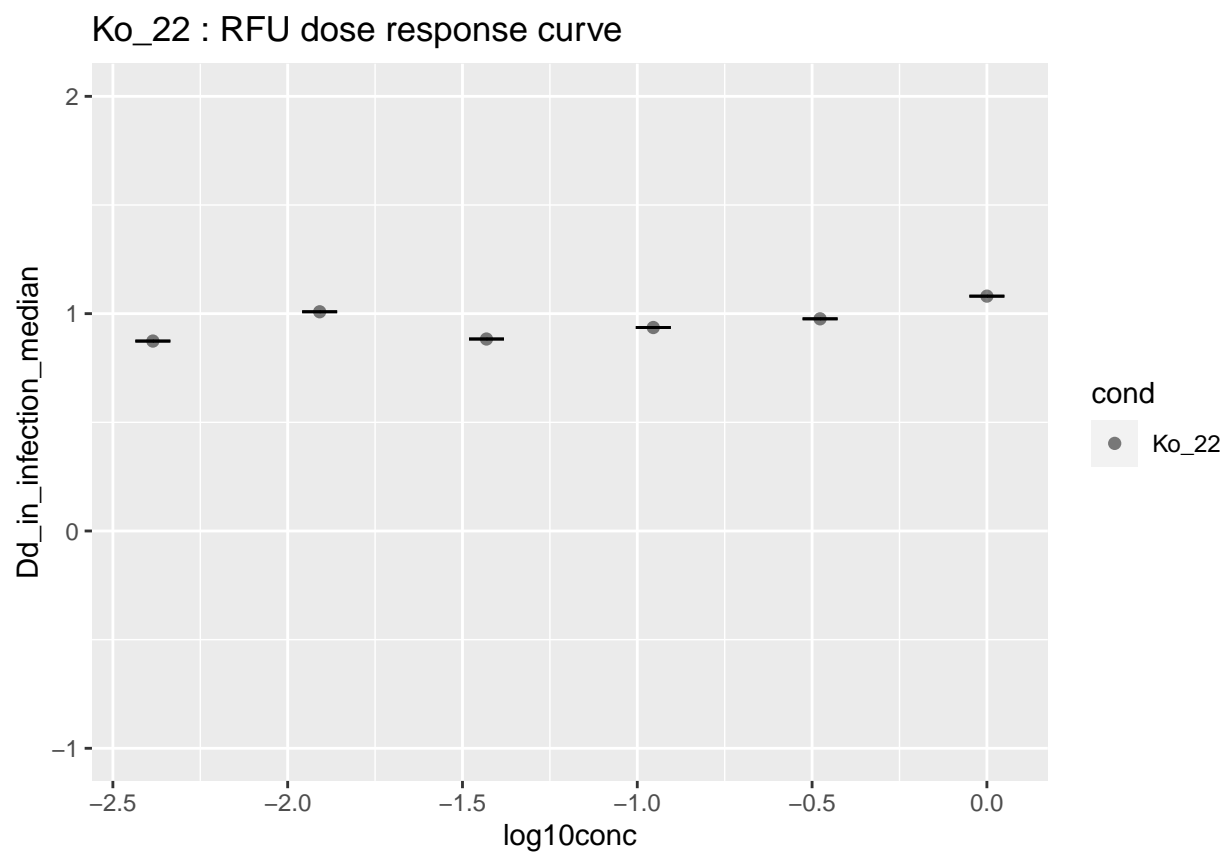


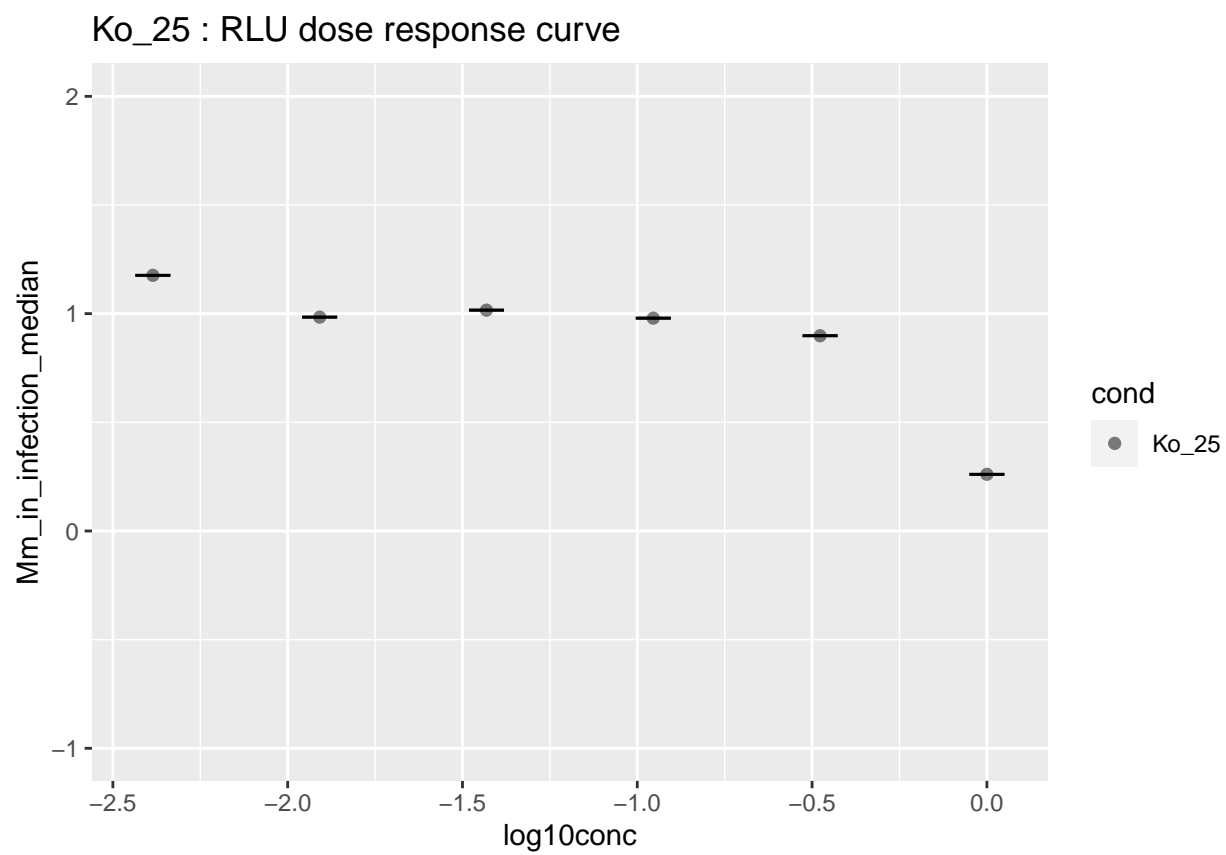


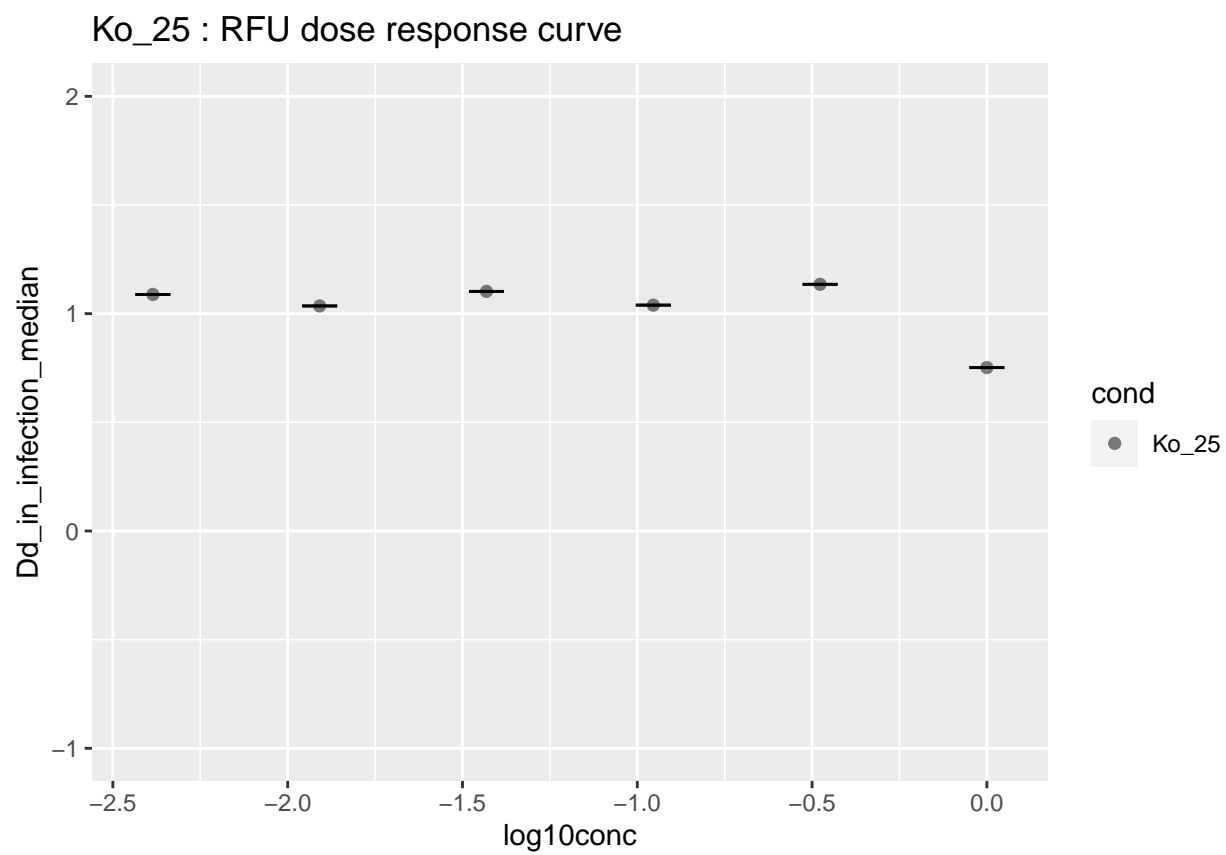


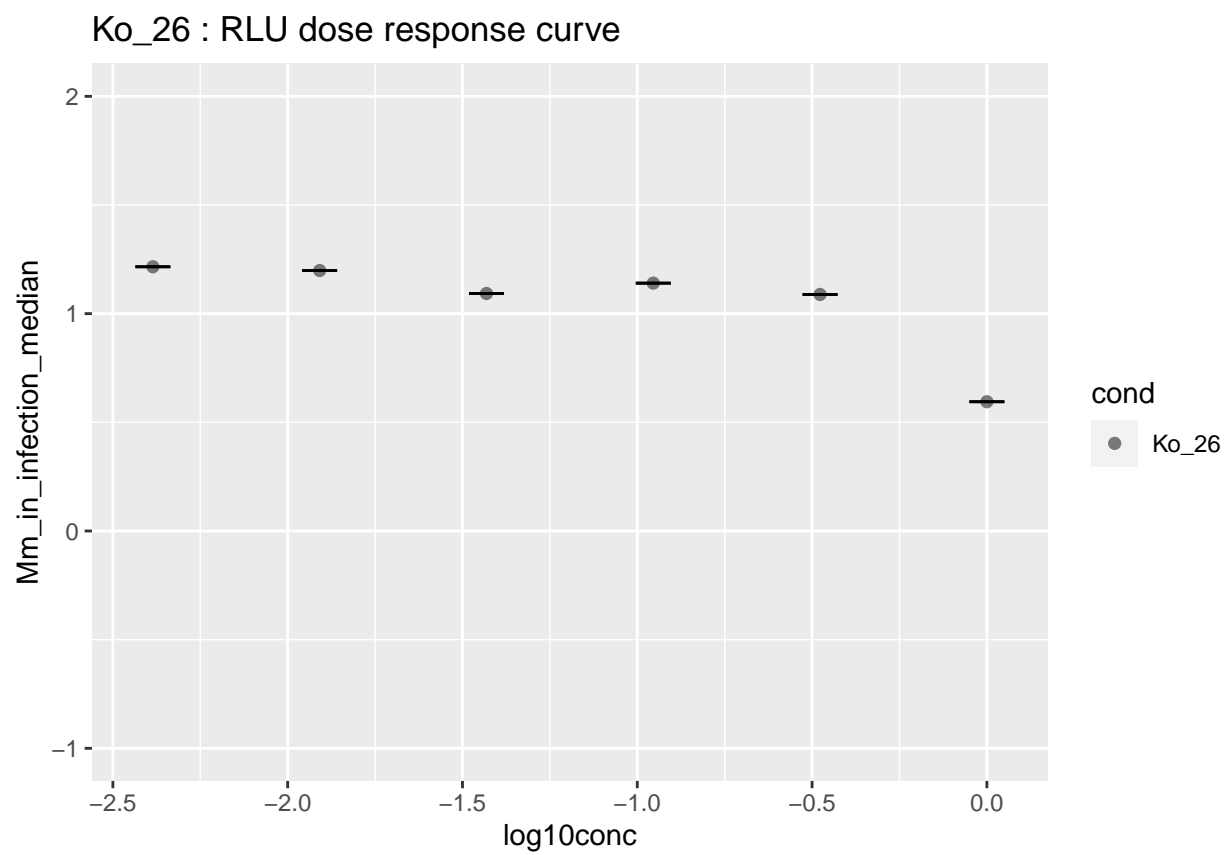


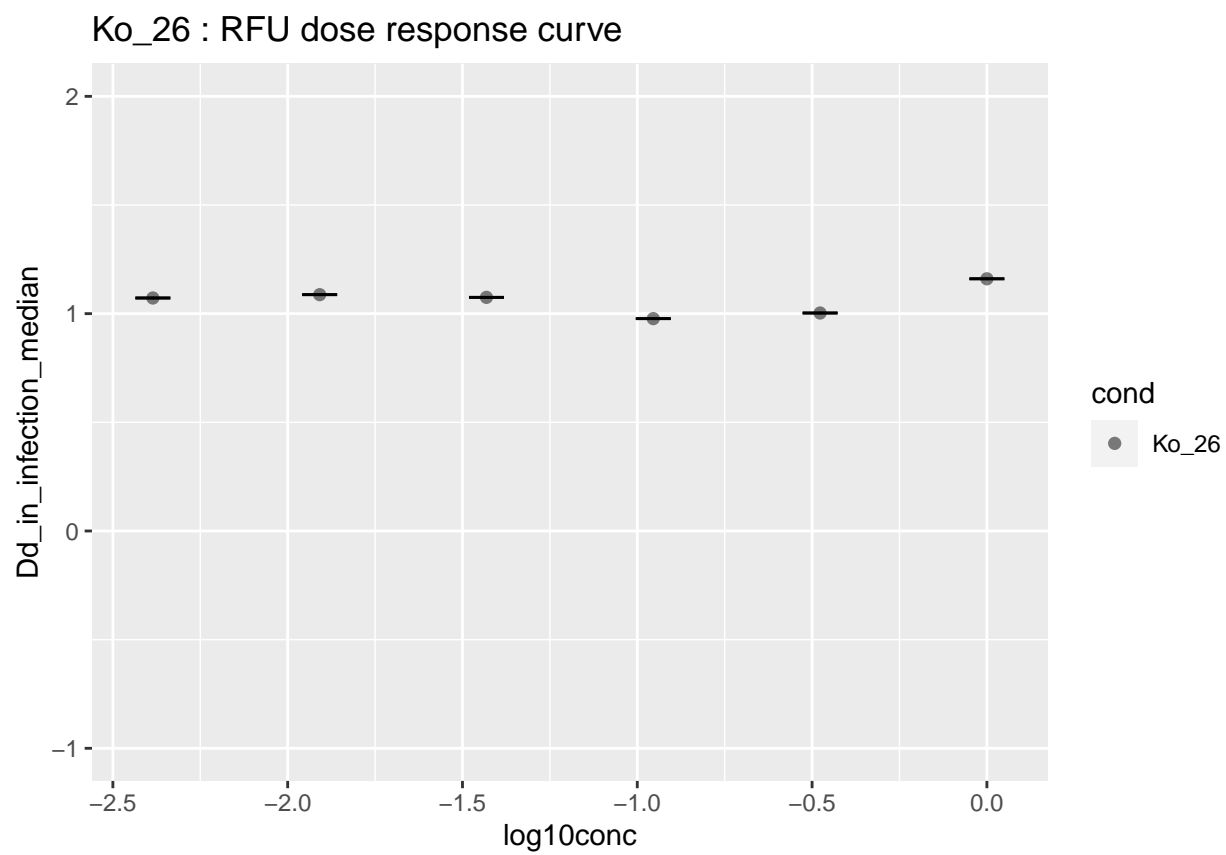


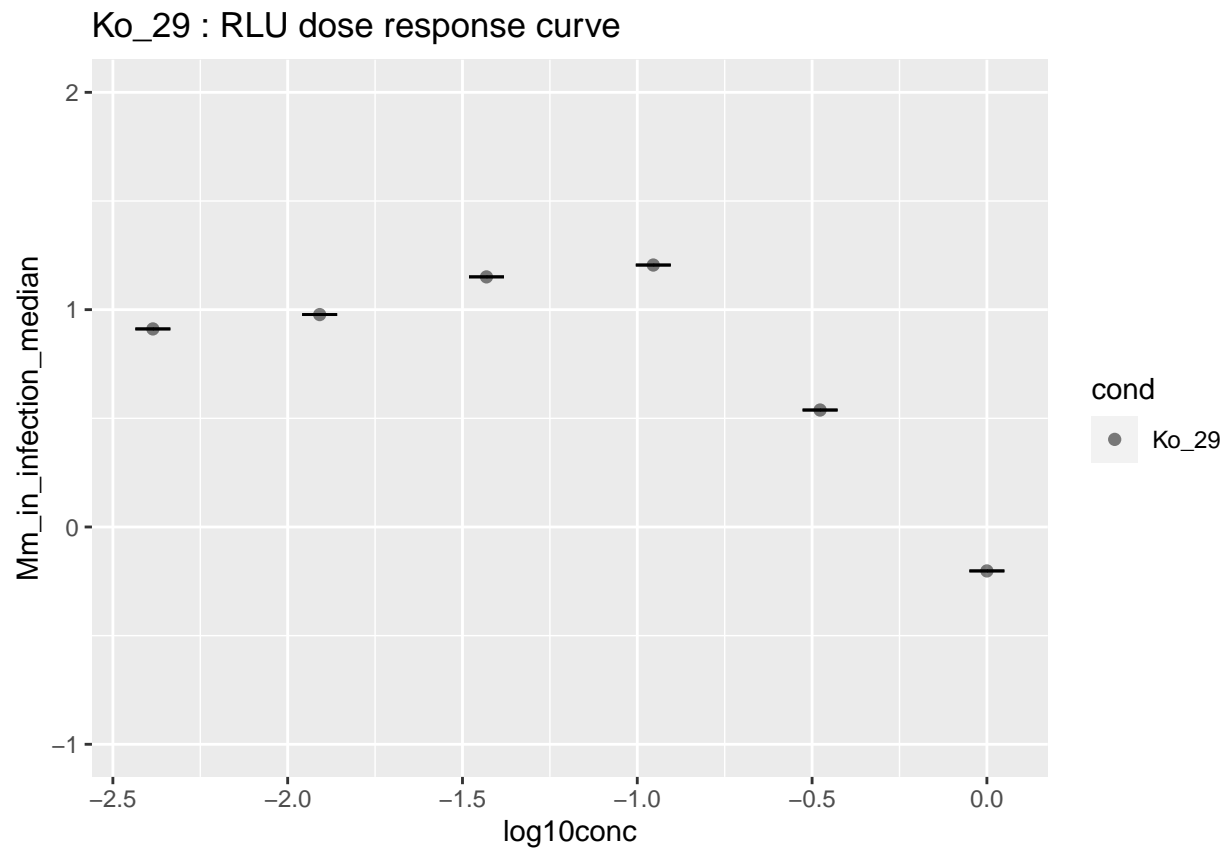


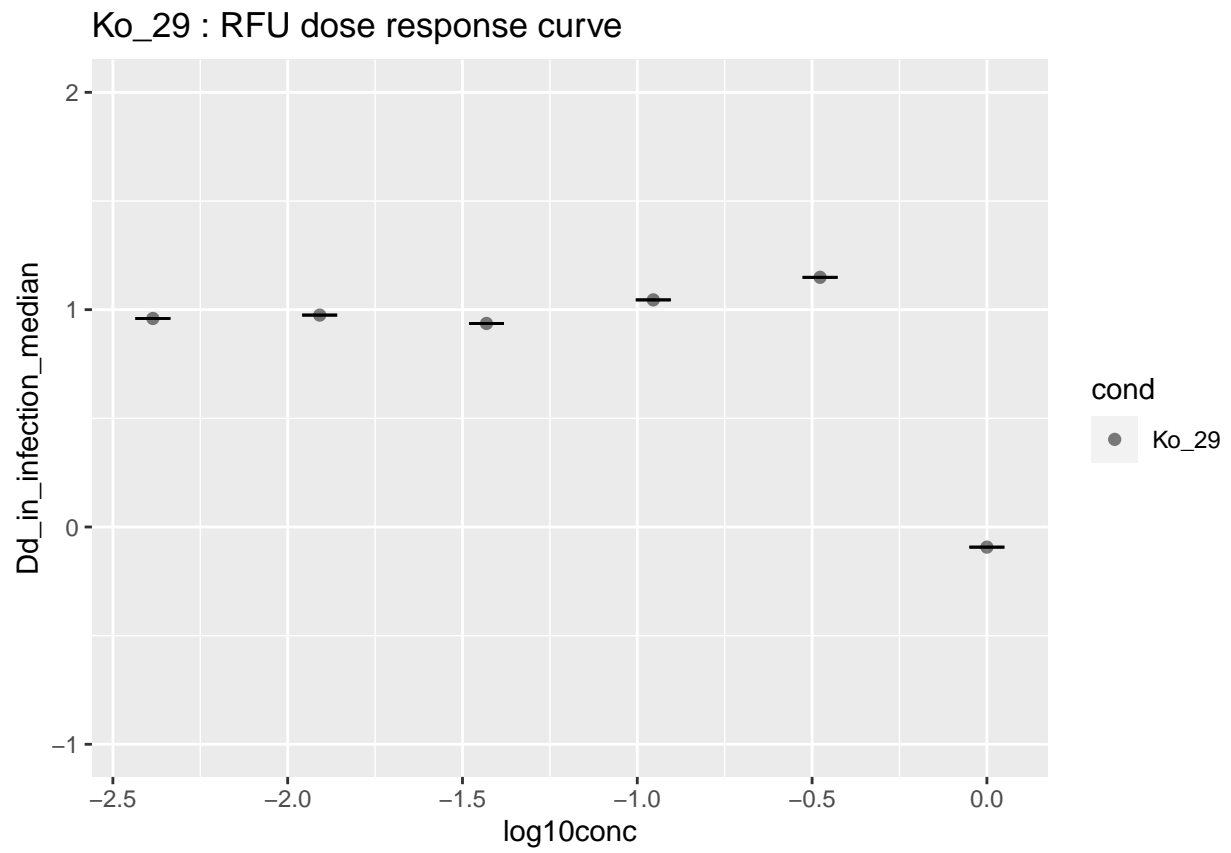


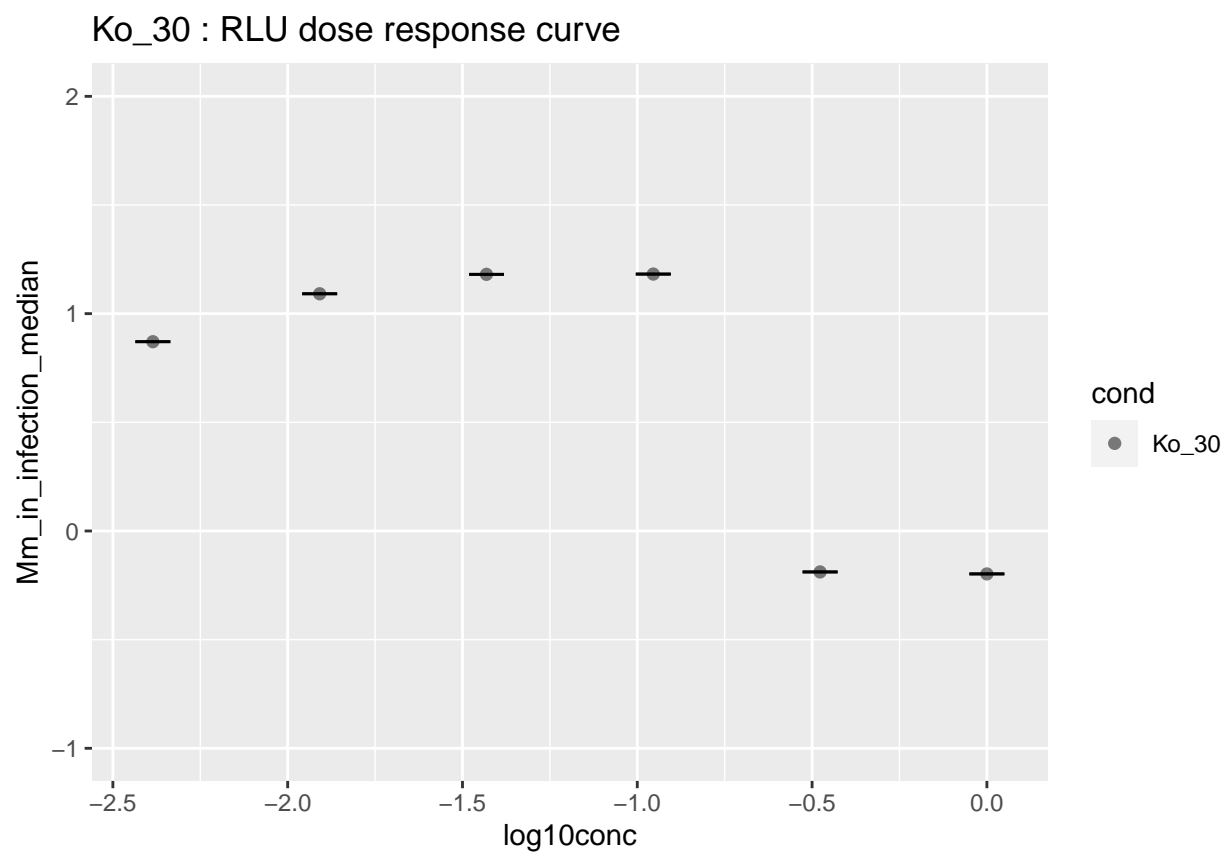


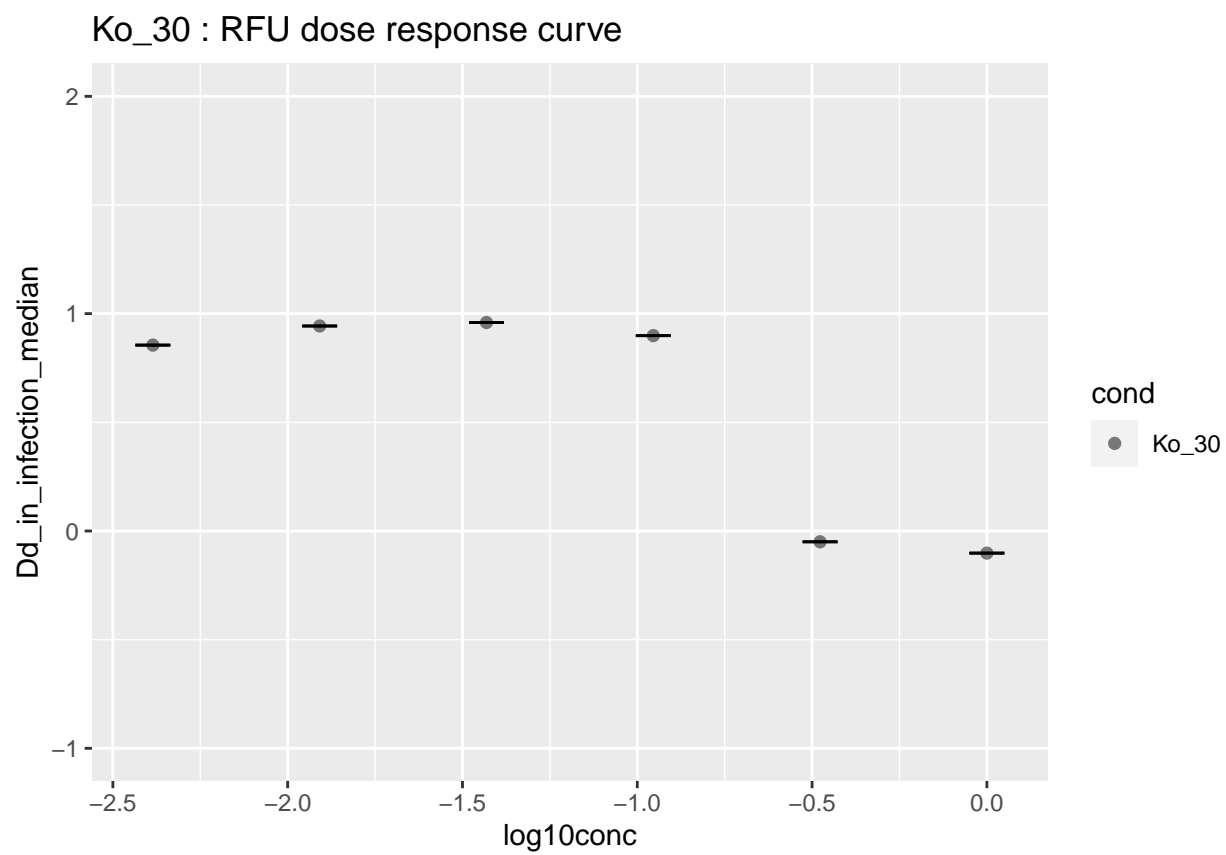


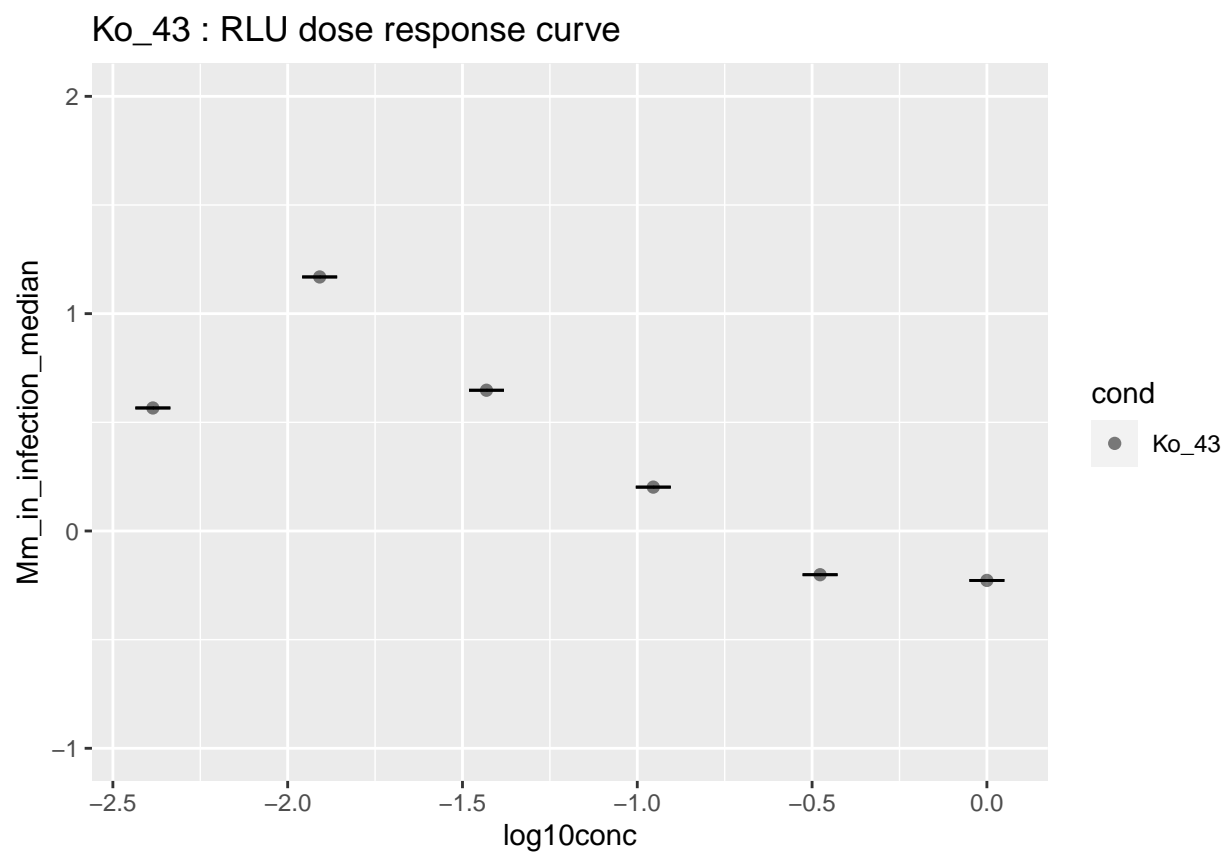


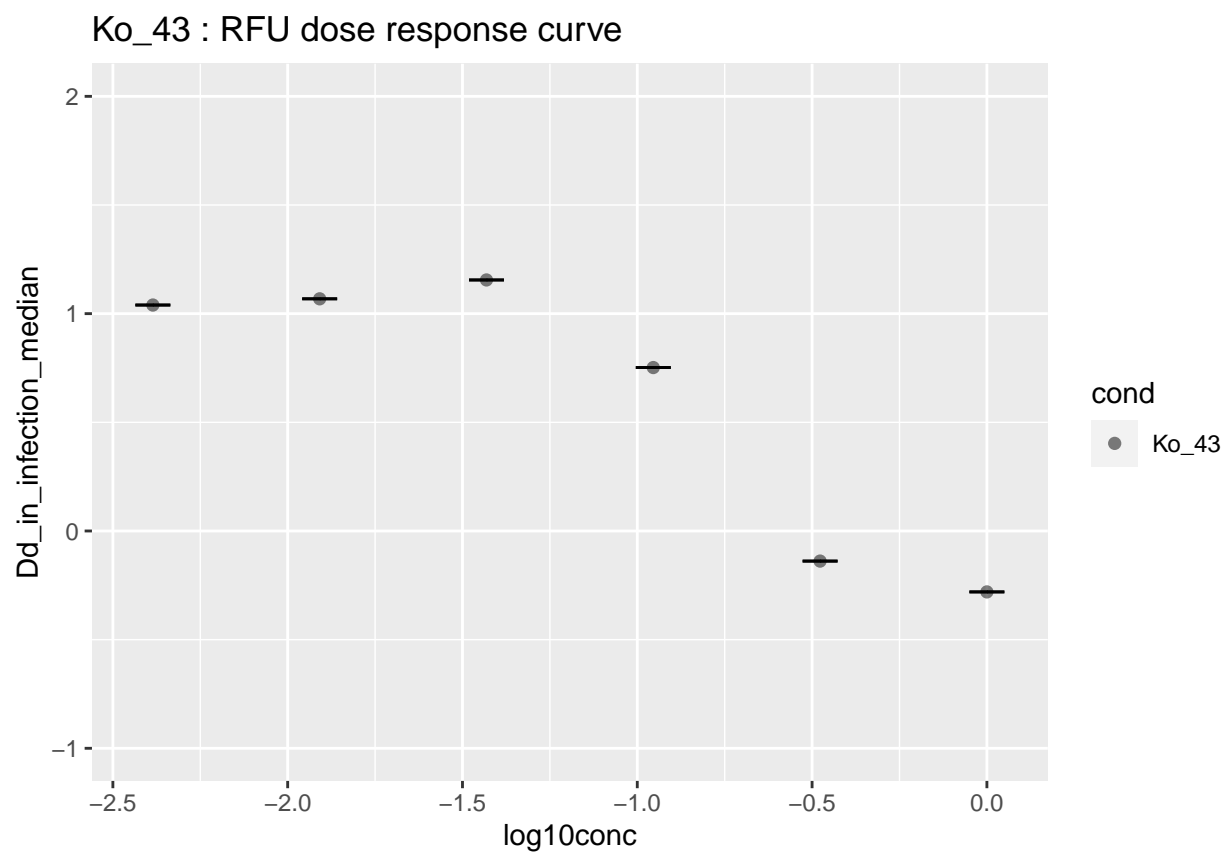


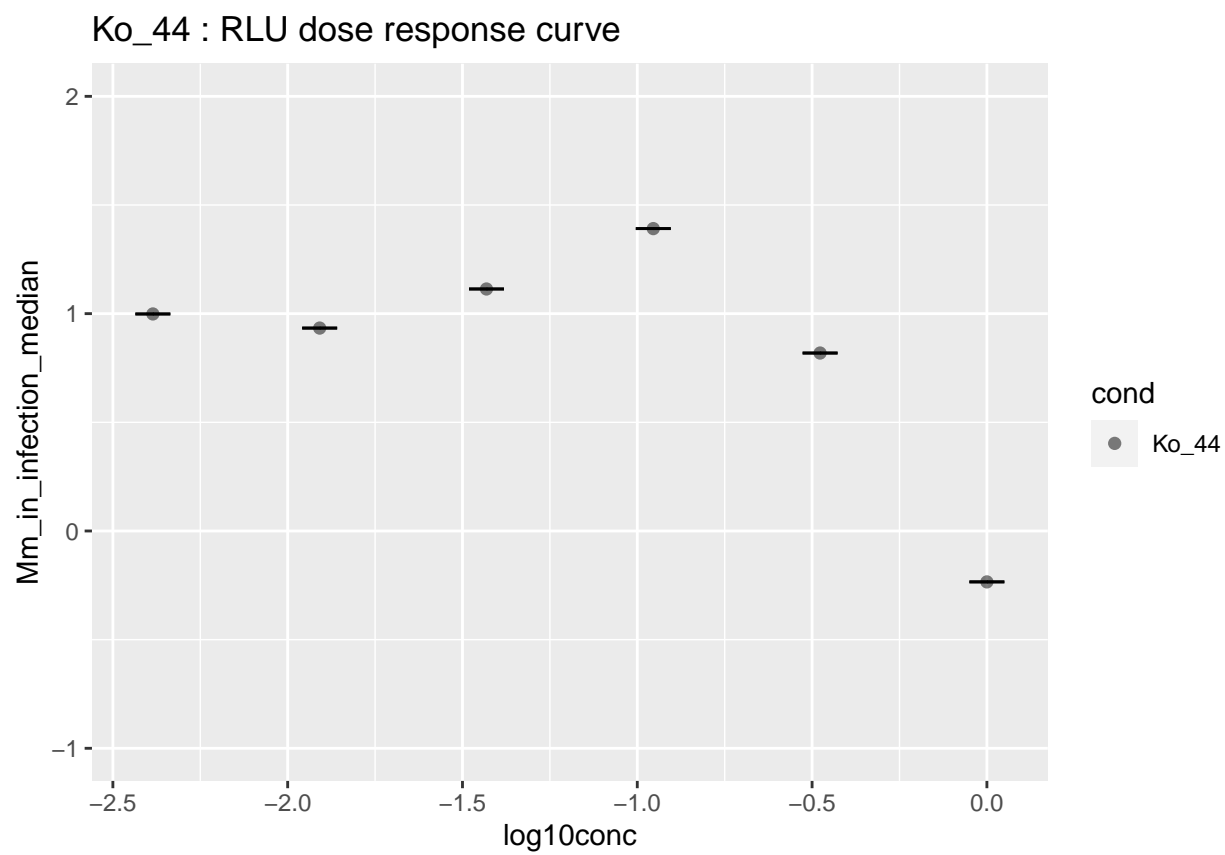


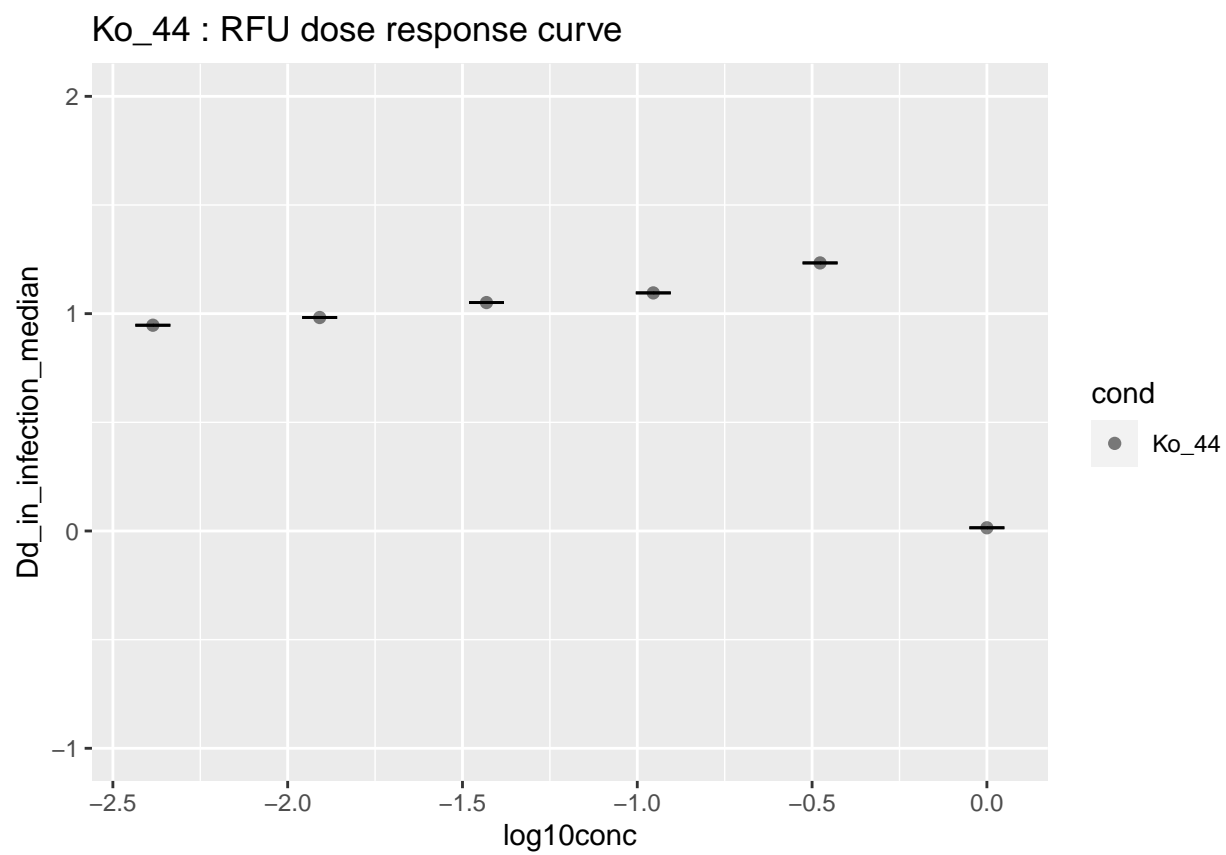


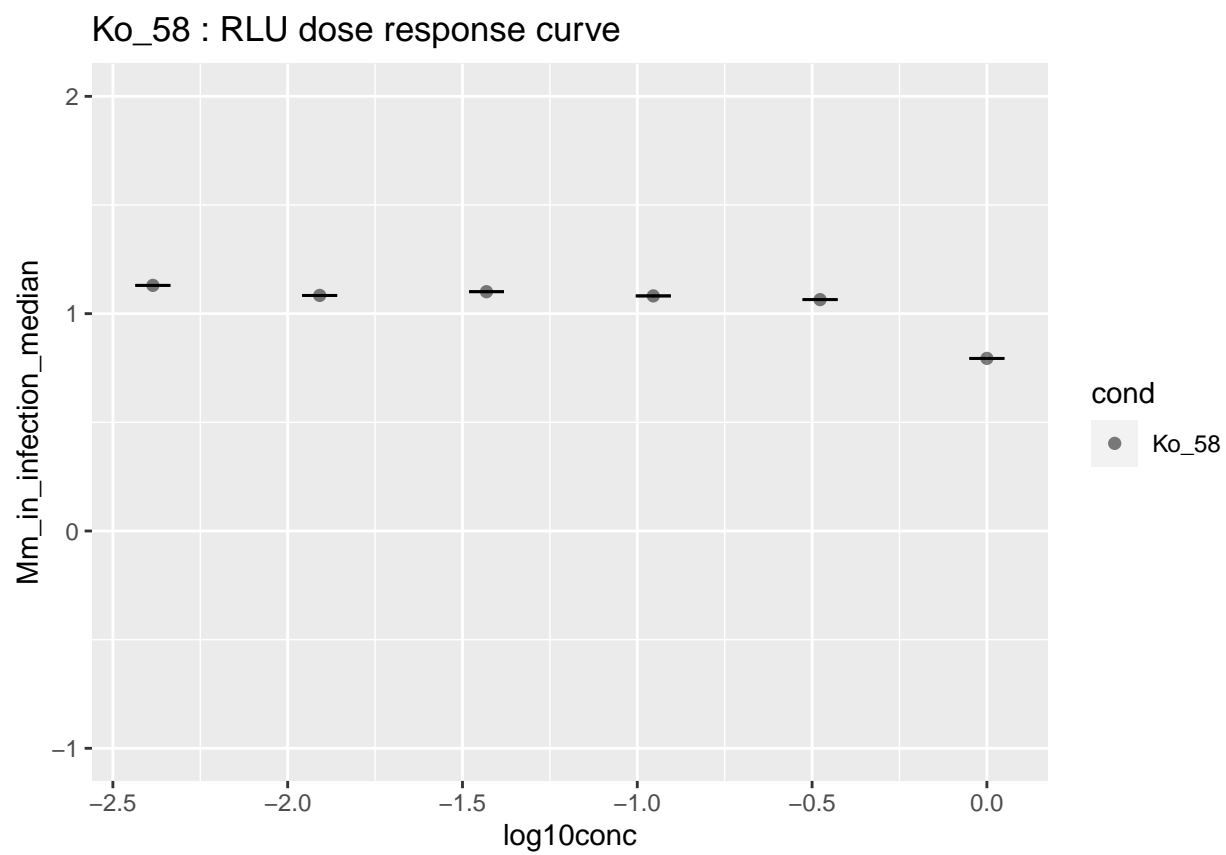


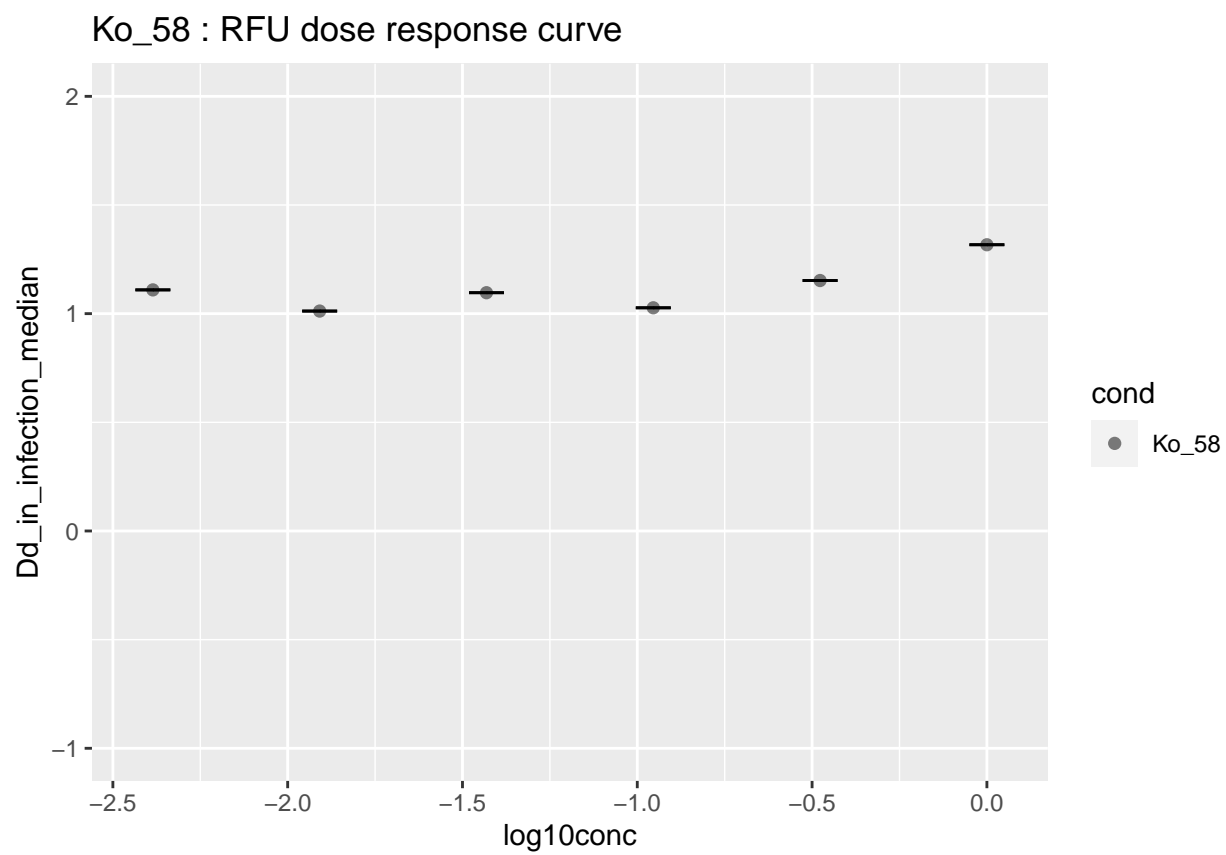


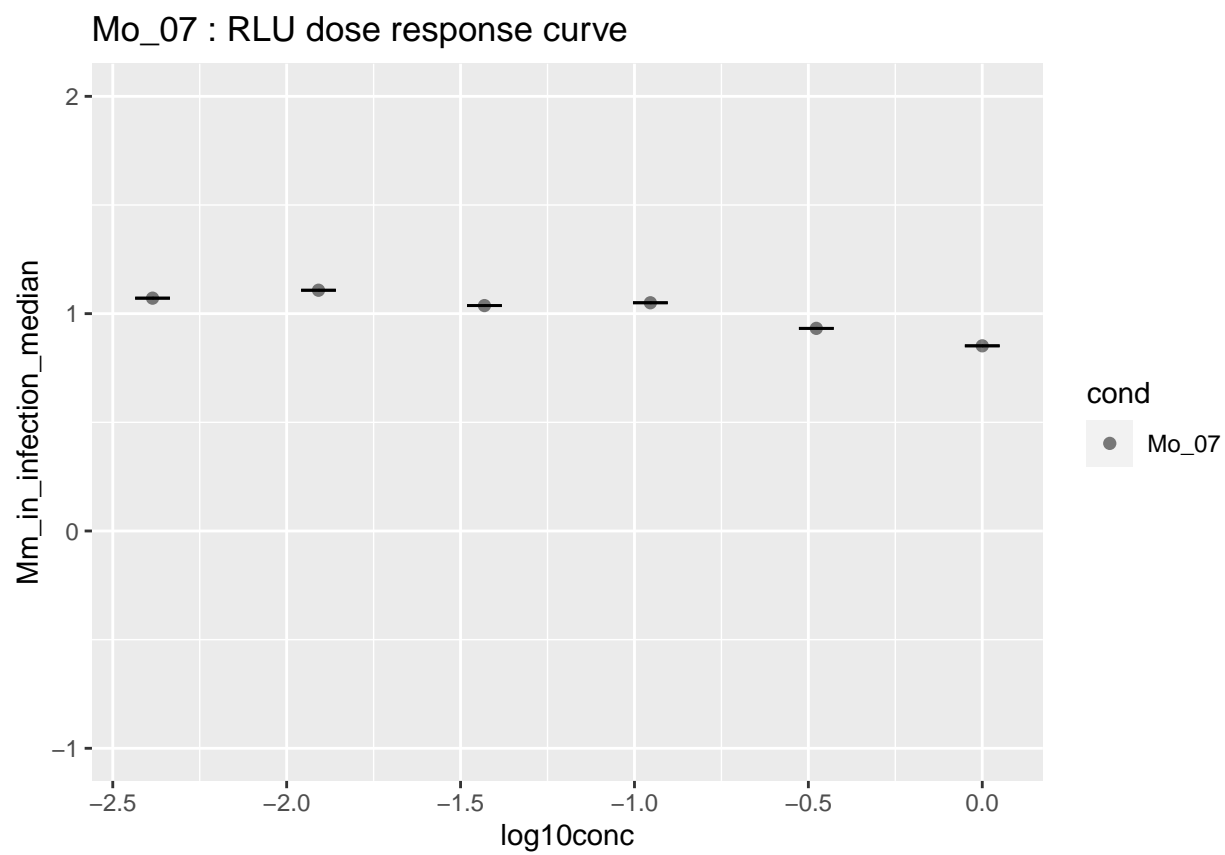


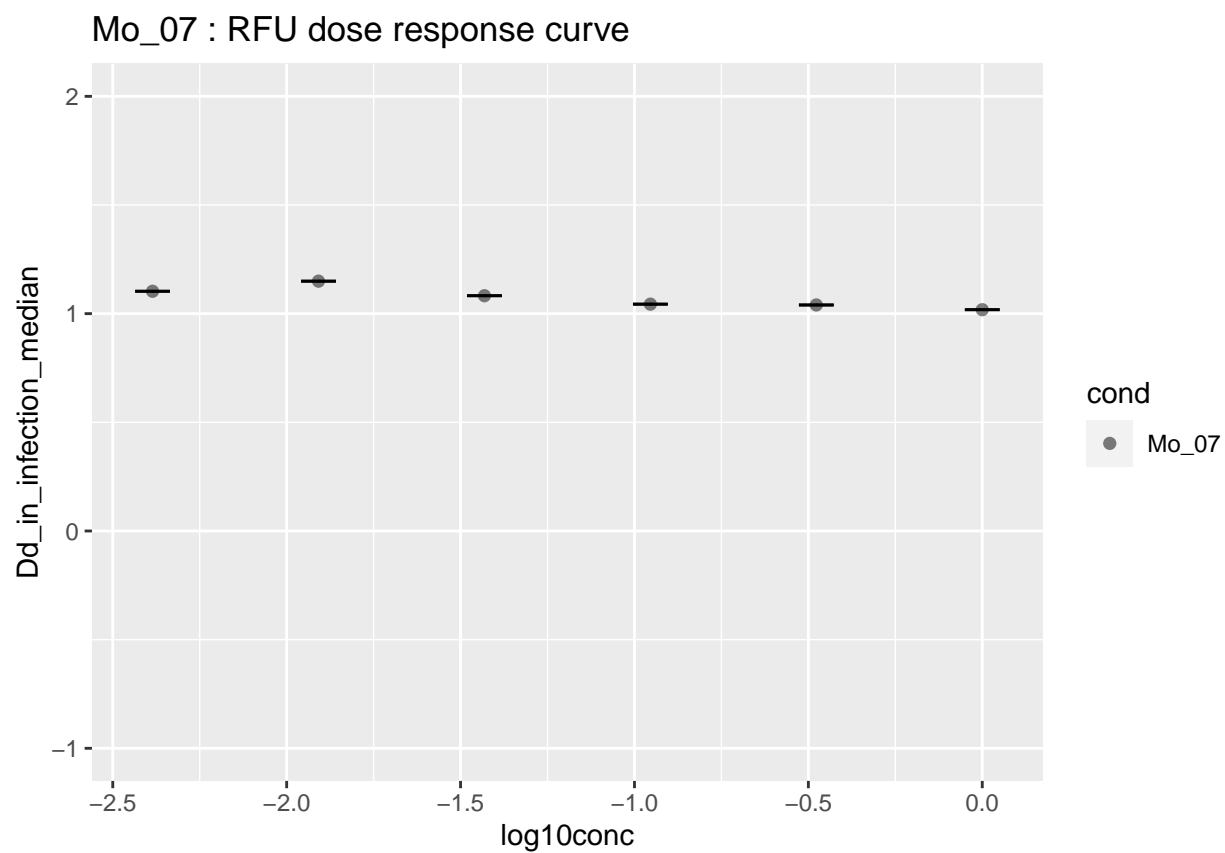


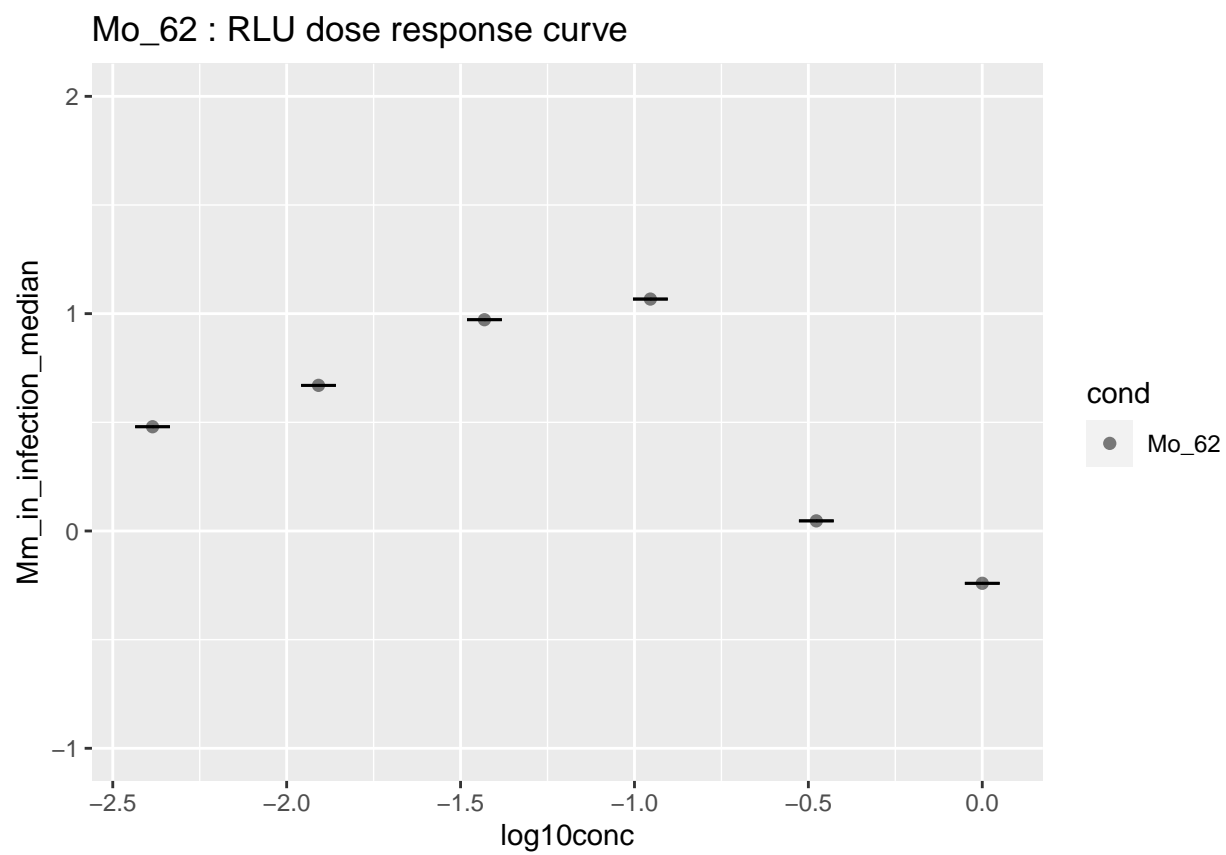


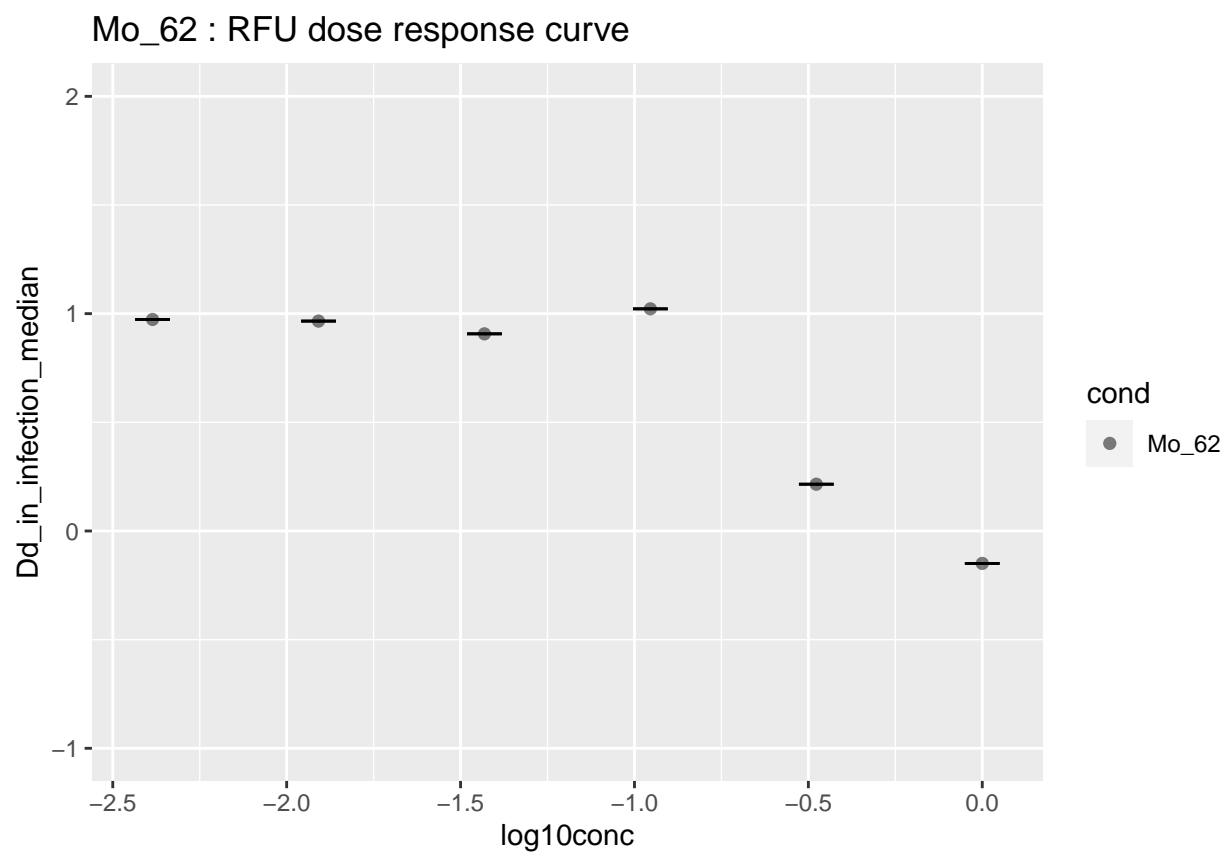


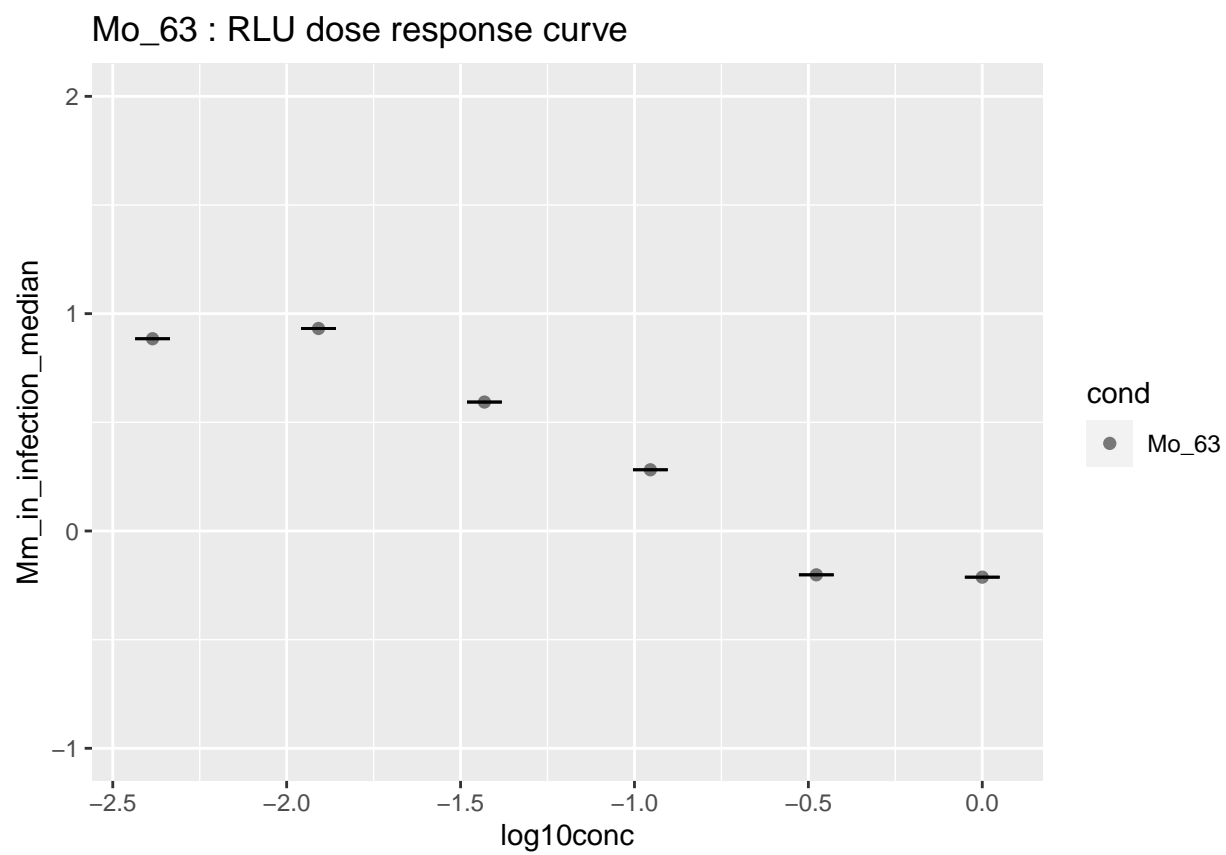


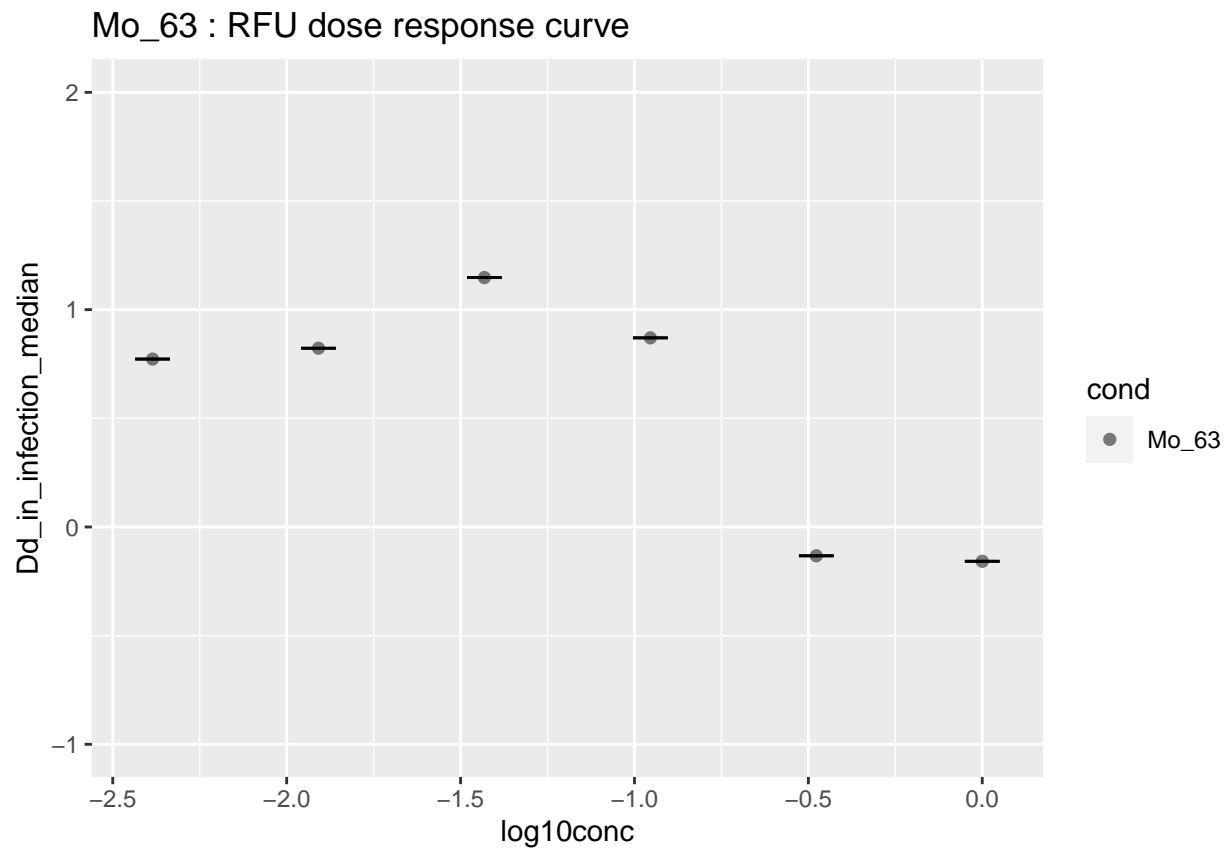


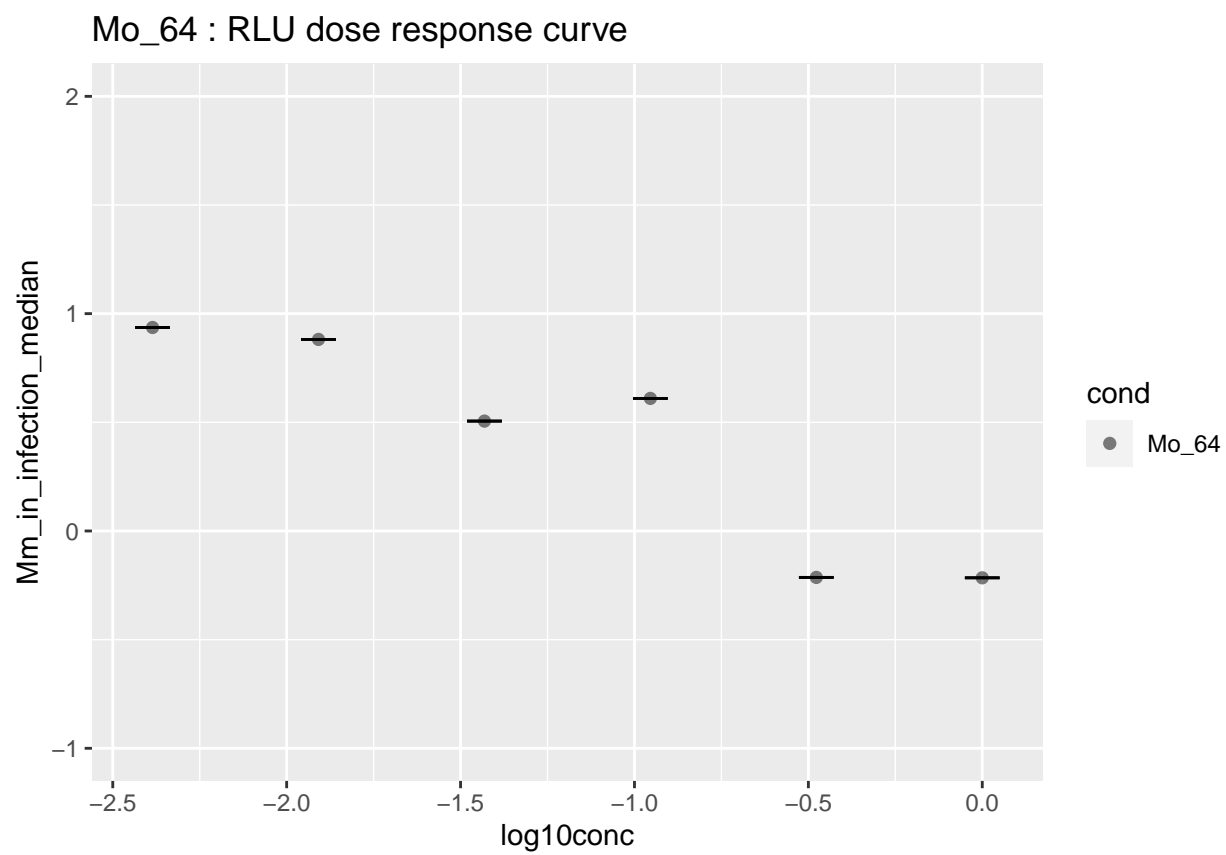


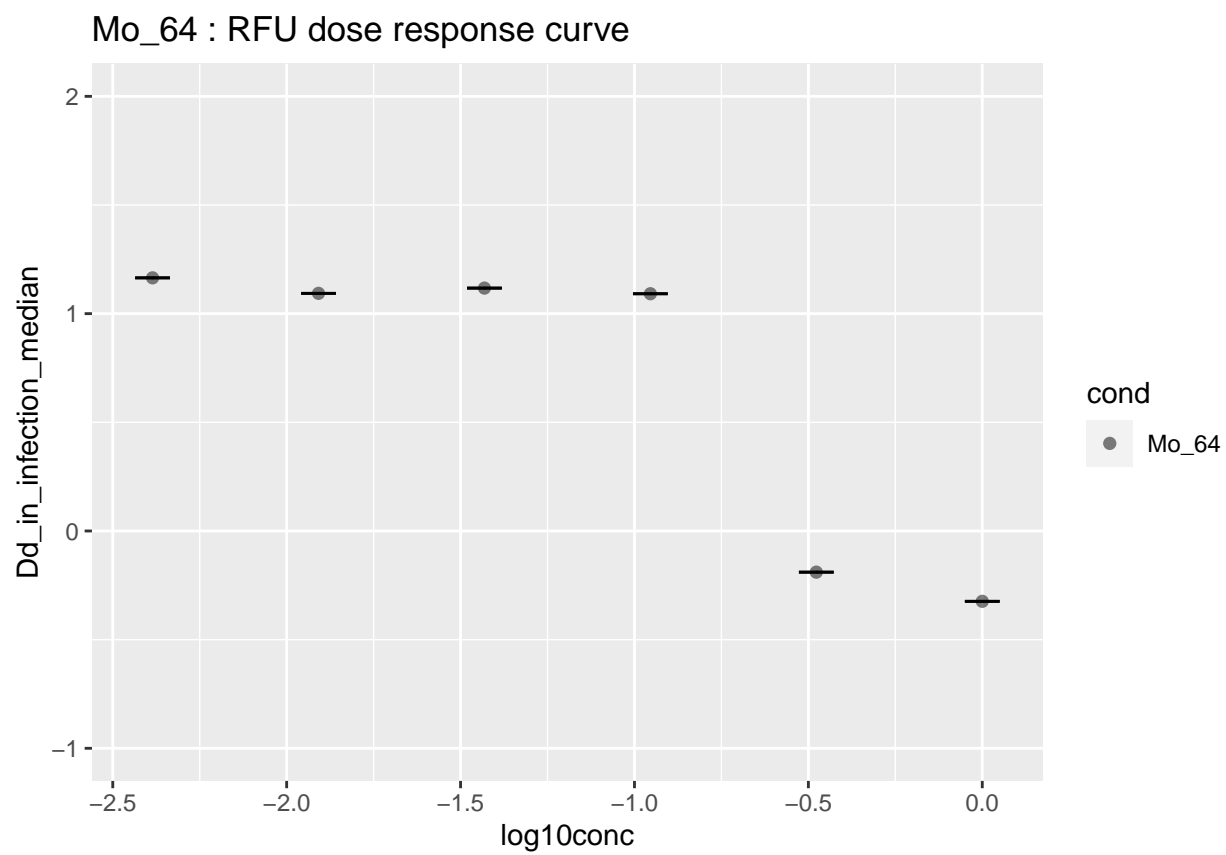


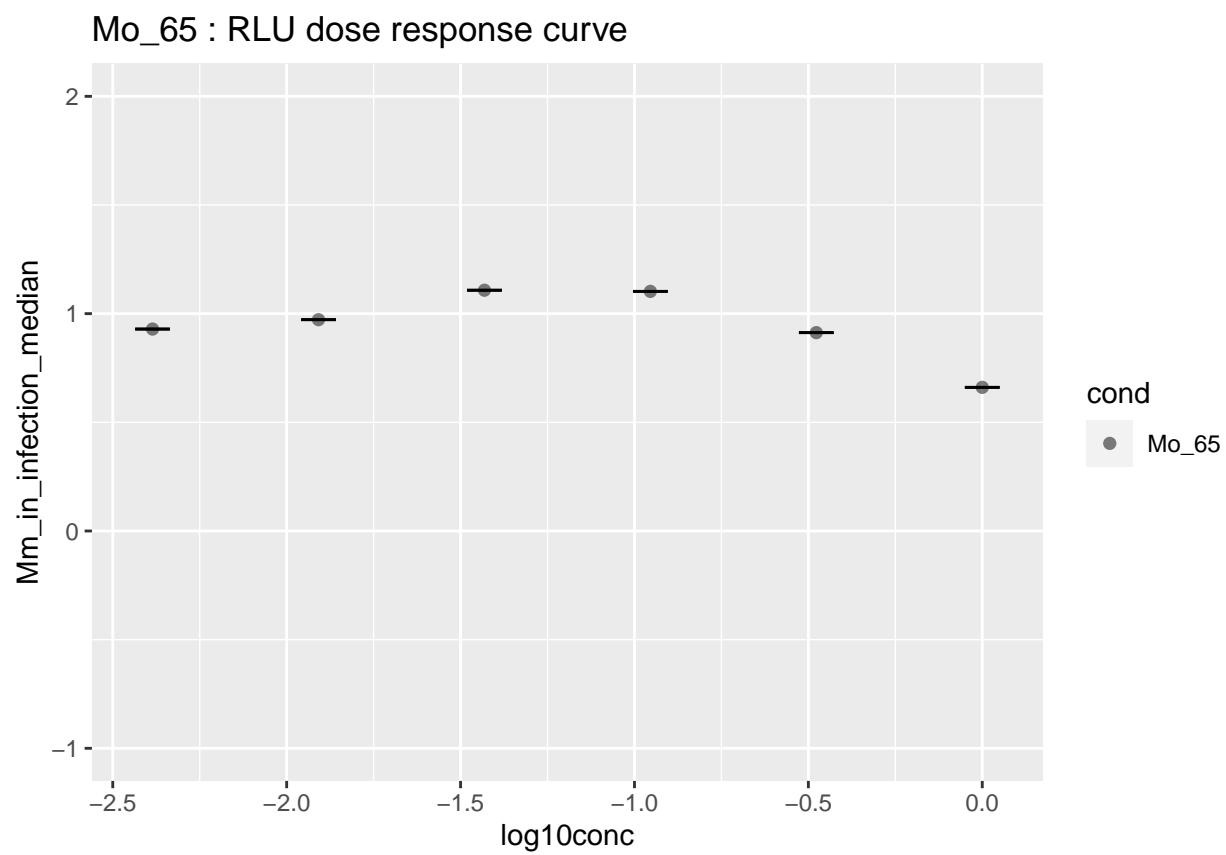


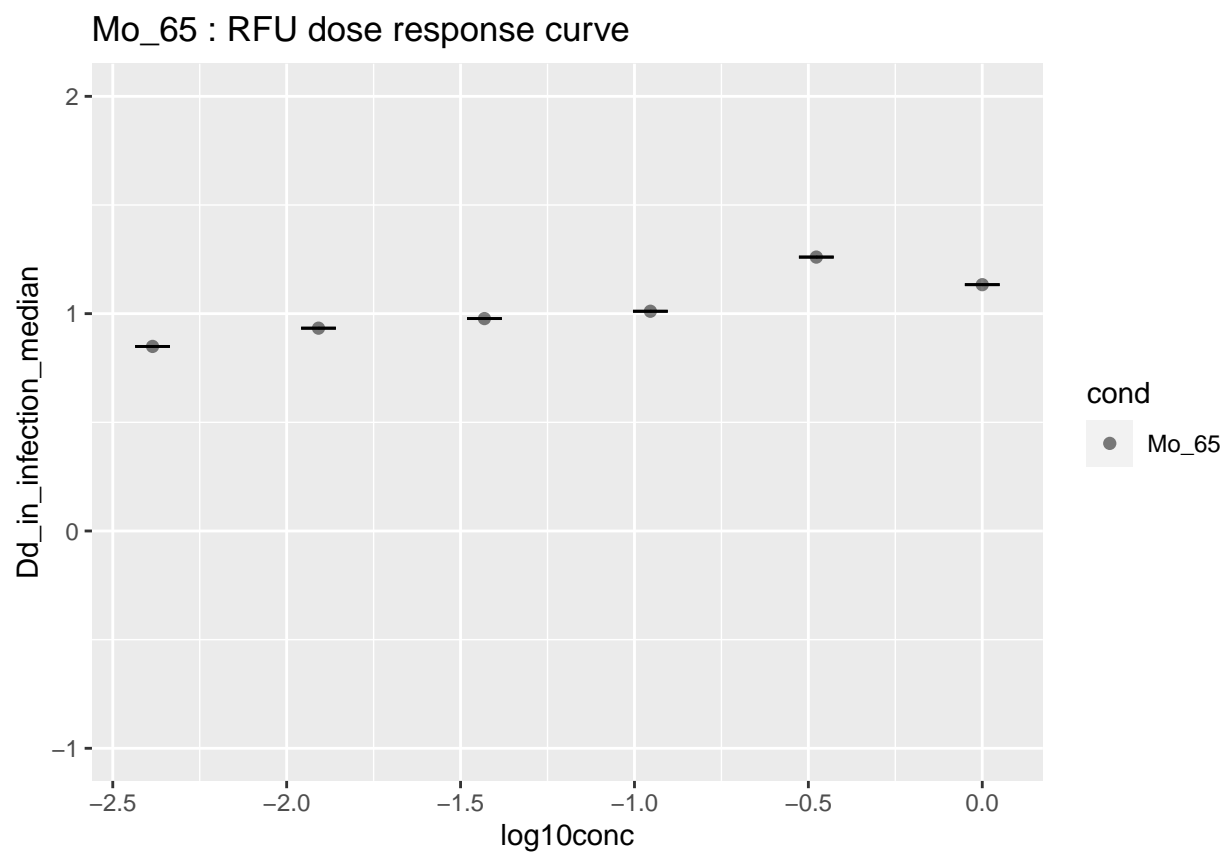


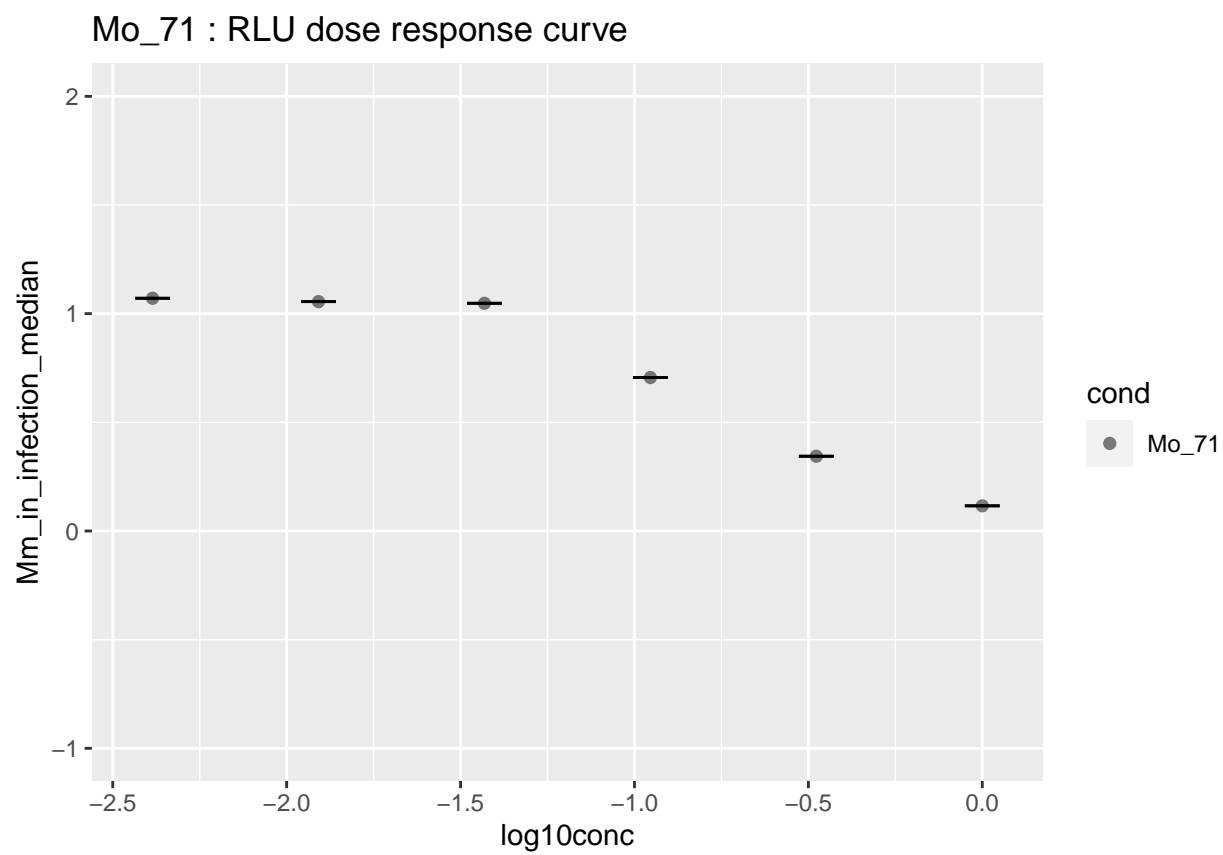


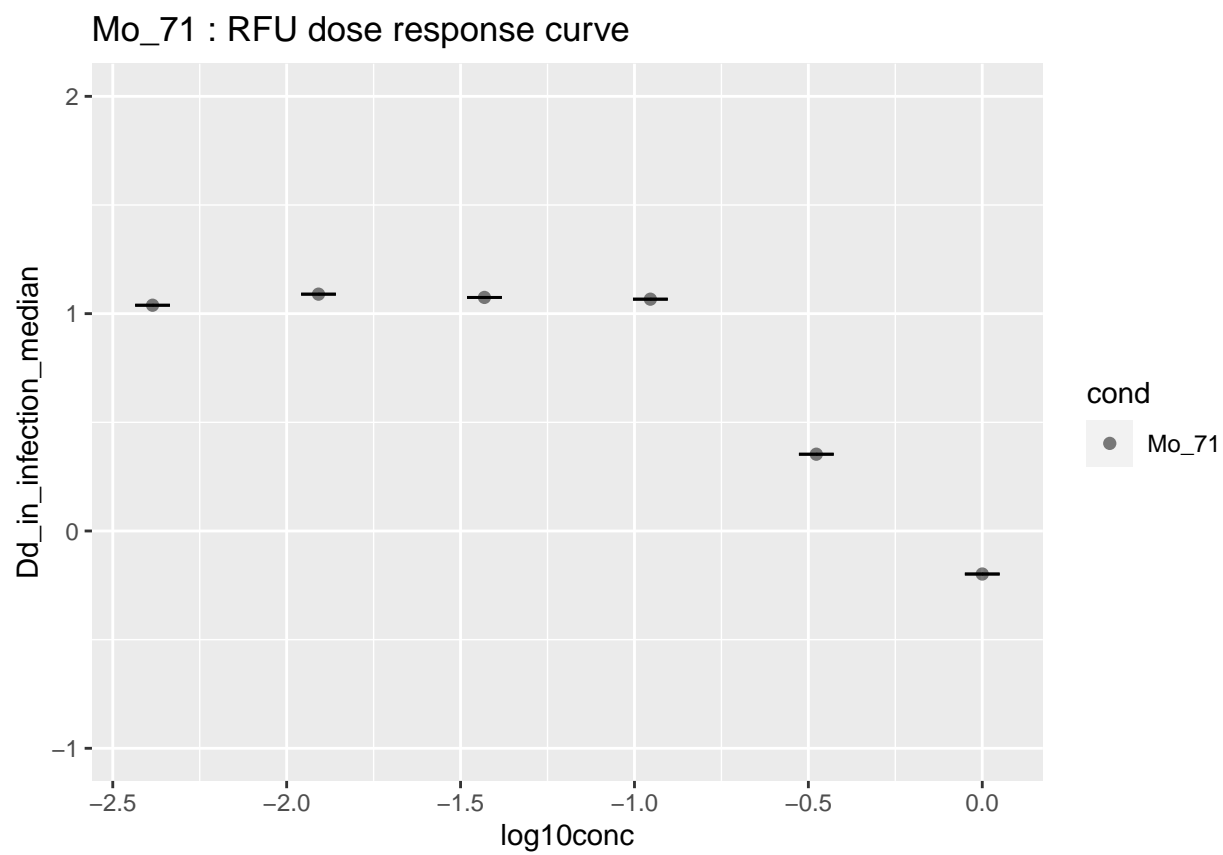


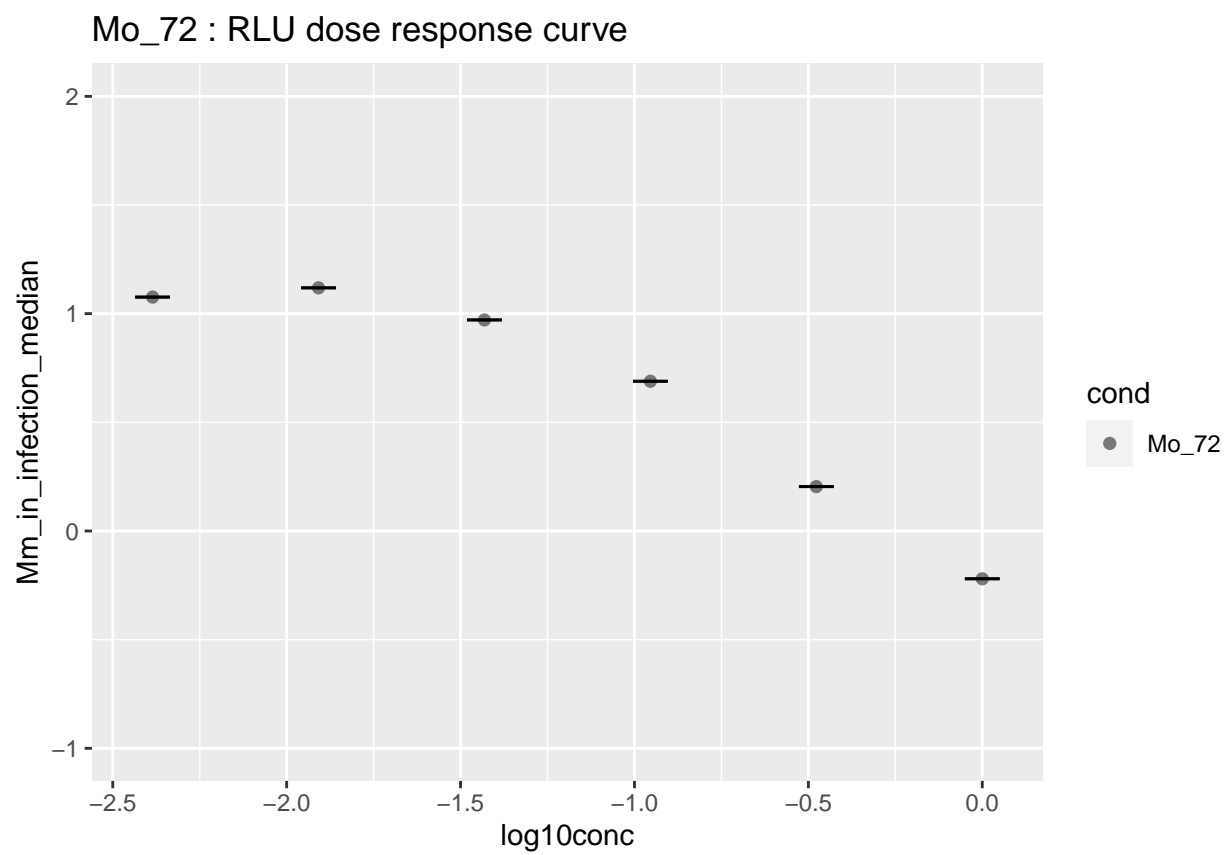


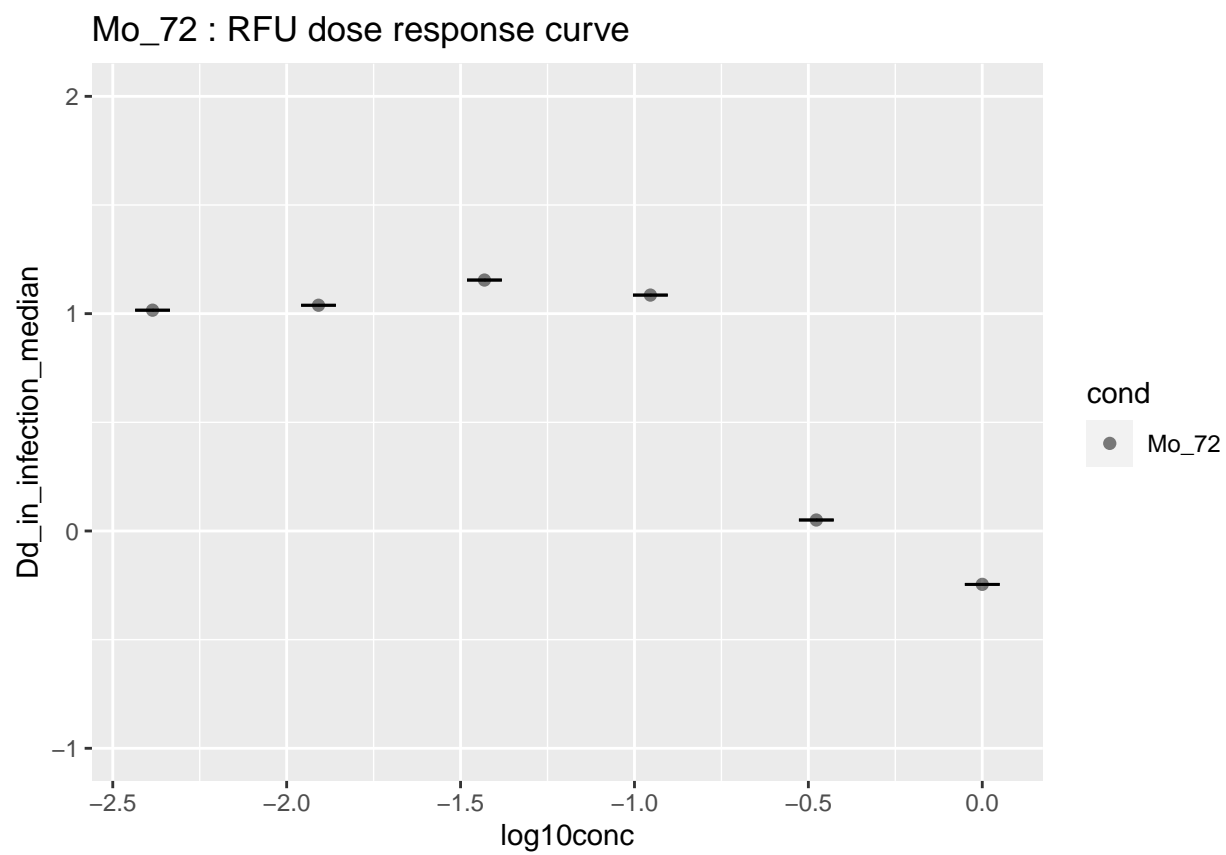


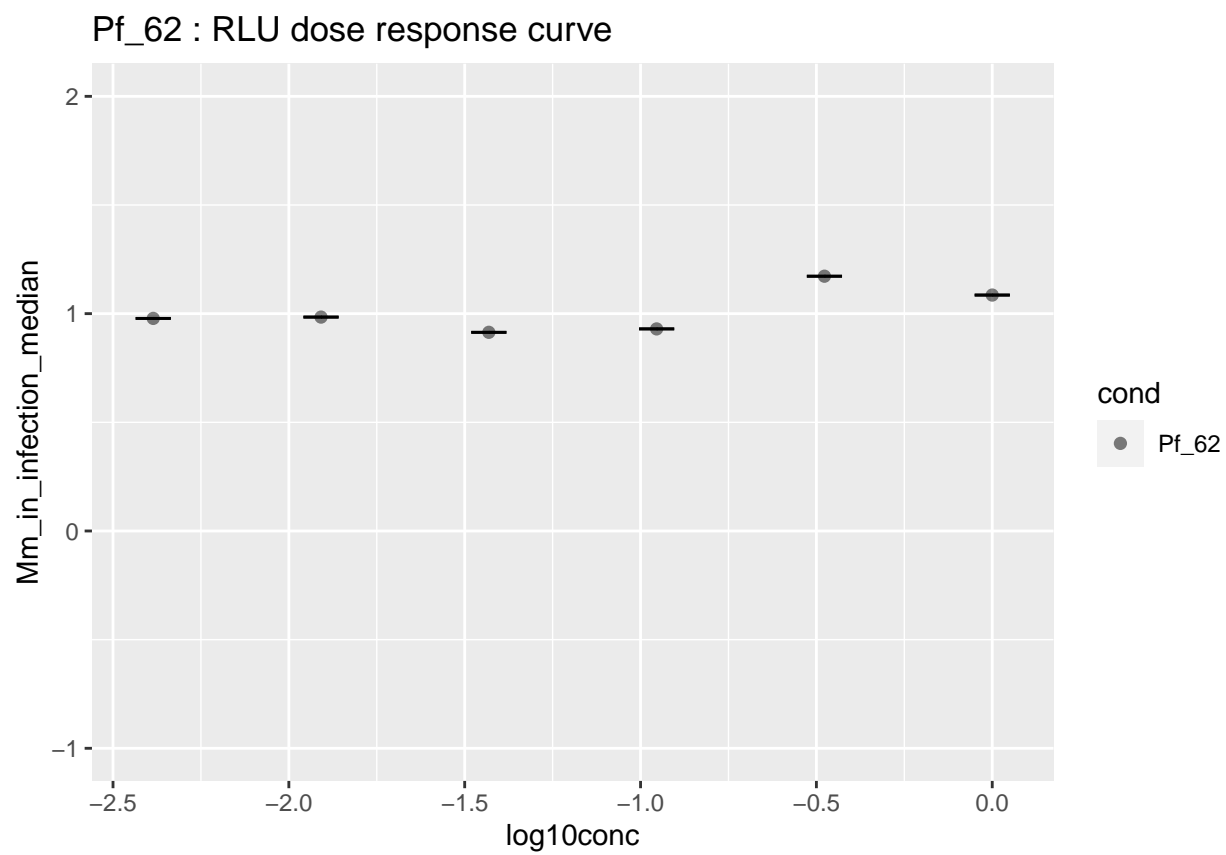


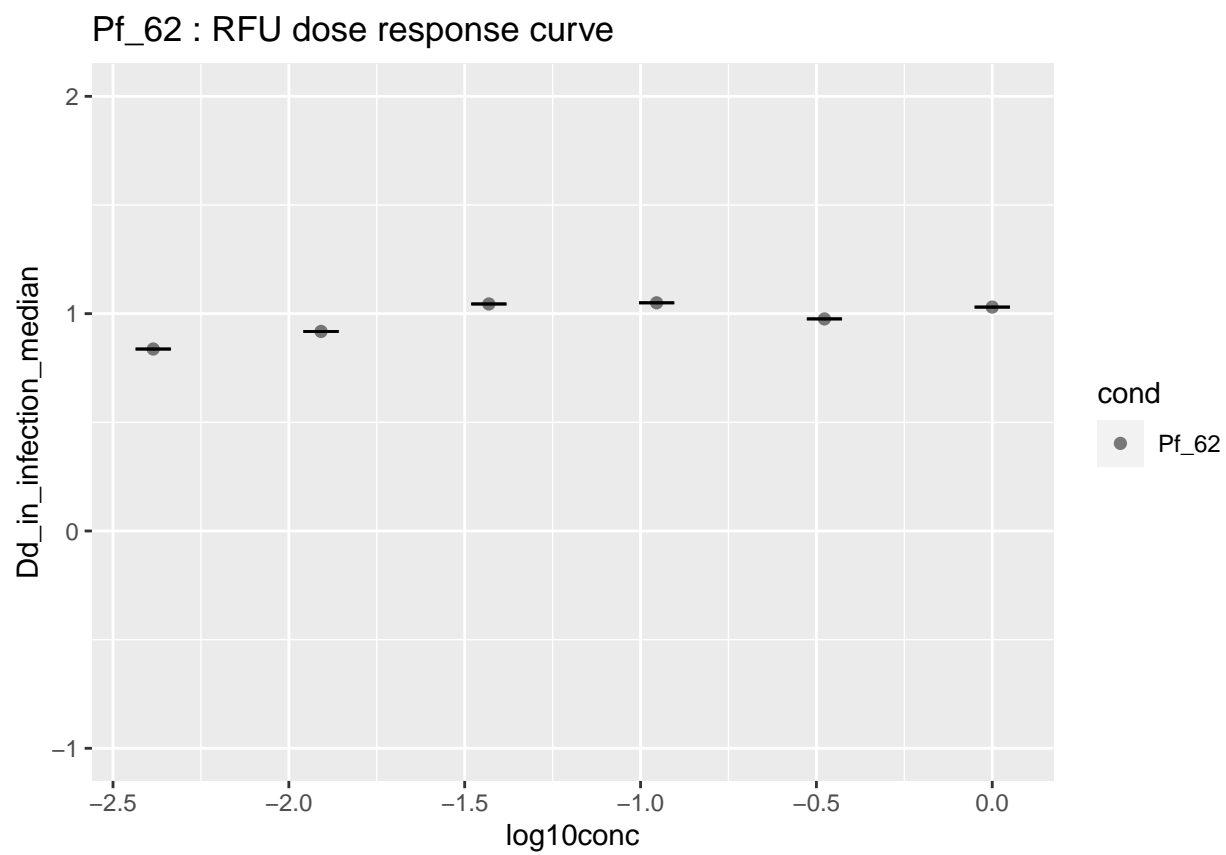




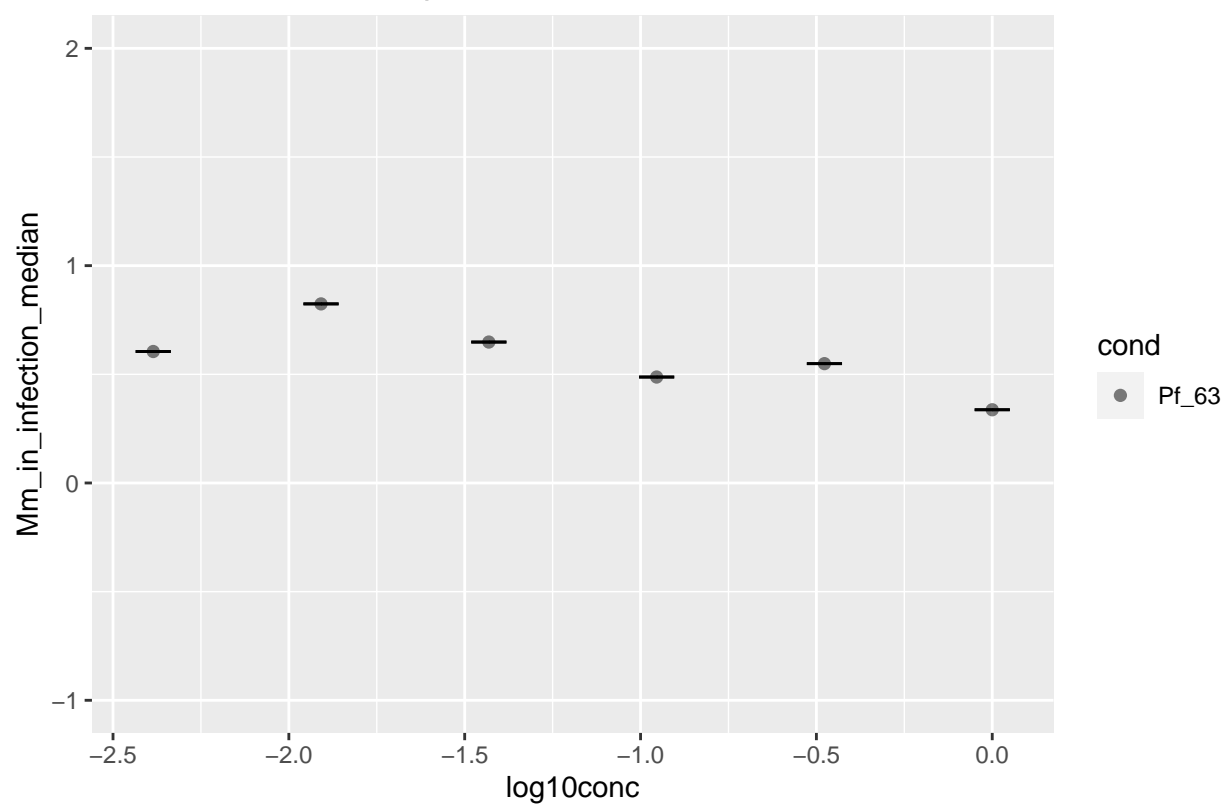


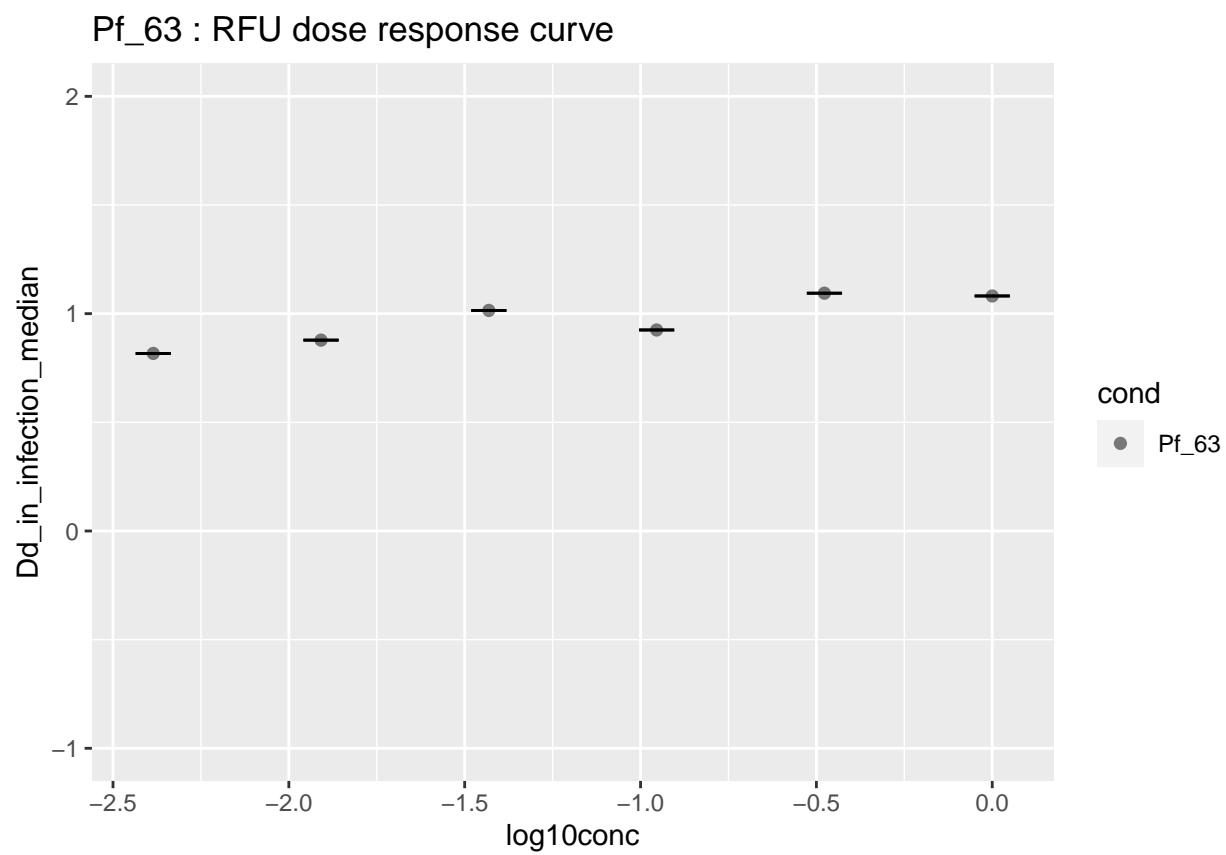




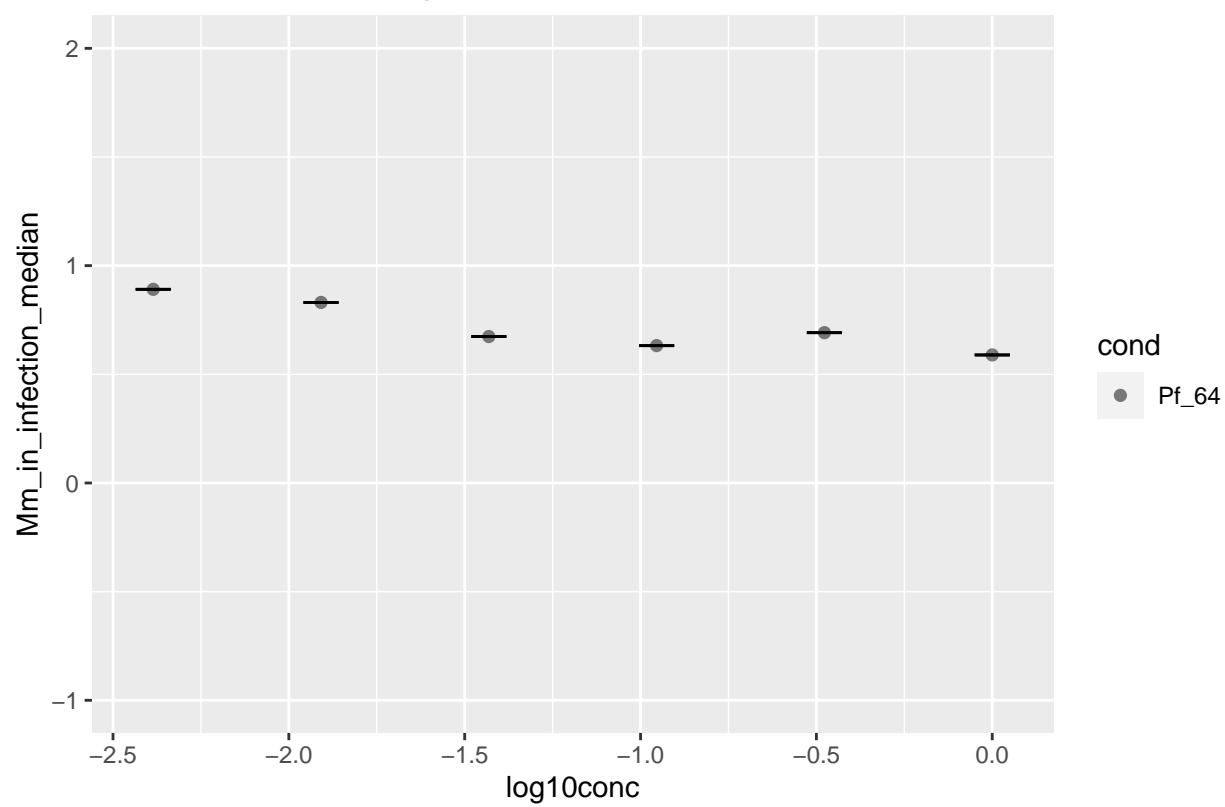


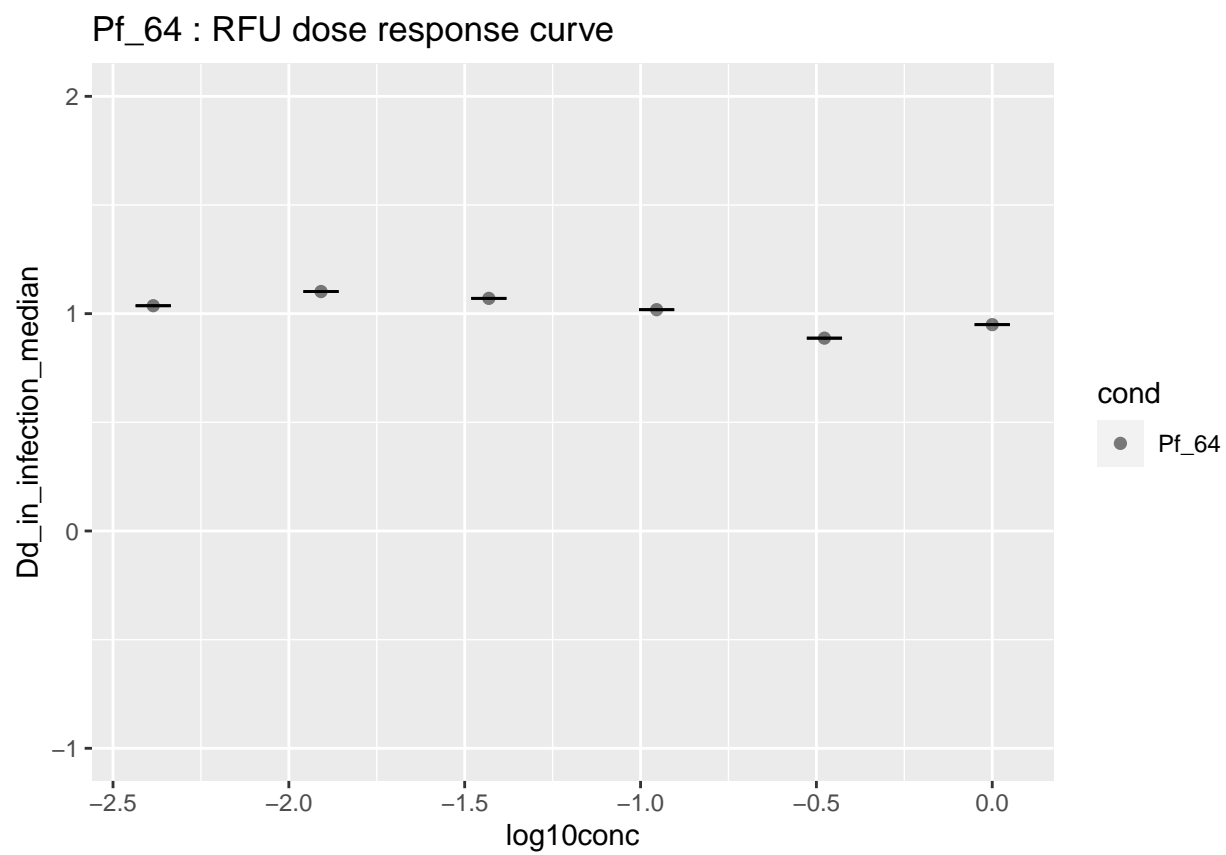
Pf_63 : RLU dose response curve

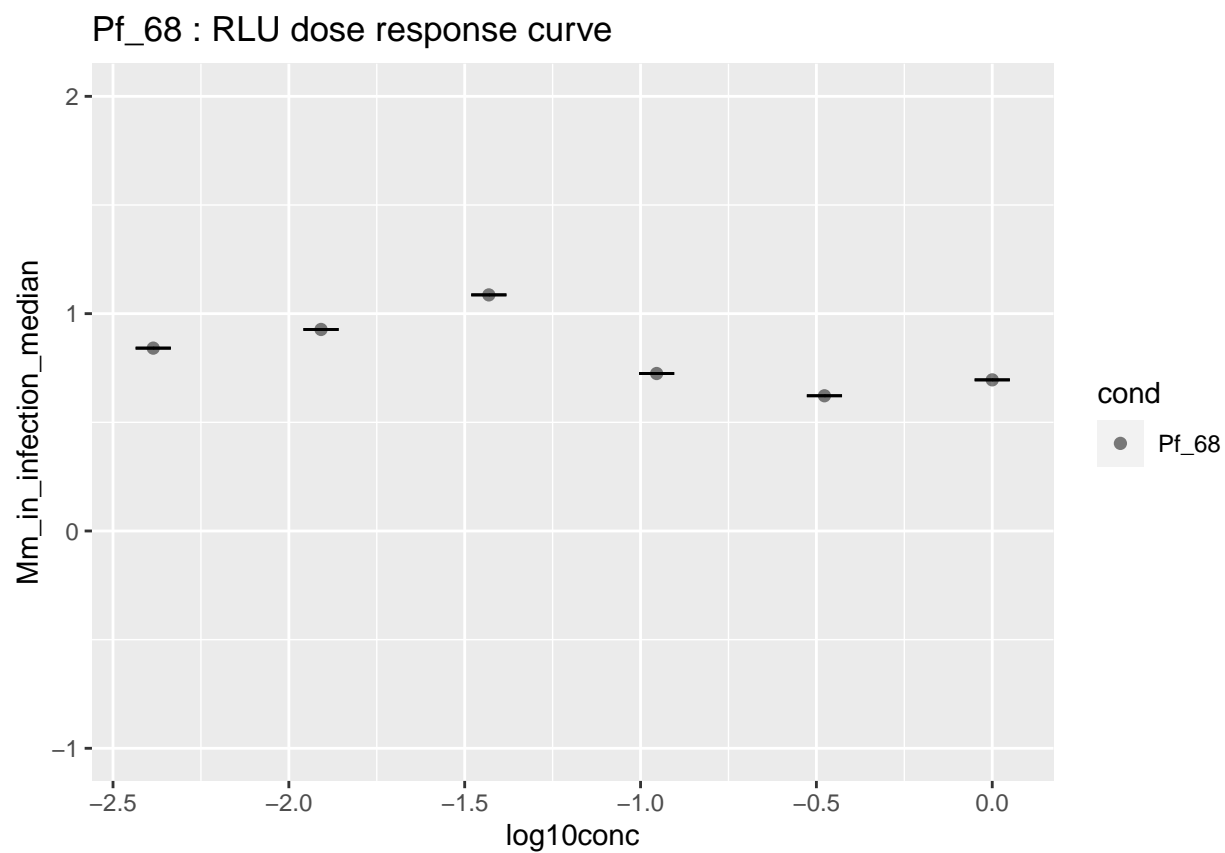


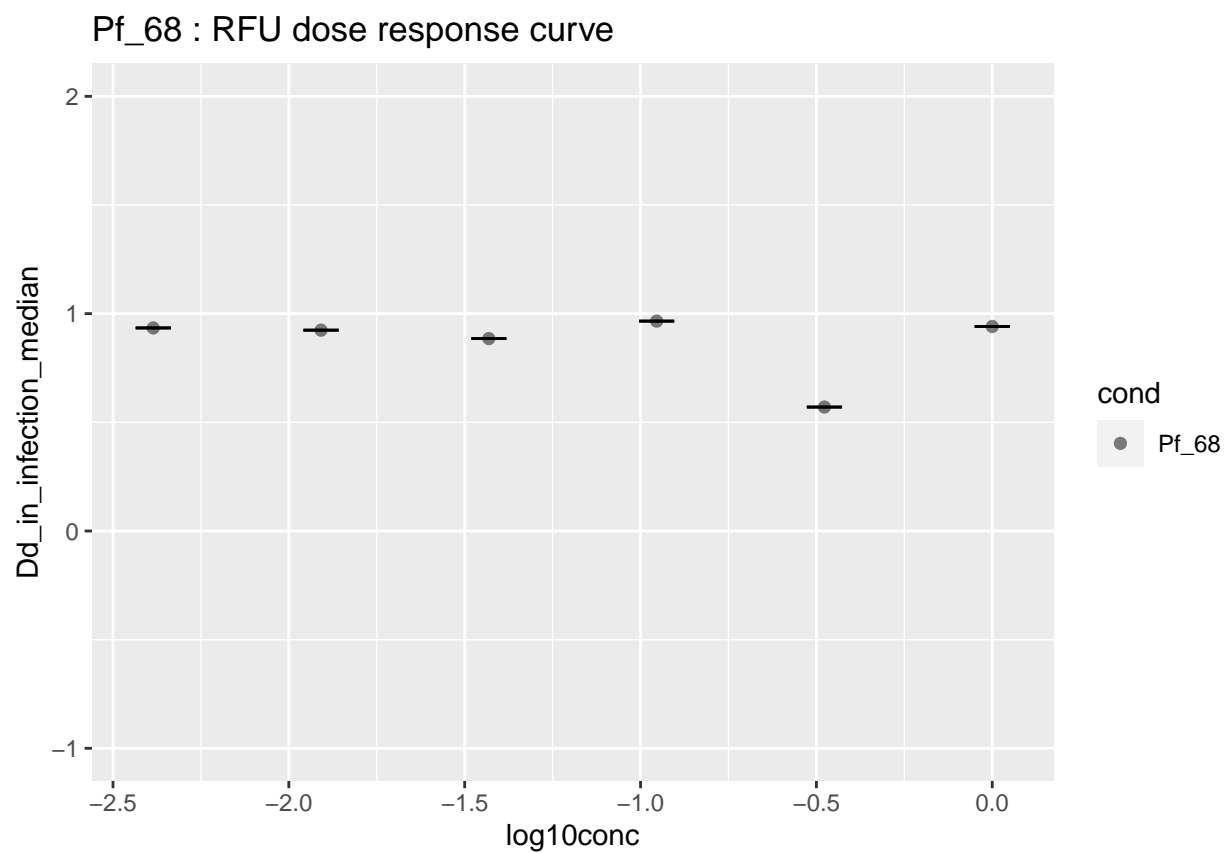


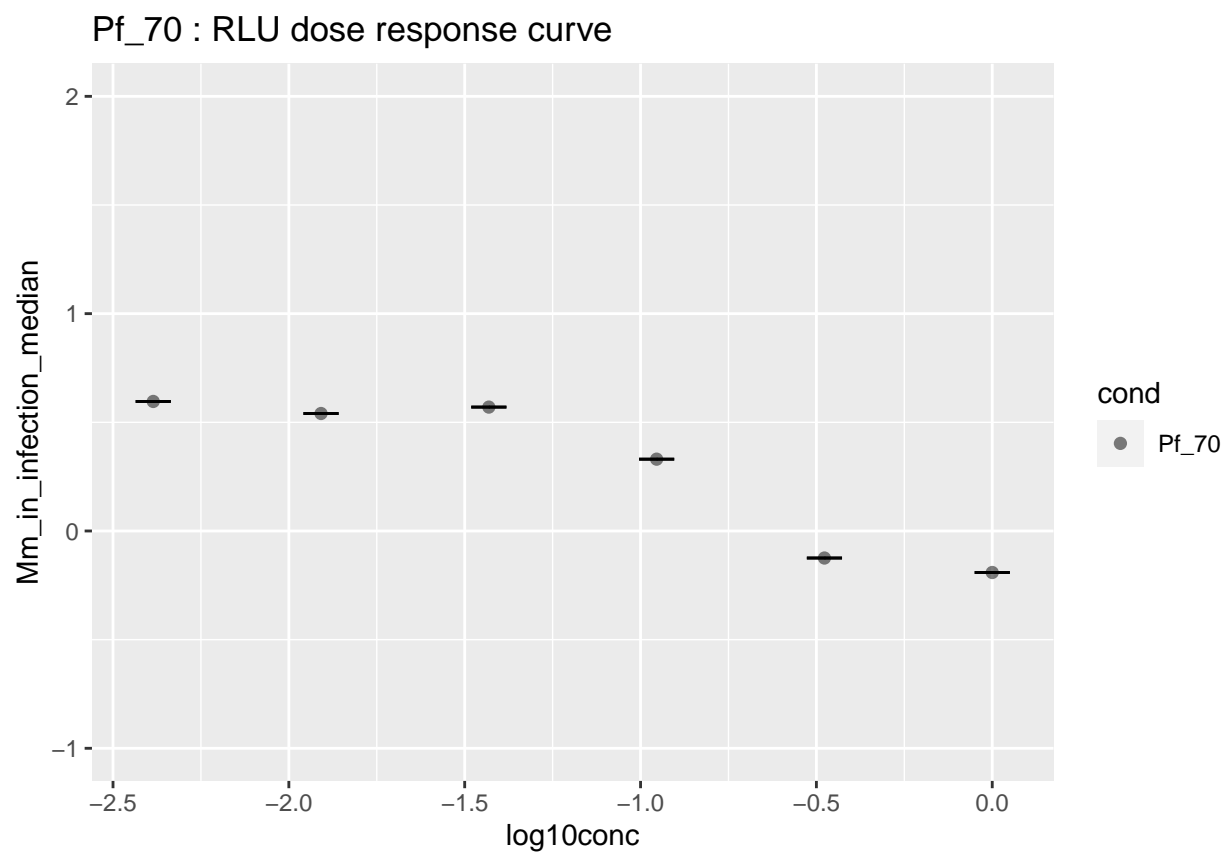
Pf_64 : RLU dose response curve

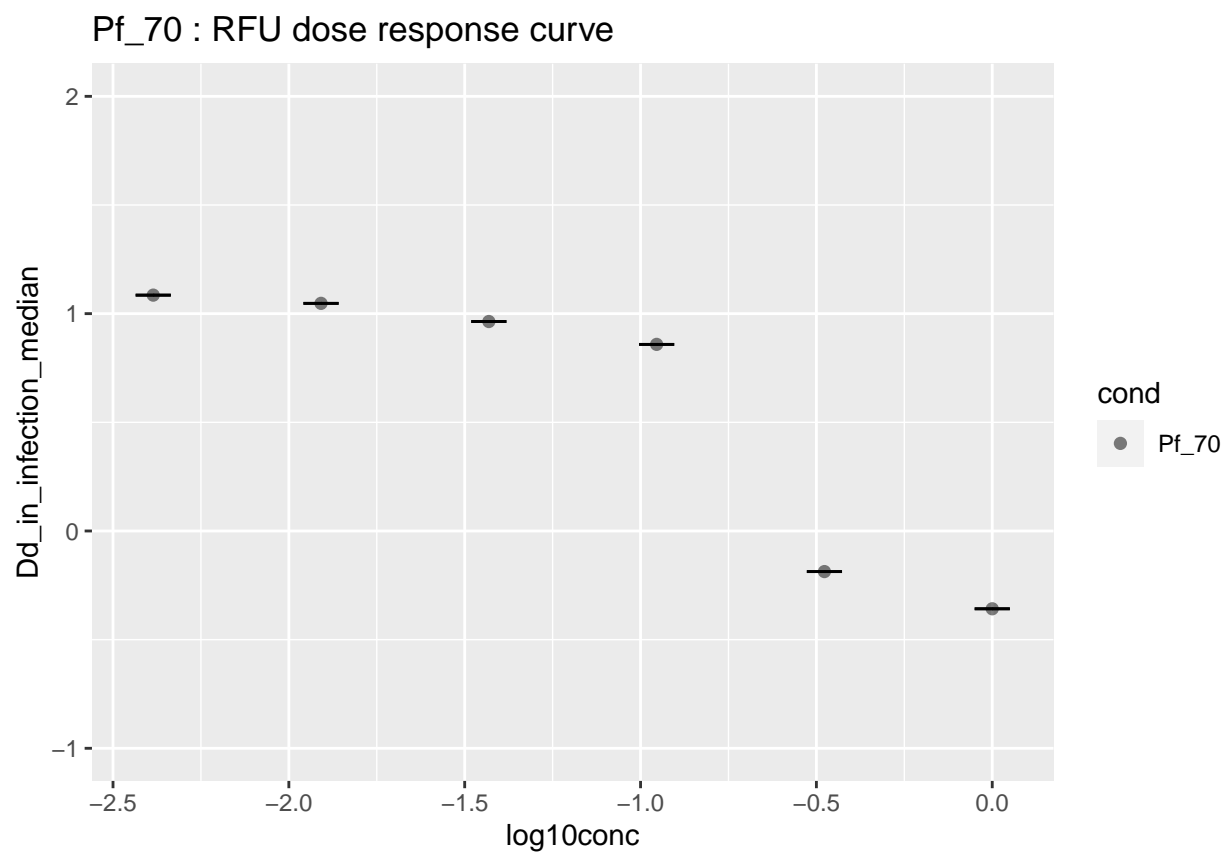


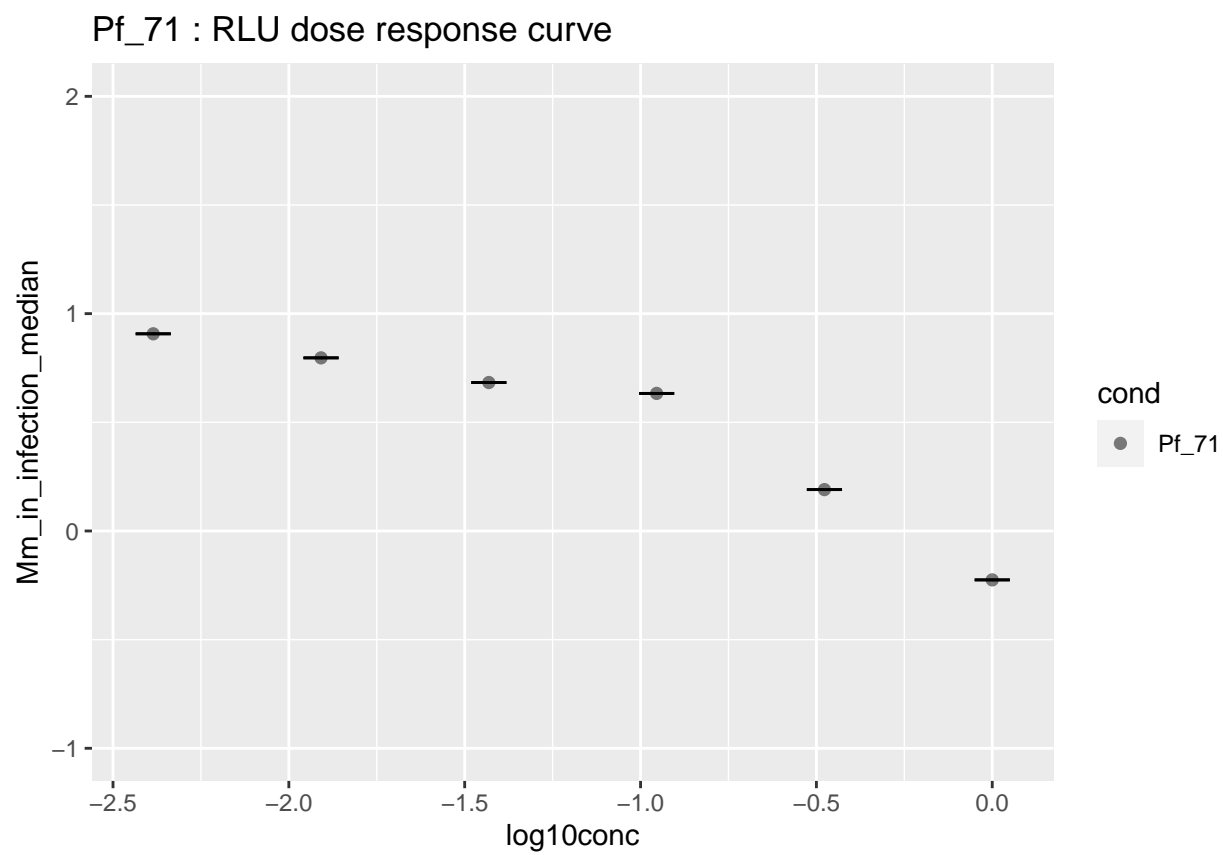


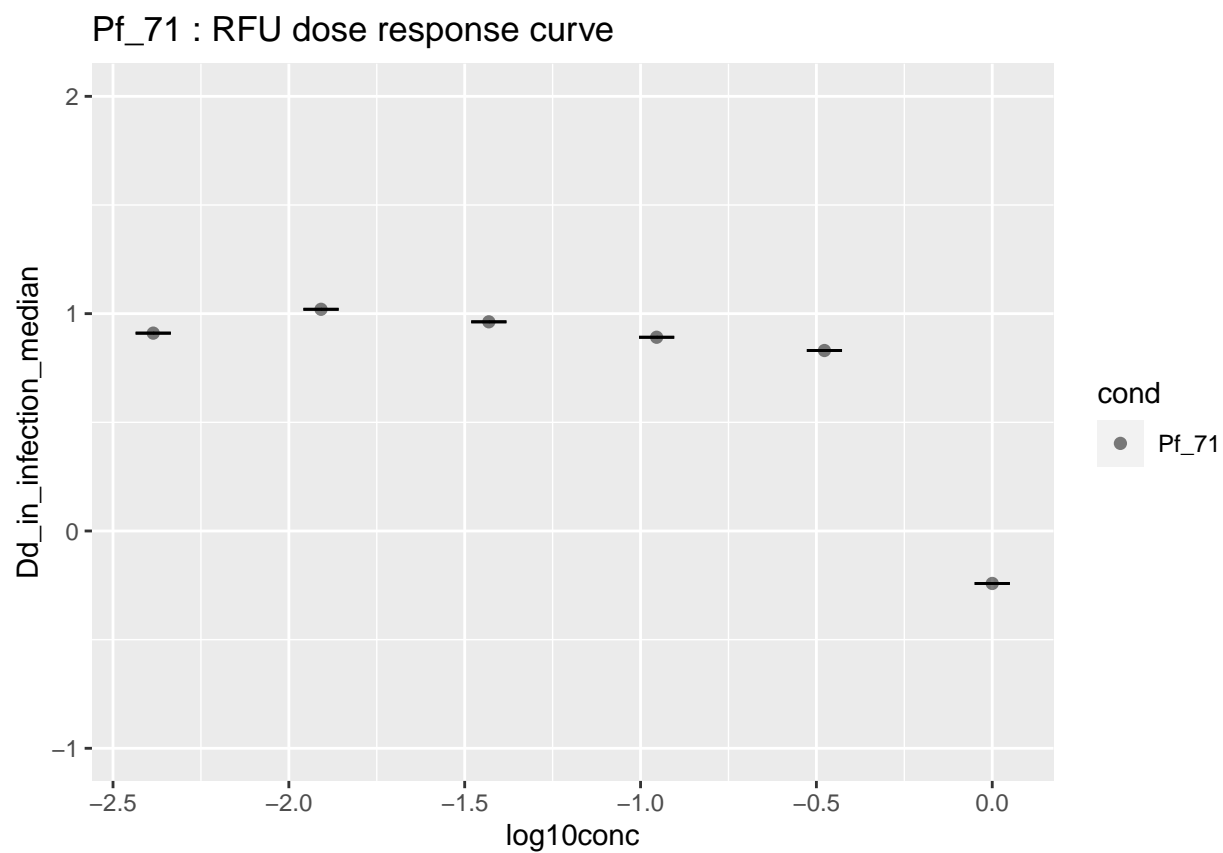


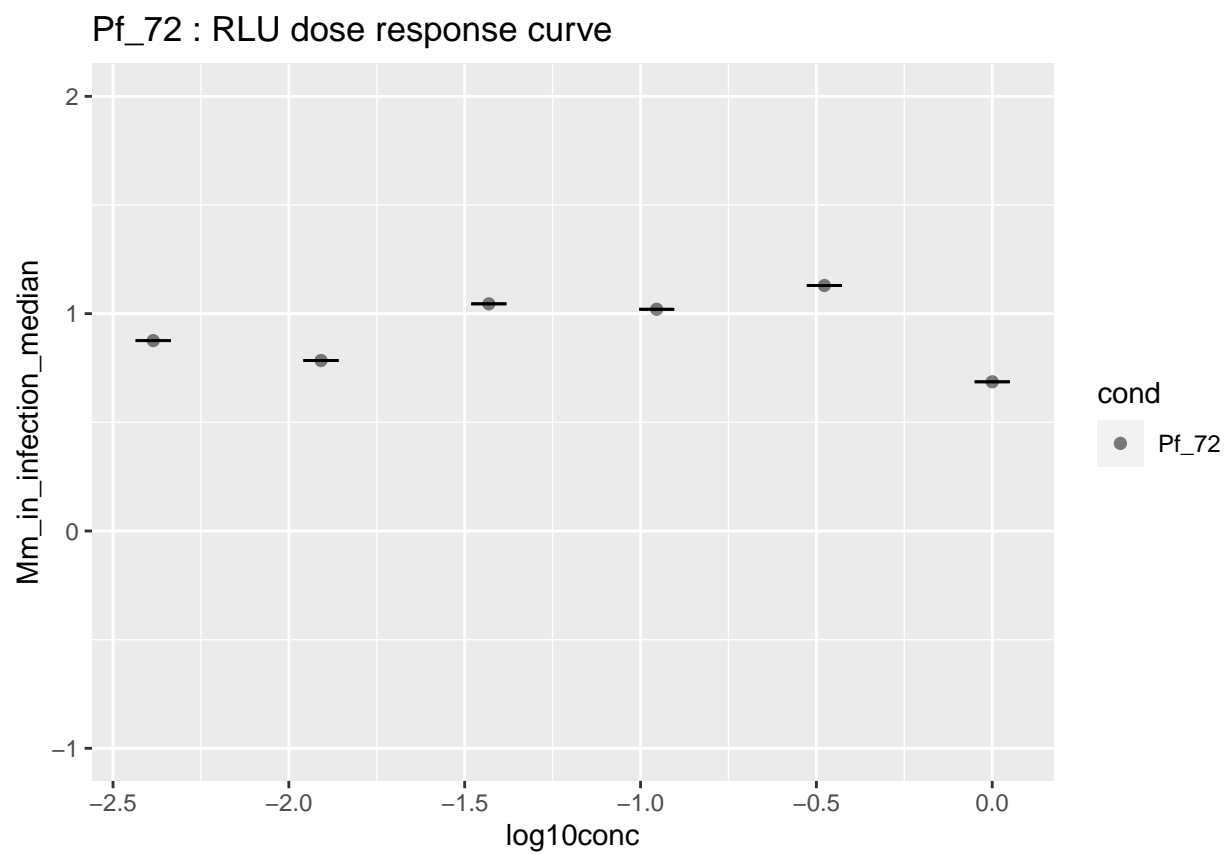


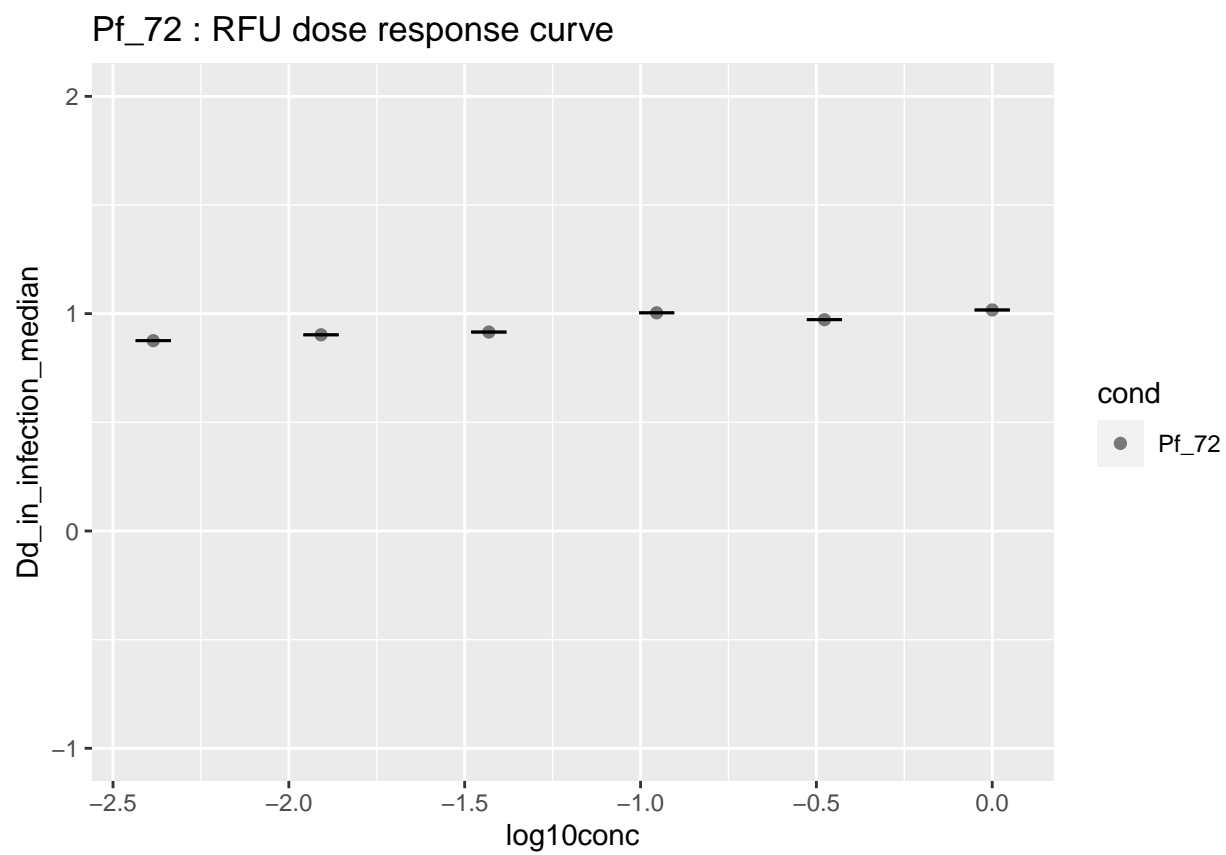


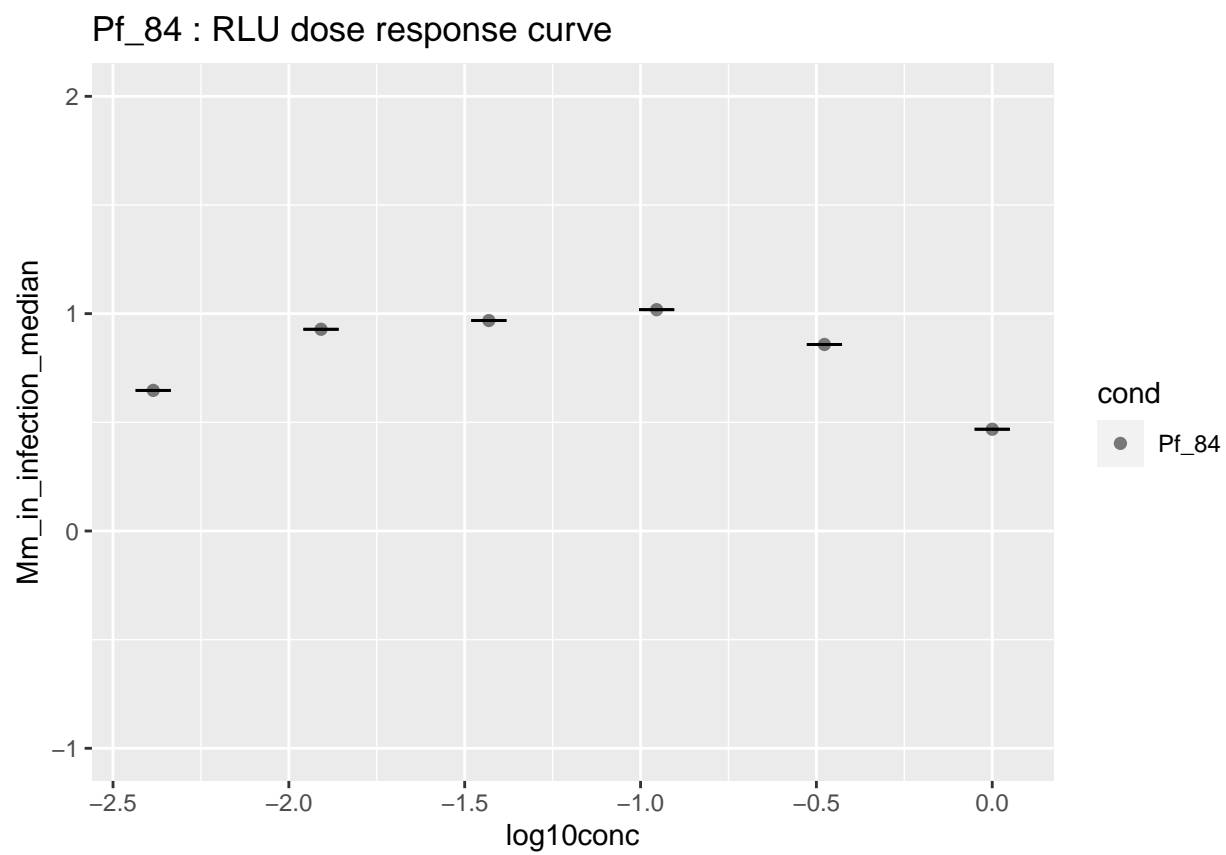


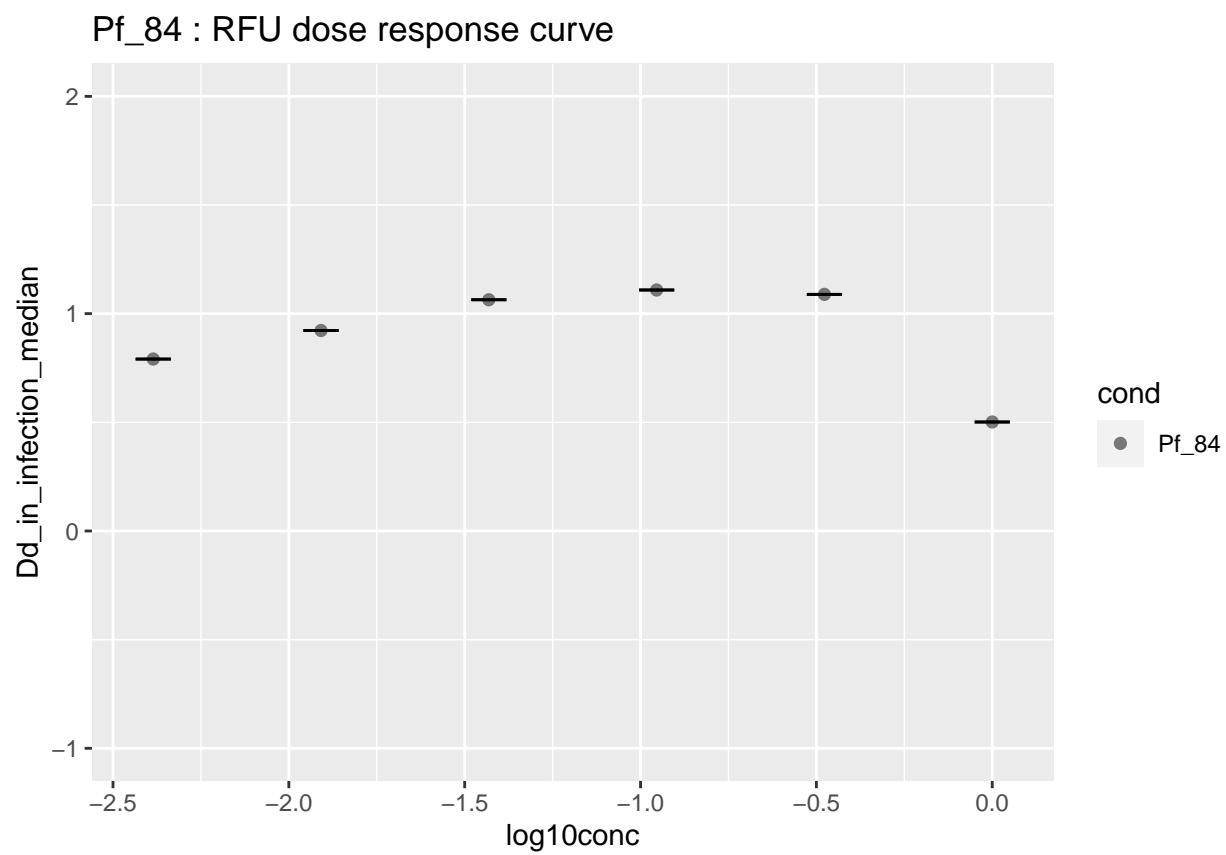


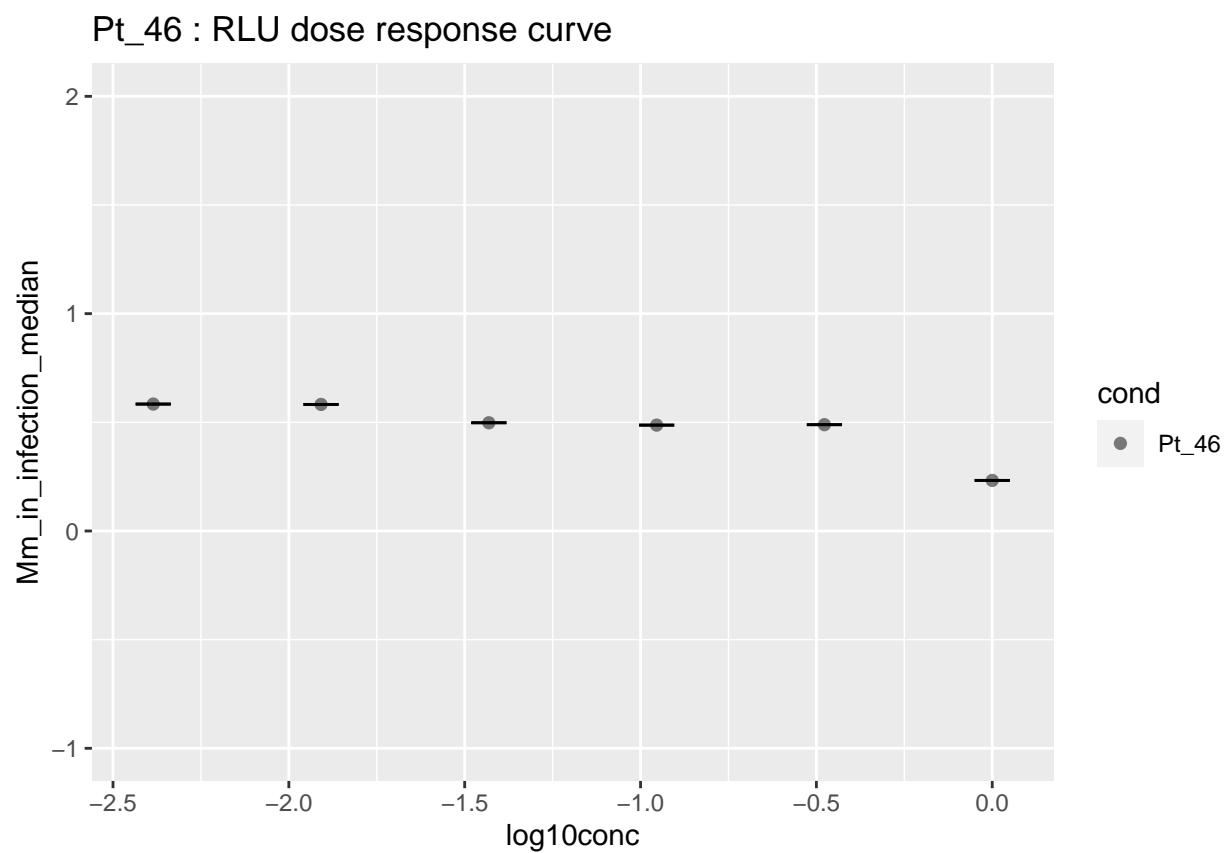


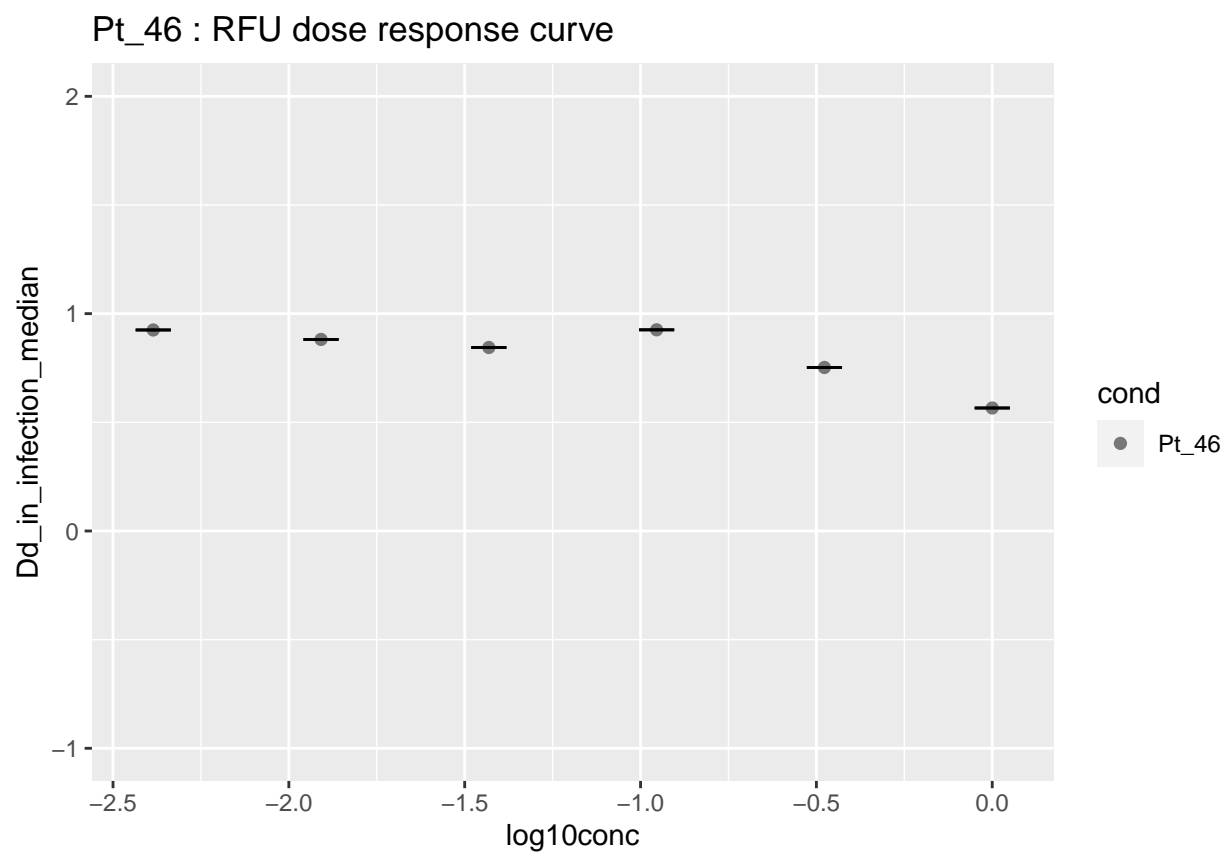


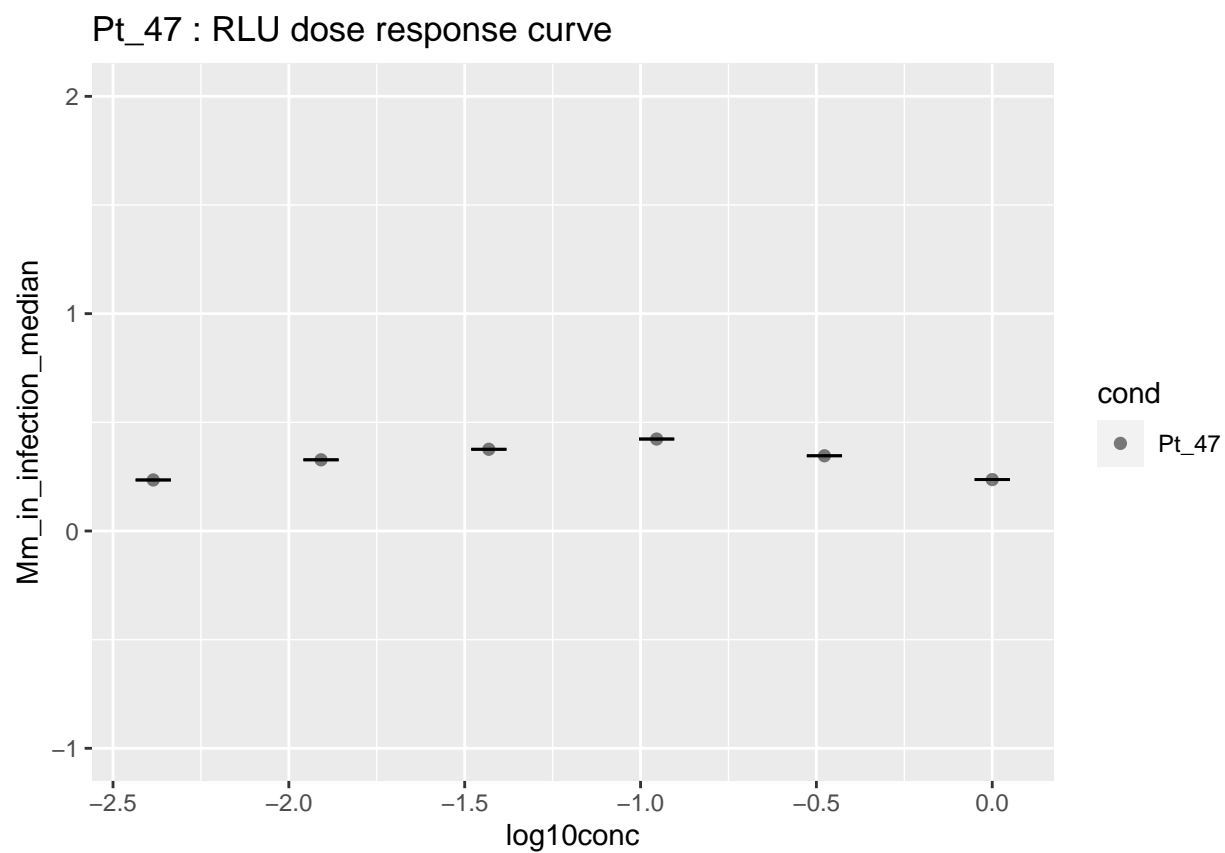


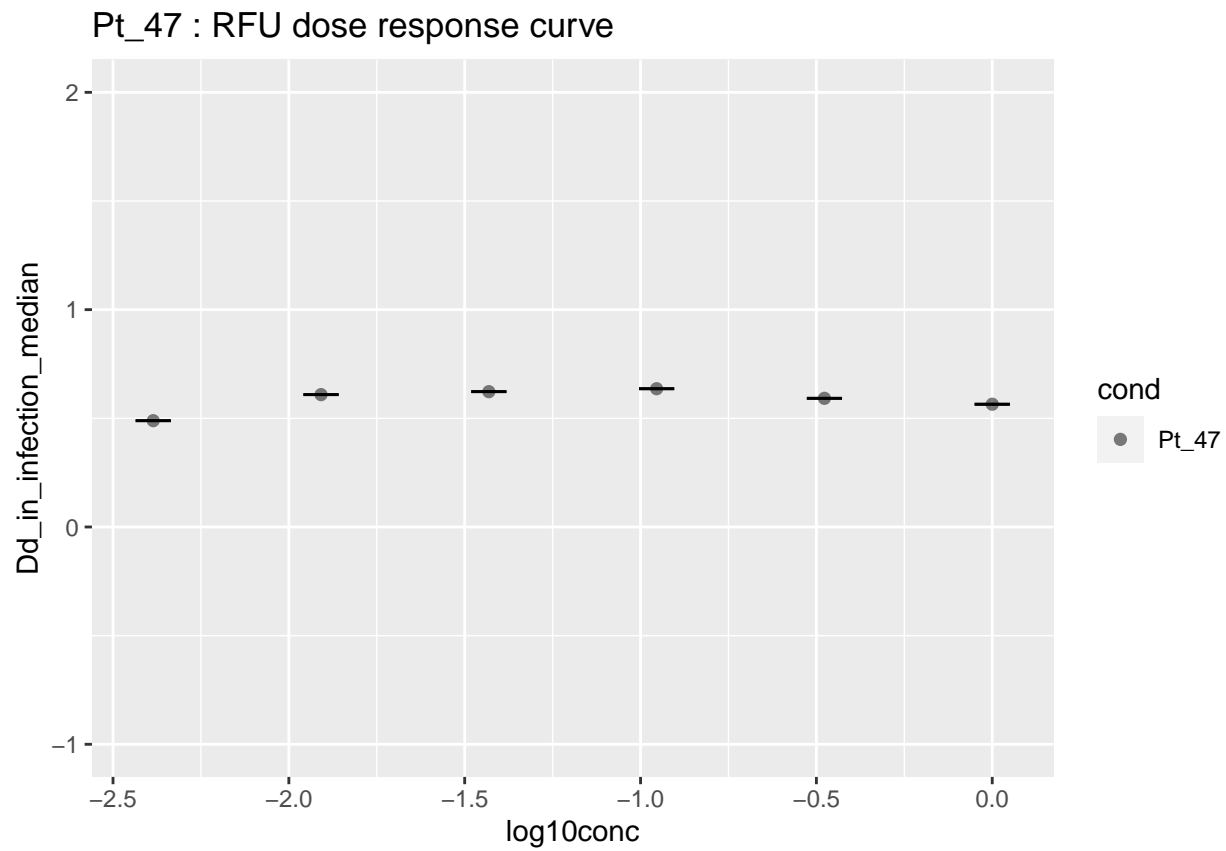




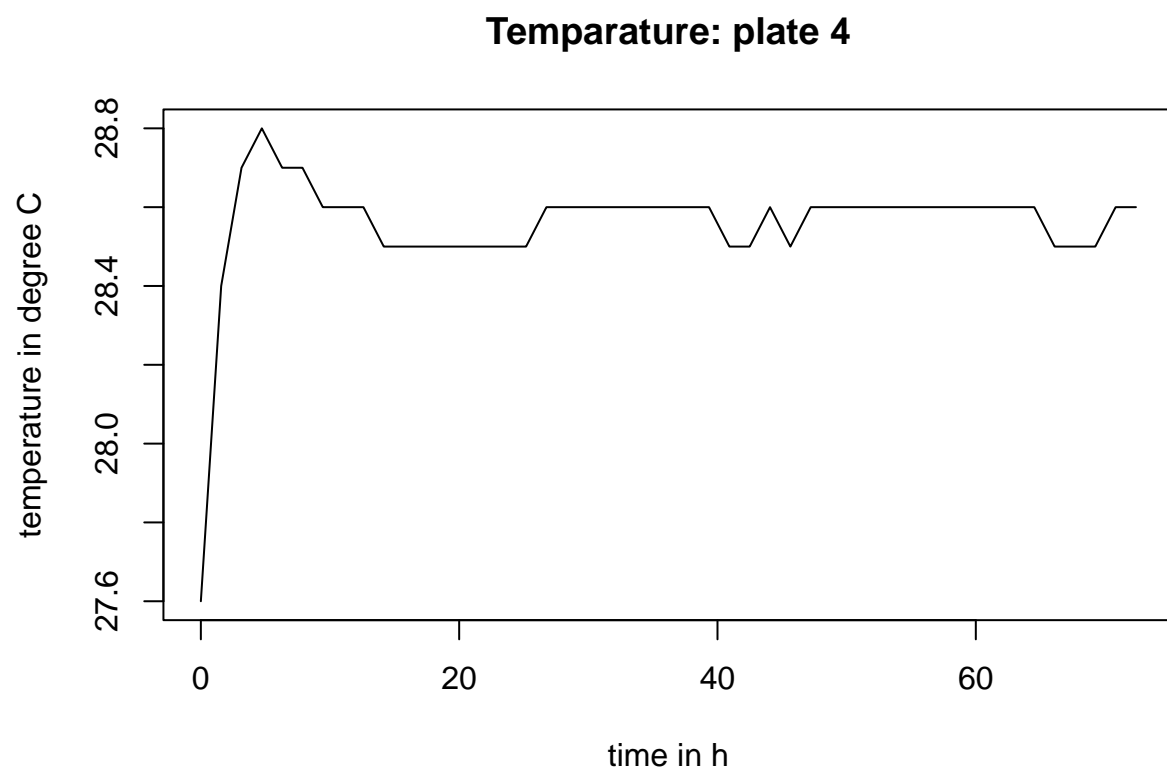




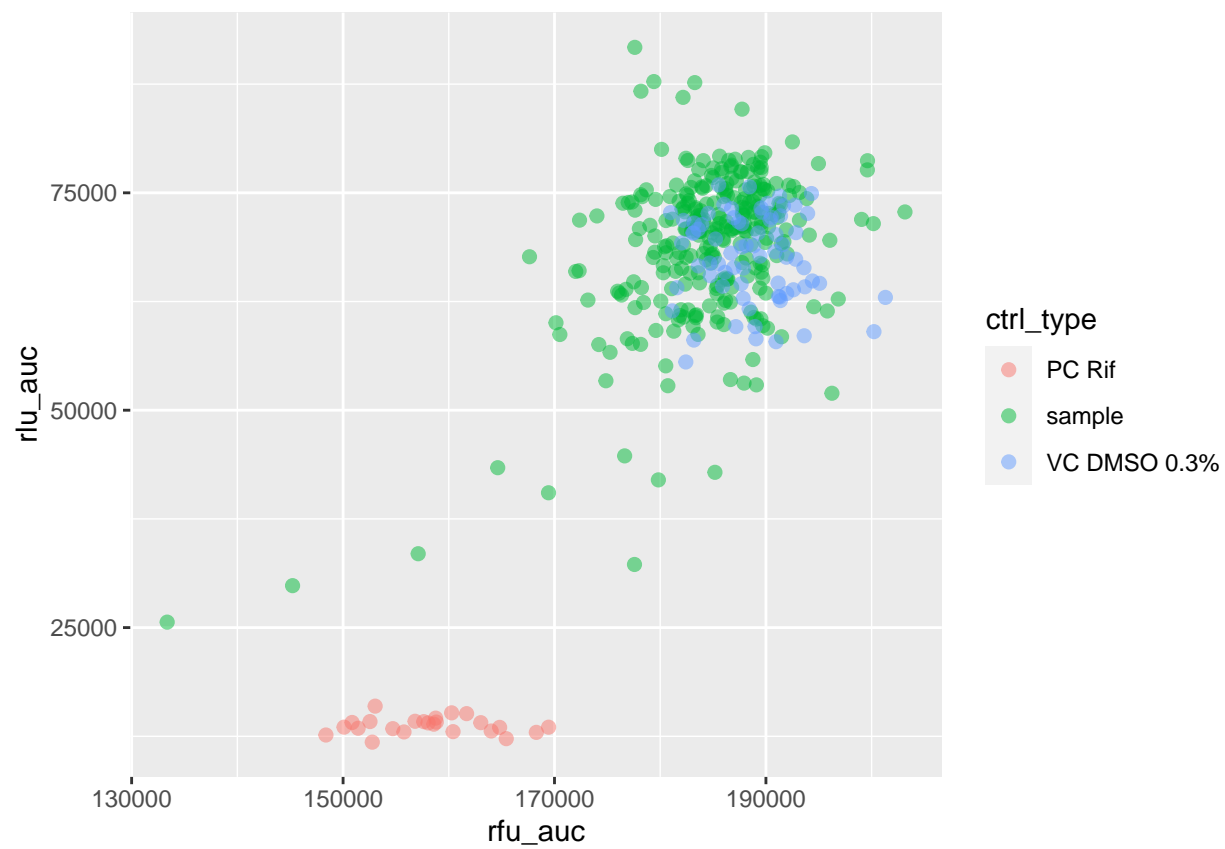


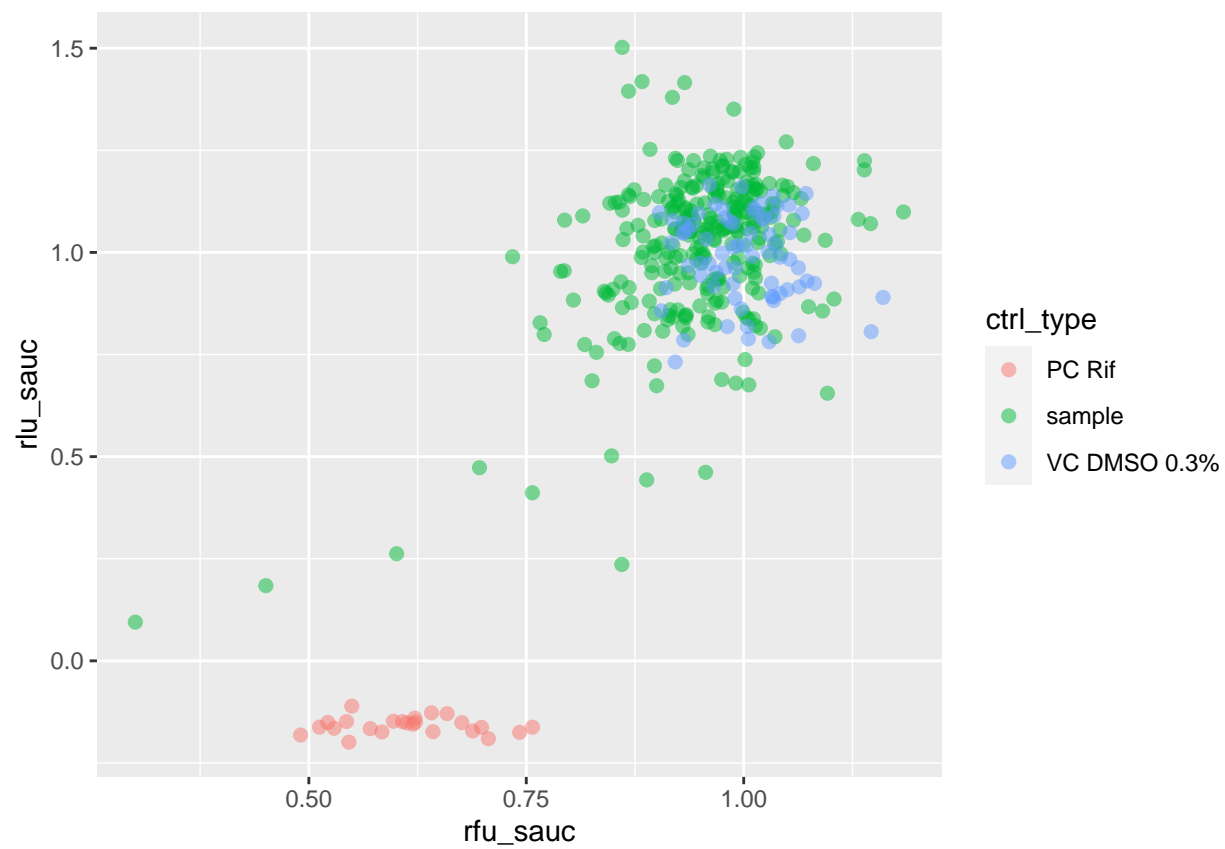


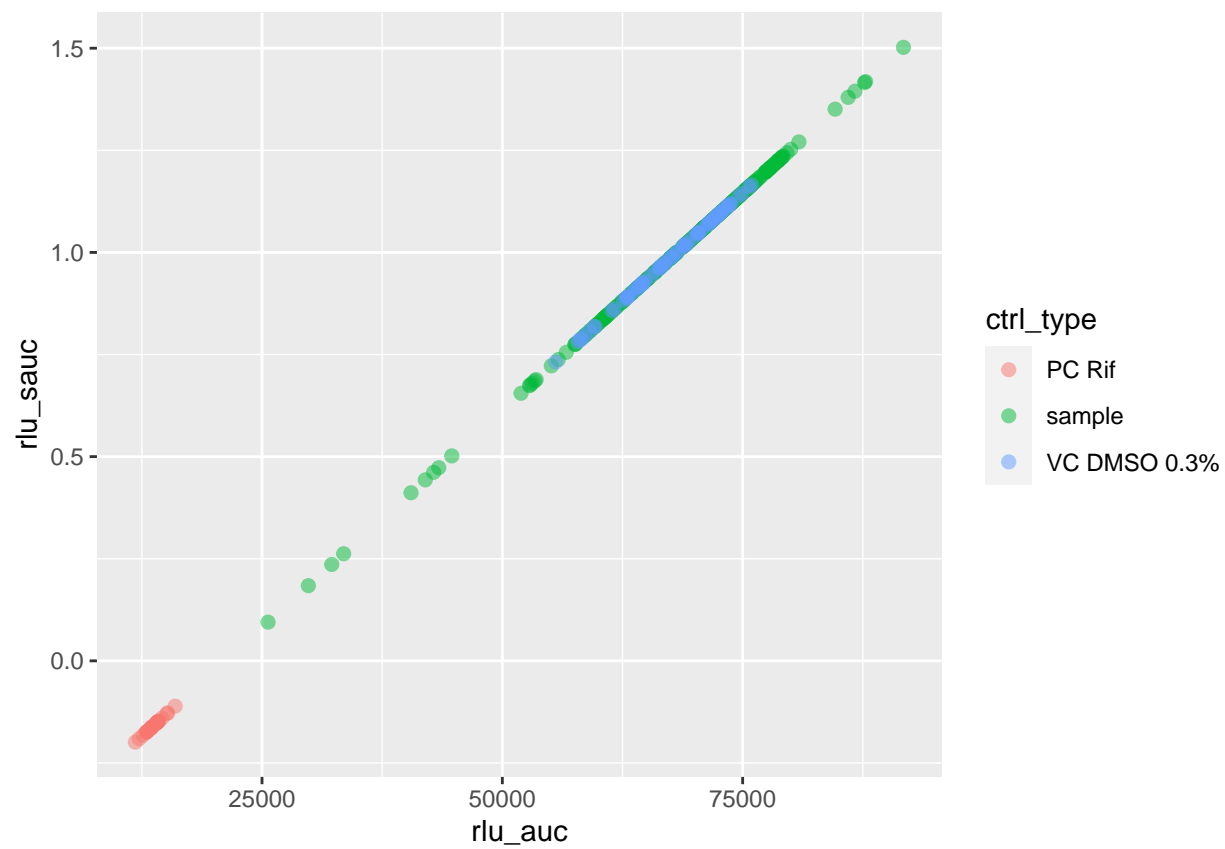
```
## [1] "analysis for plate05.xlsx"
```

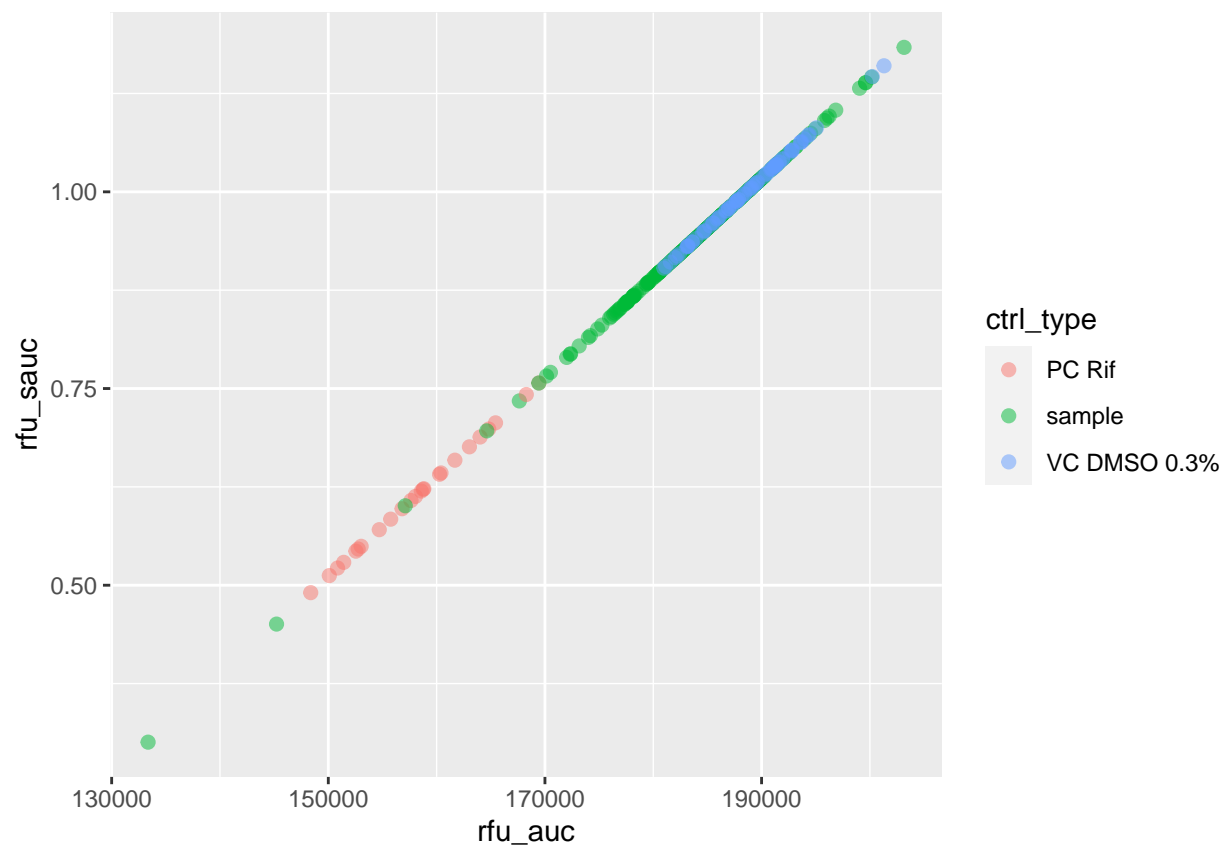


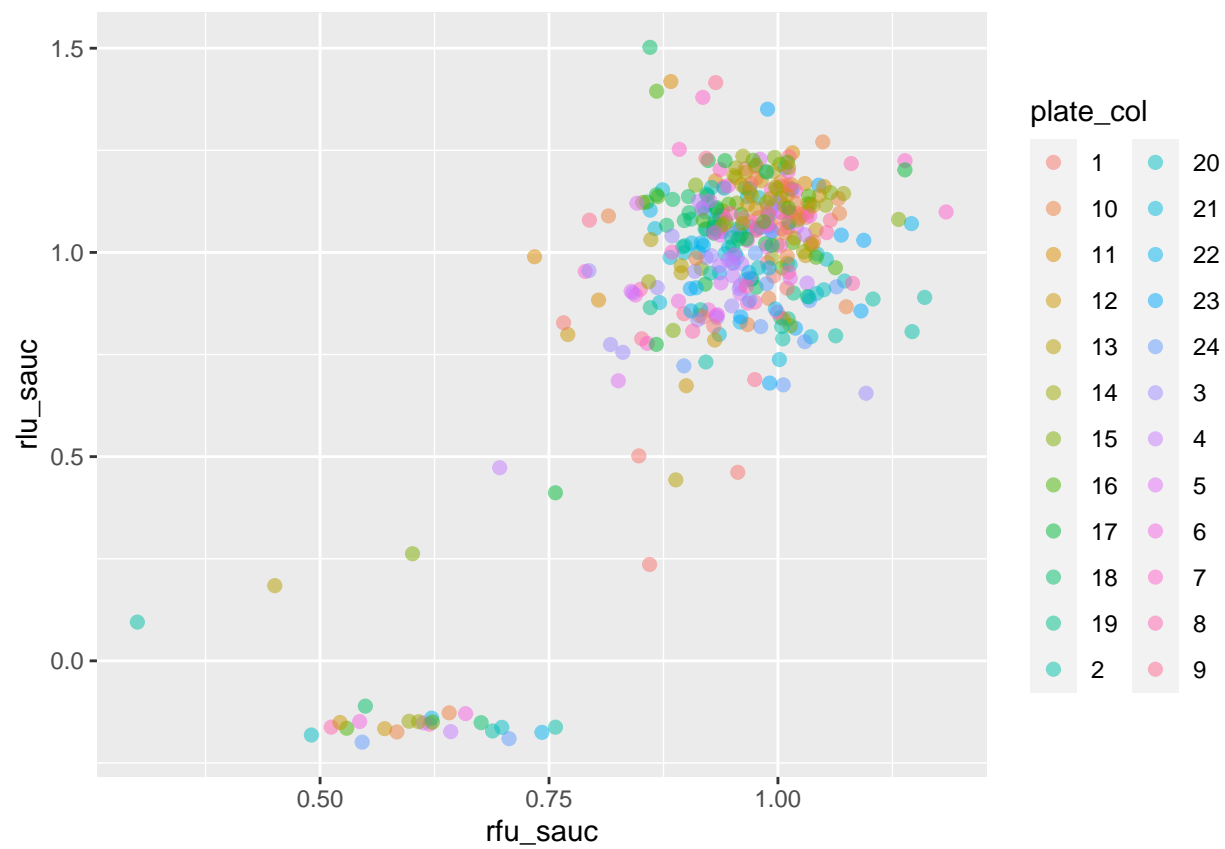
```
## [1] "VC DMSO 0.3% Robust z'-factor of rlu for plate plate05.xlsx, biorep 2 : "  
## [1] 0.64
```

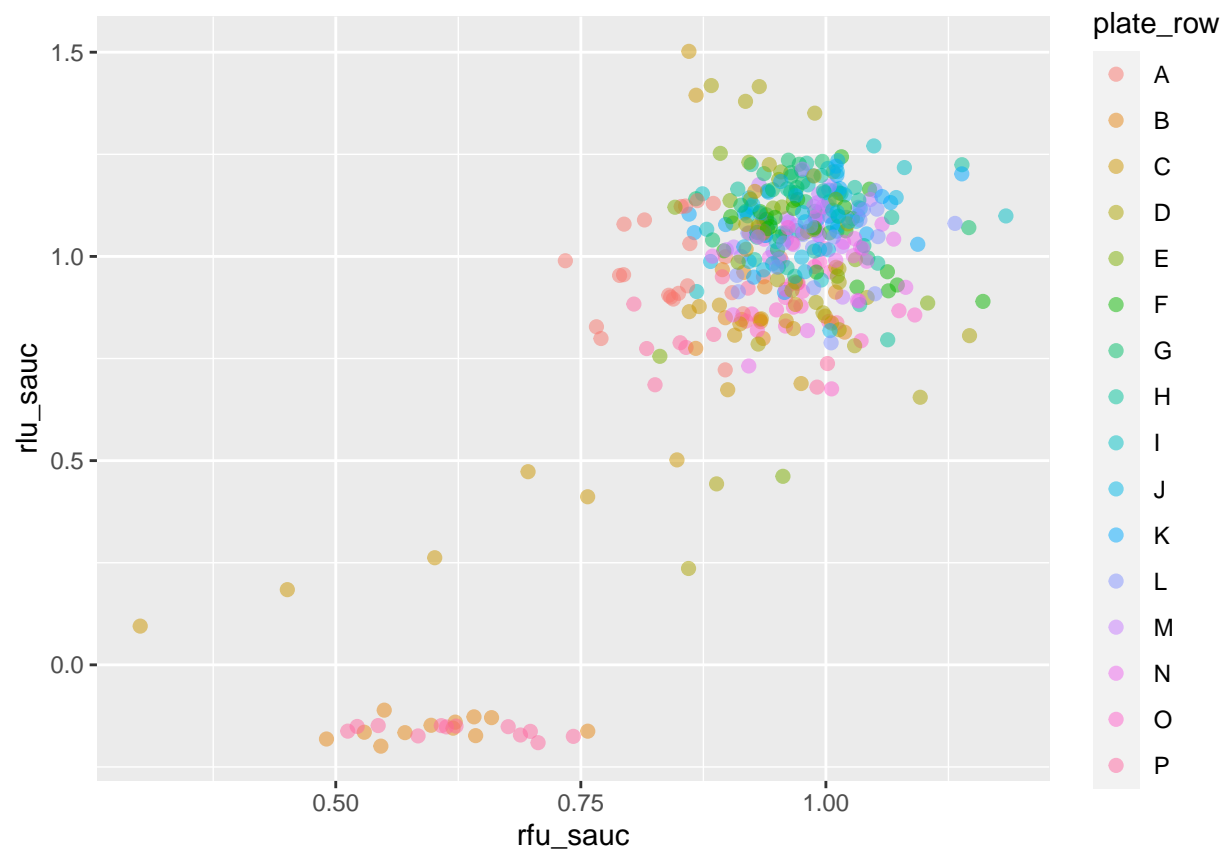





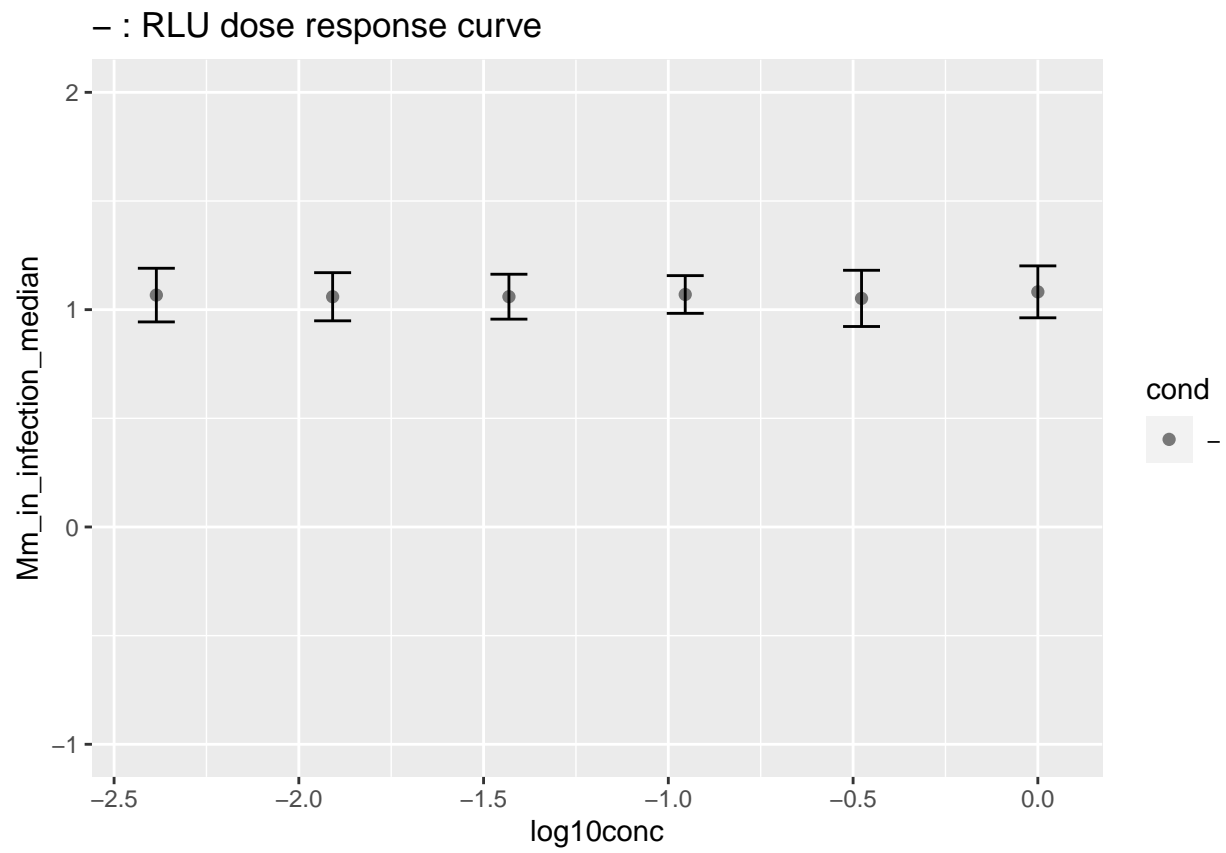


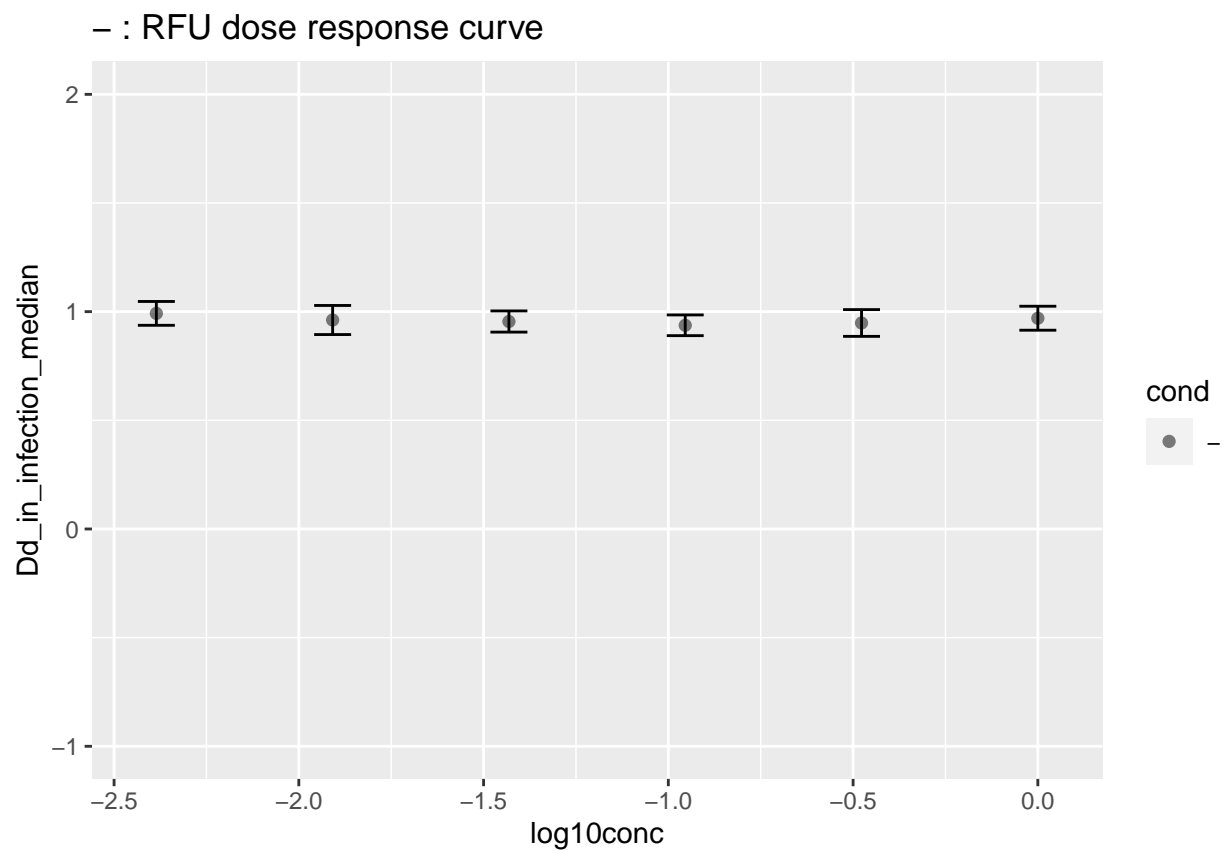




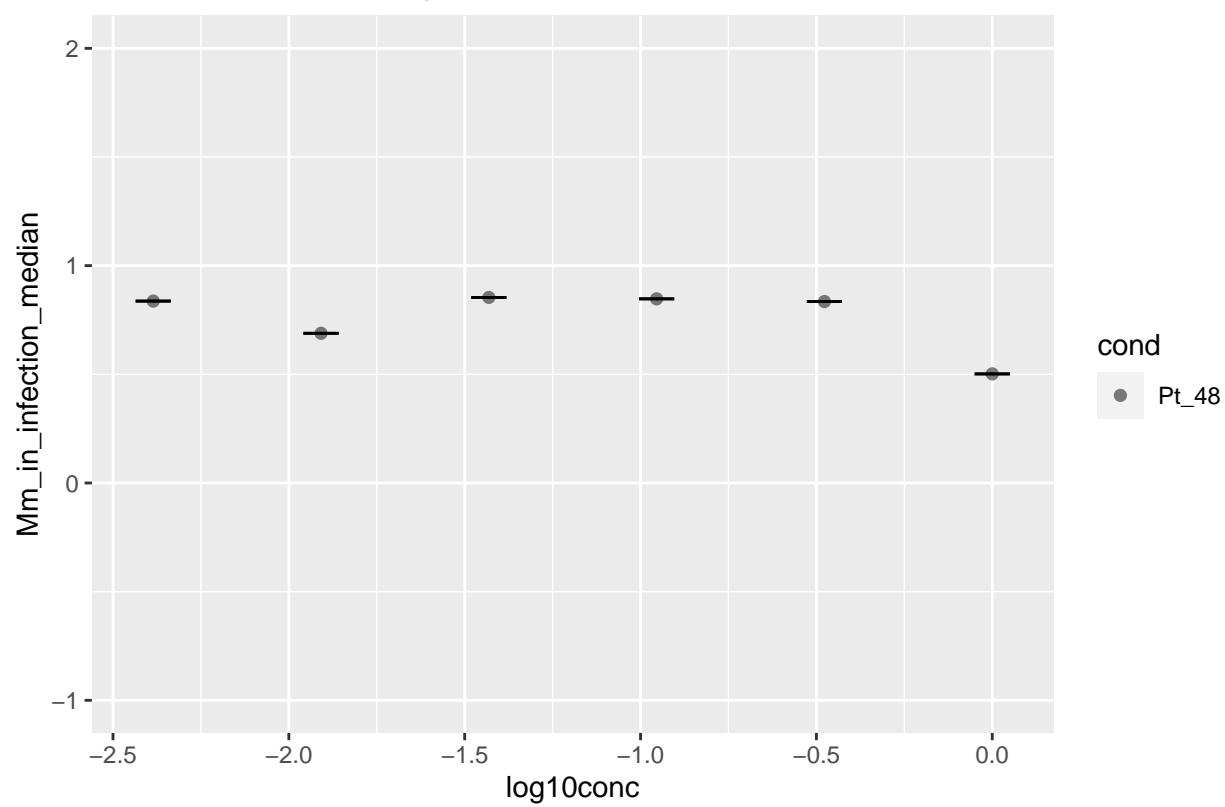


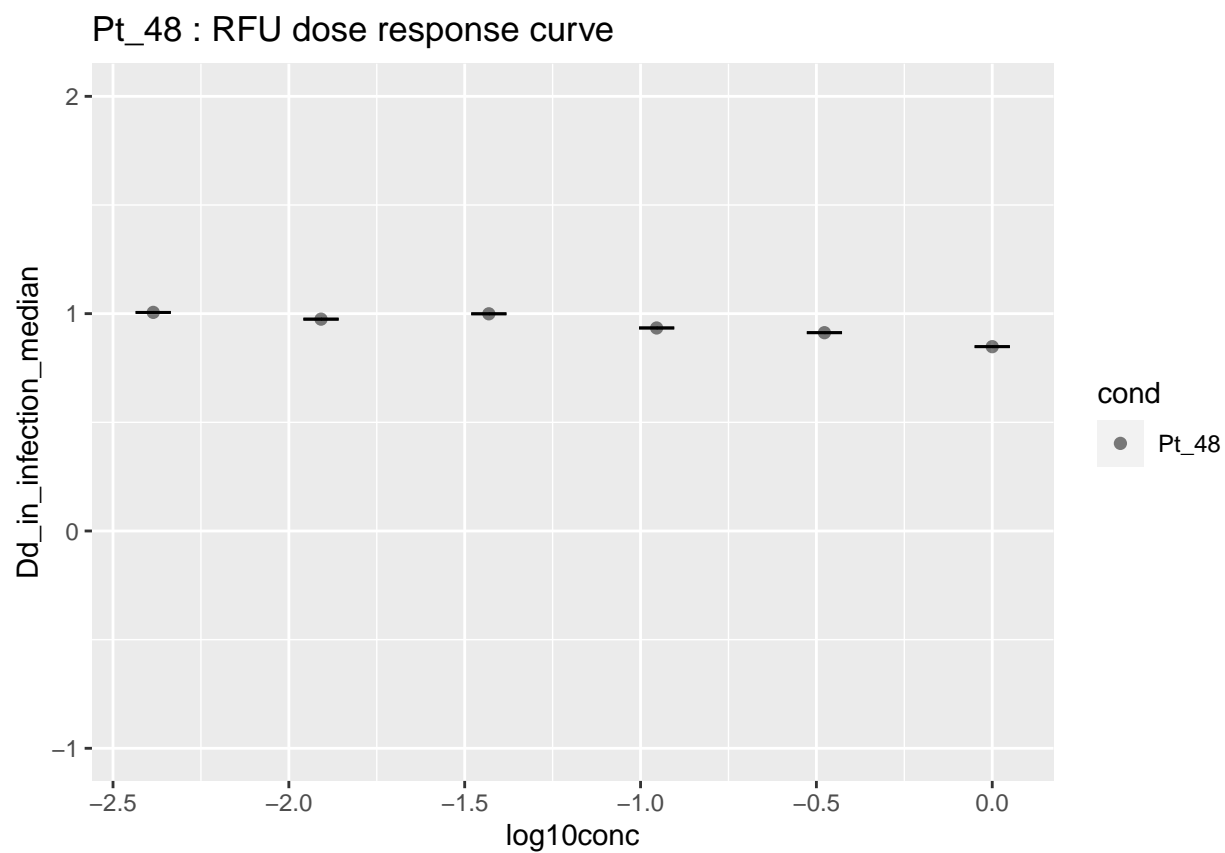
```
## [1] "Dose response curves over all bioreps within this plate"
```

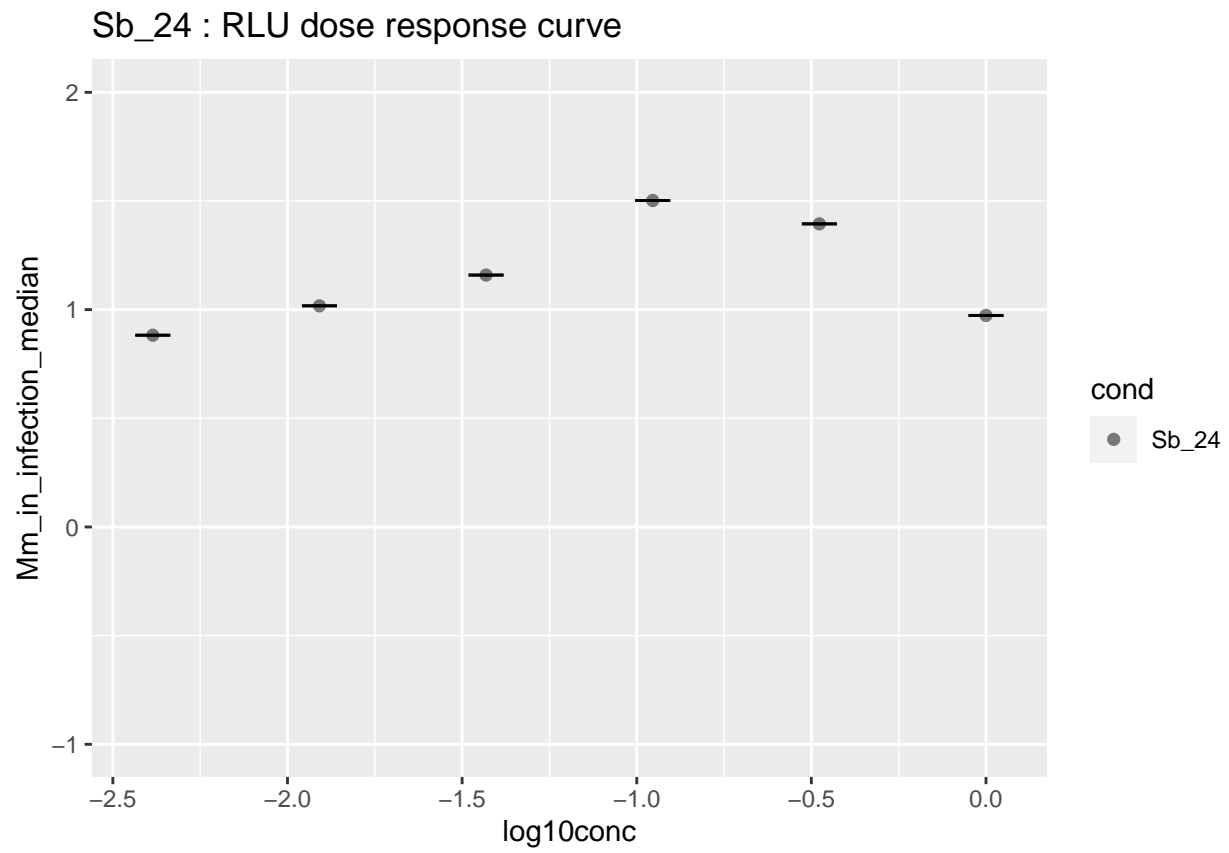


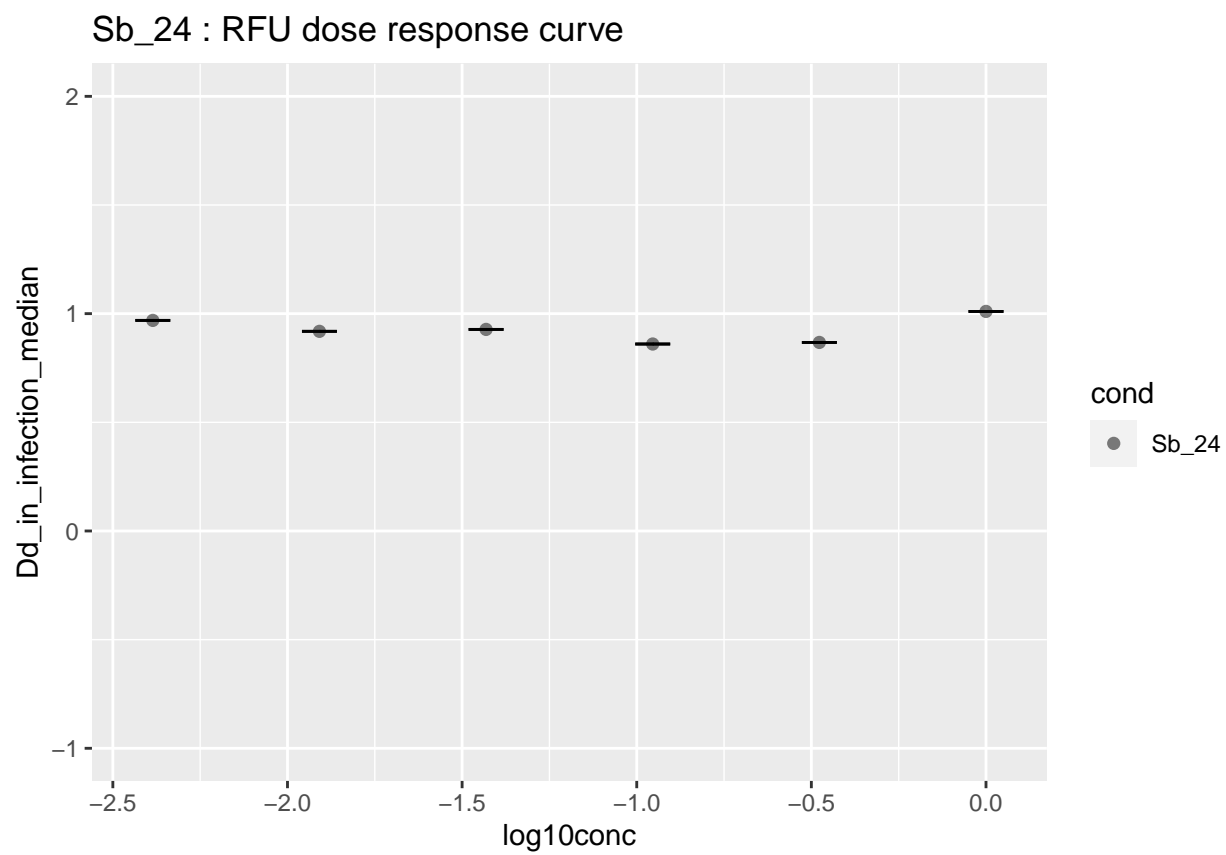


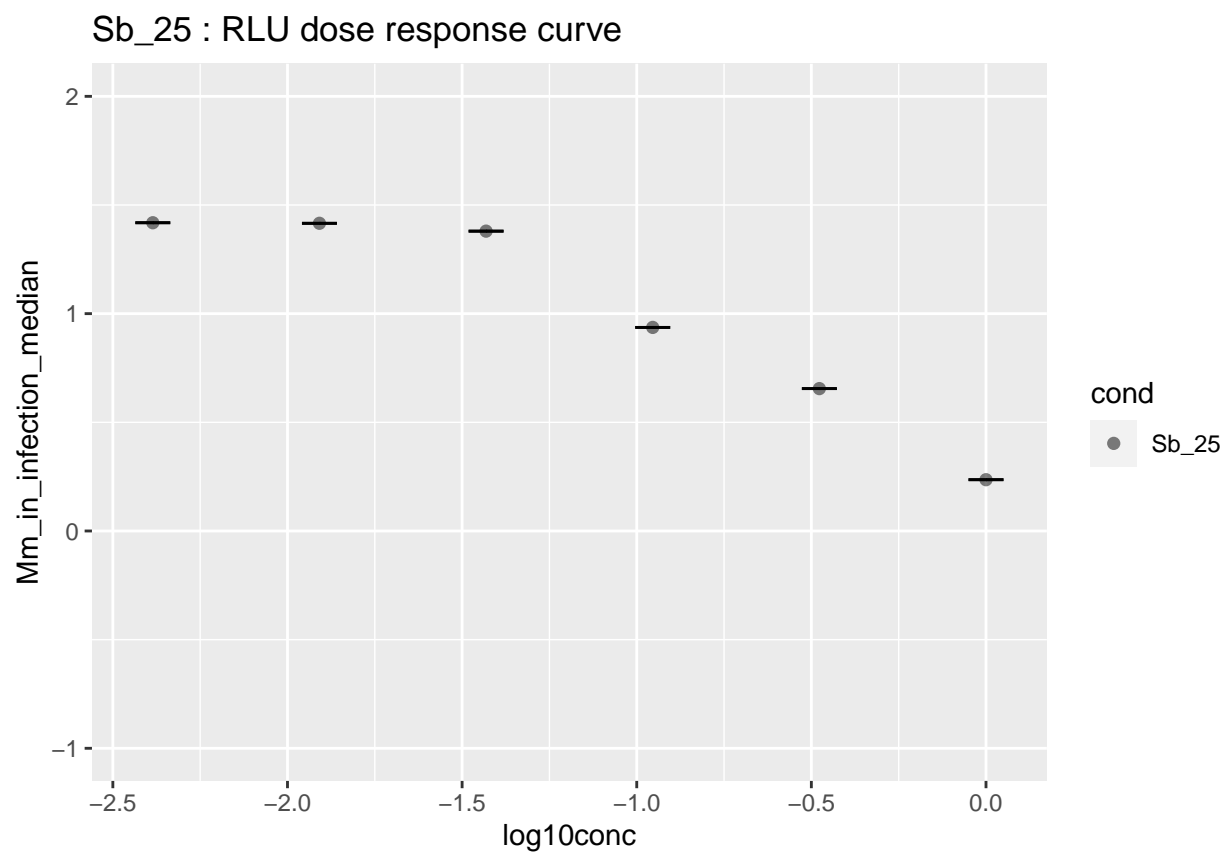
Pt_48 : RLU dose response curve

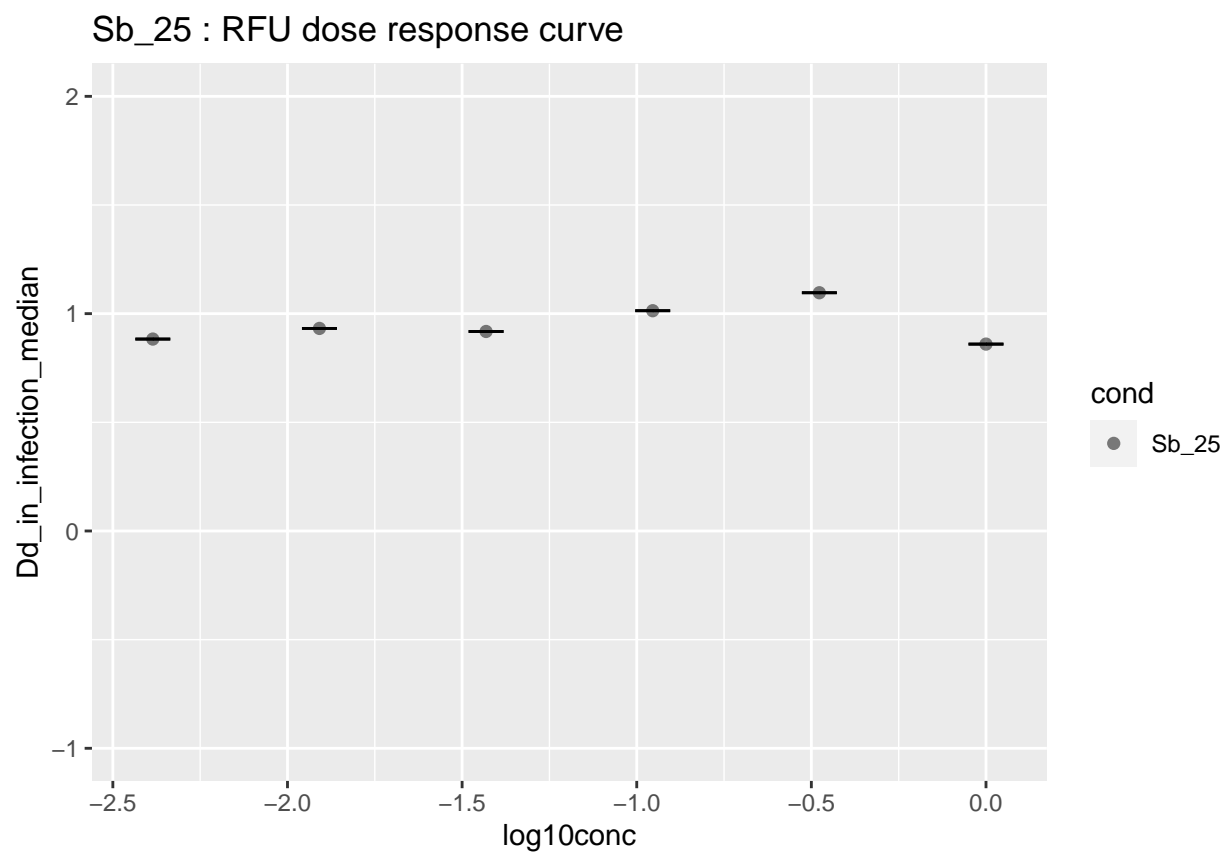


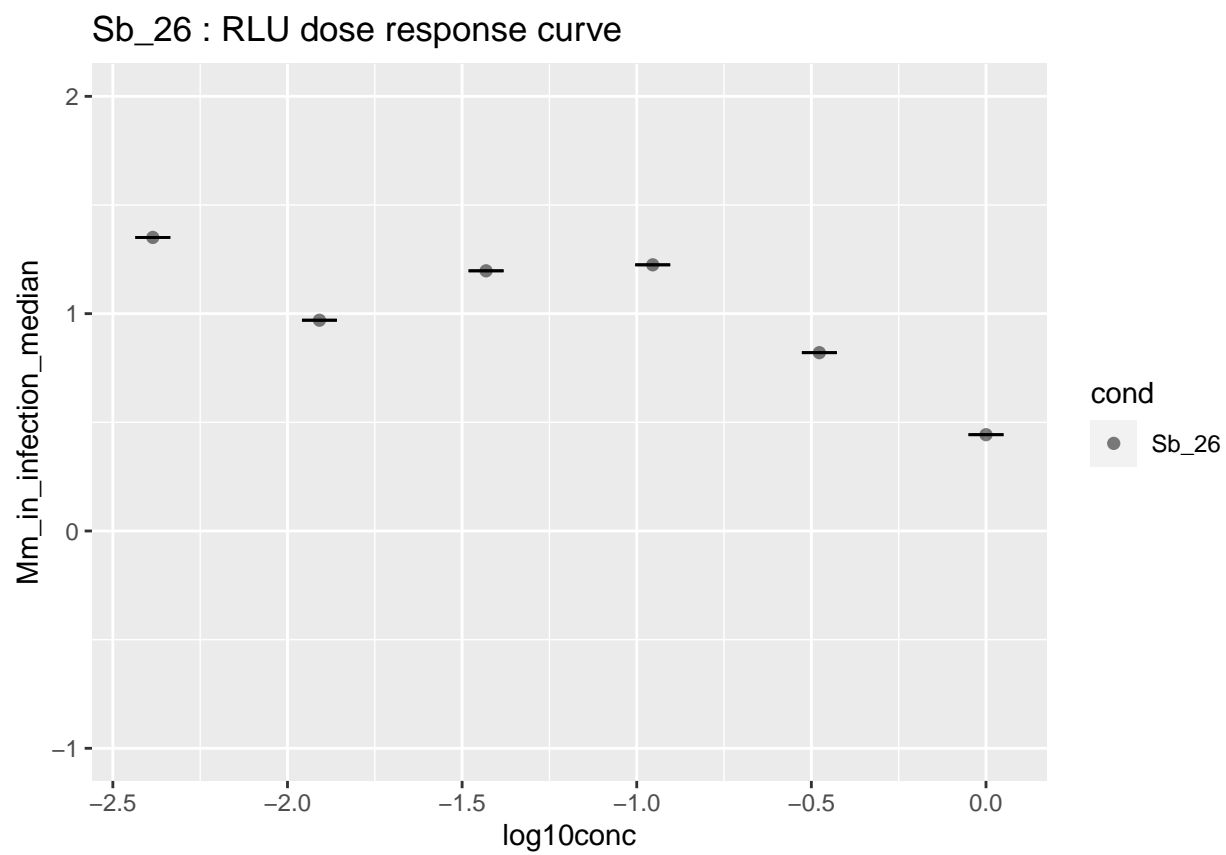


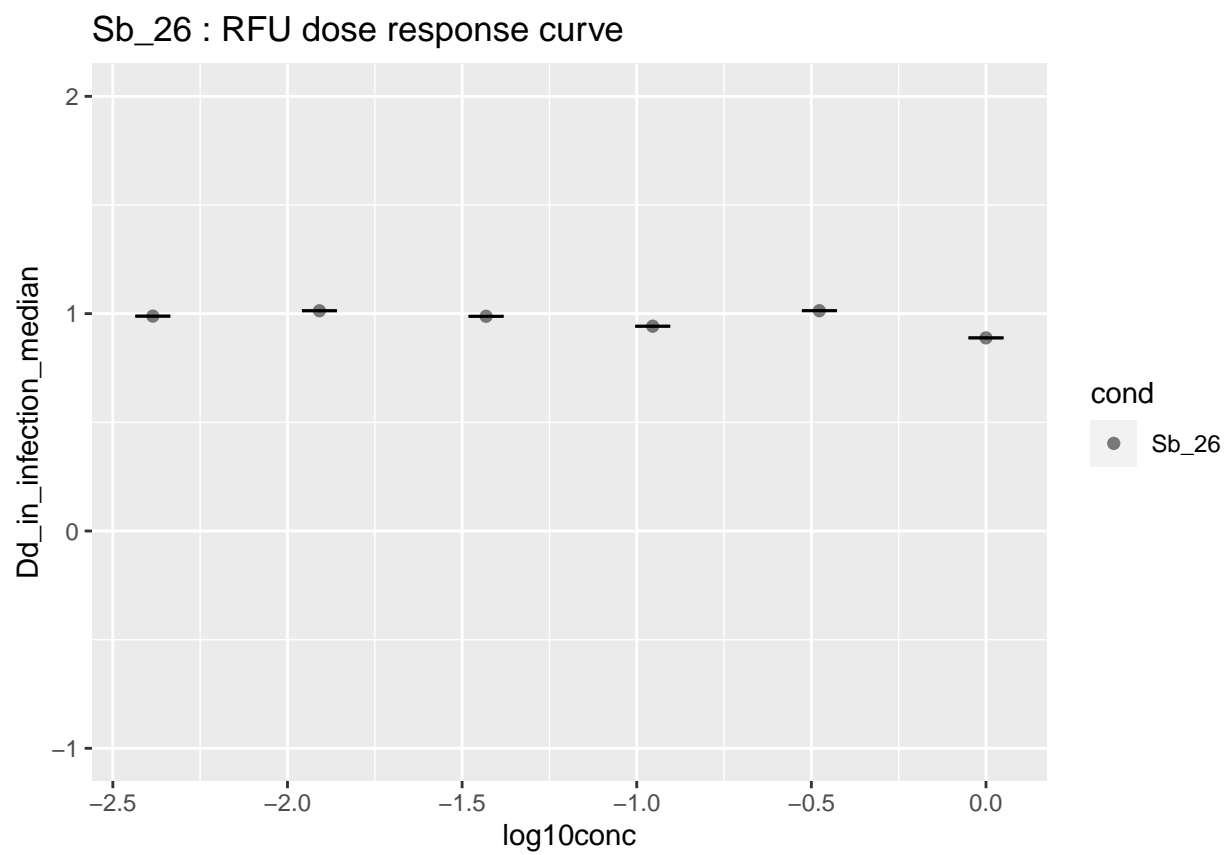


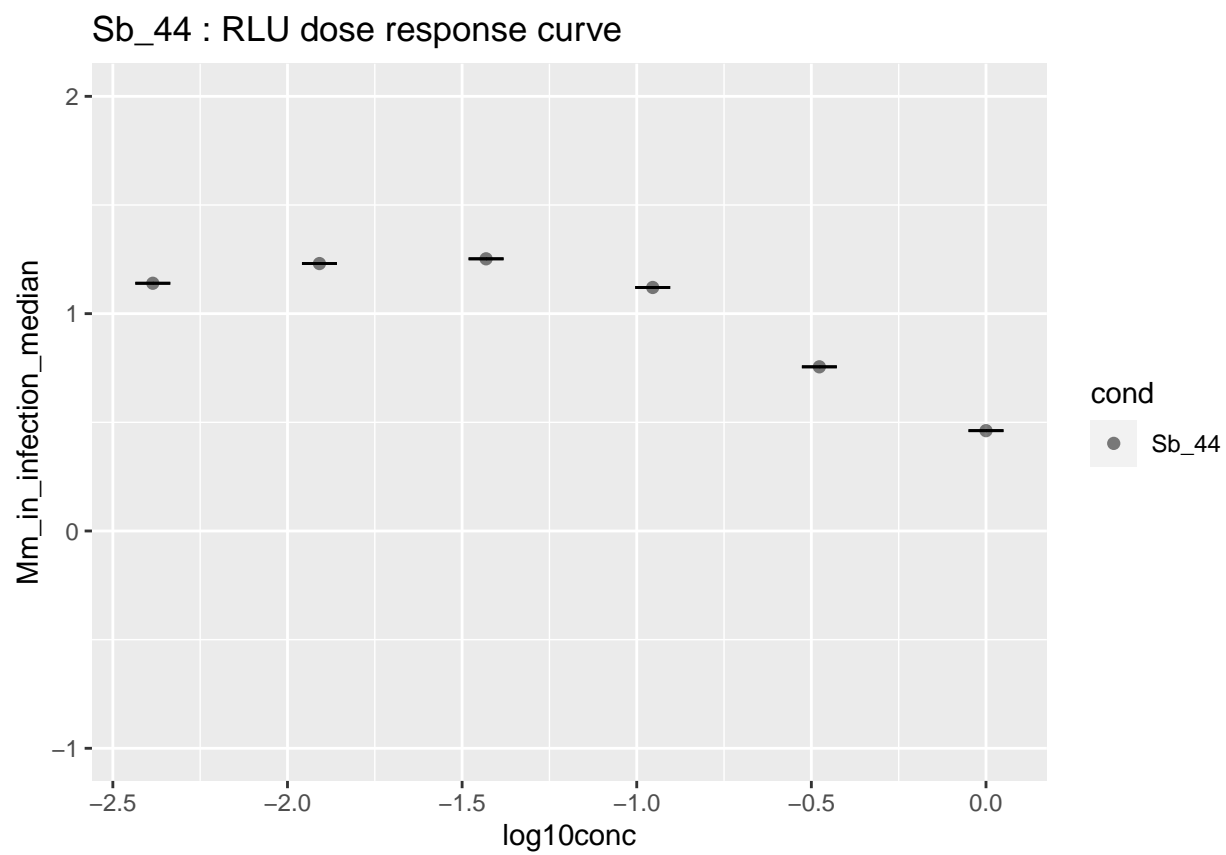


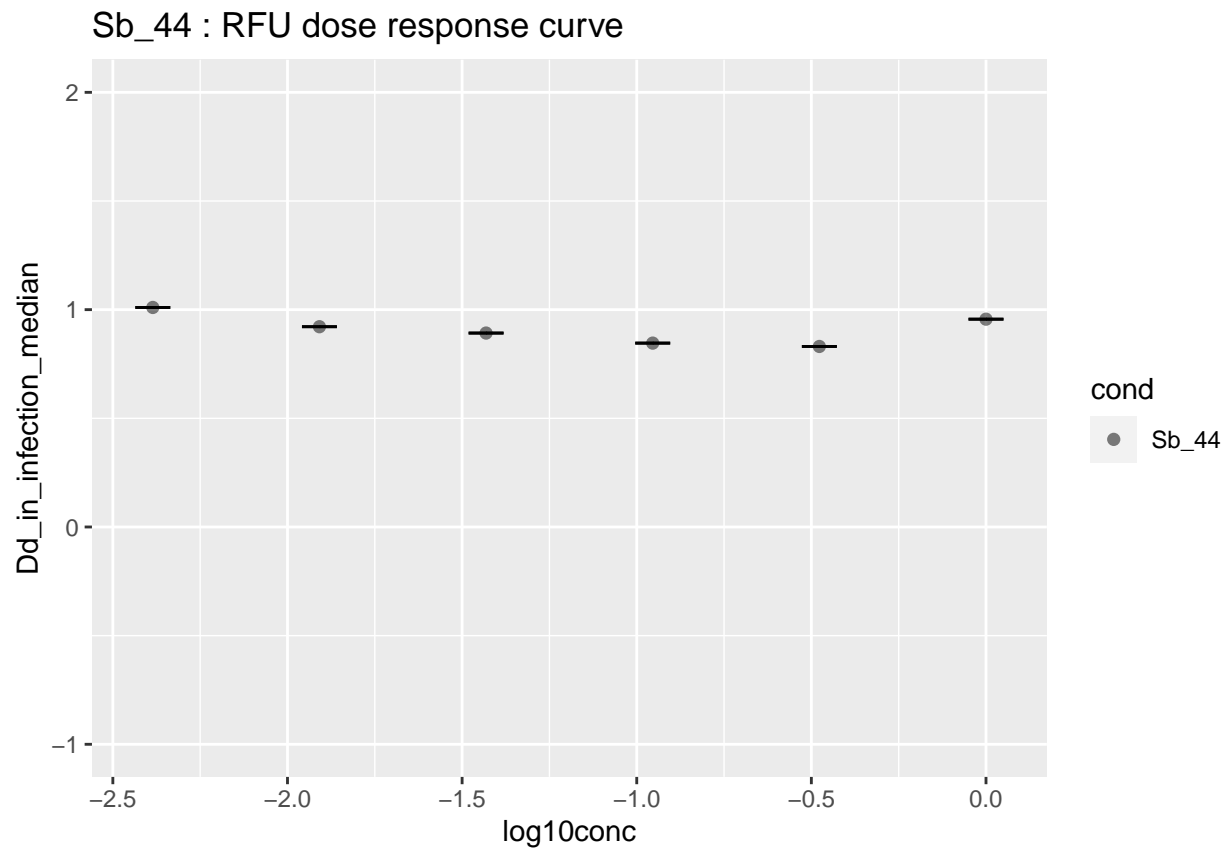


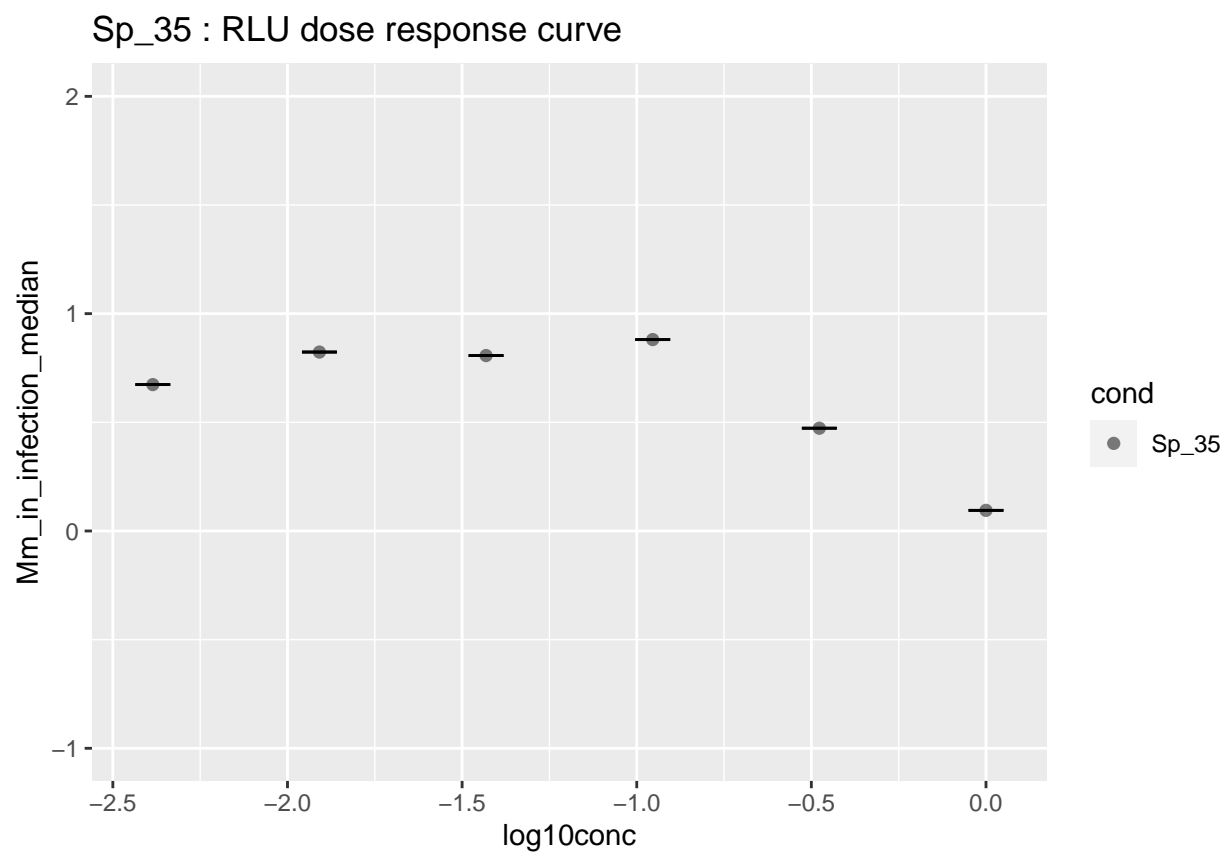


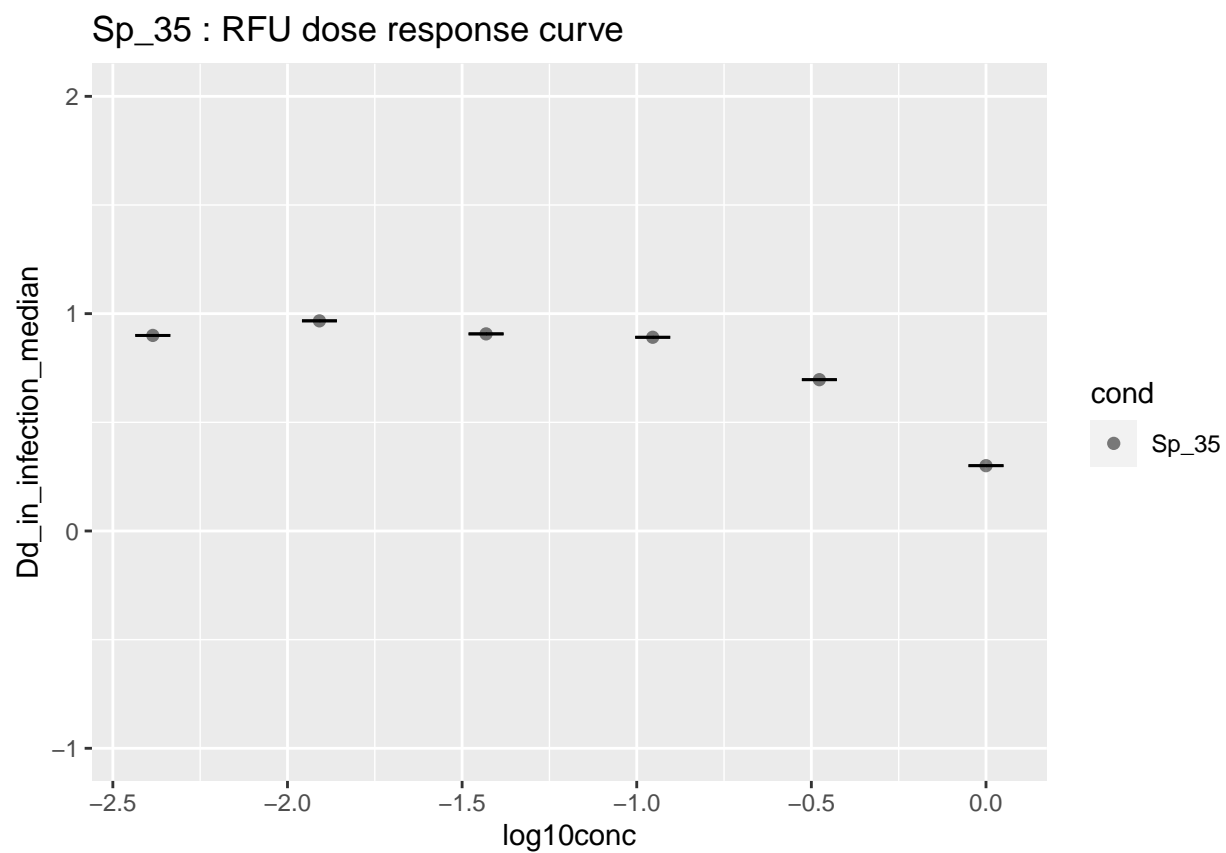


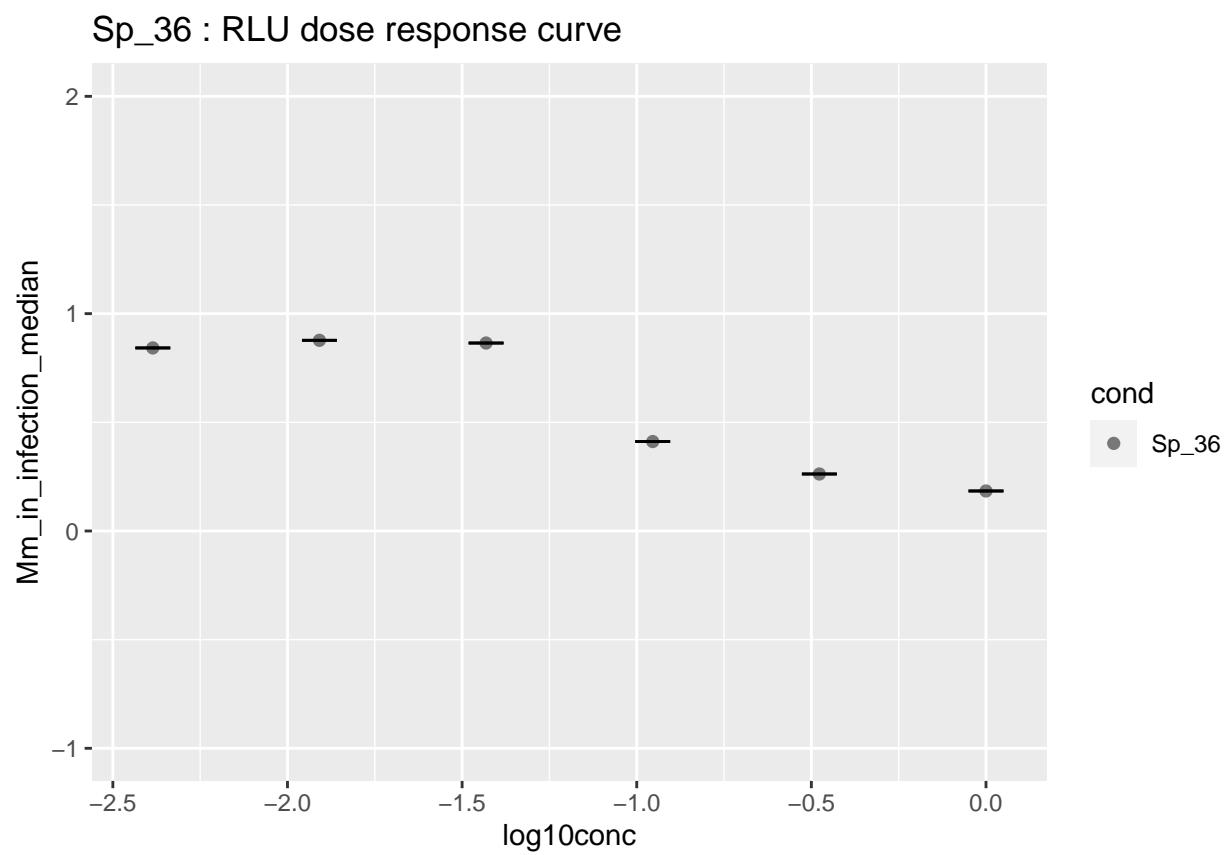


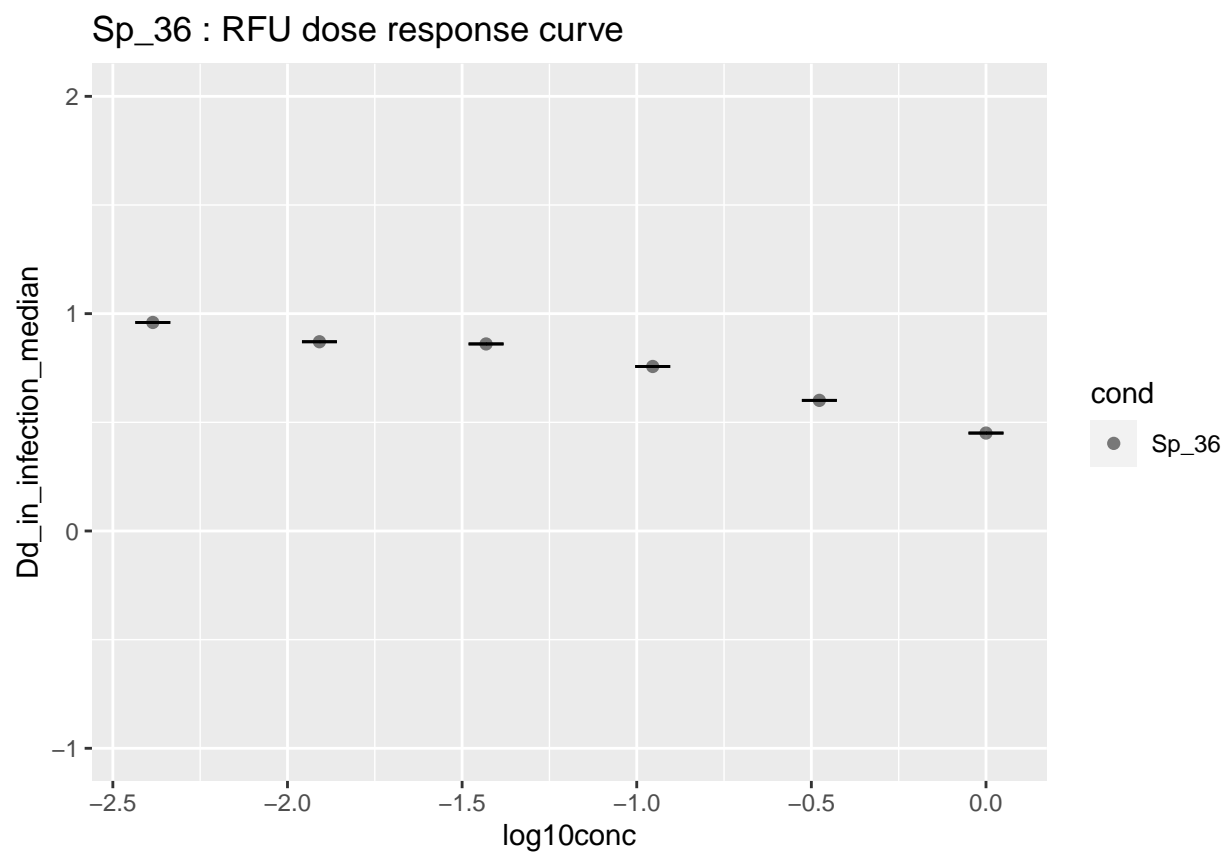


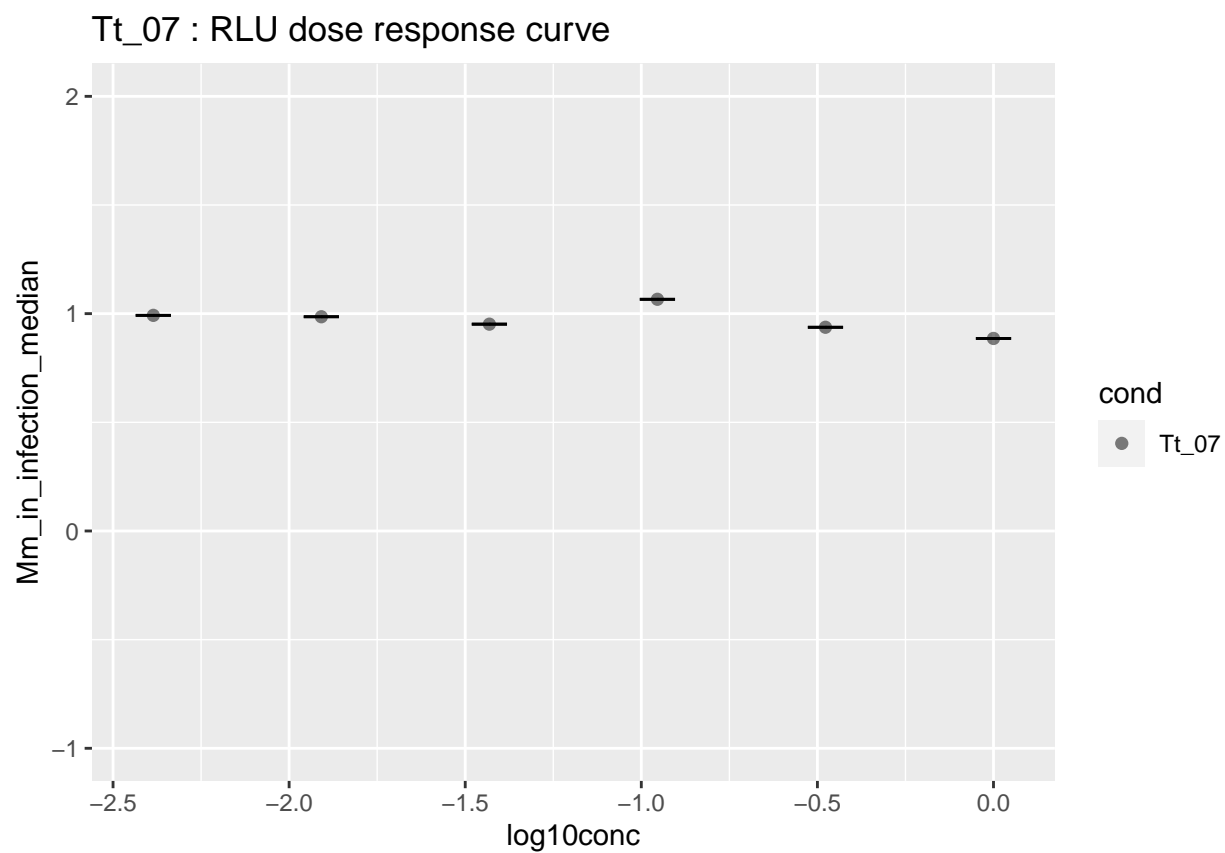


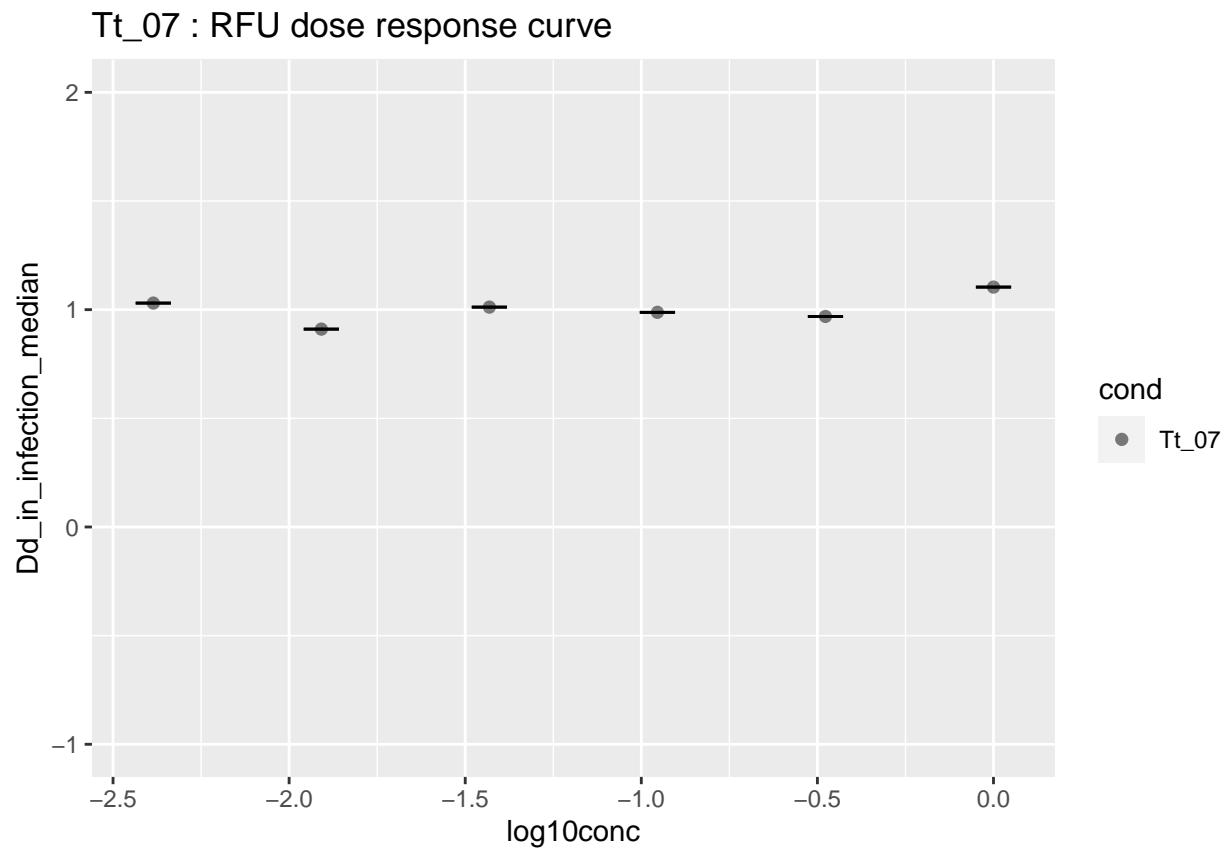




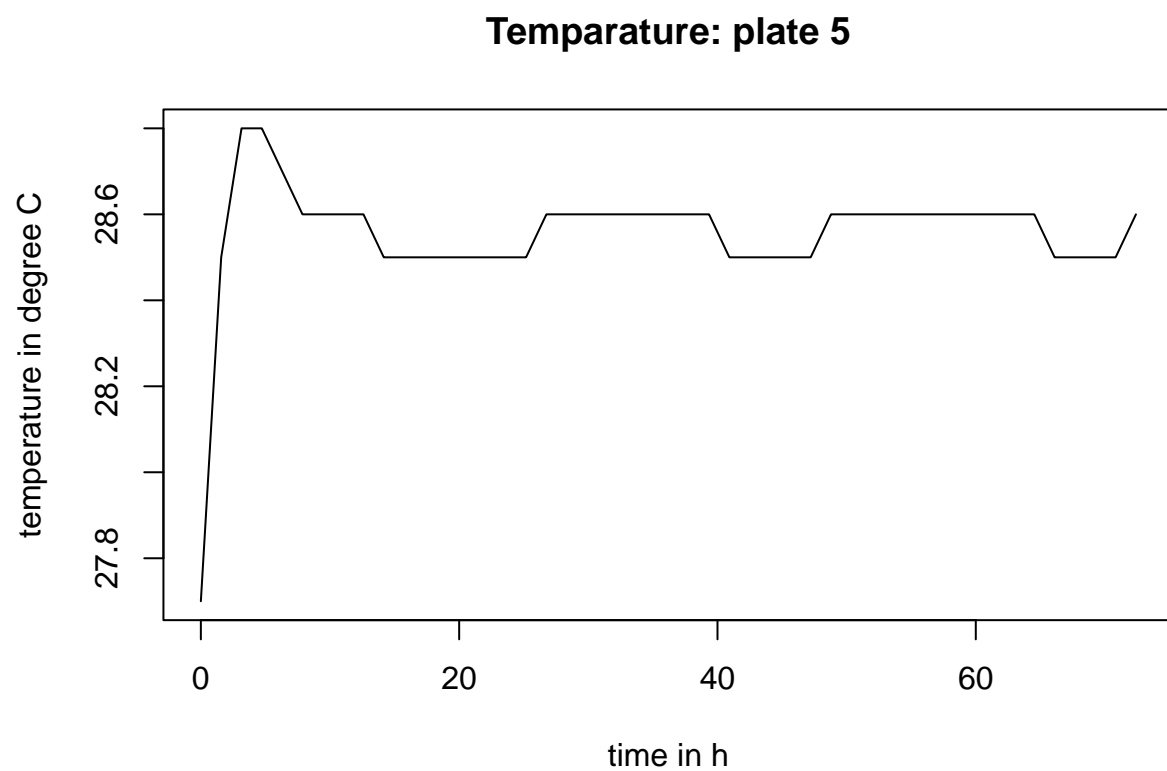




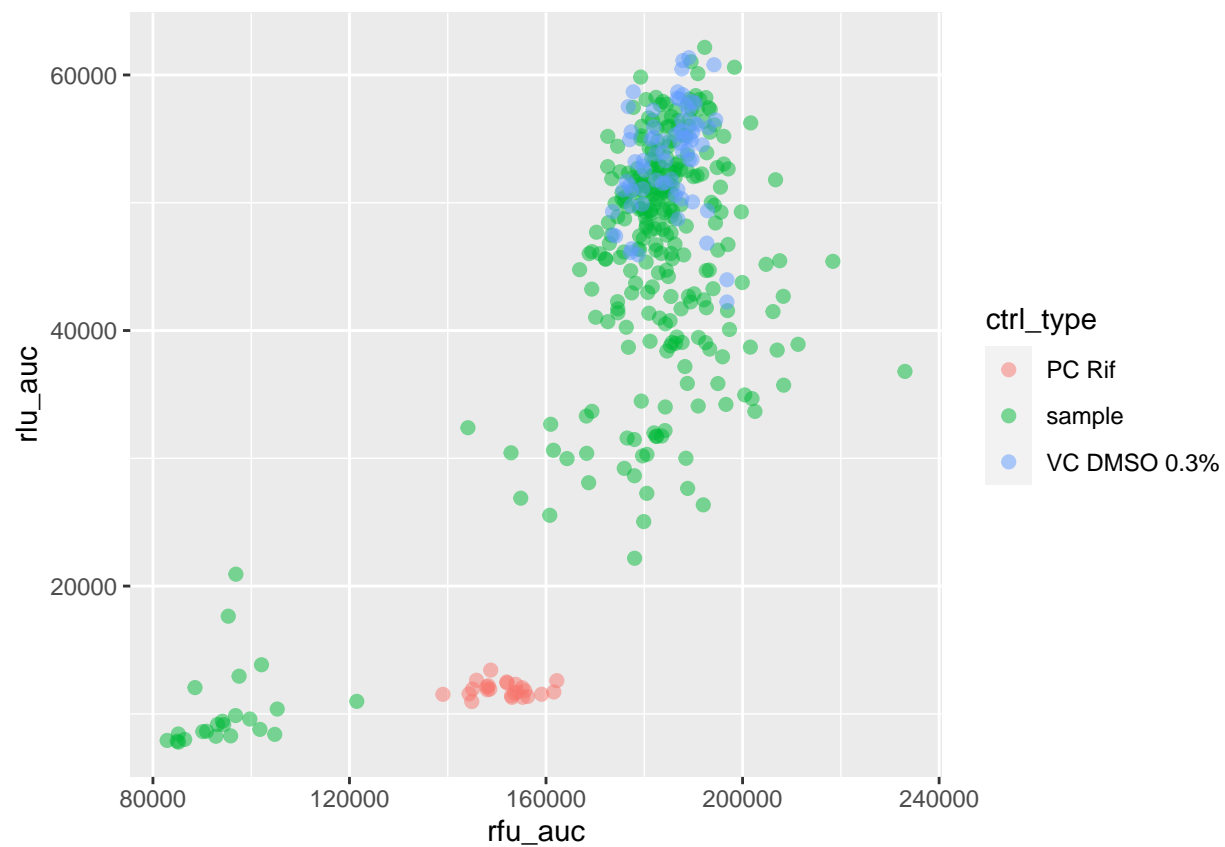


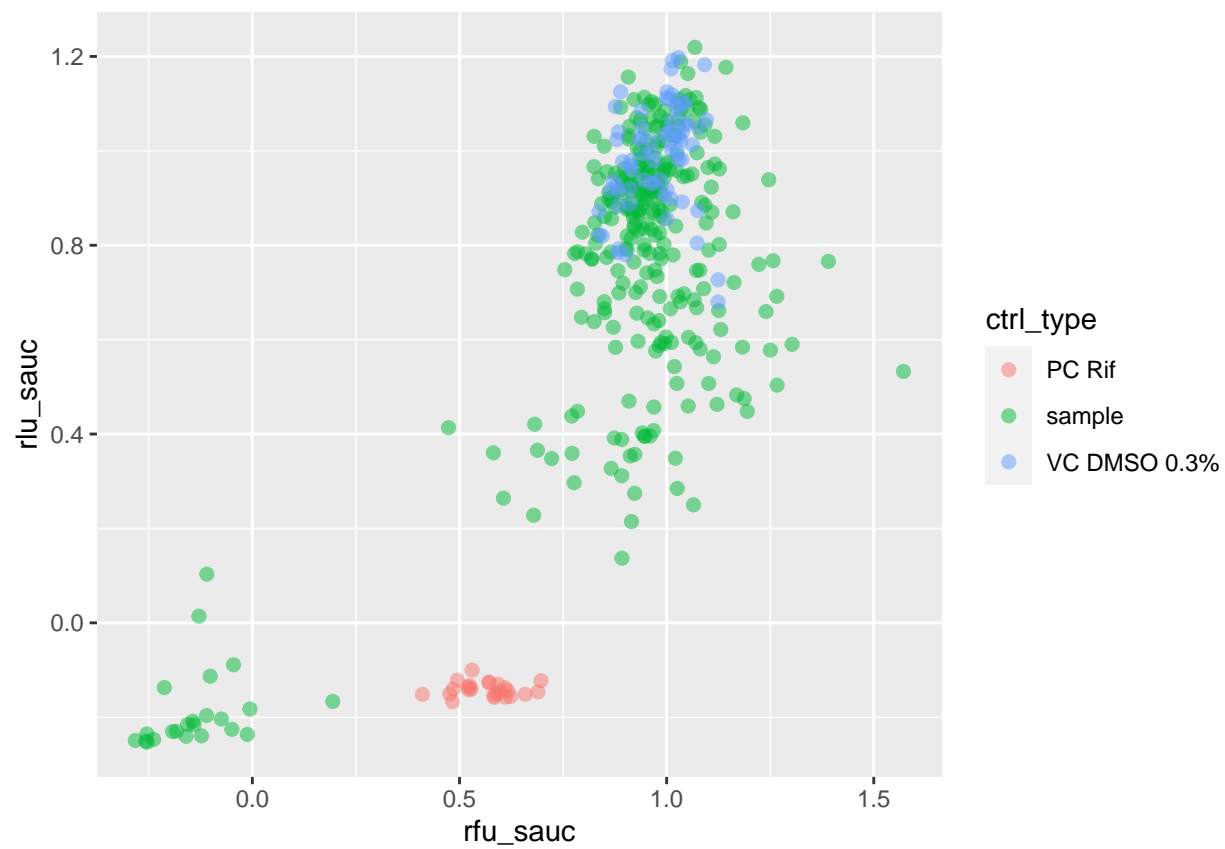


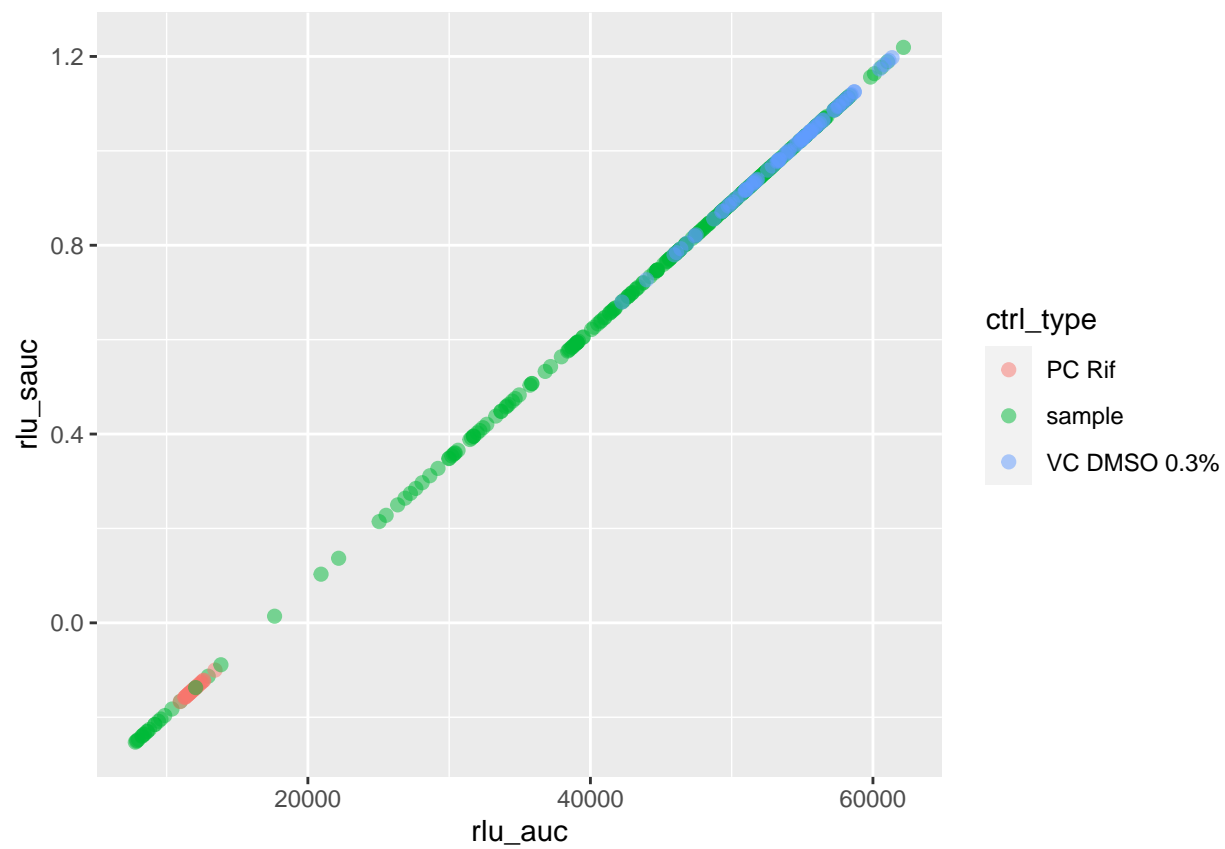
```
## [1] "analysis for plate06.xlsx"
```

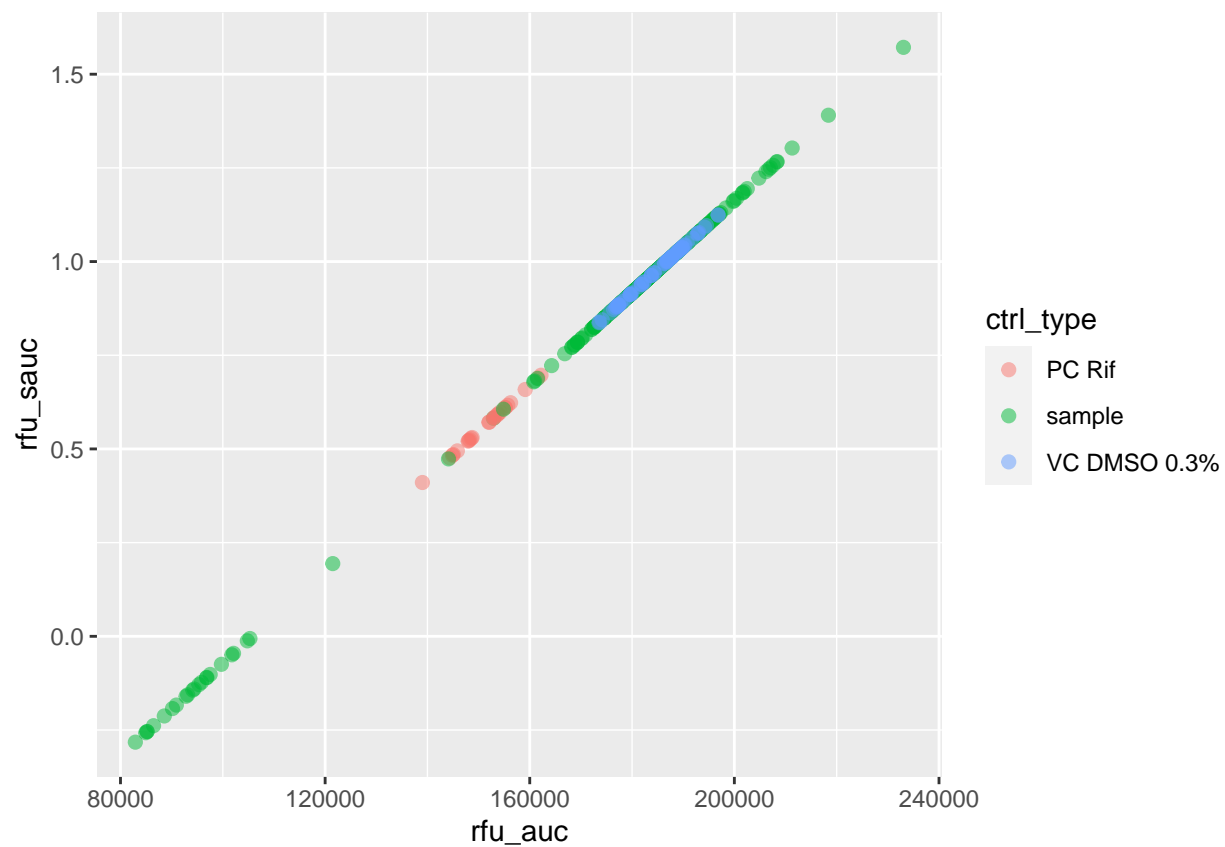



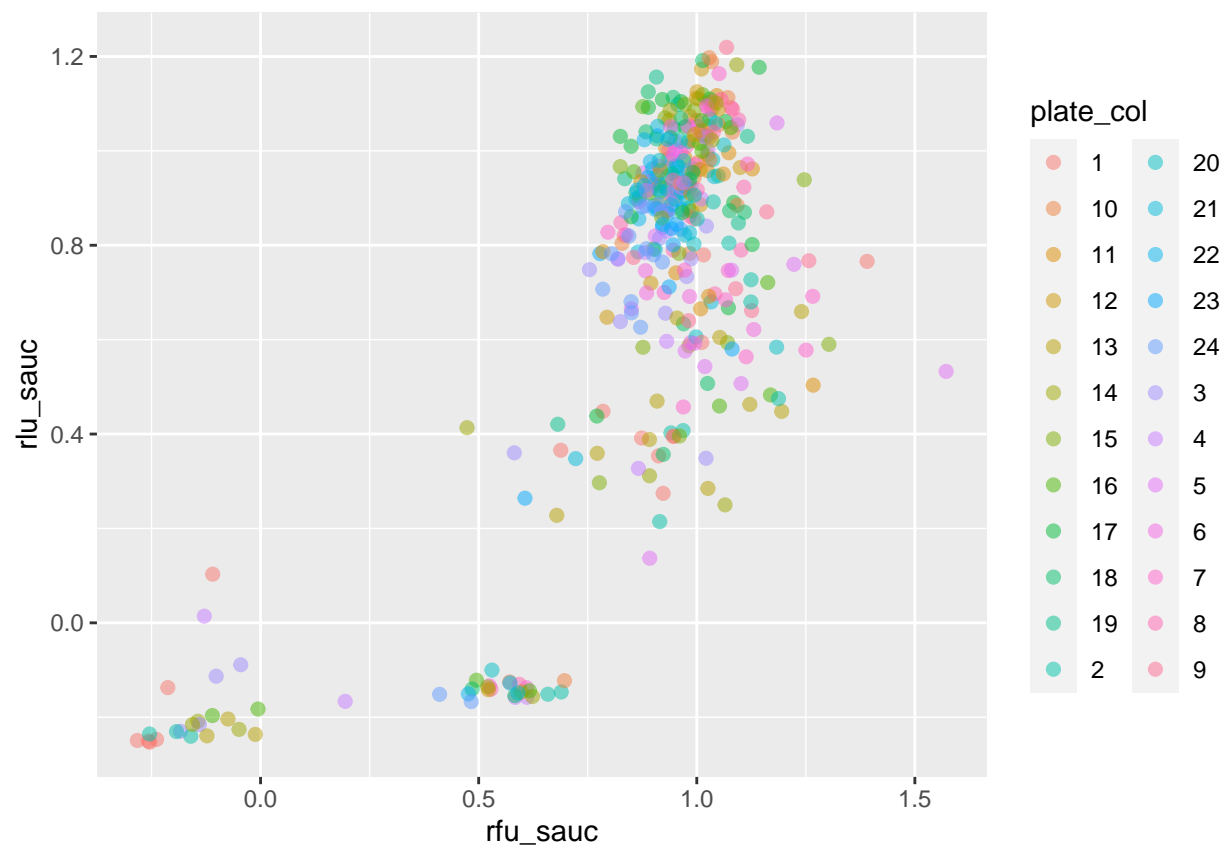
```
## [1] "VC DMSO 0.3% Robust z'-factor of rlu for plate plate06.xlsx, biorep 3 : "  
## [1] 0.69
```

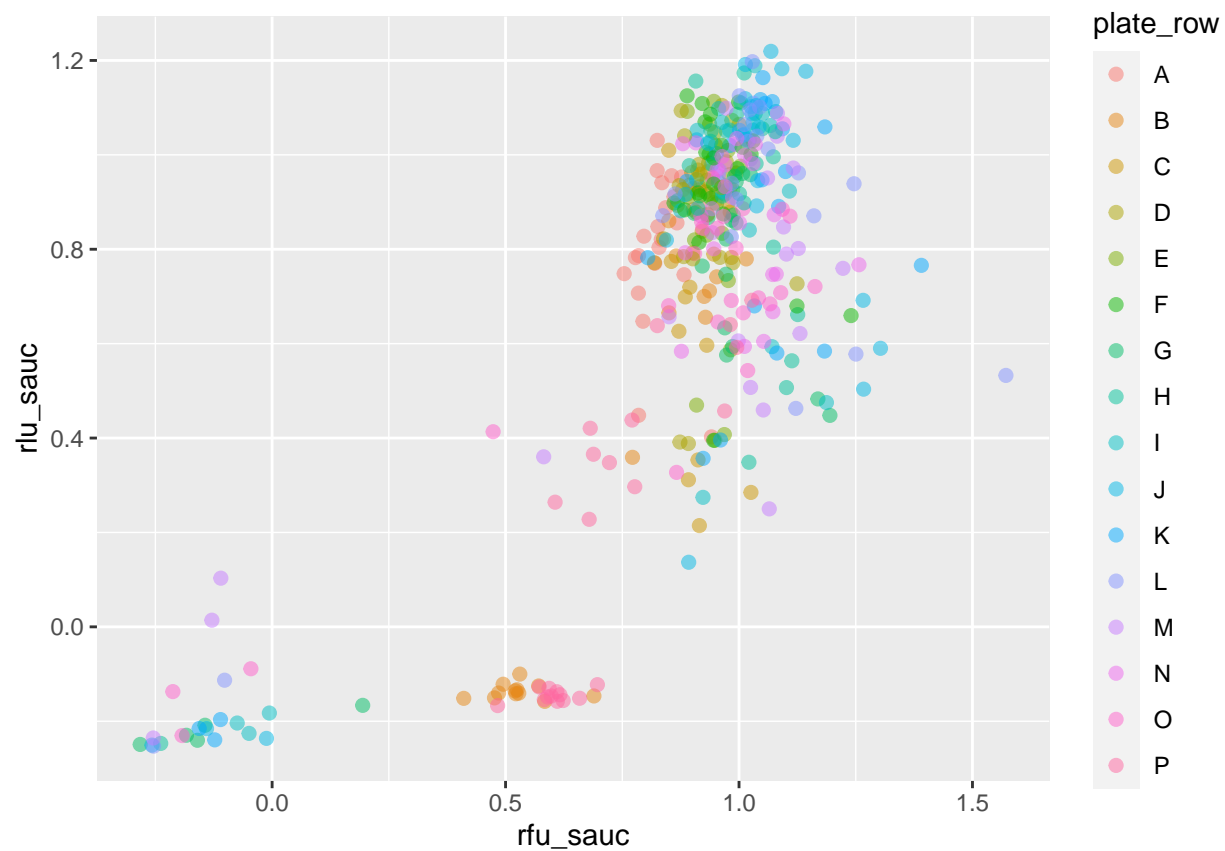




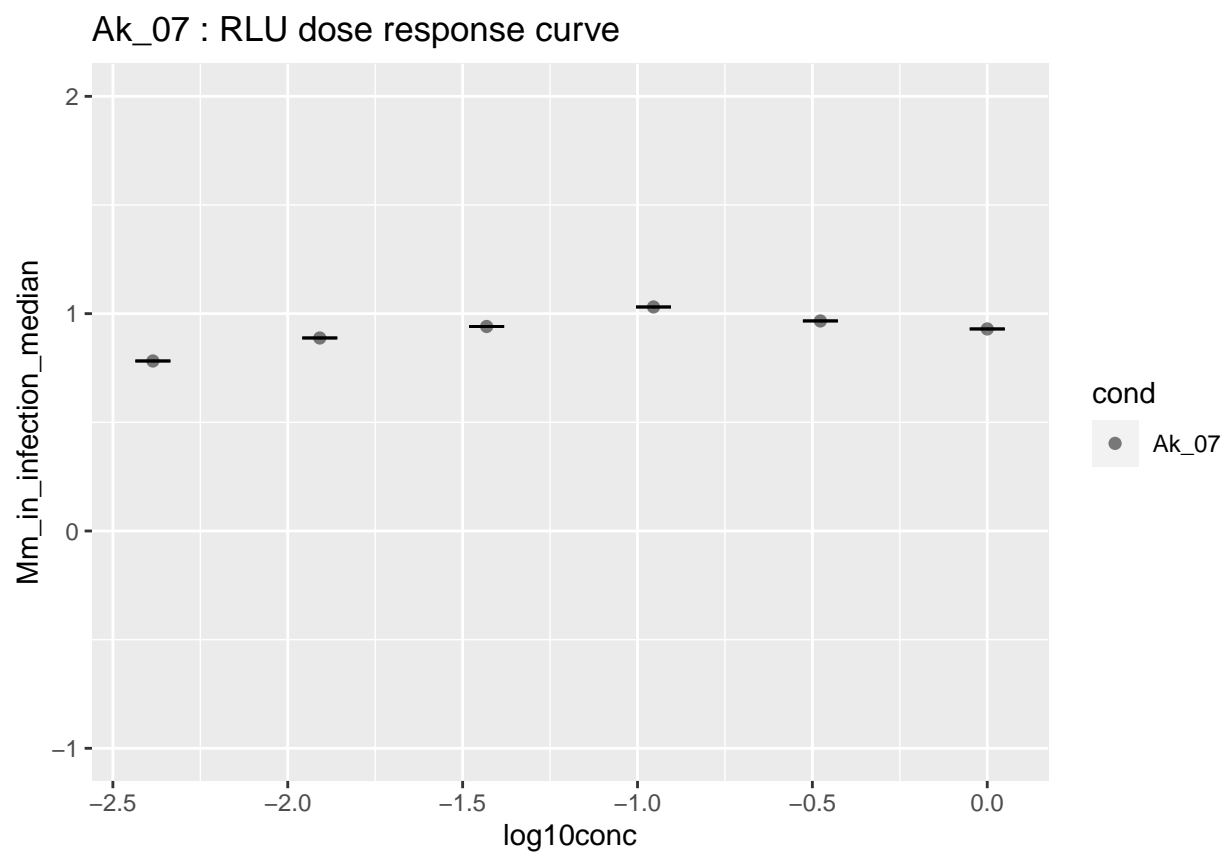


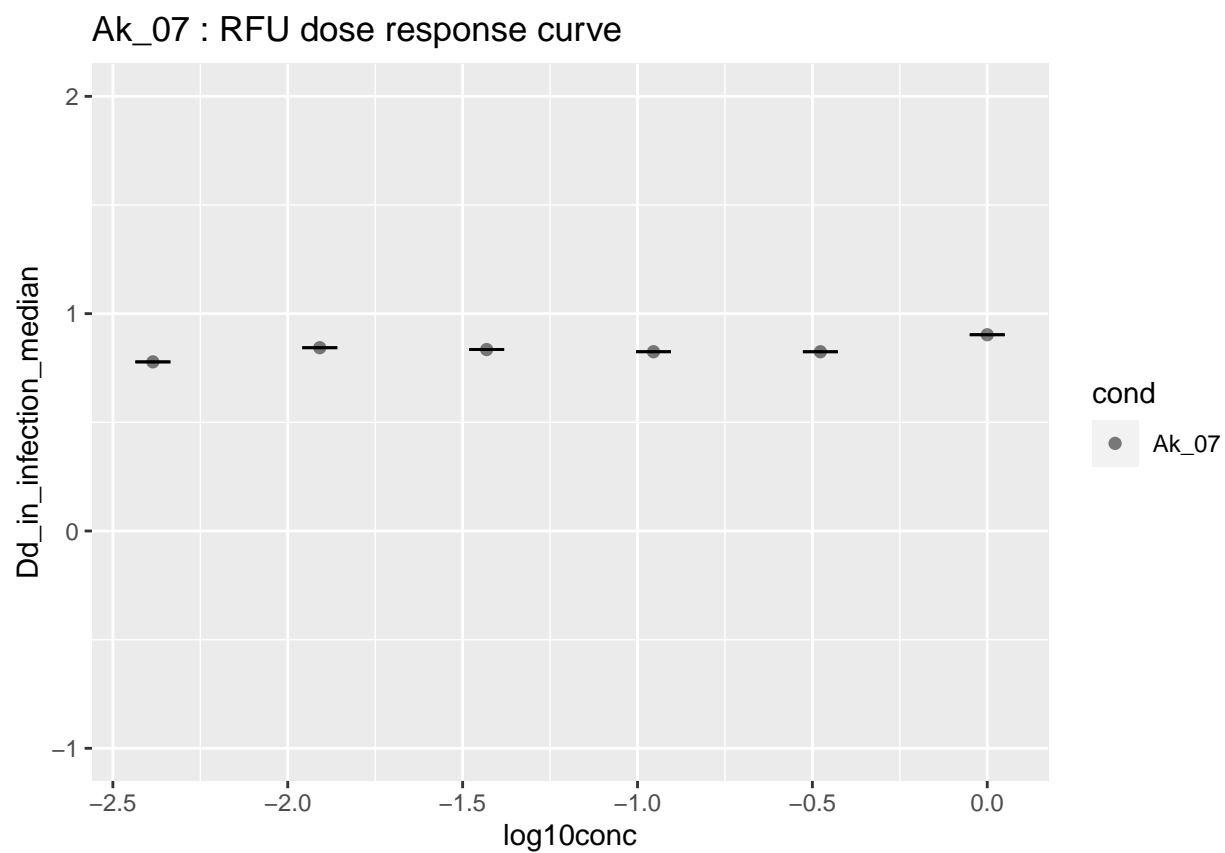


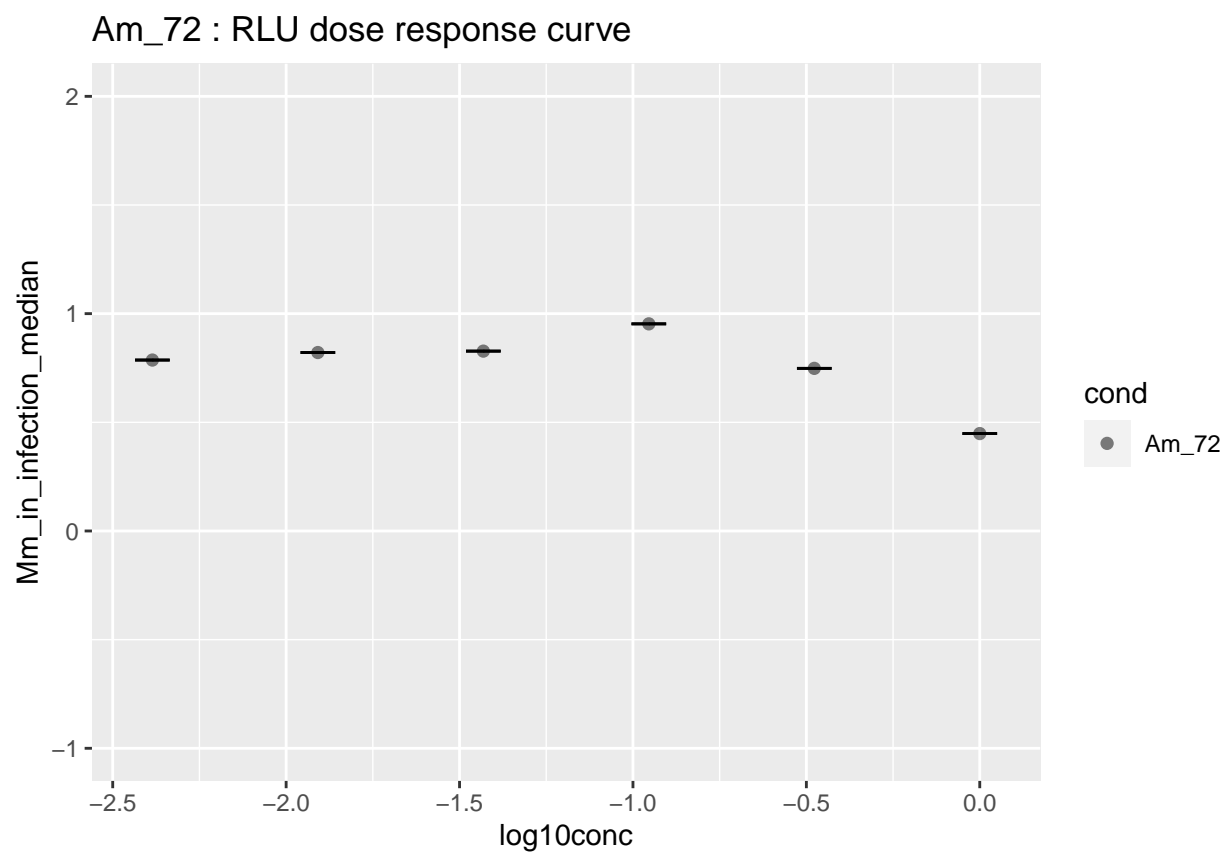


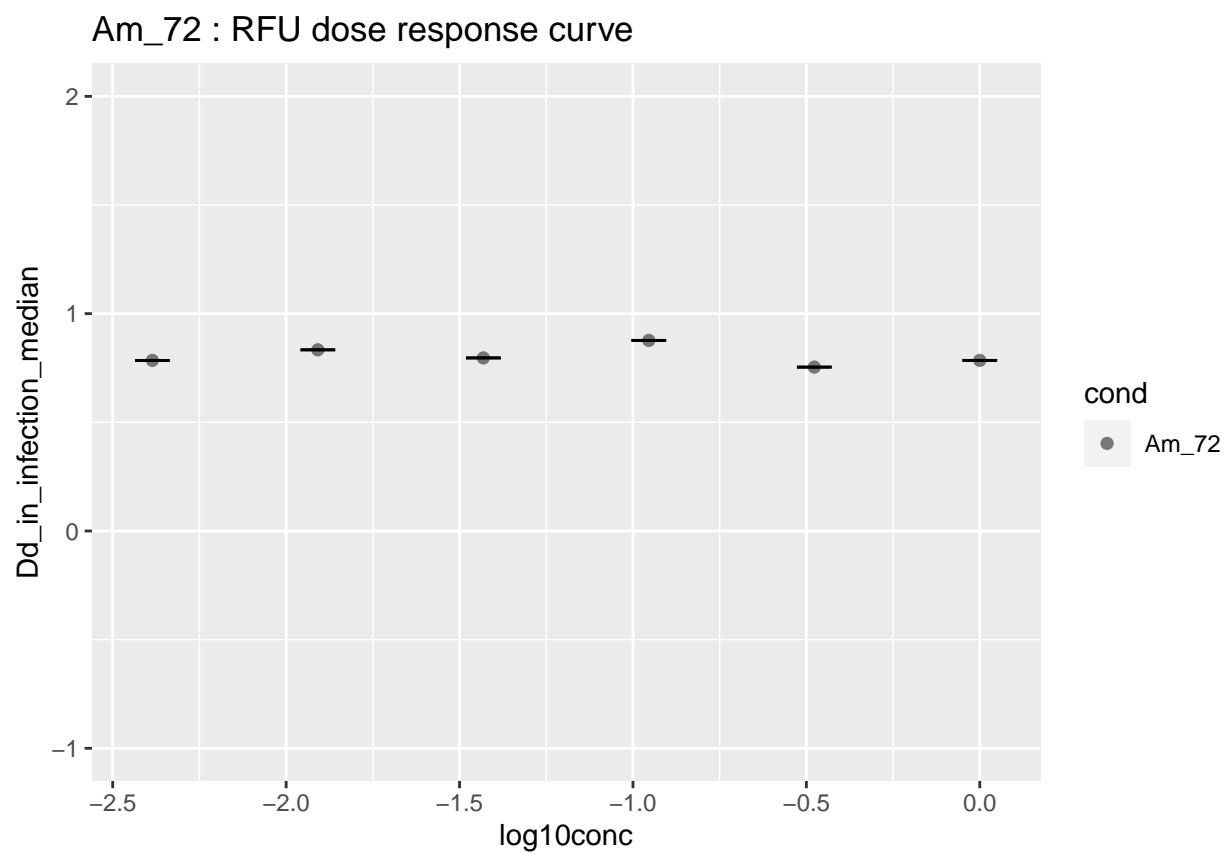


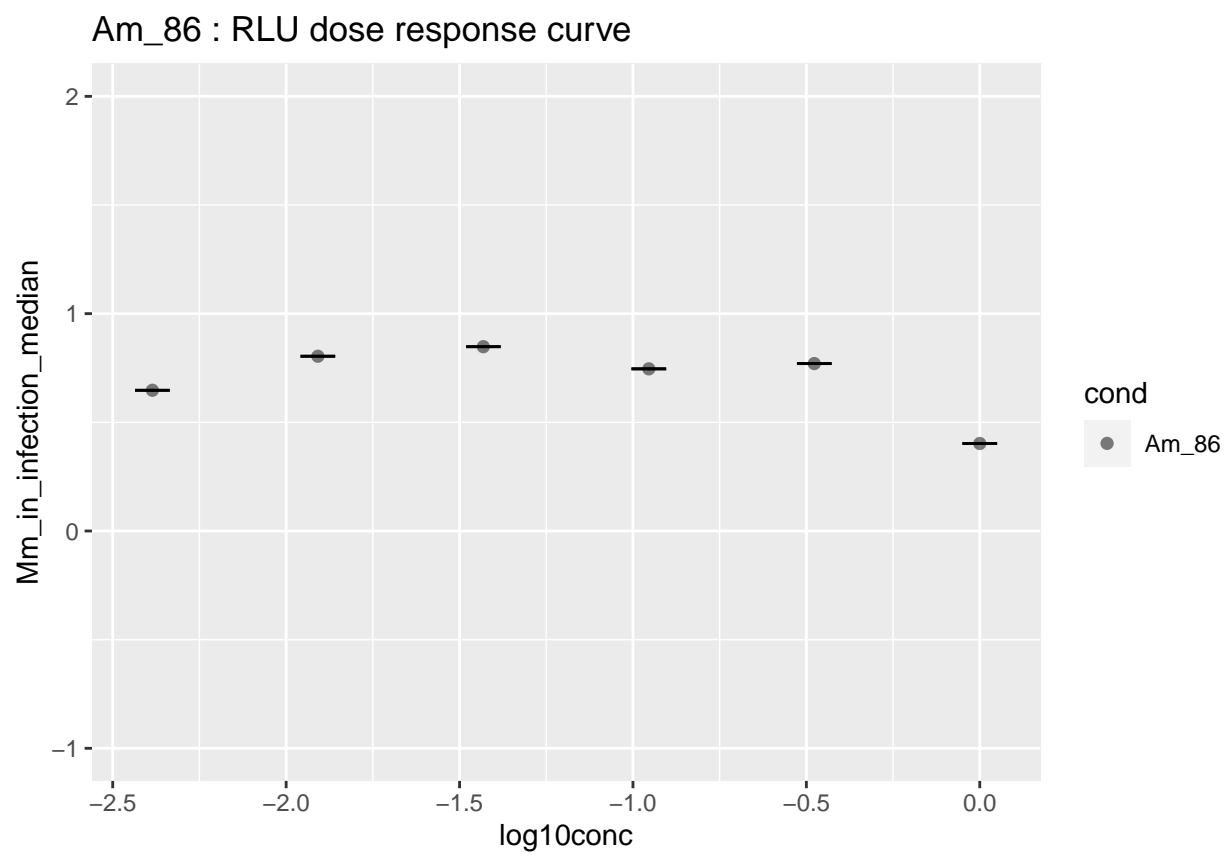
```
## [1] "Dose response curves over all bioreps within this plate"
```

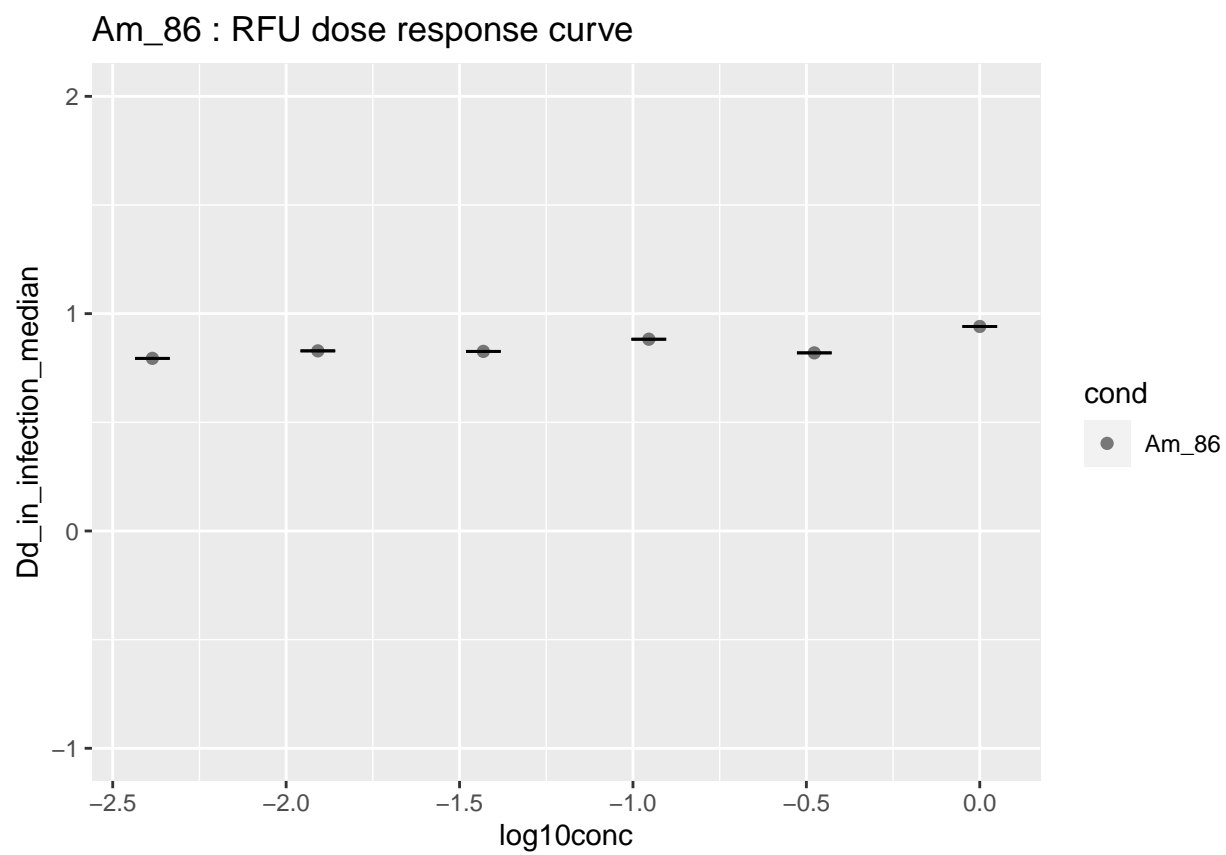




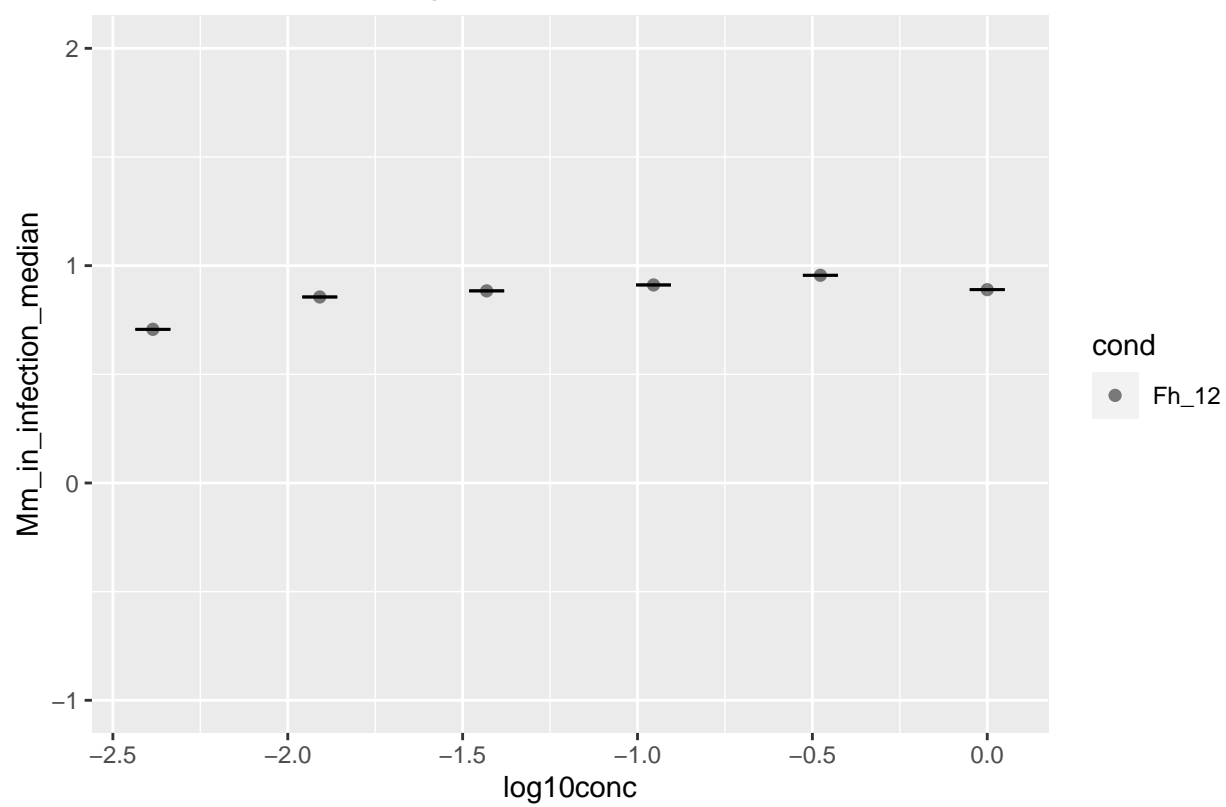




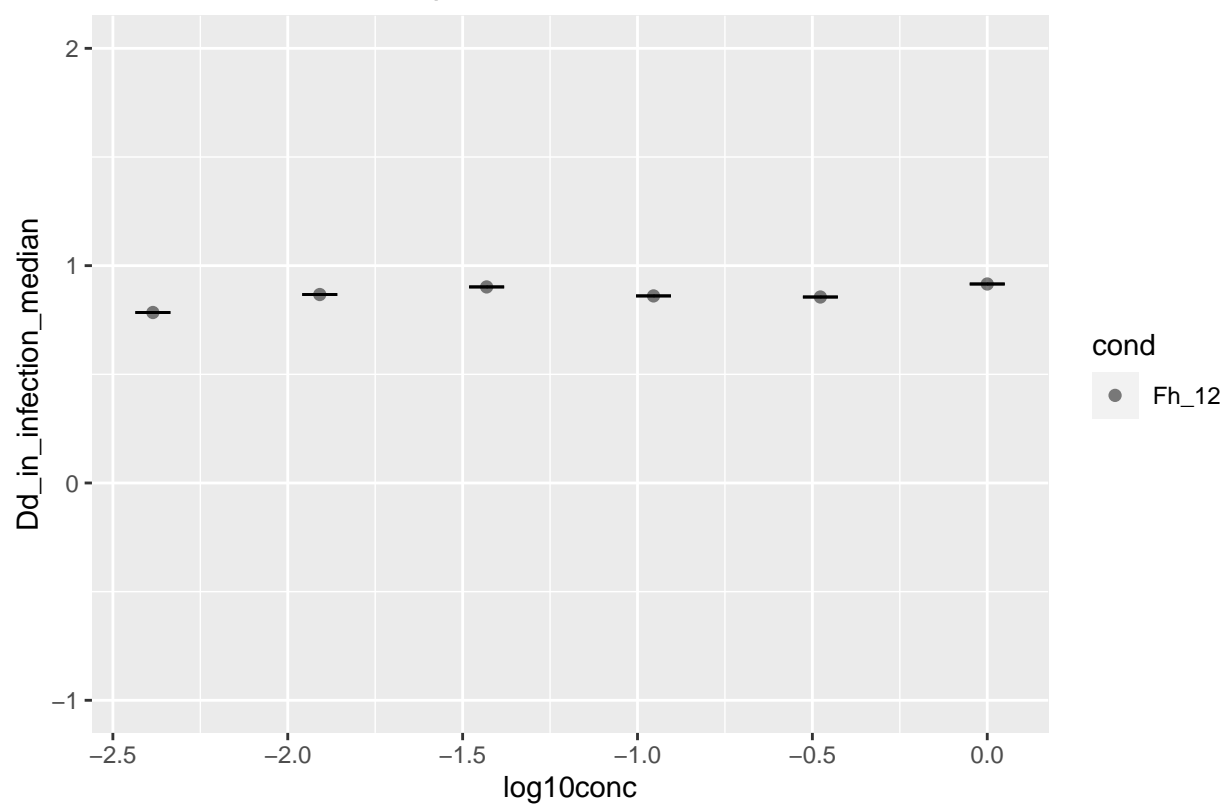


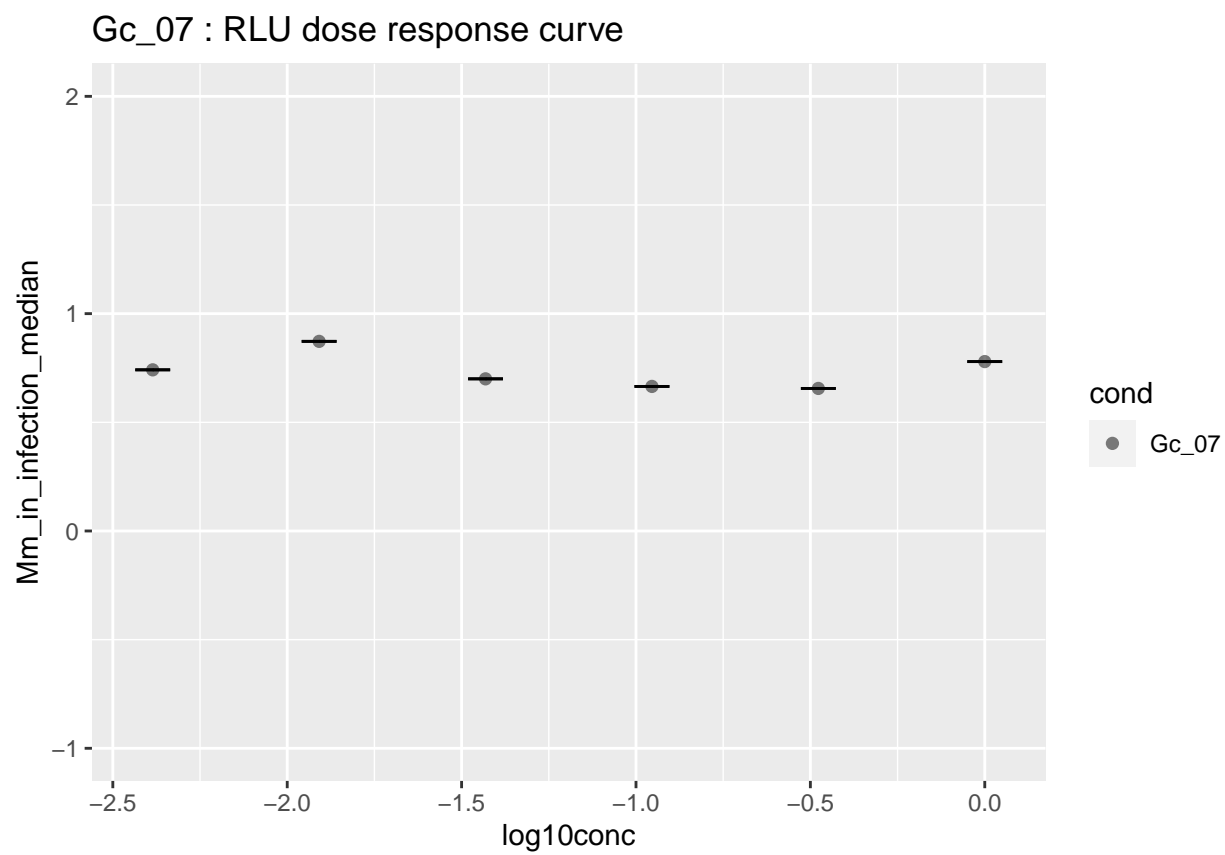


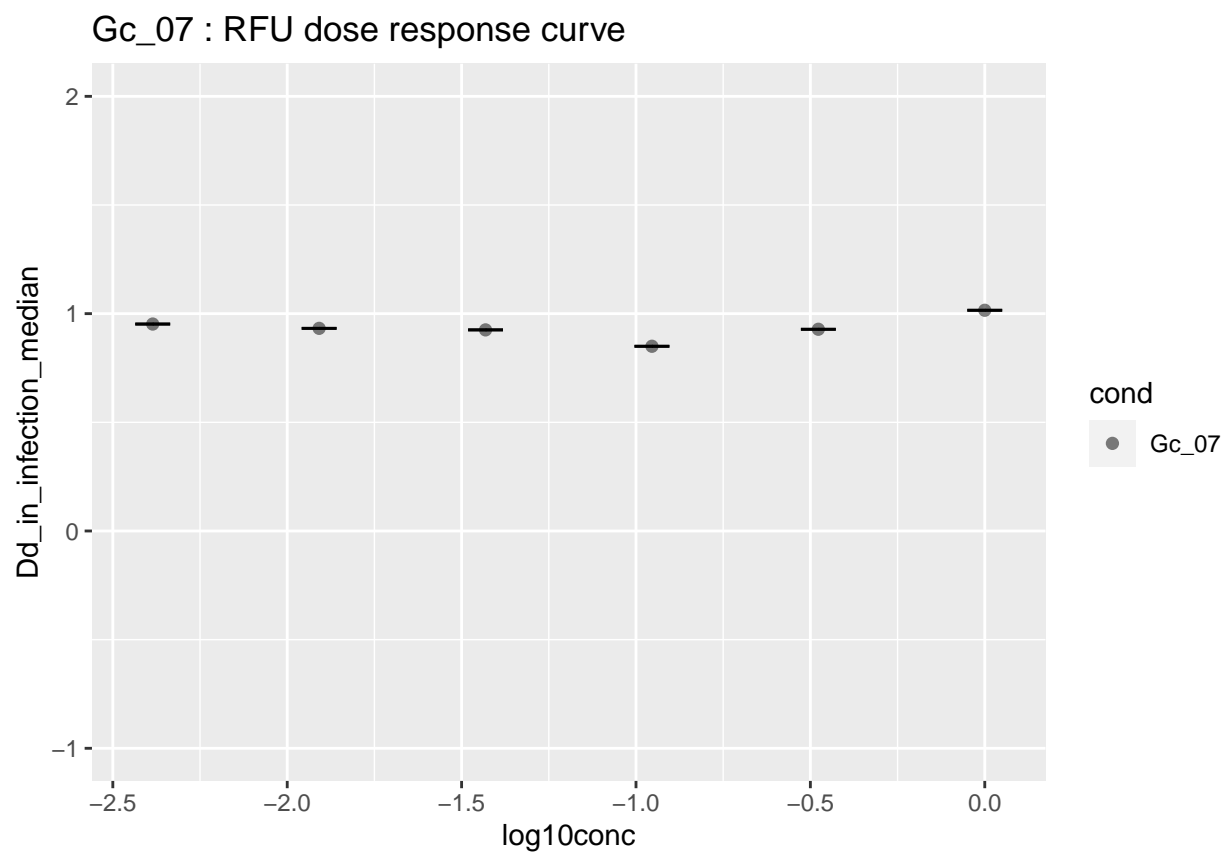
Fh_12 : RLU dose response curve

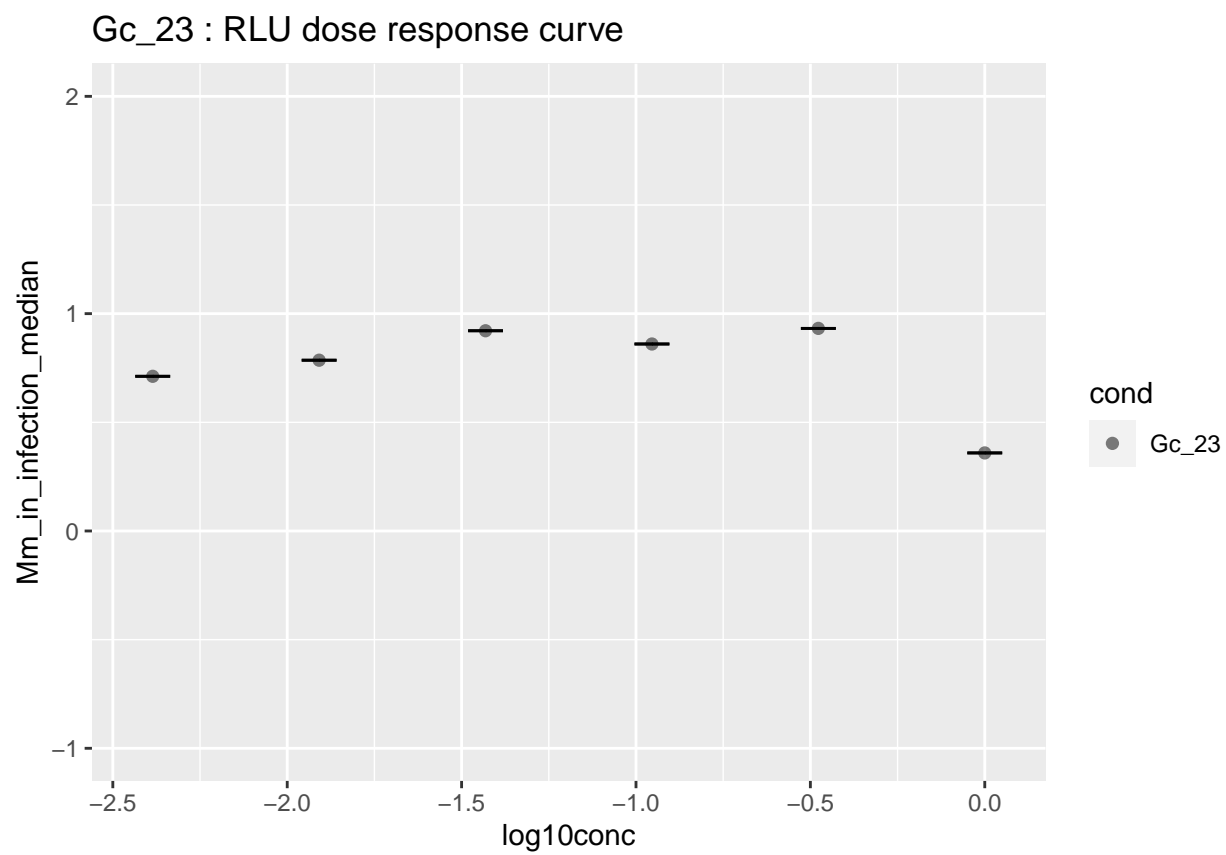


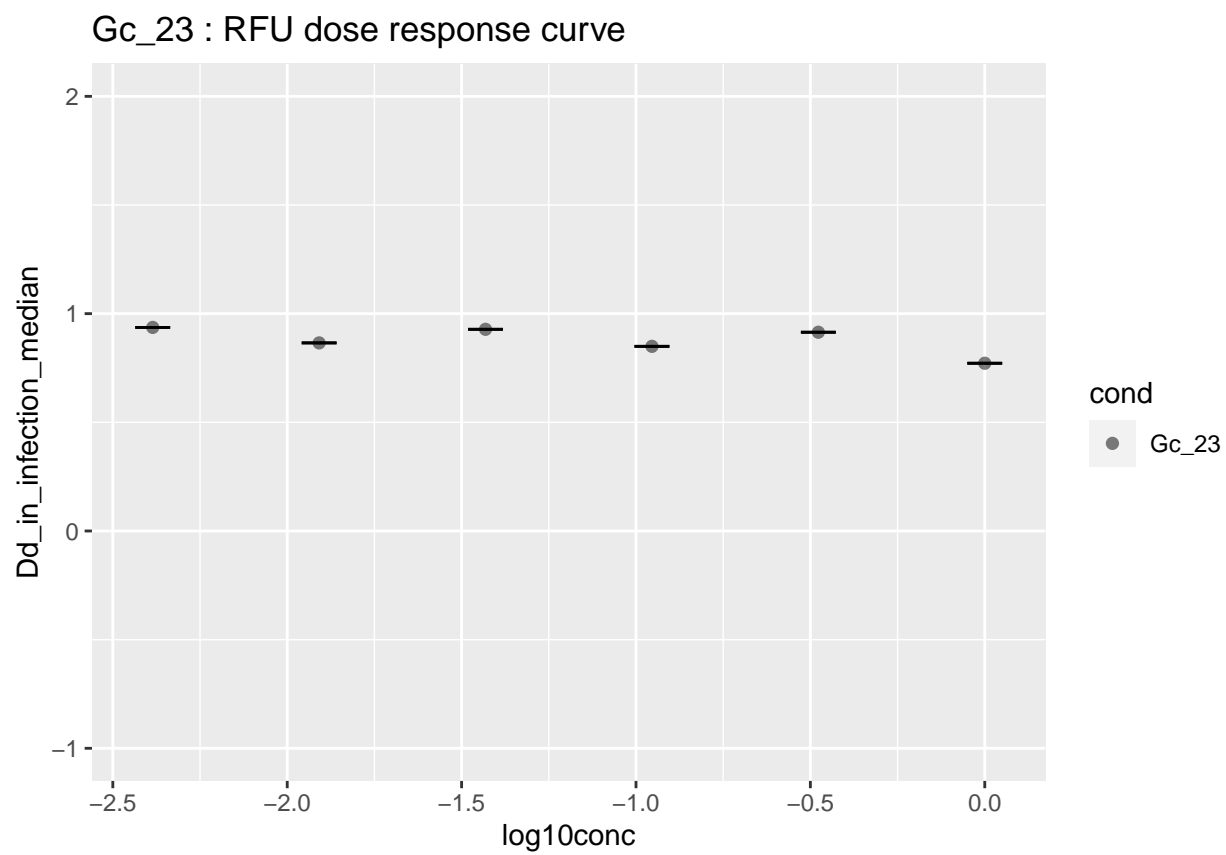
Fh_12 : RFU dose response curve

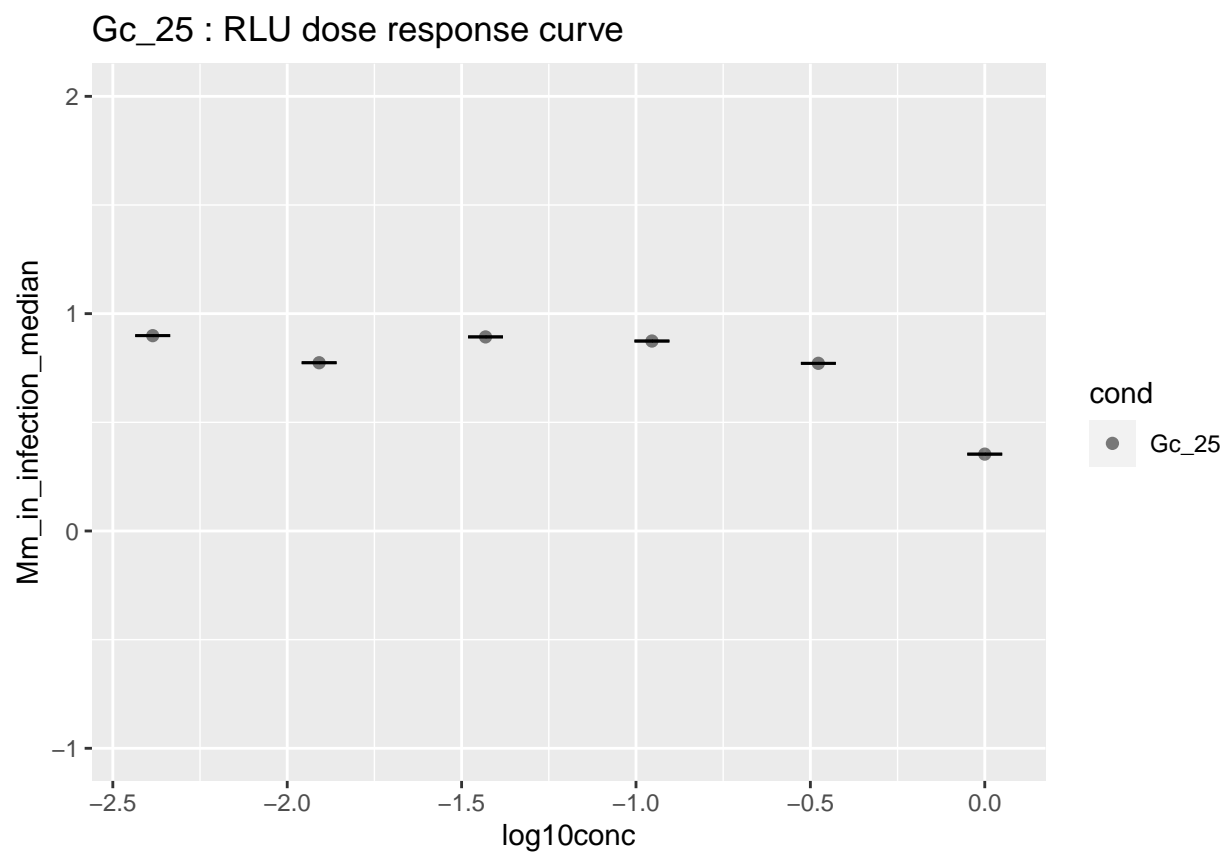


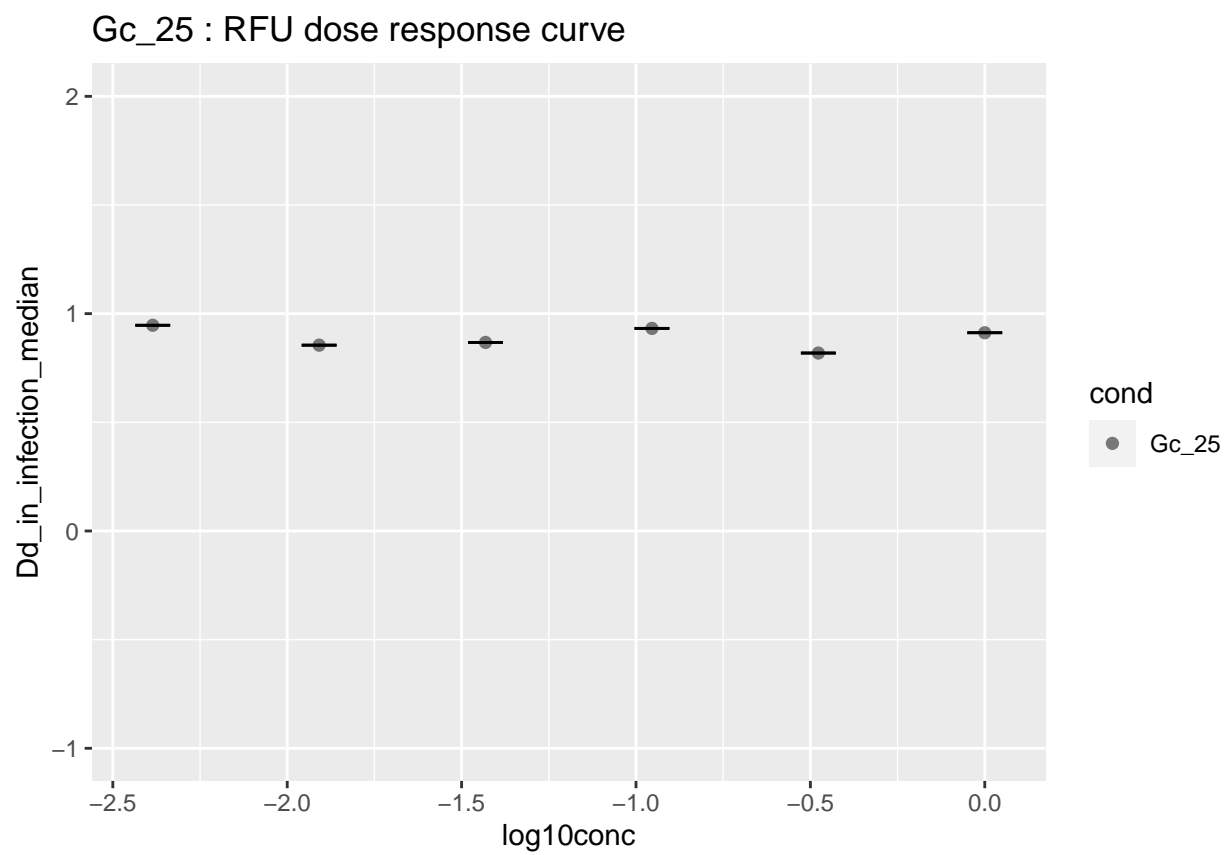


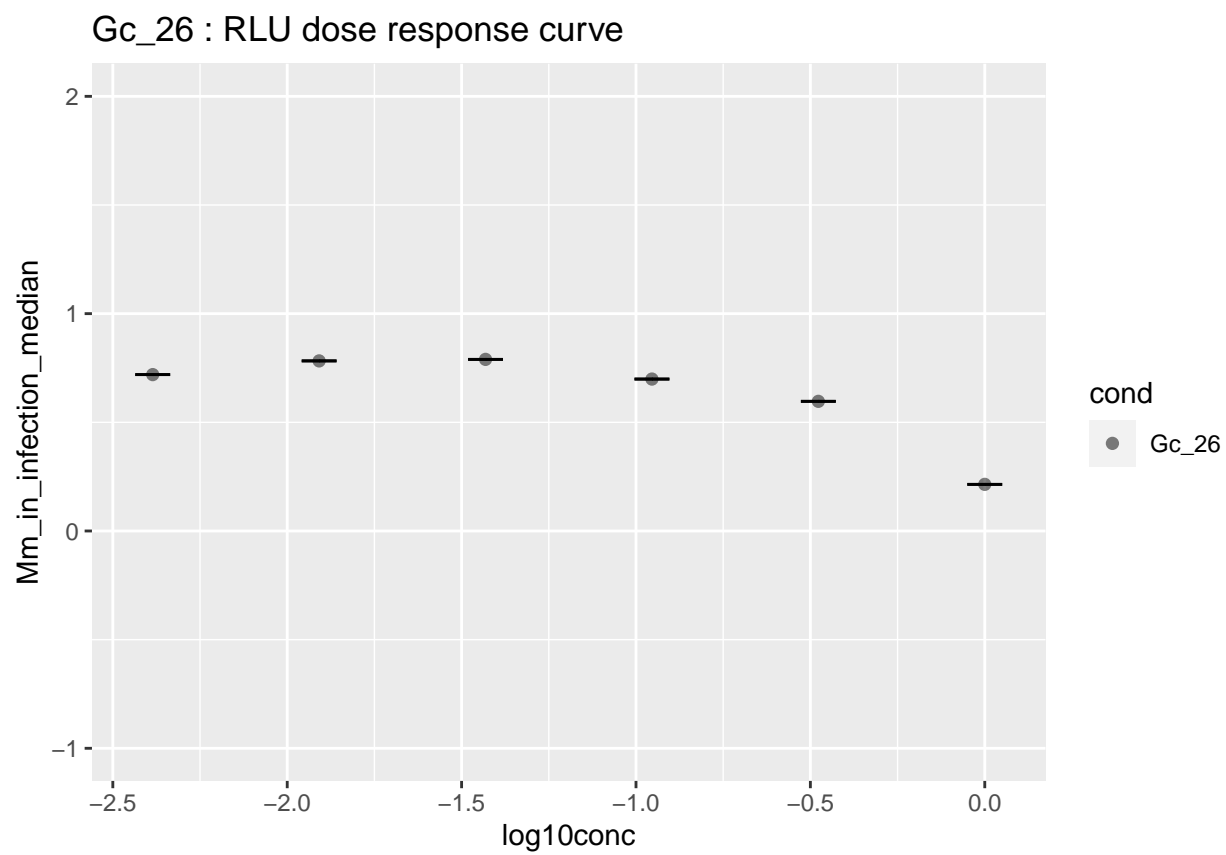


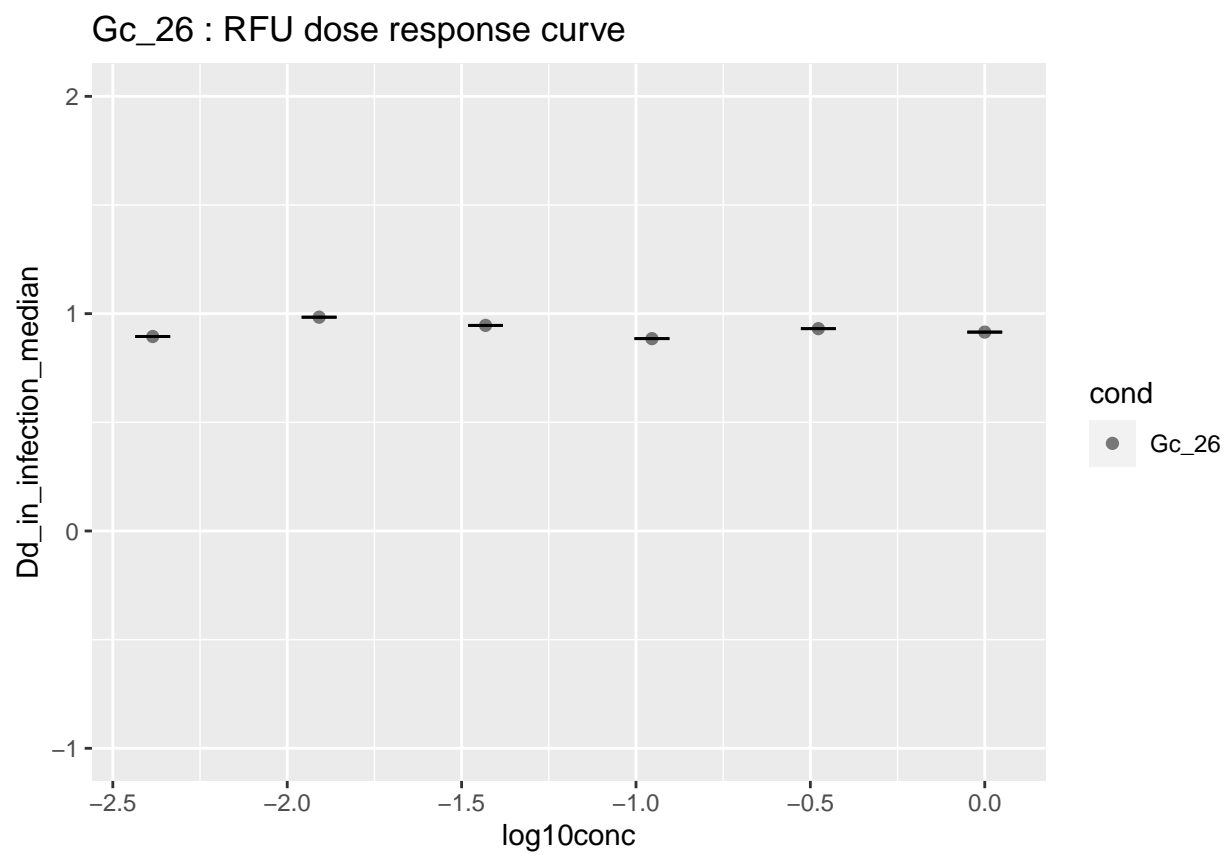


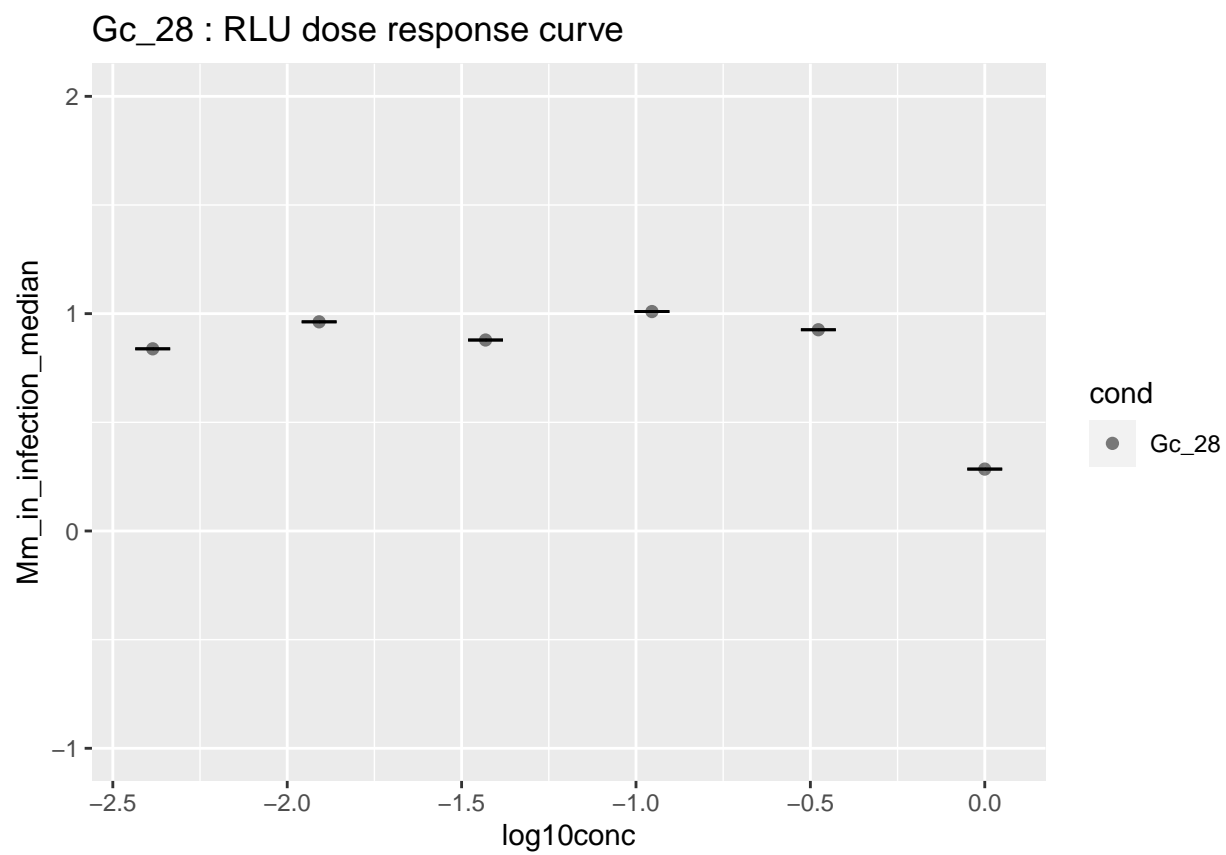


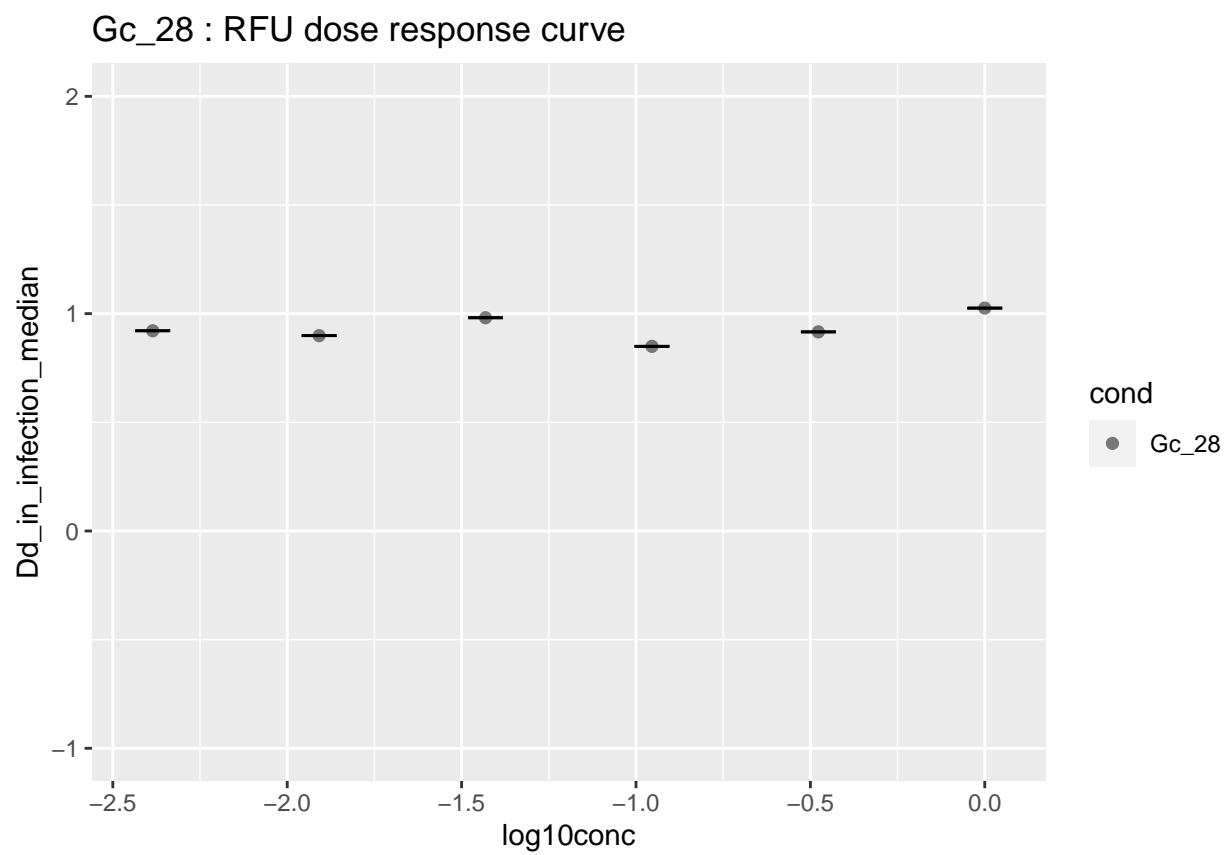


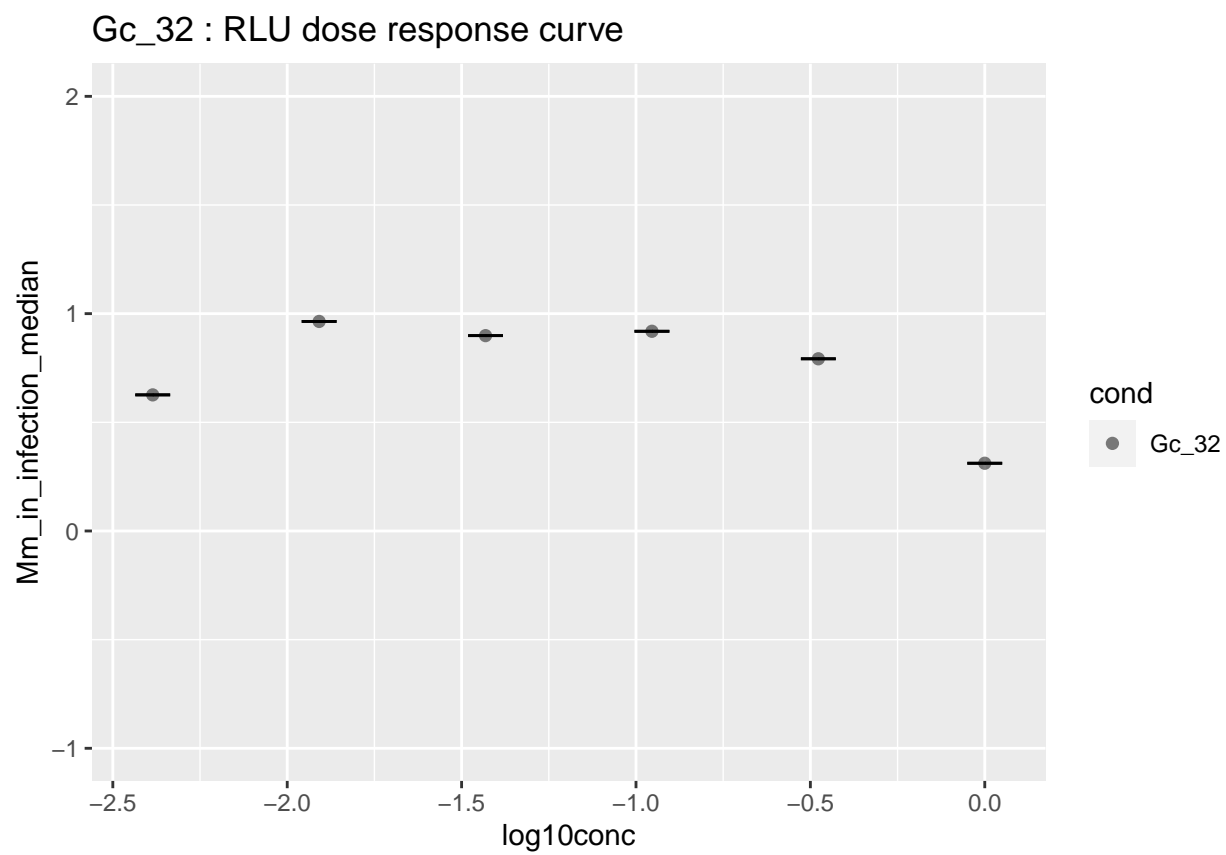


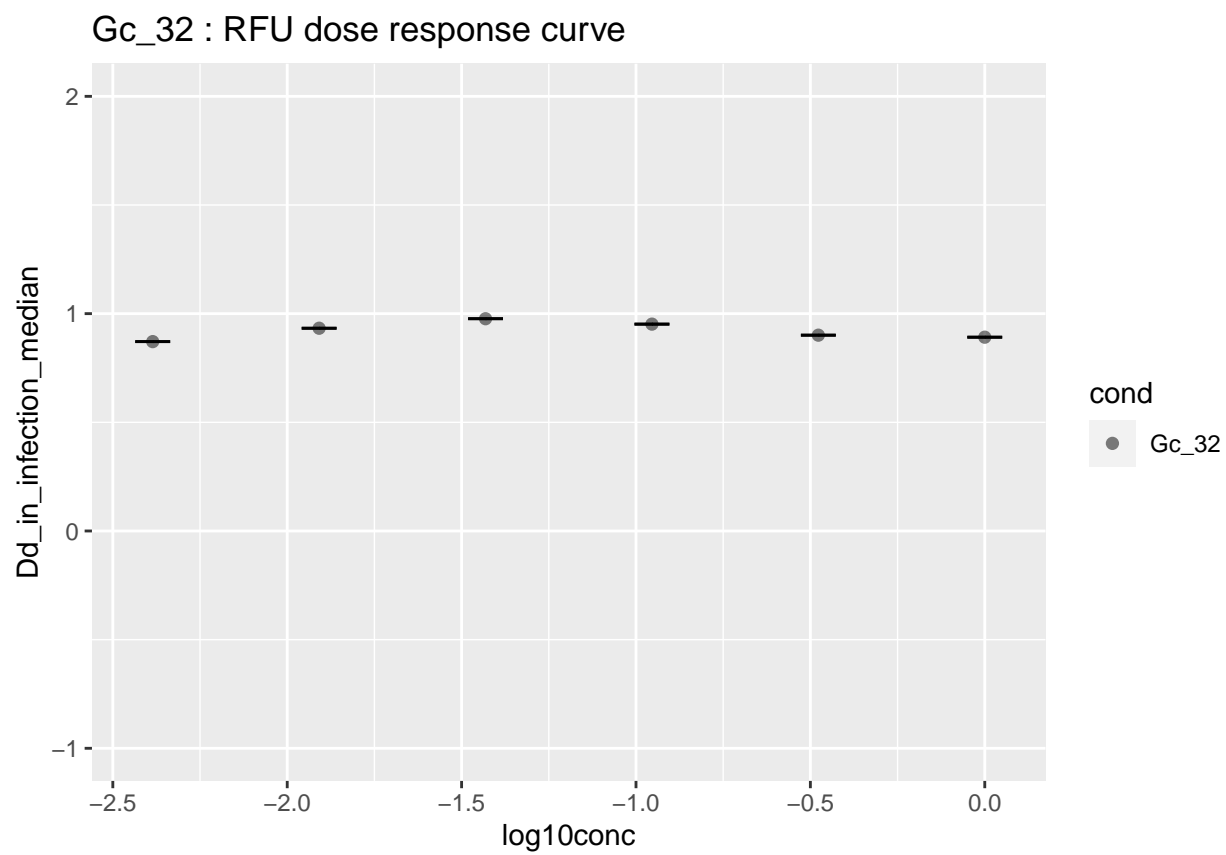


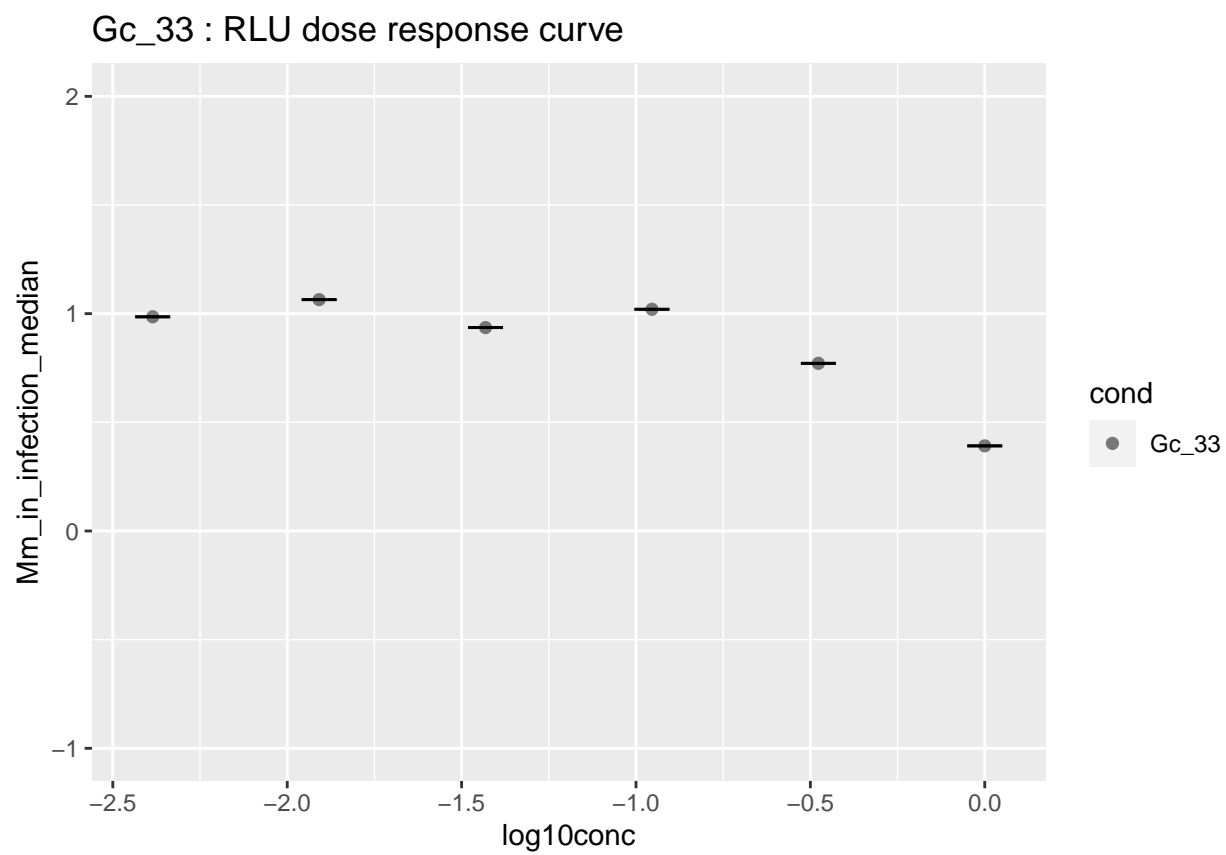


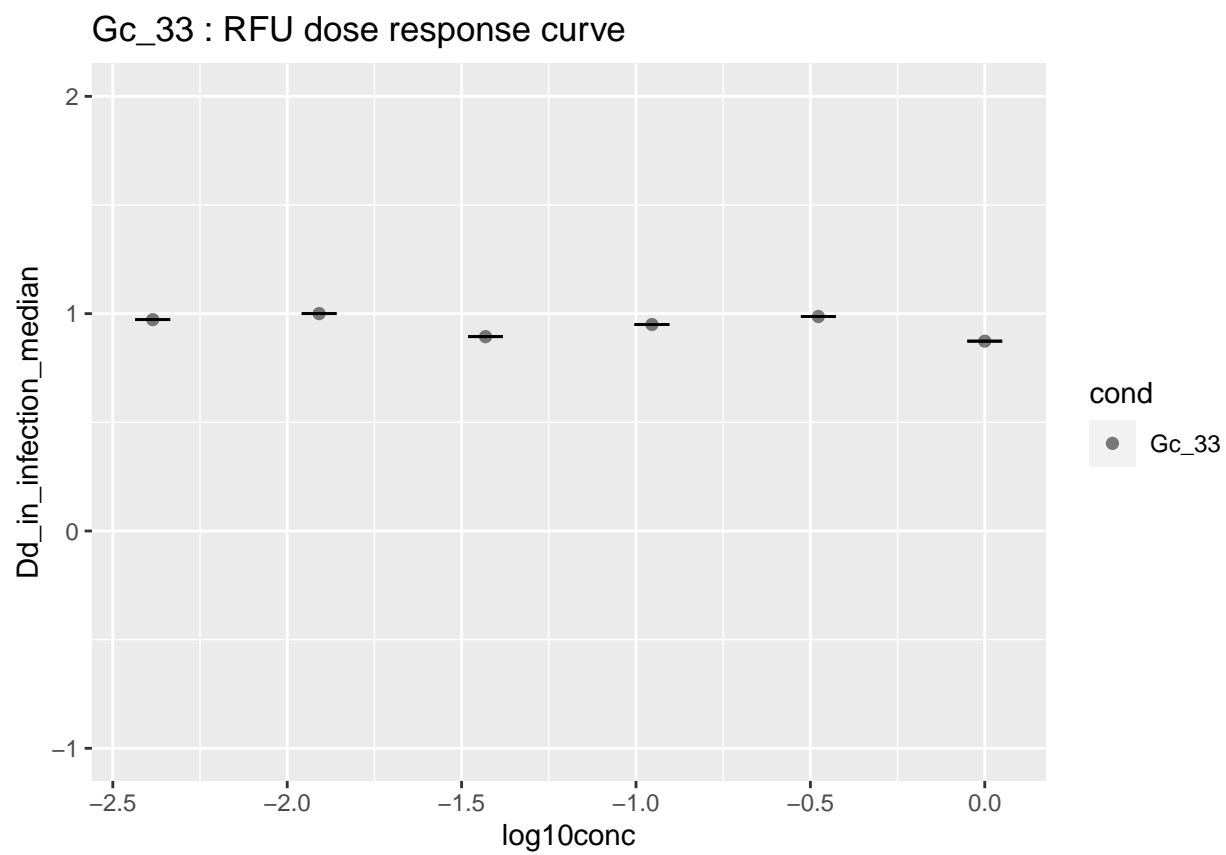


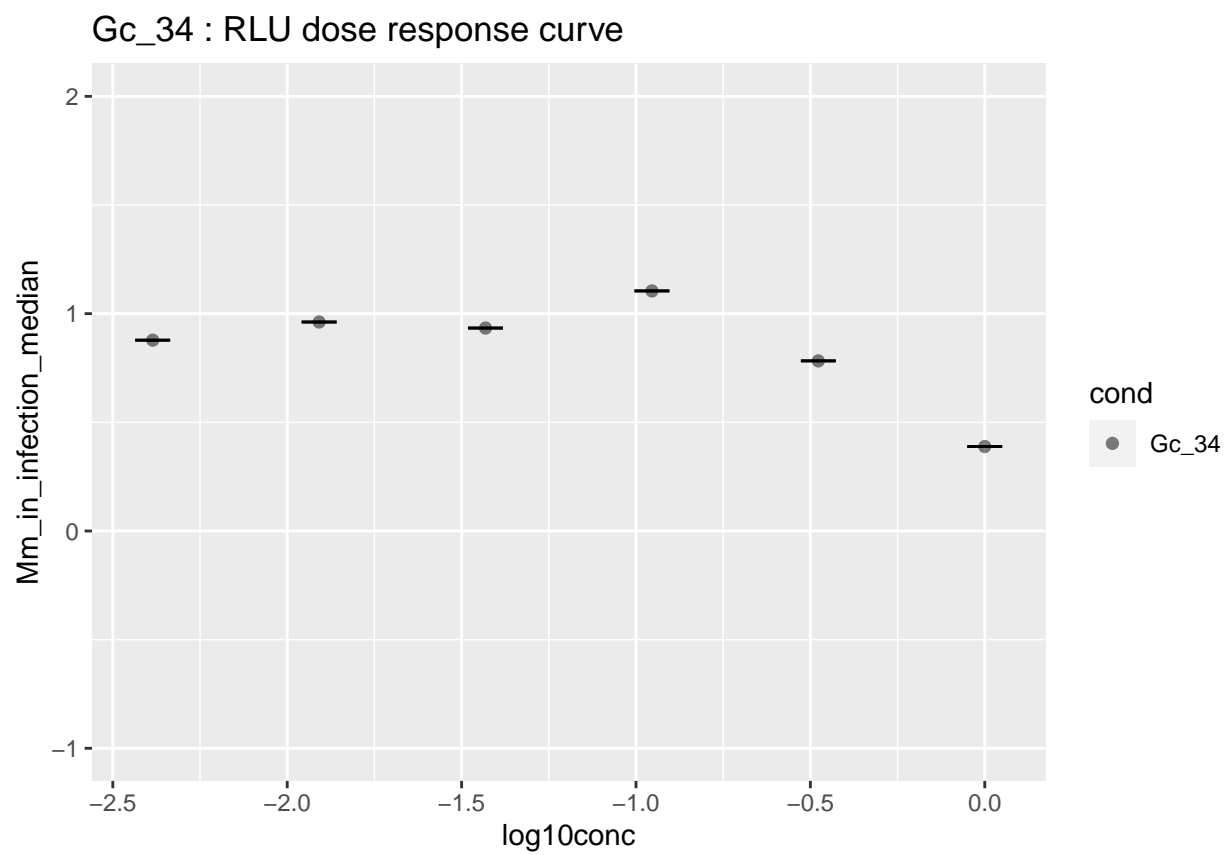


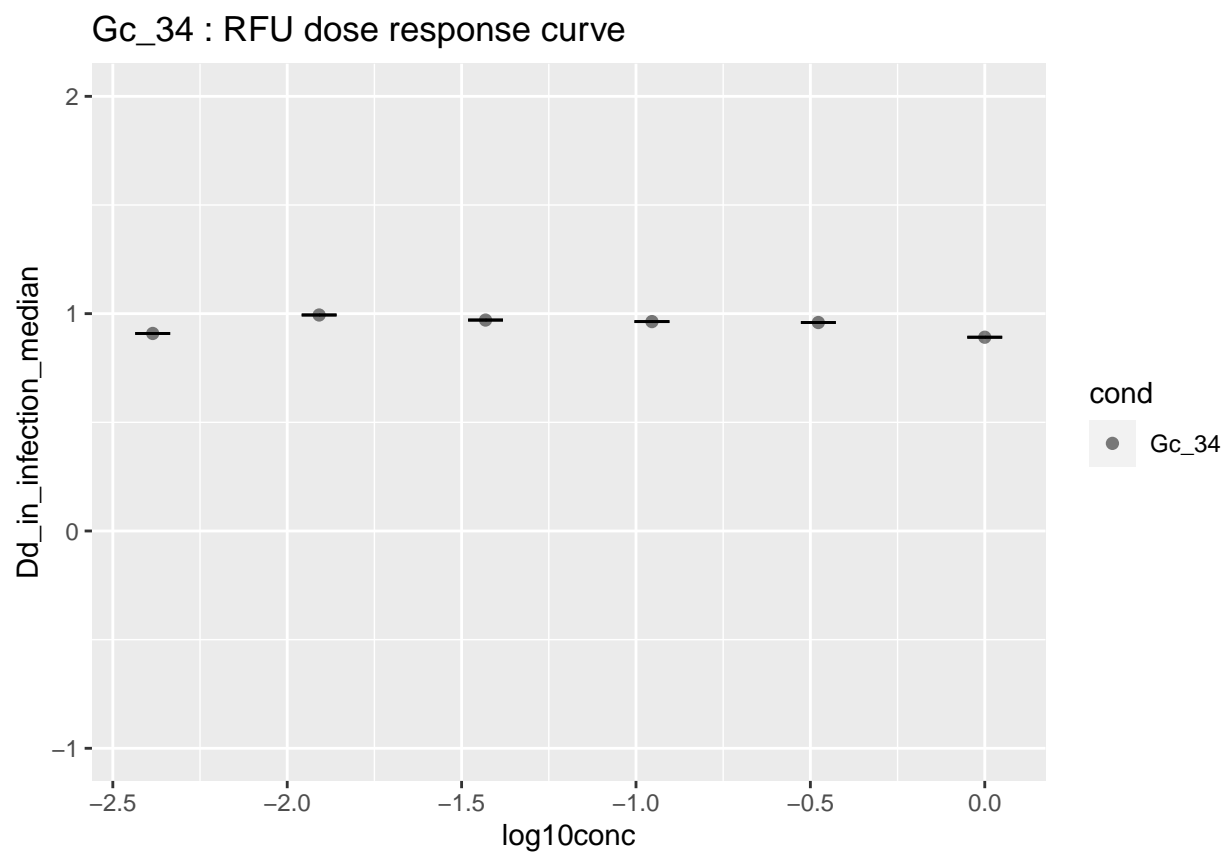


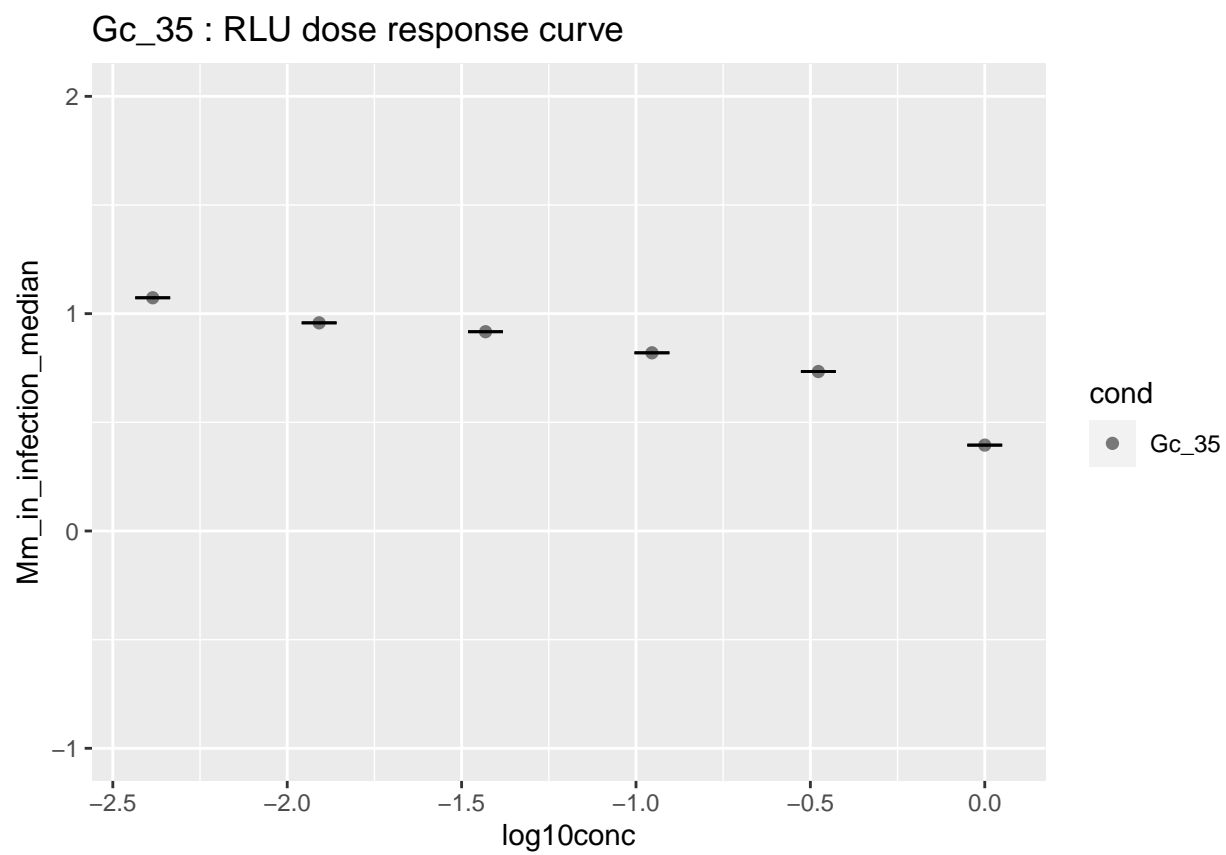


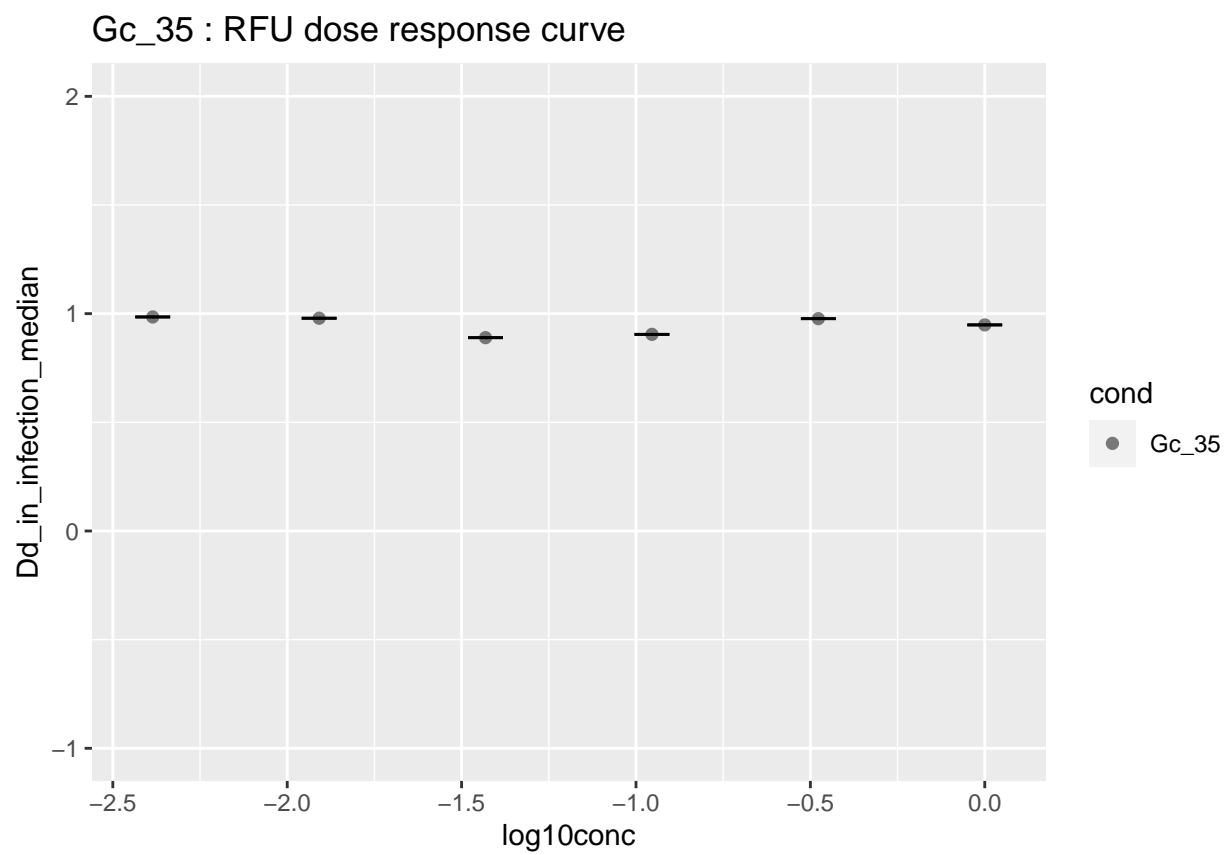


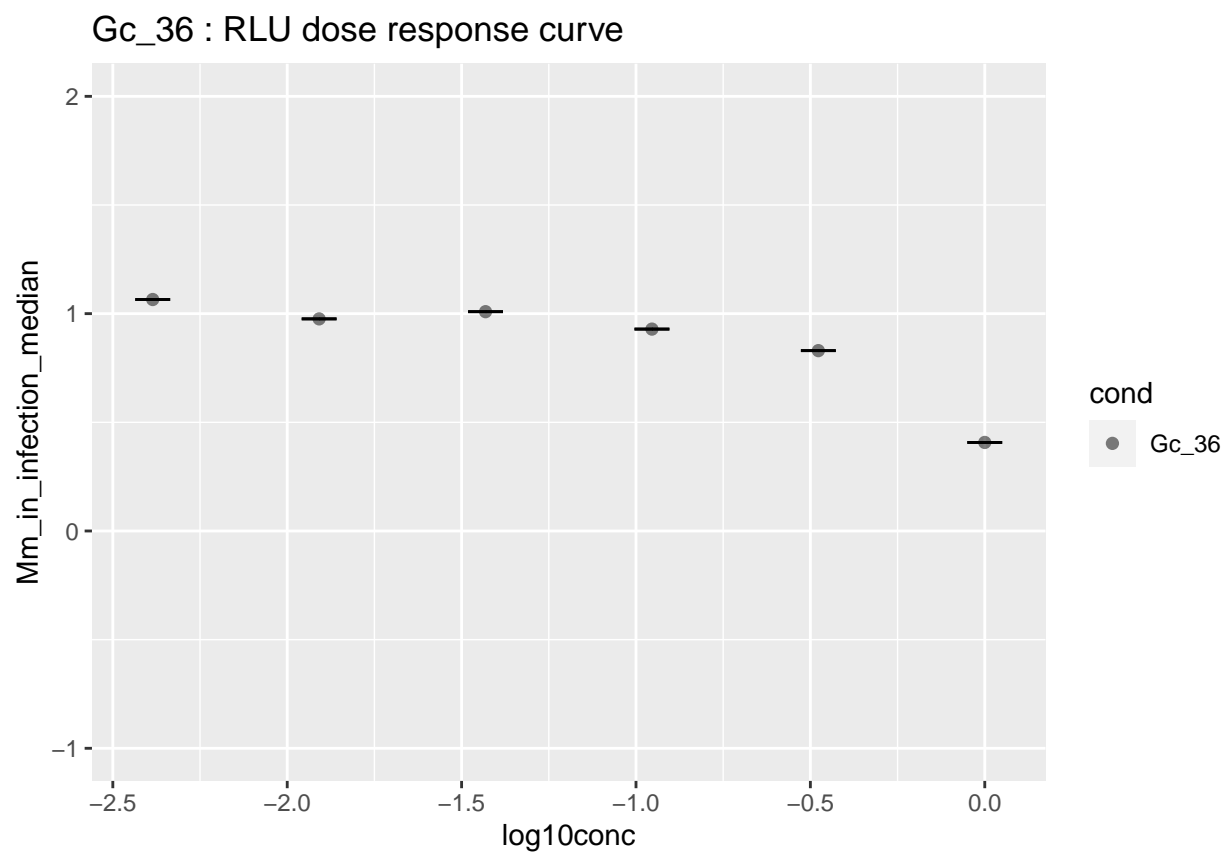


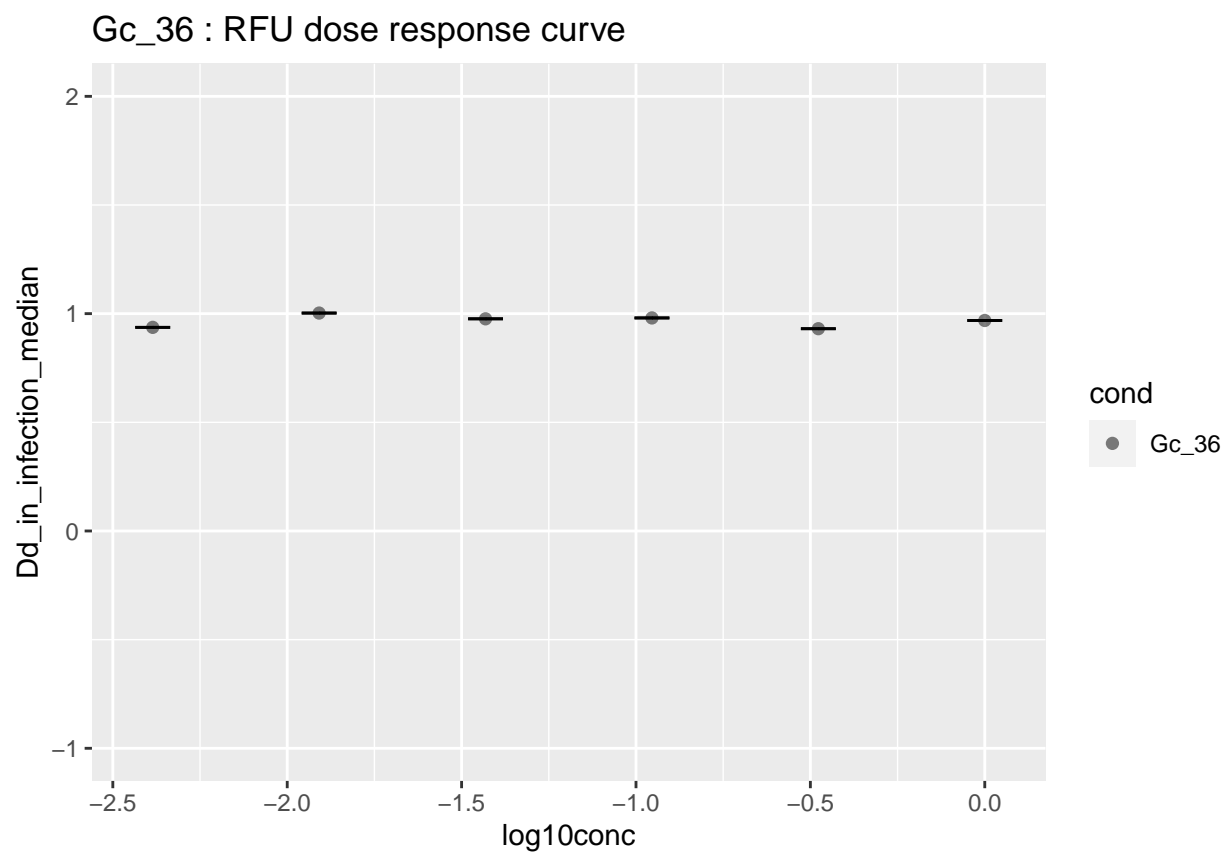


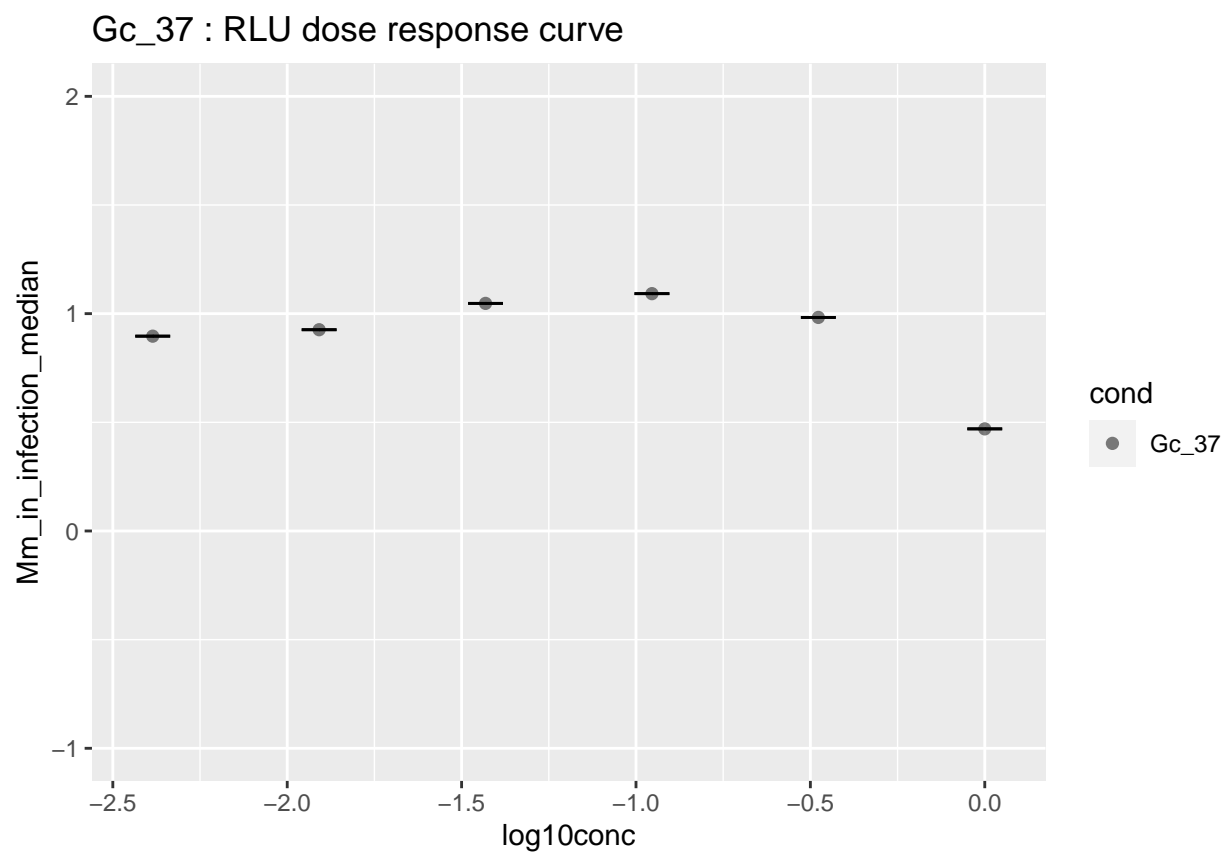


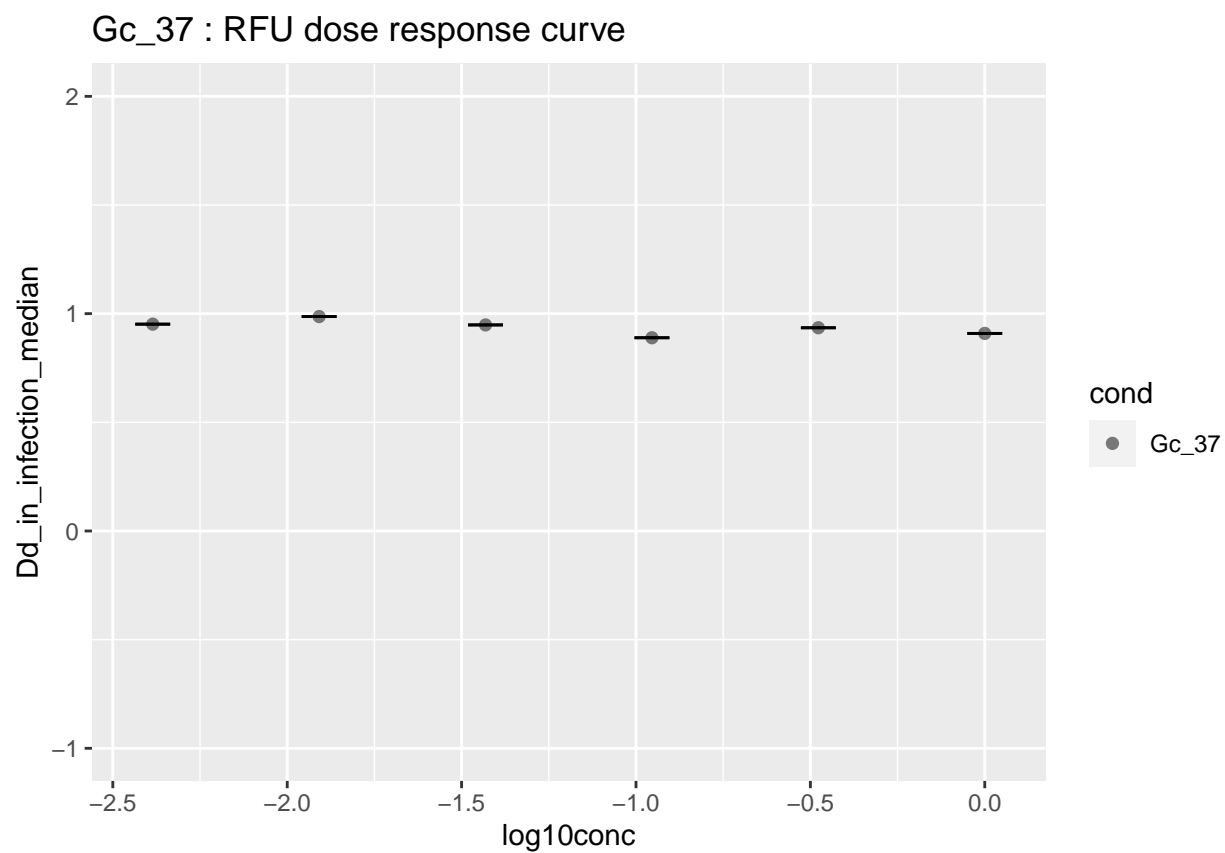


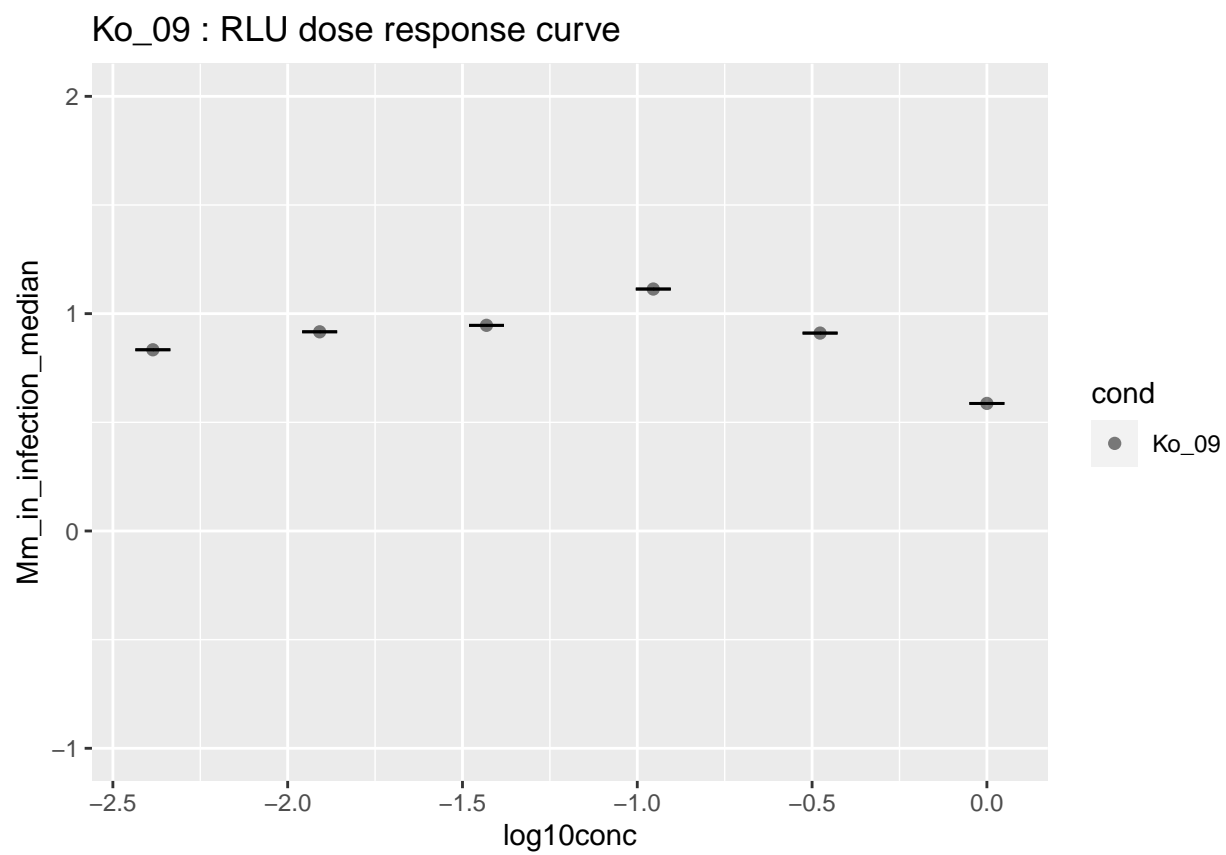


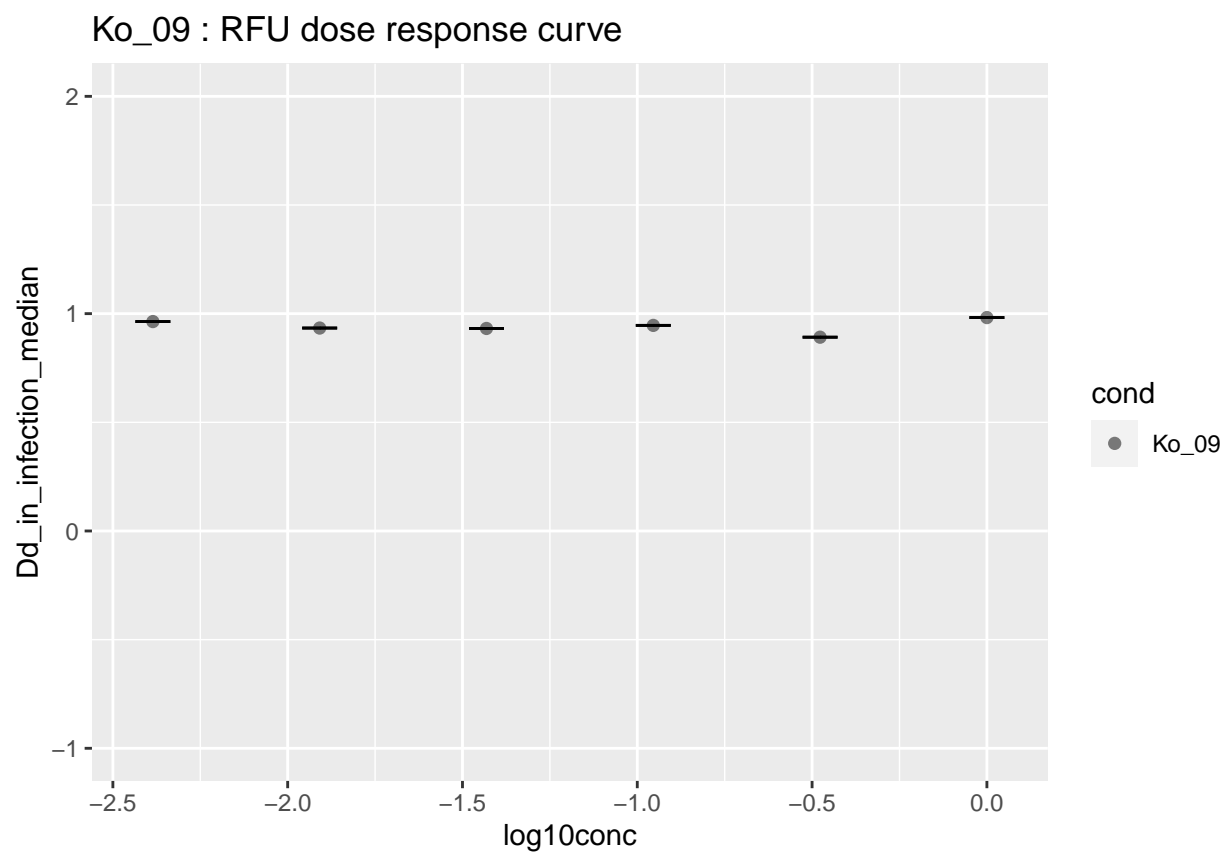


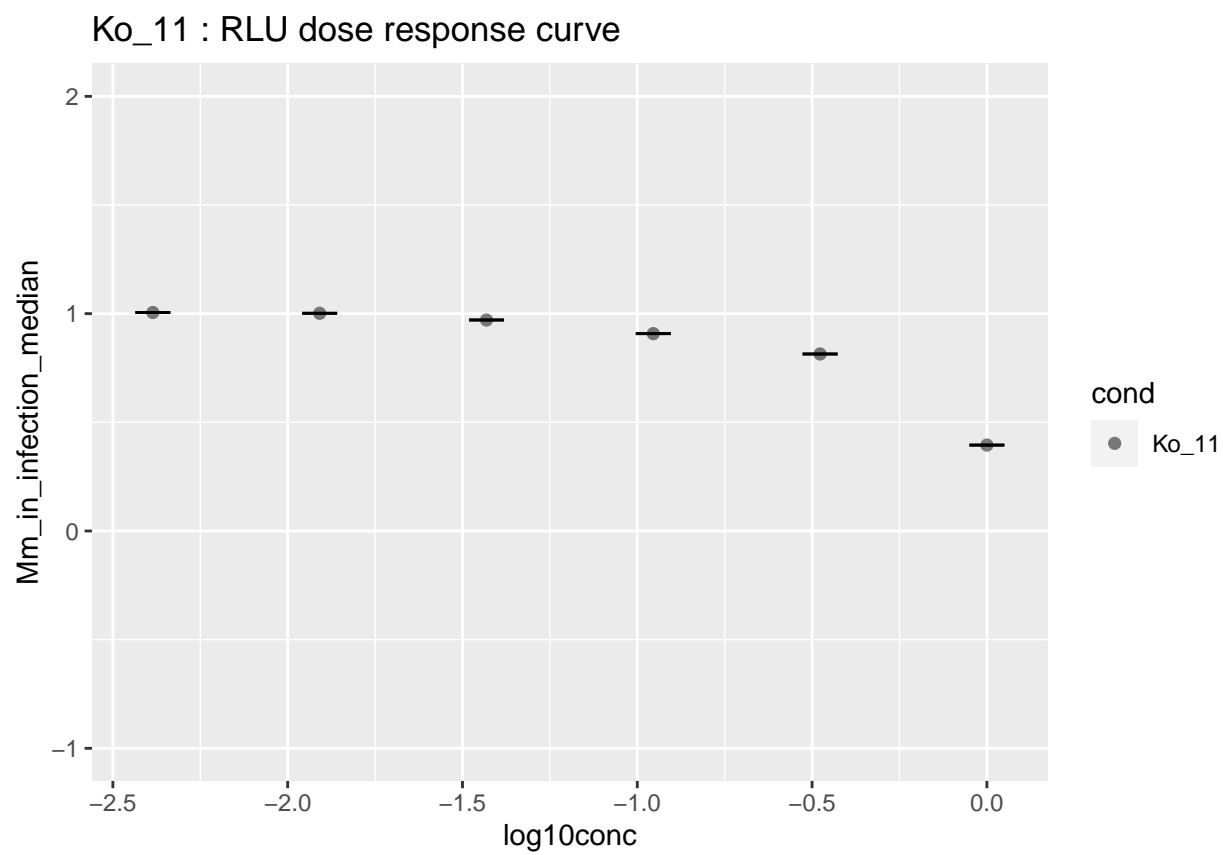


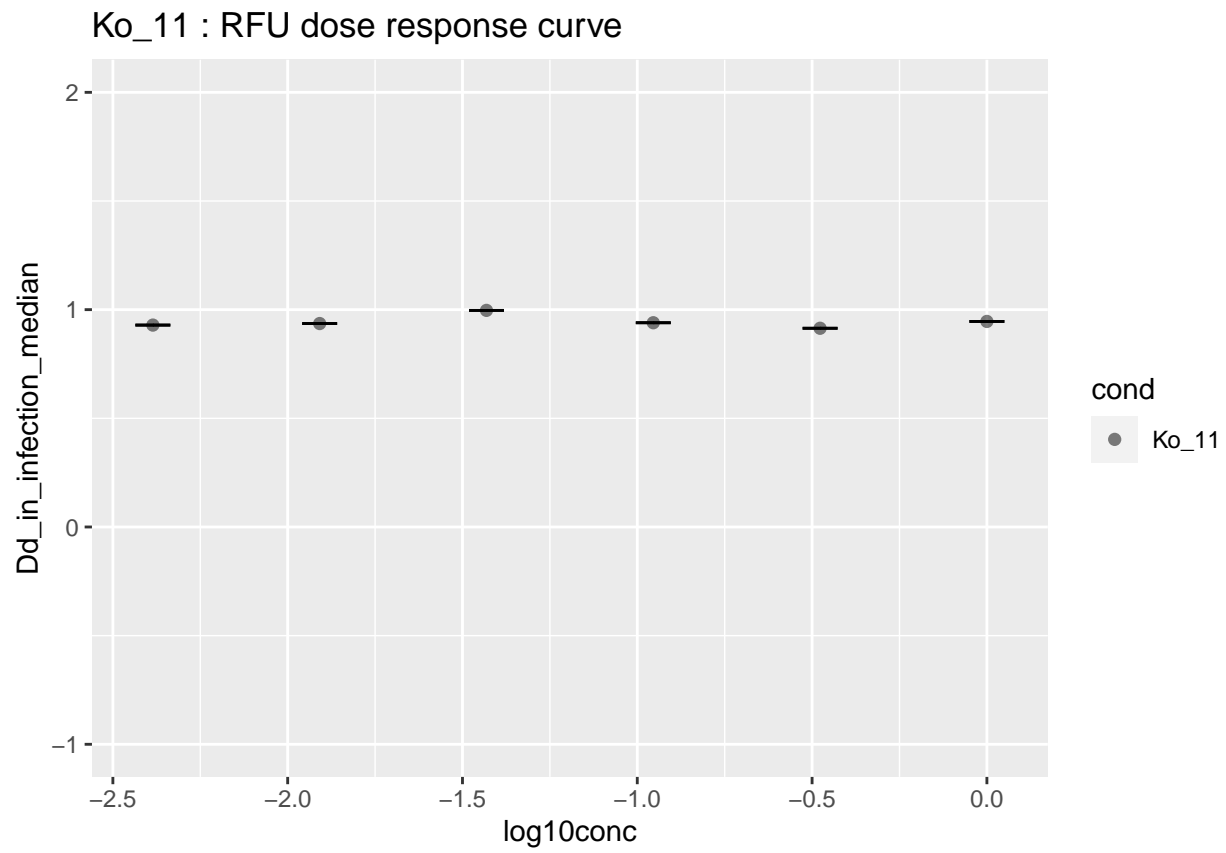


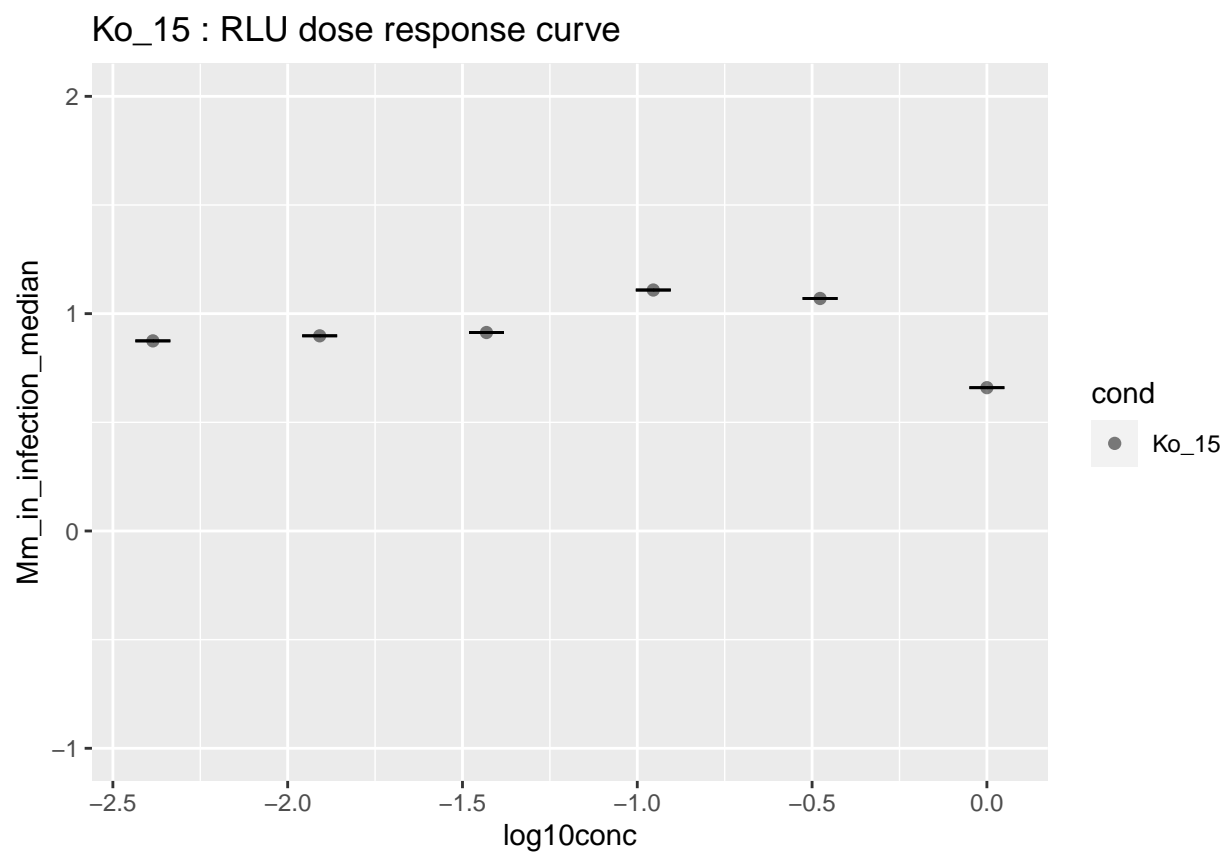


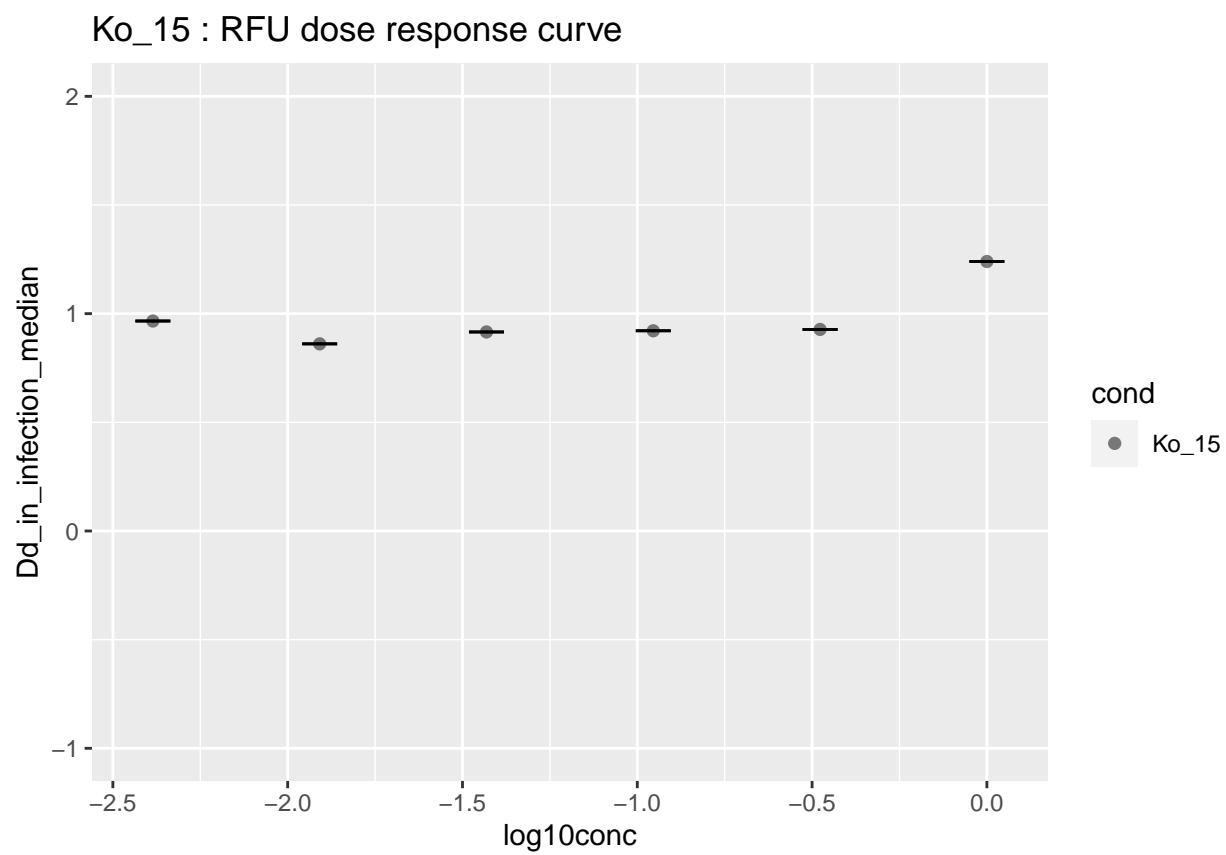


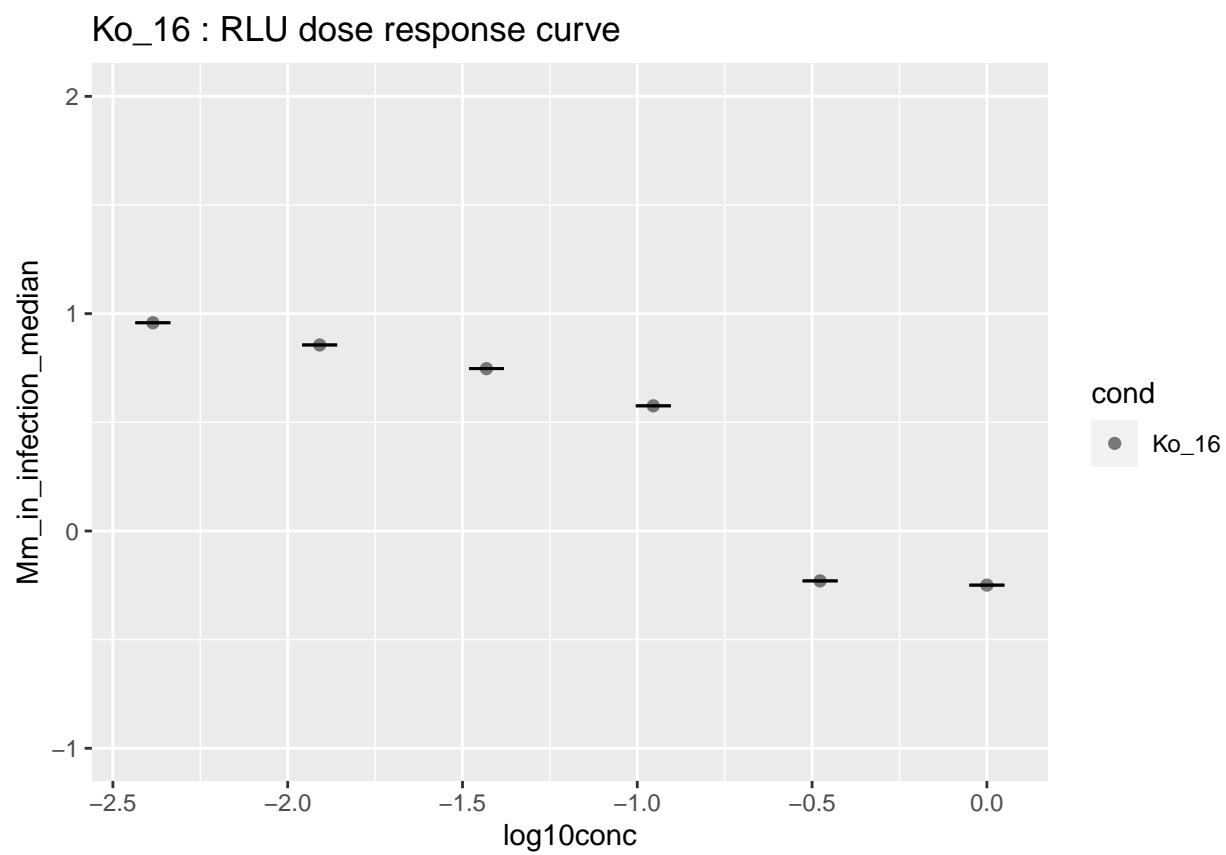


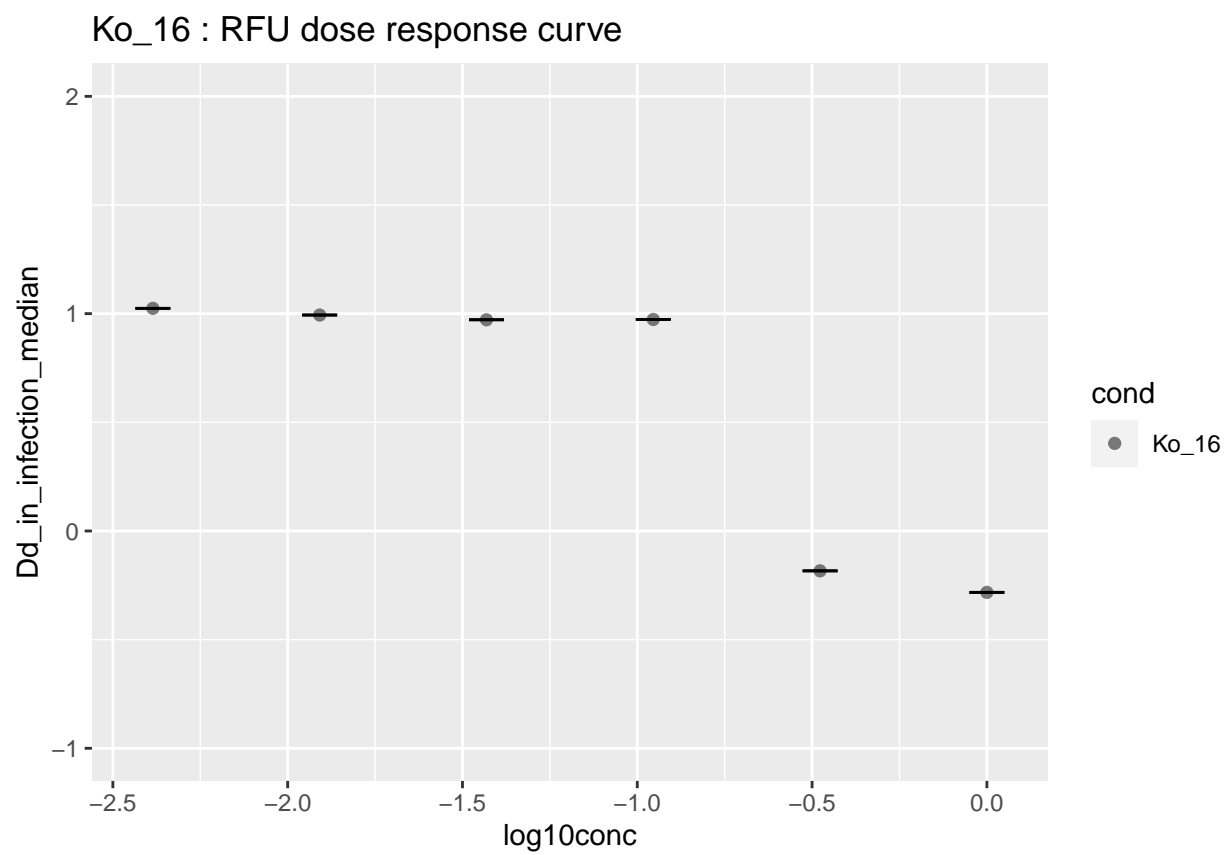


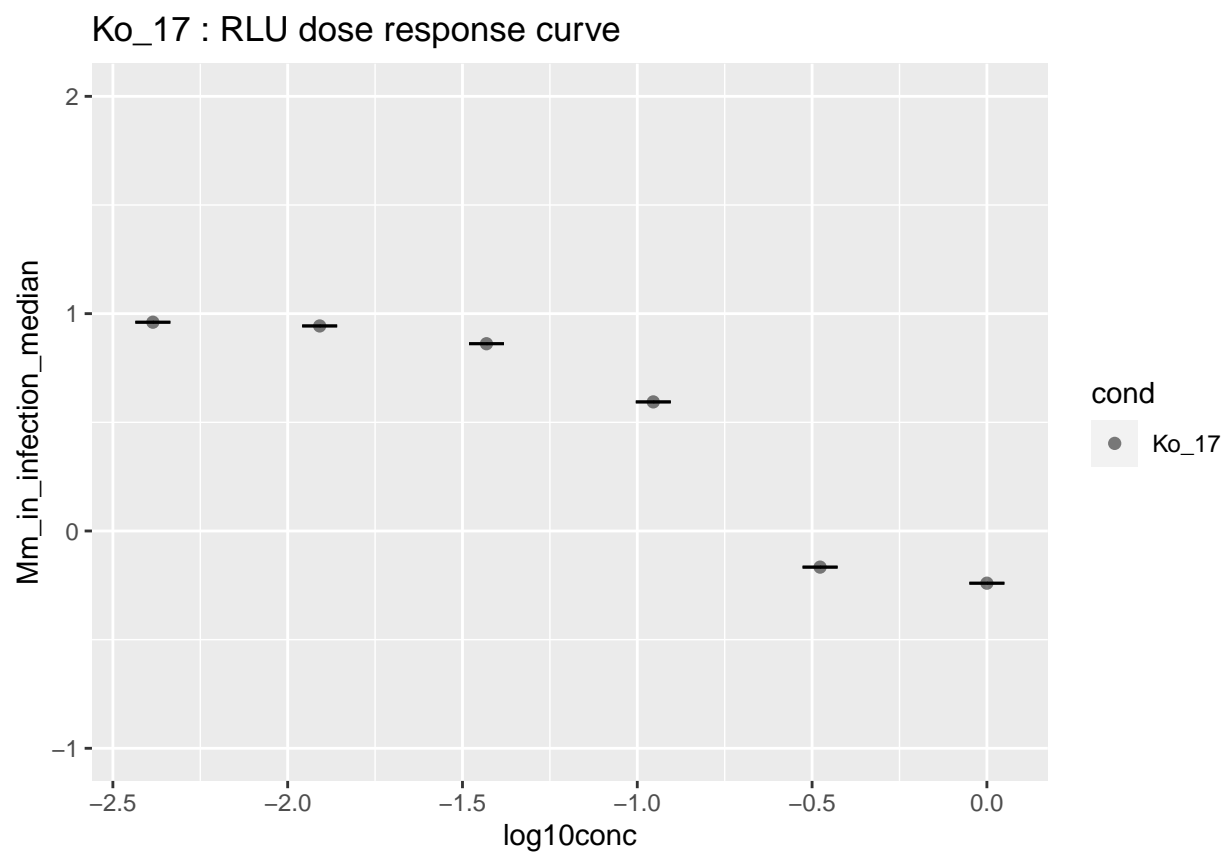


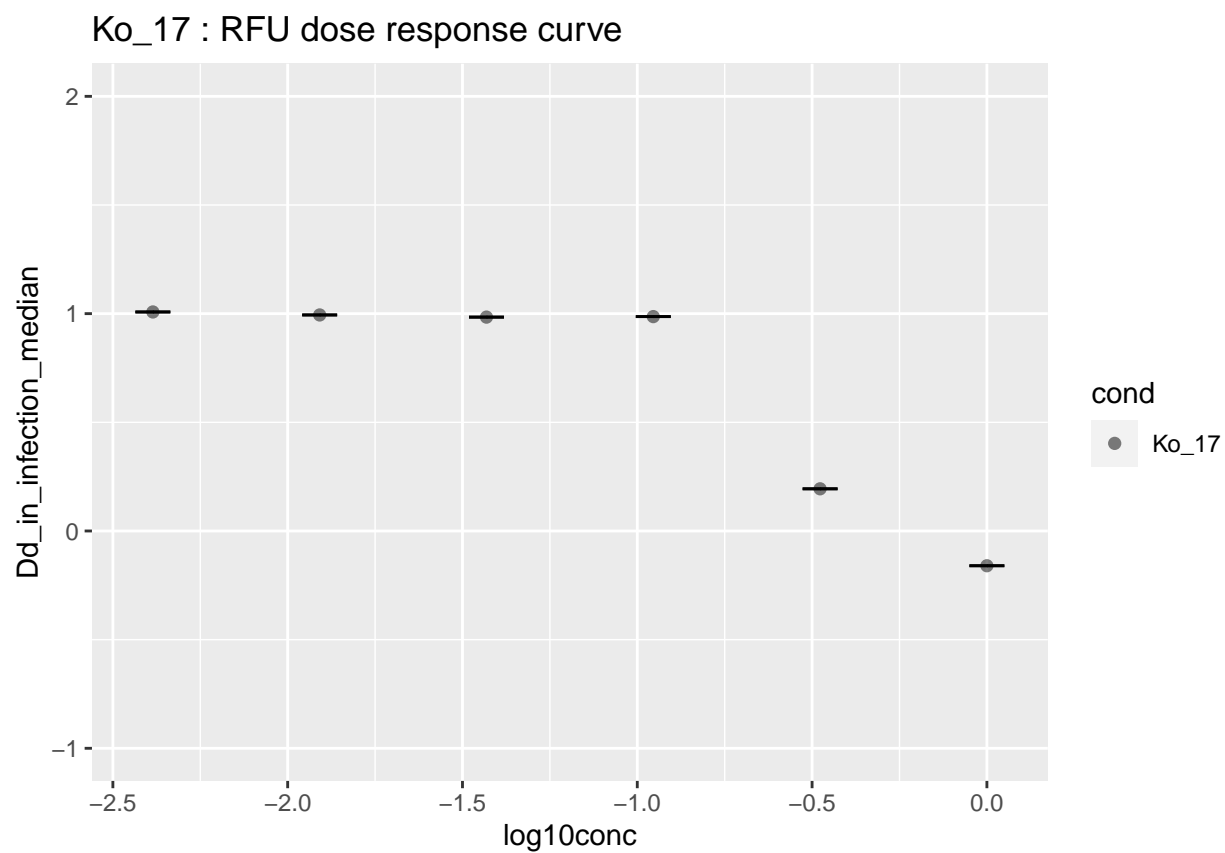


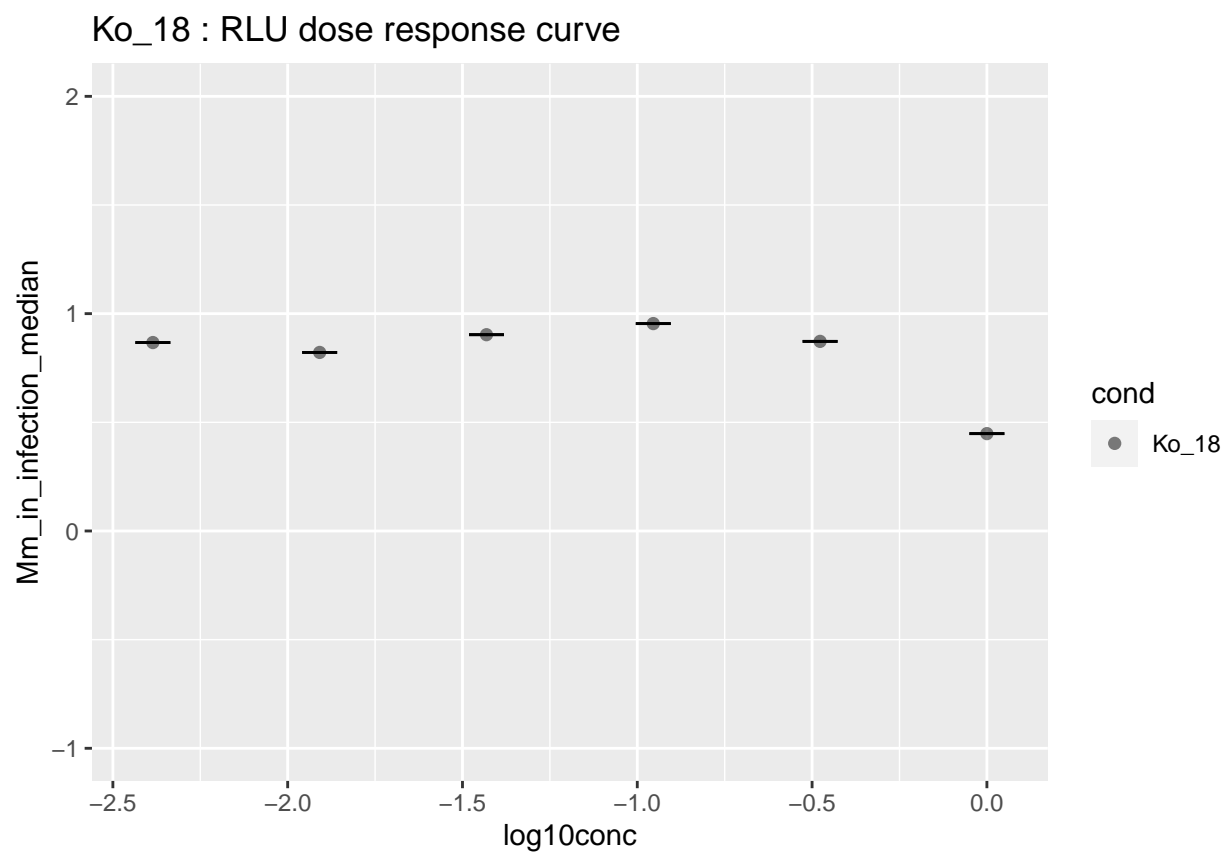


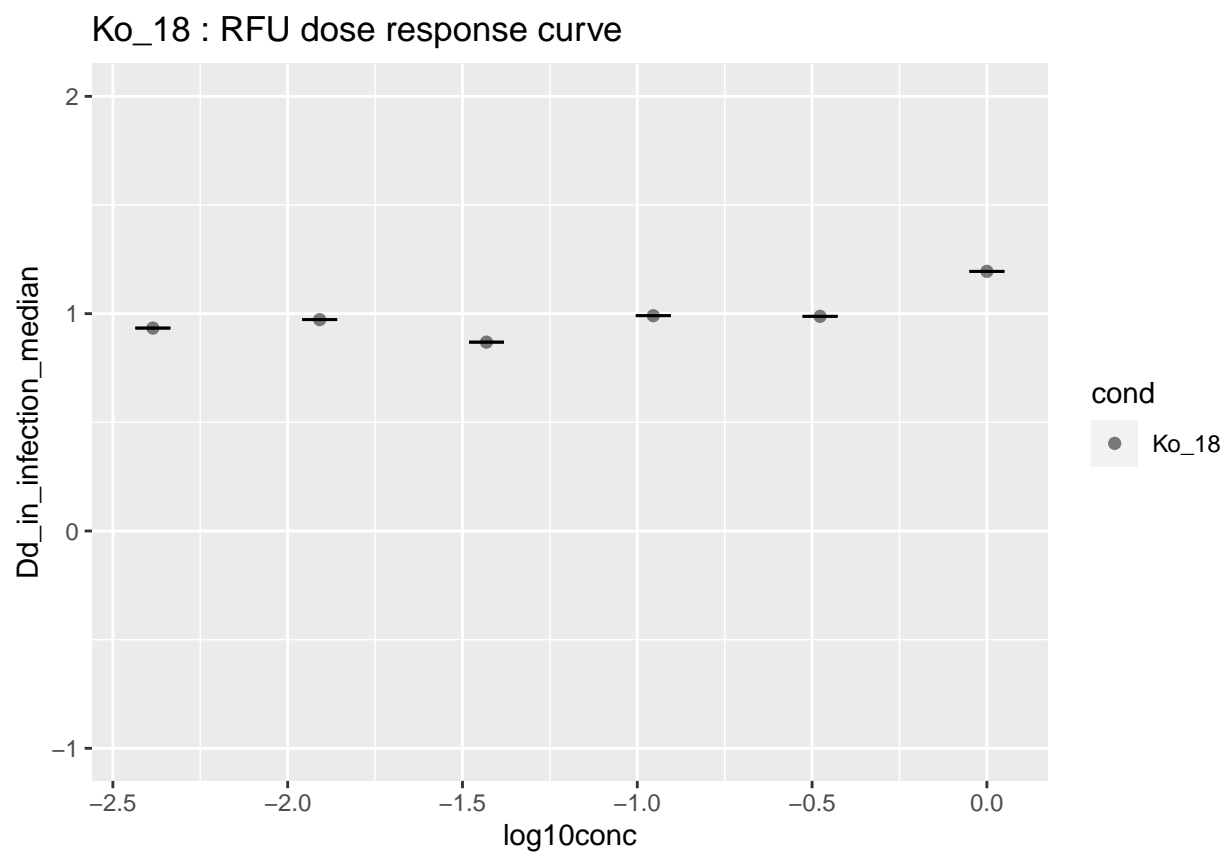


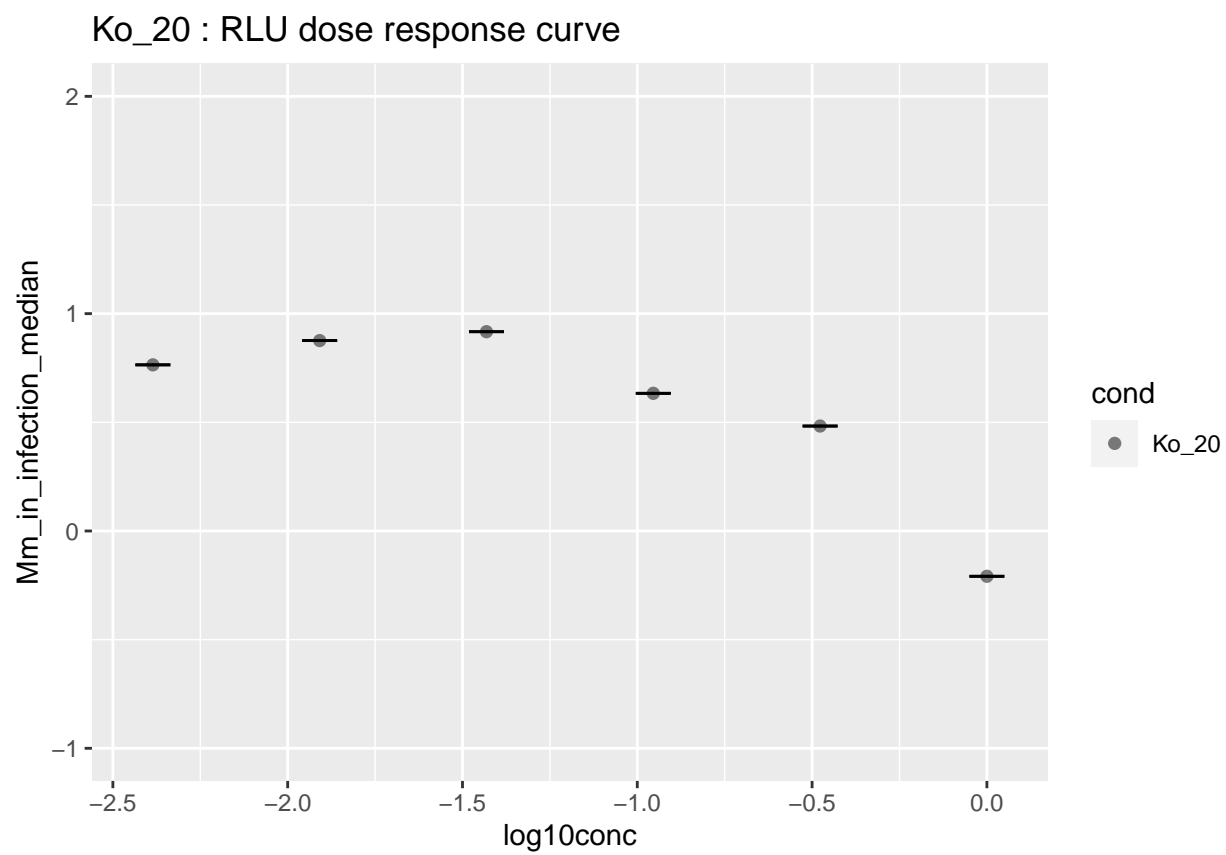


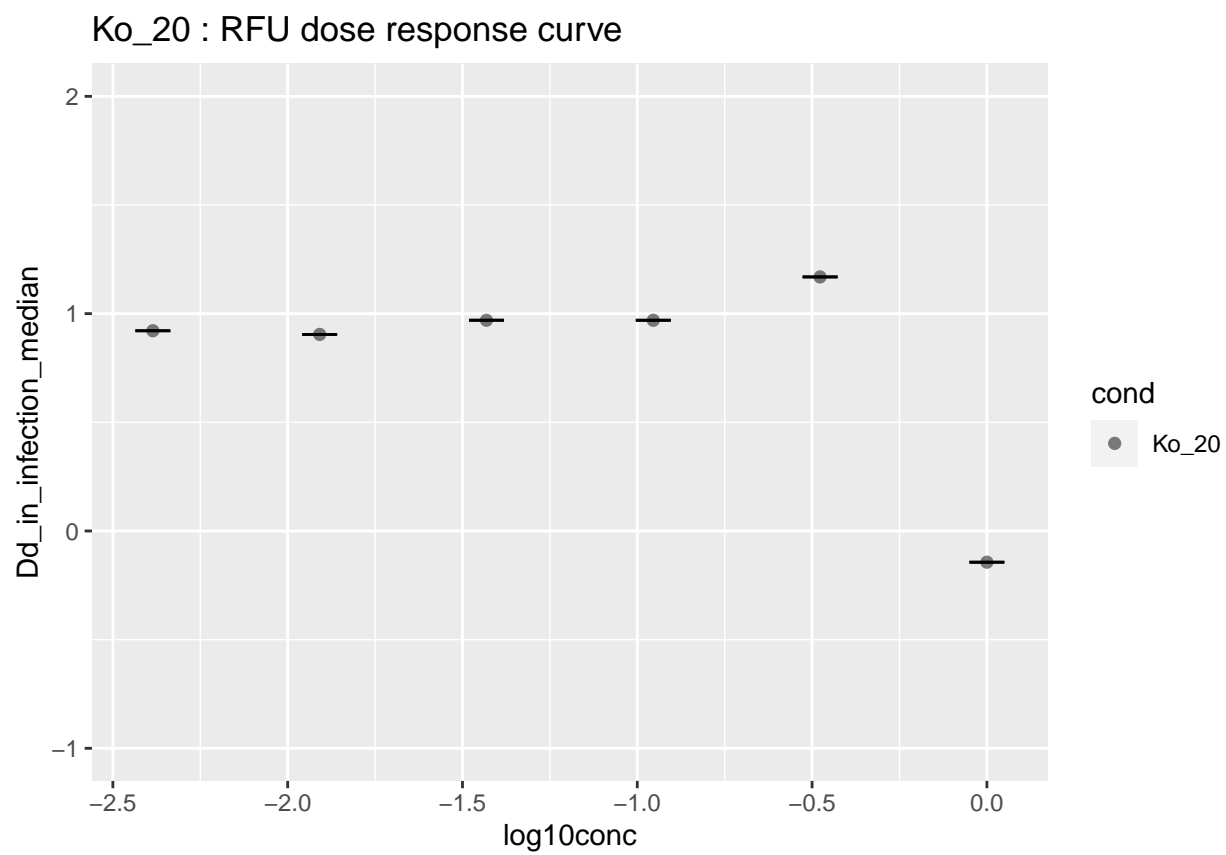


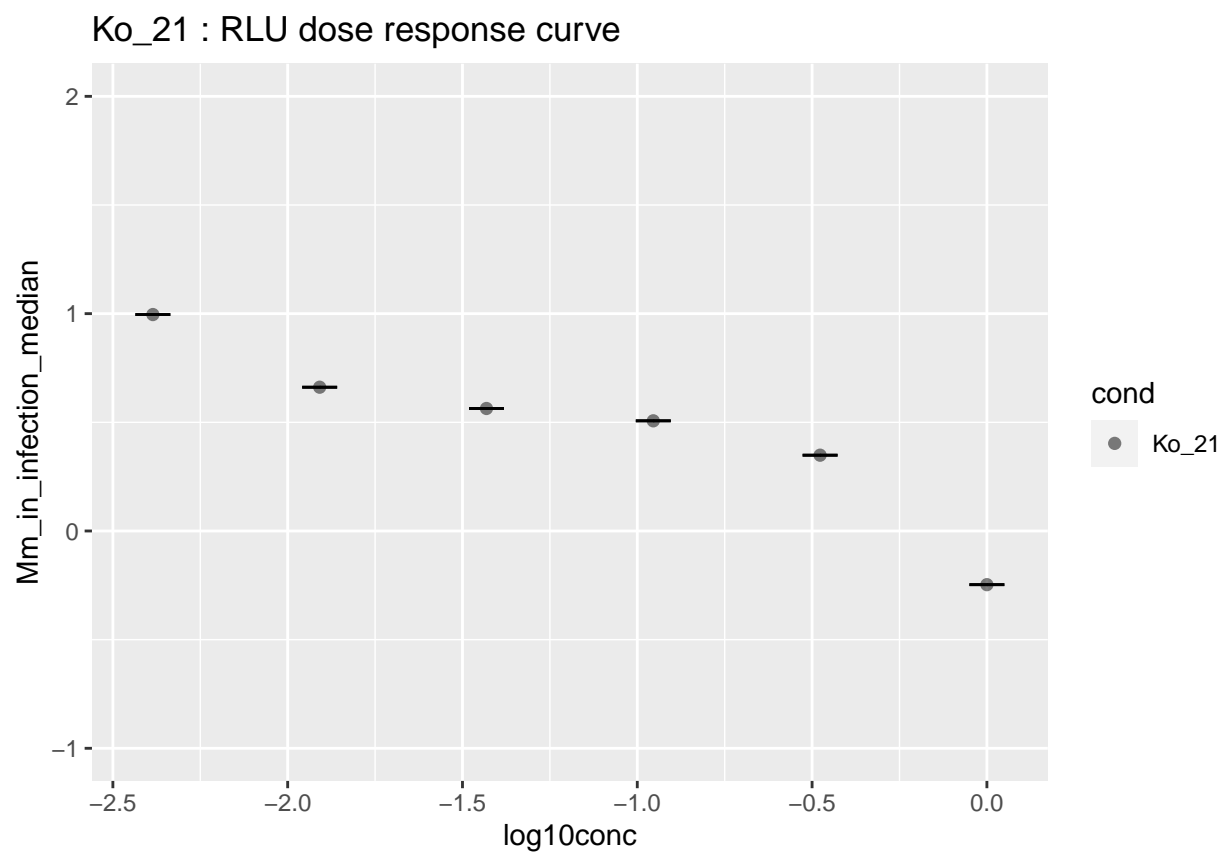


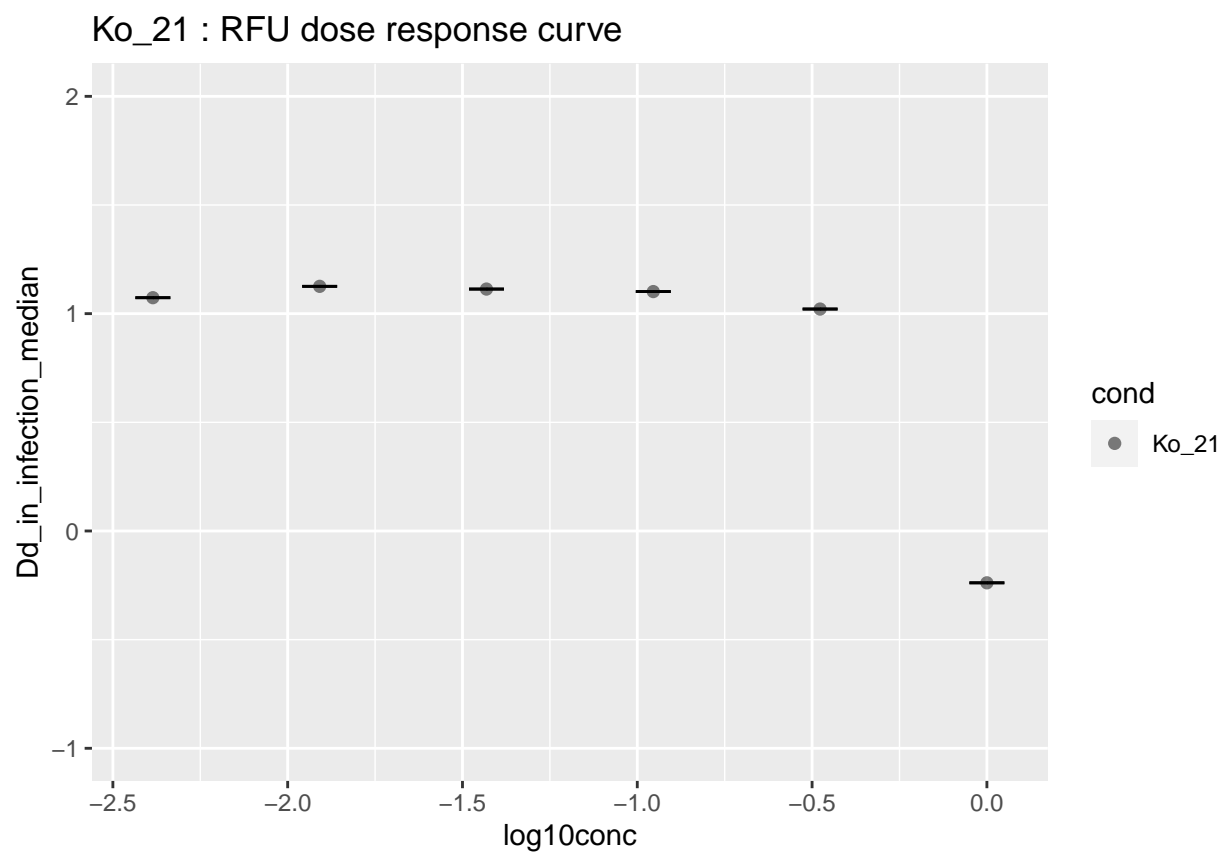


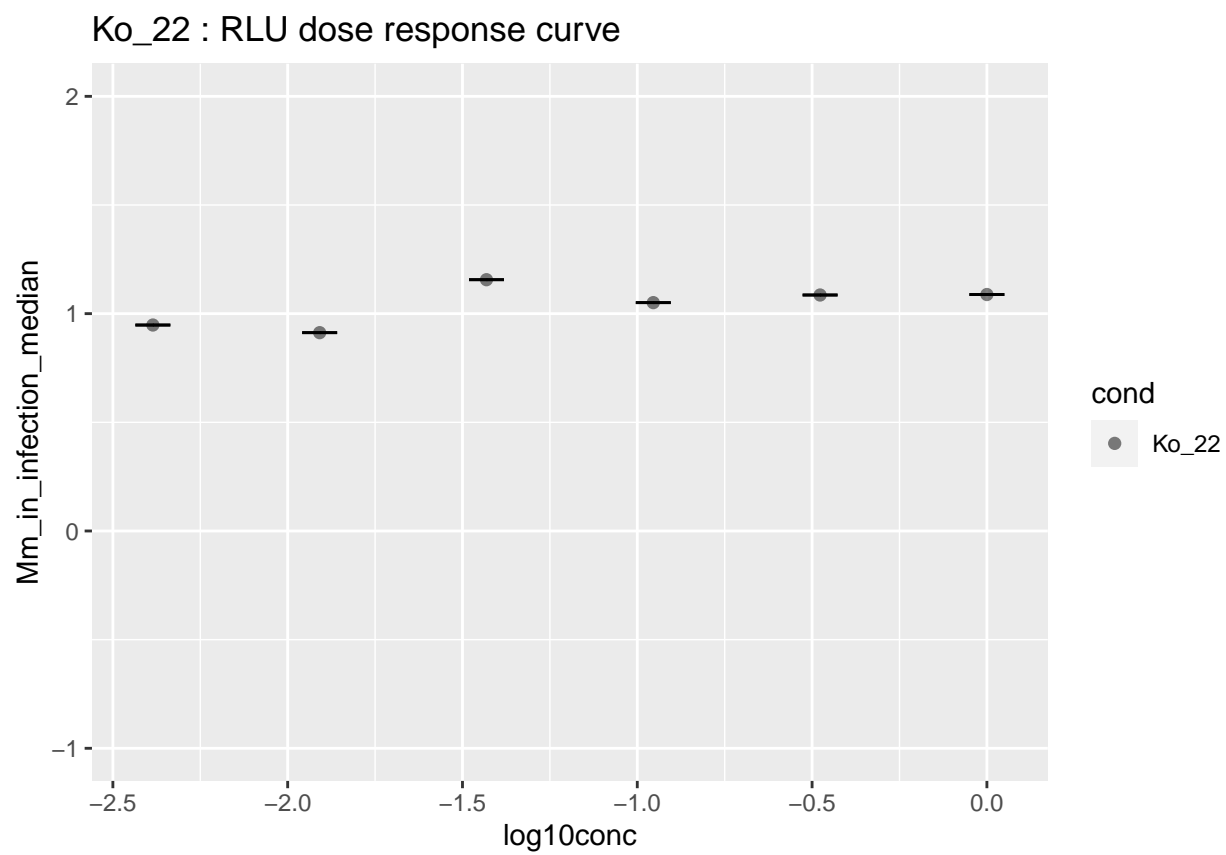


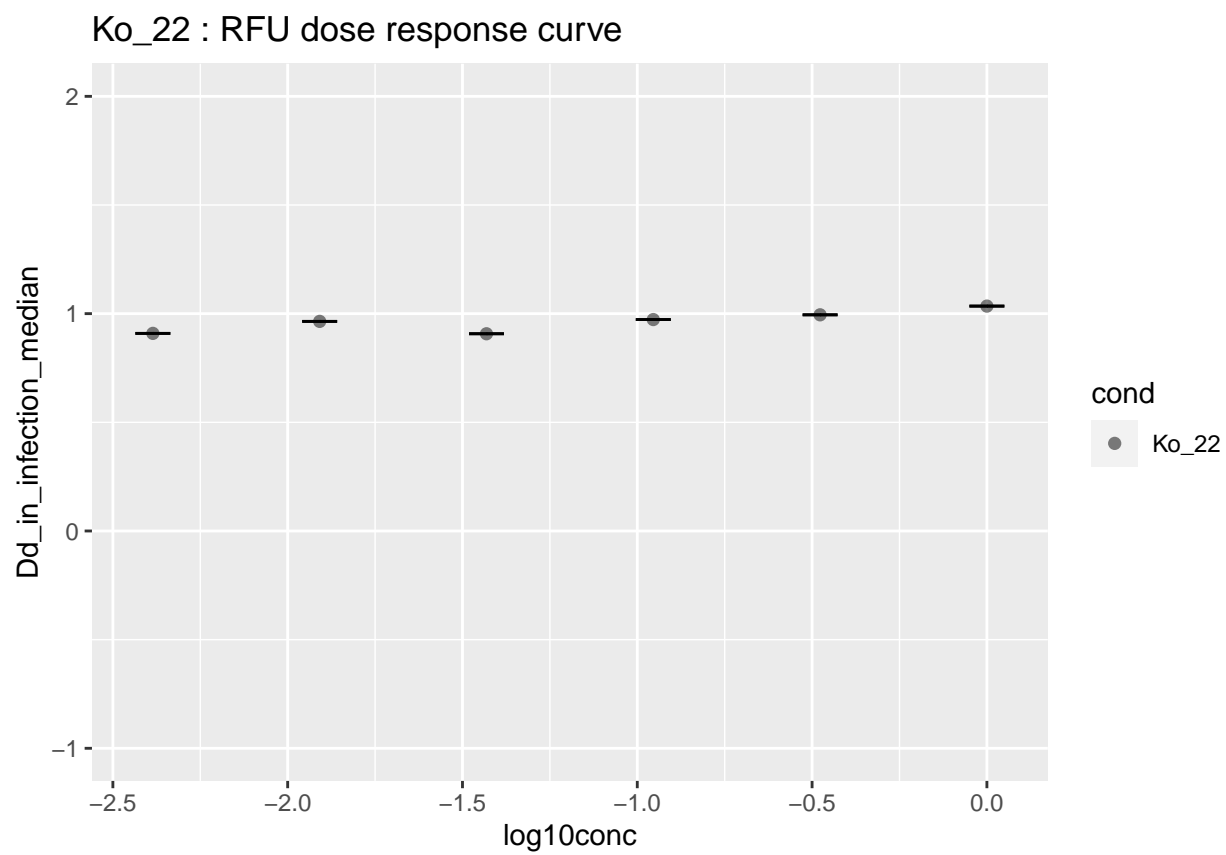


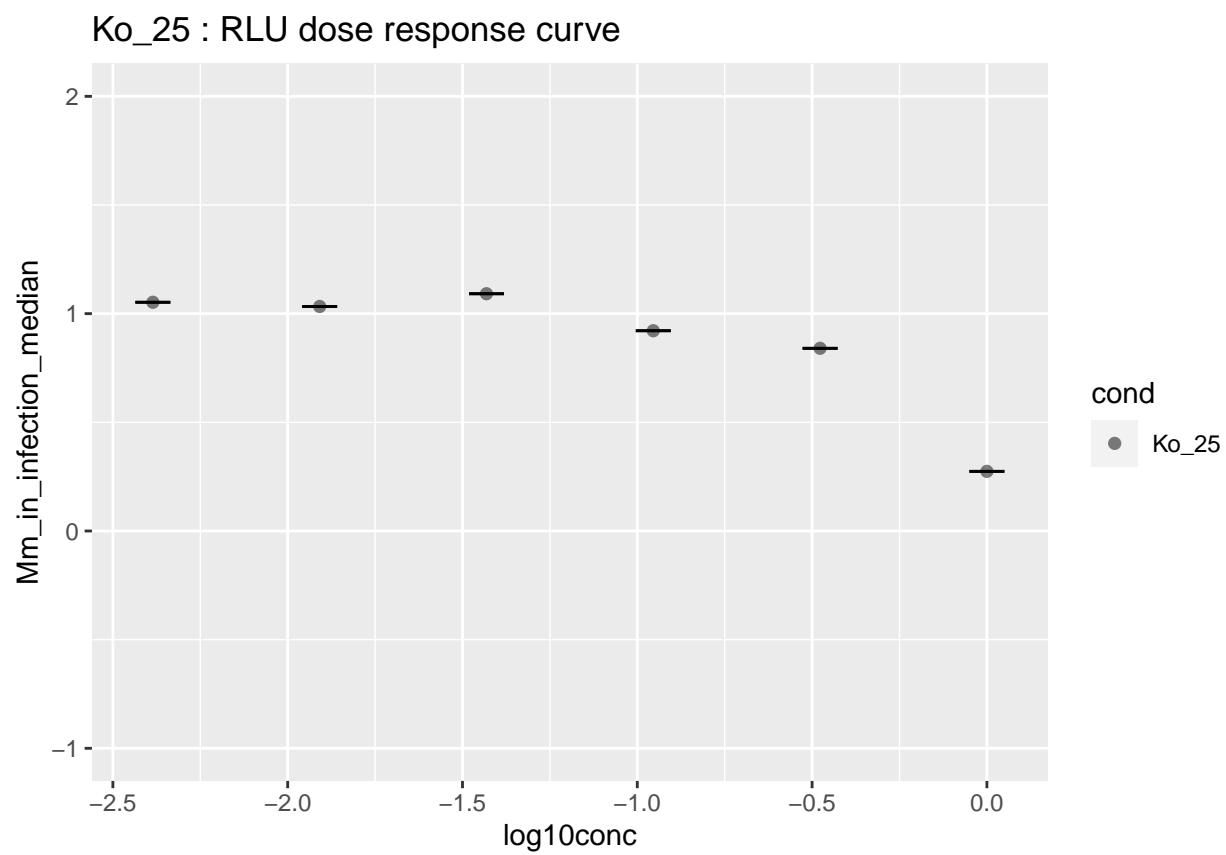


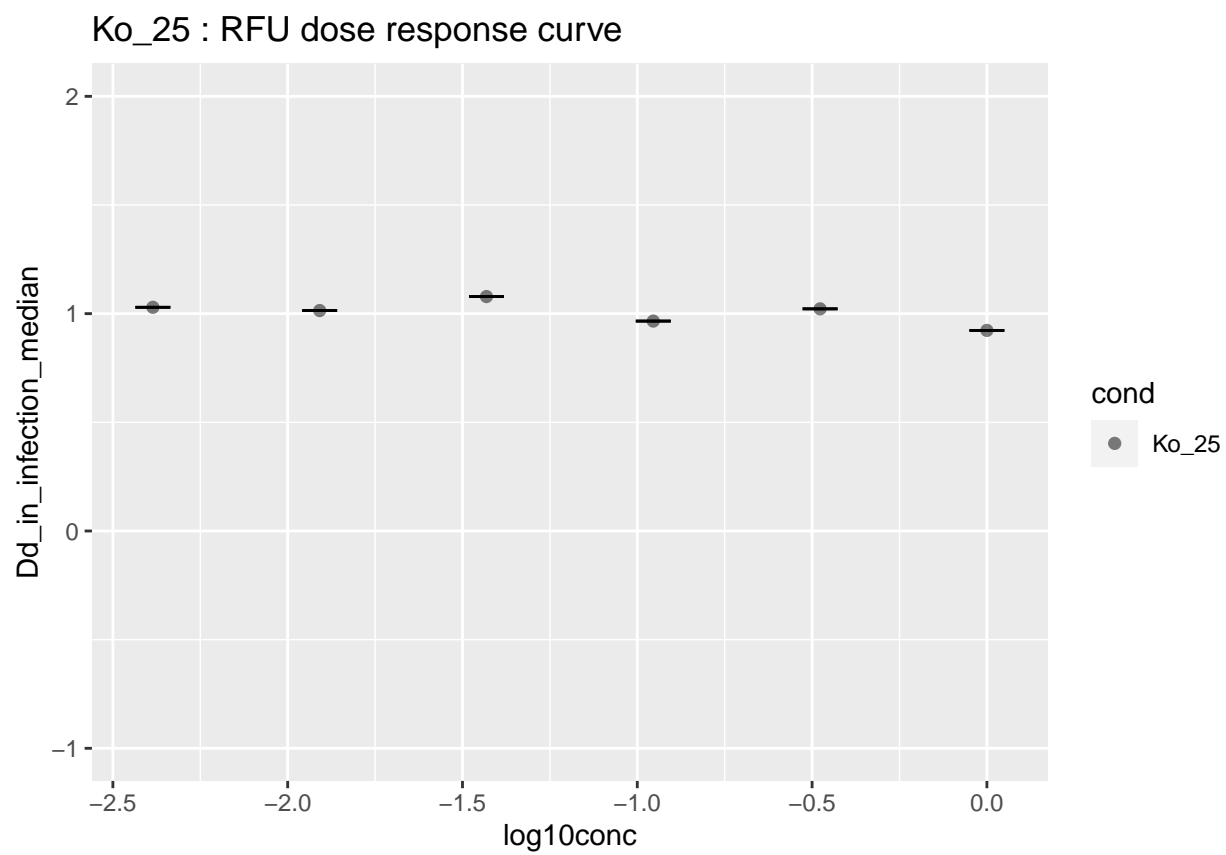


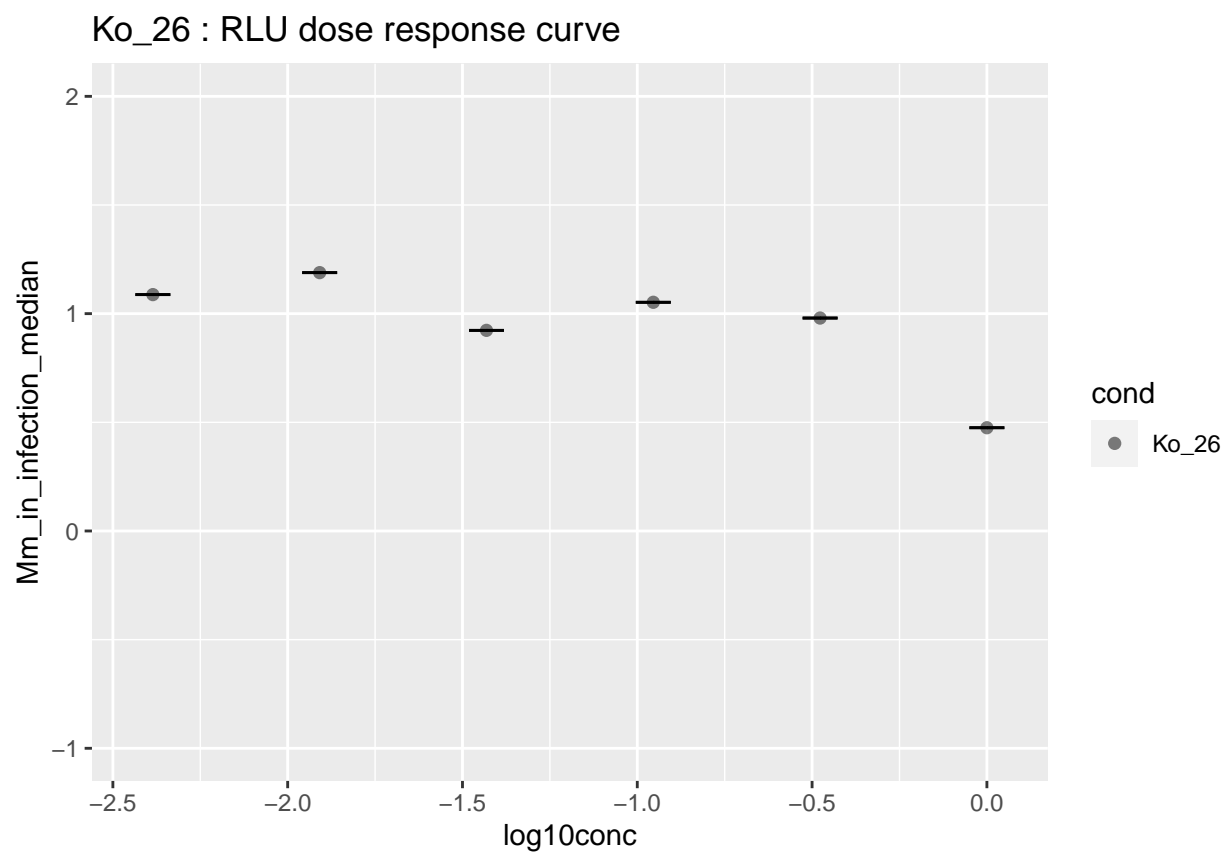


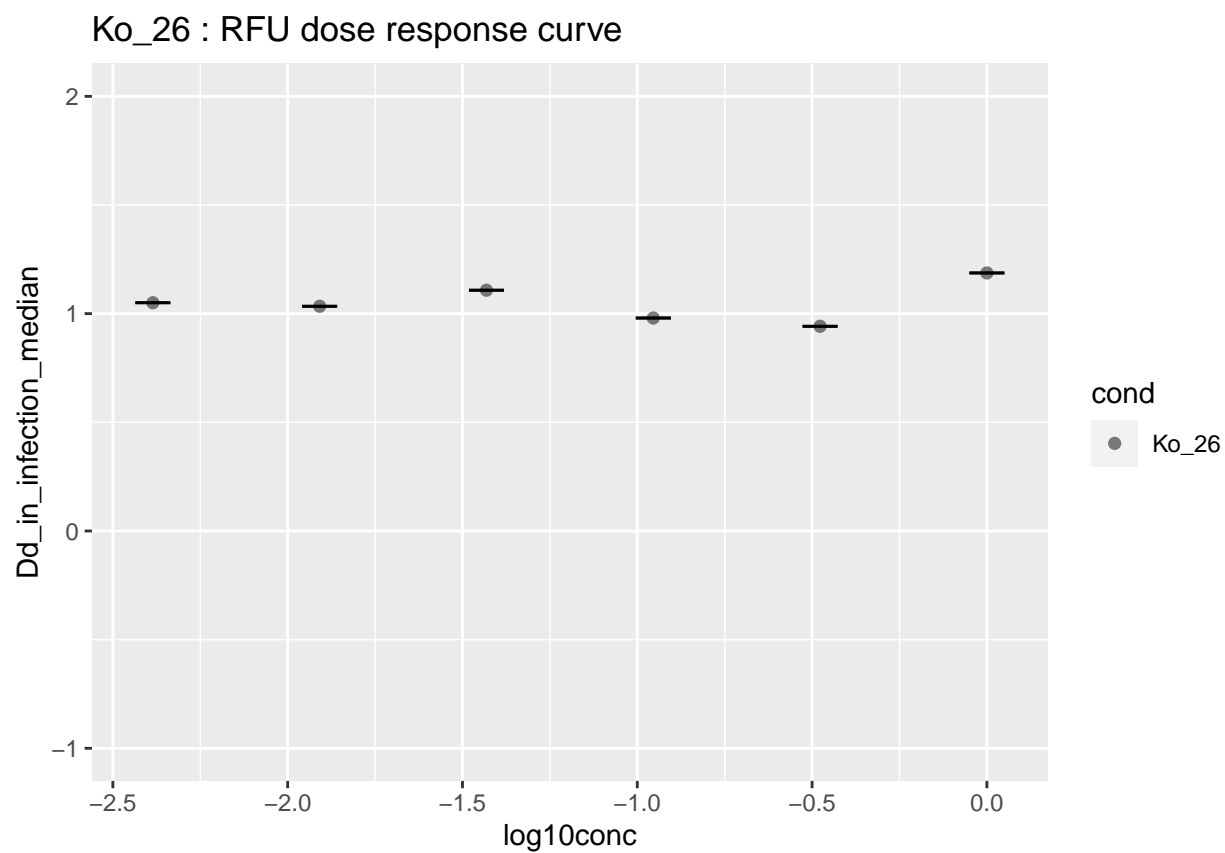


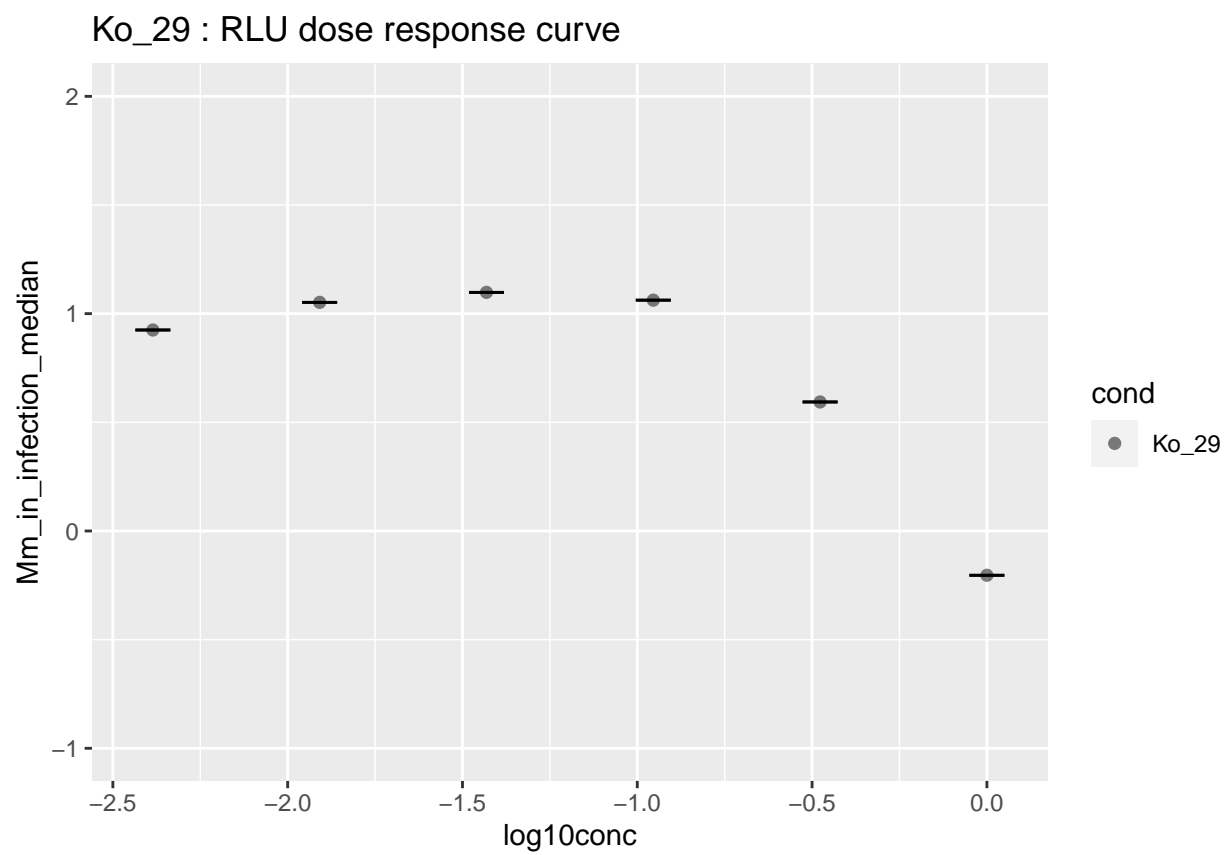


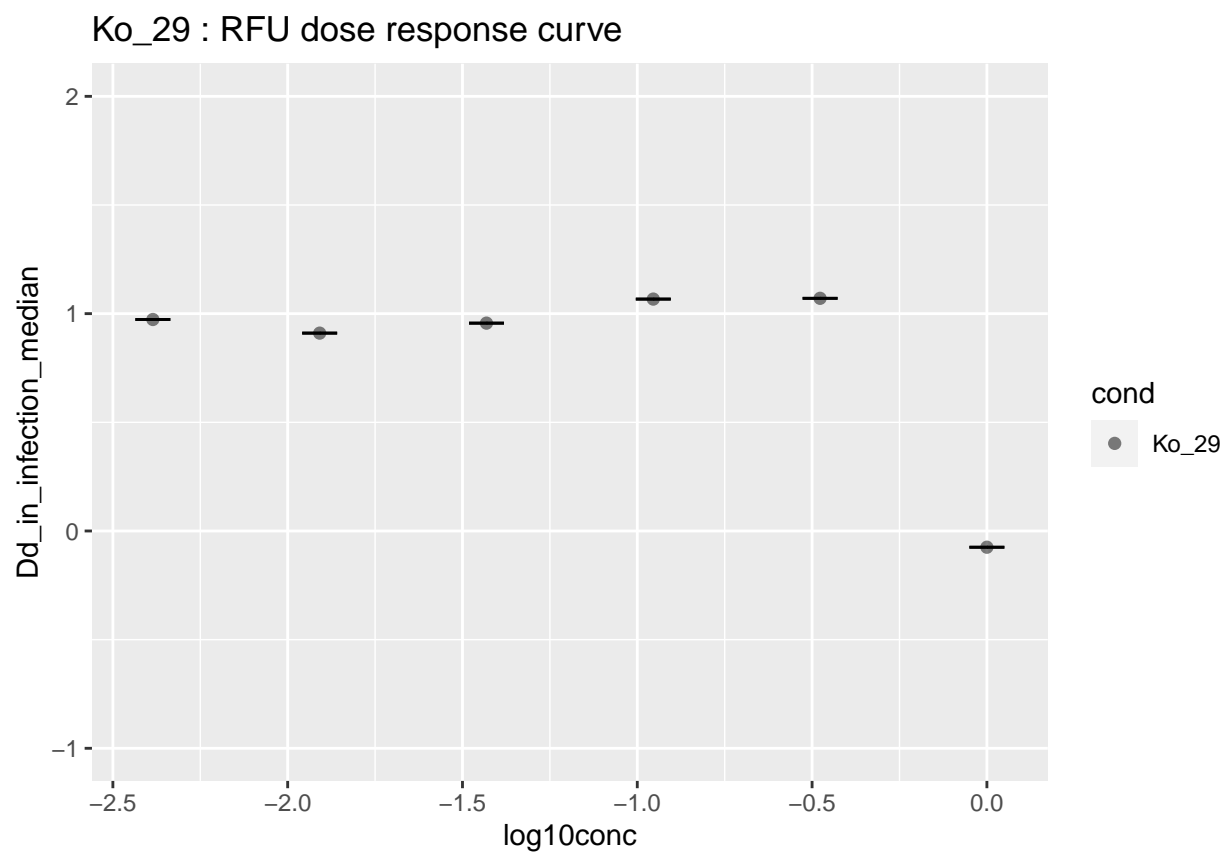


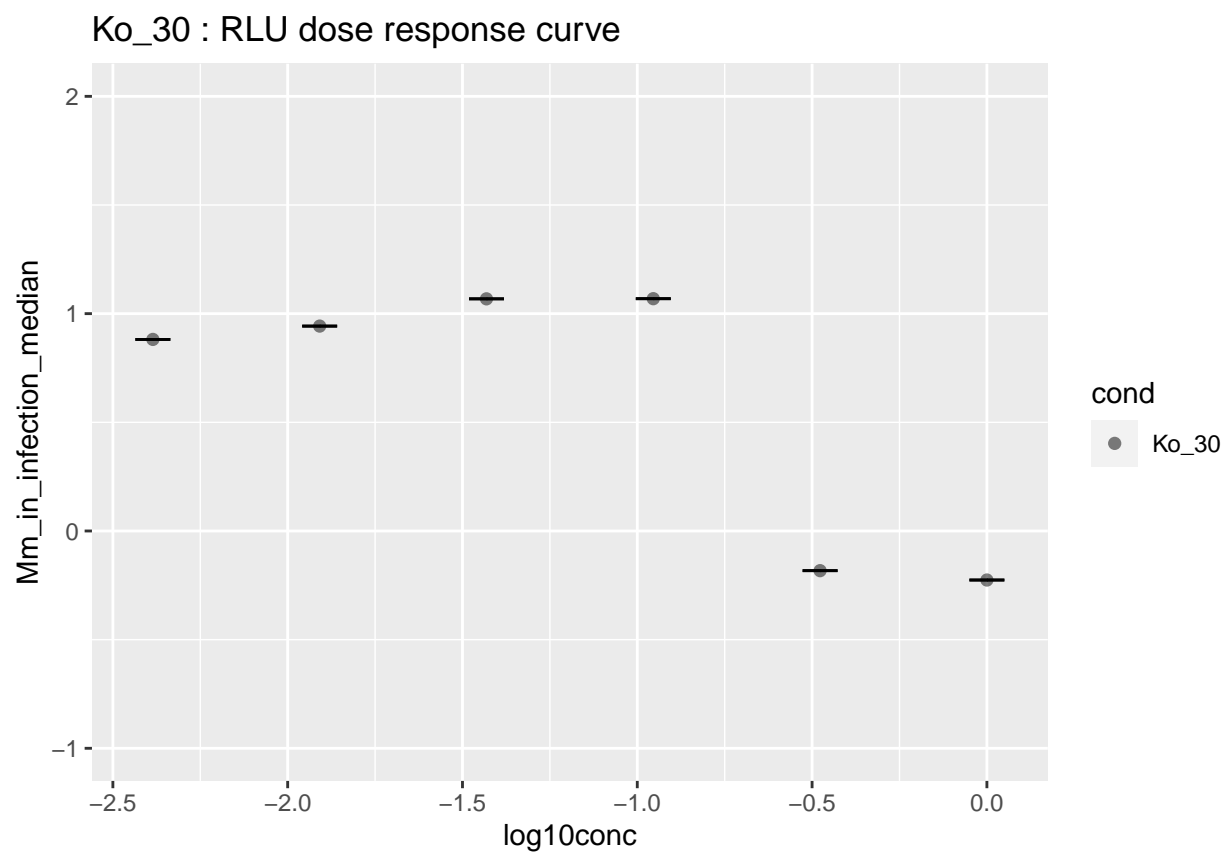


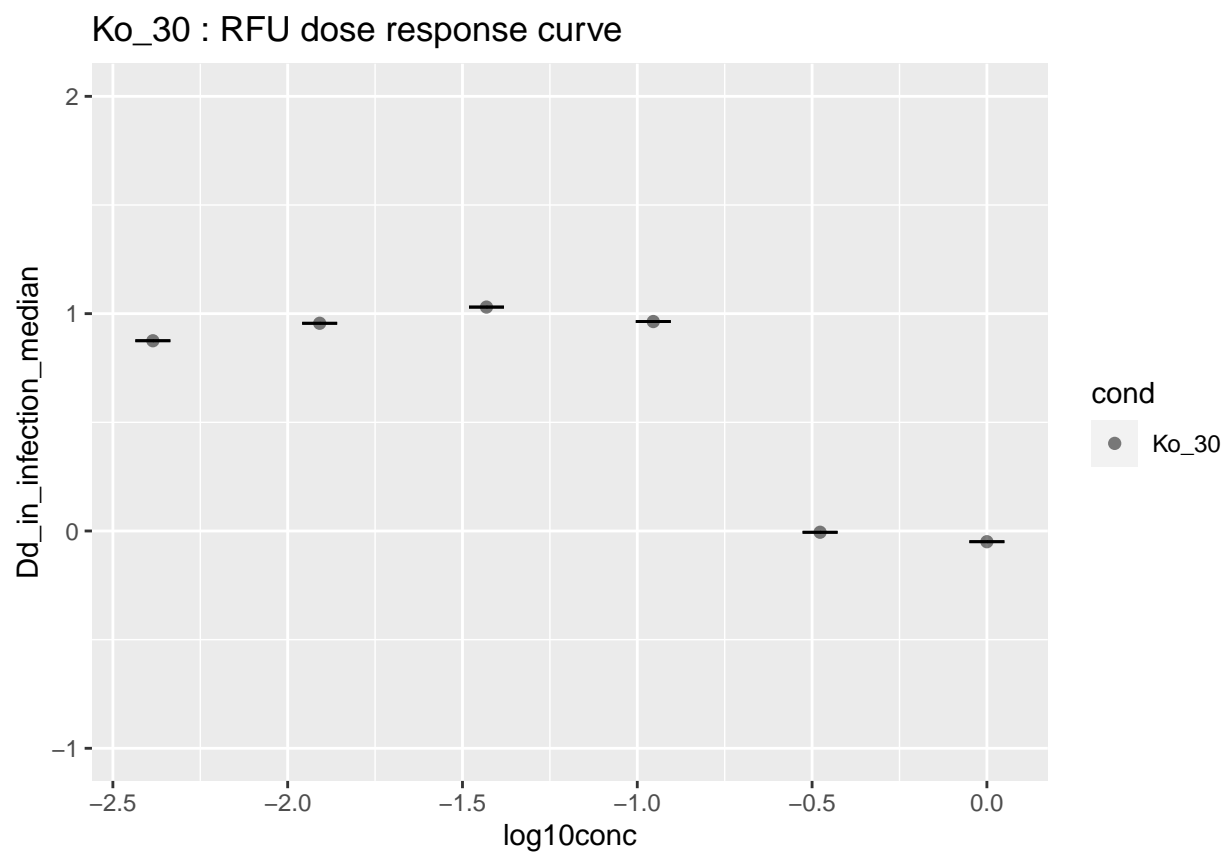


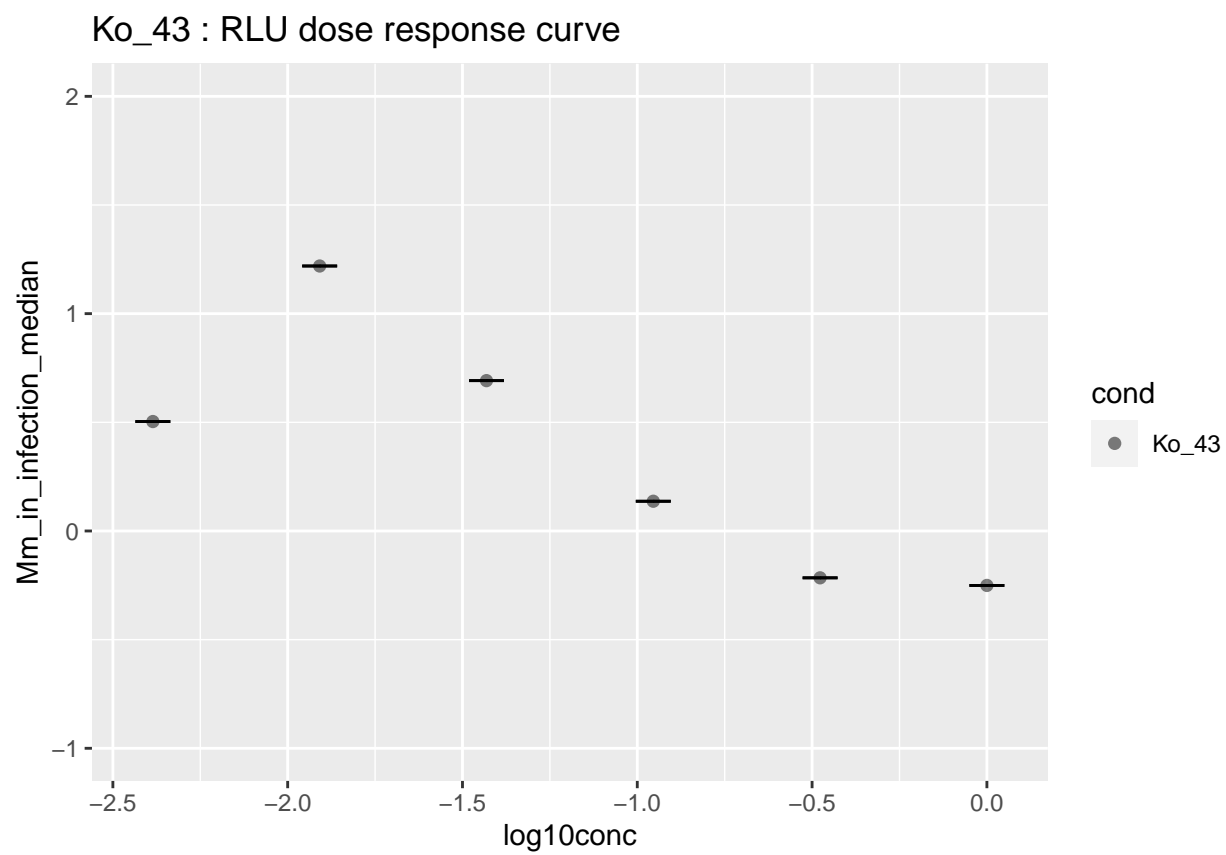


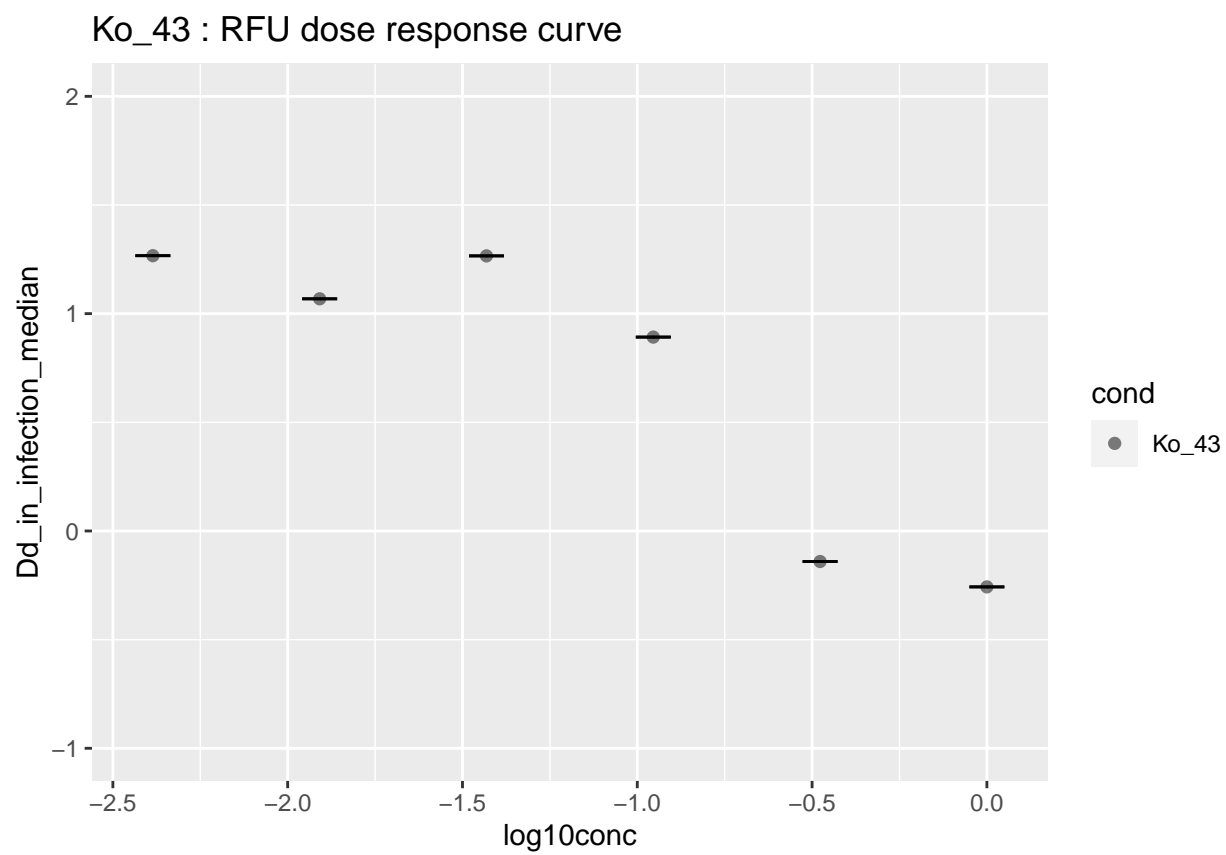


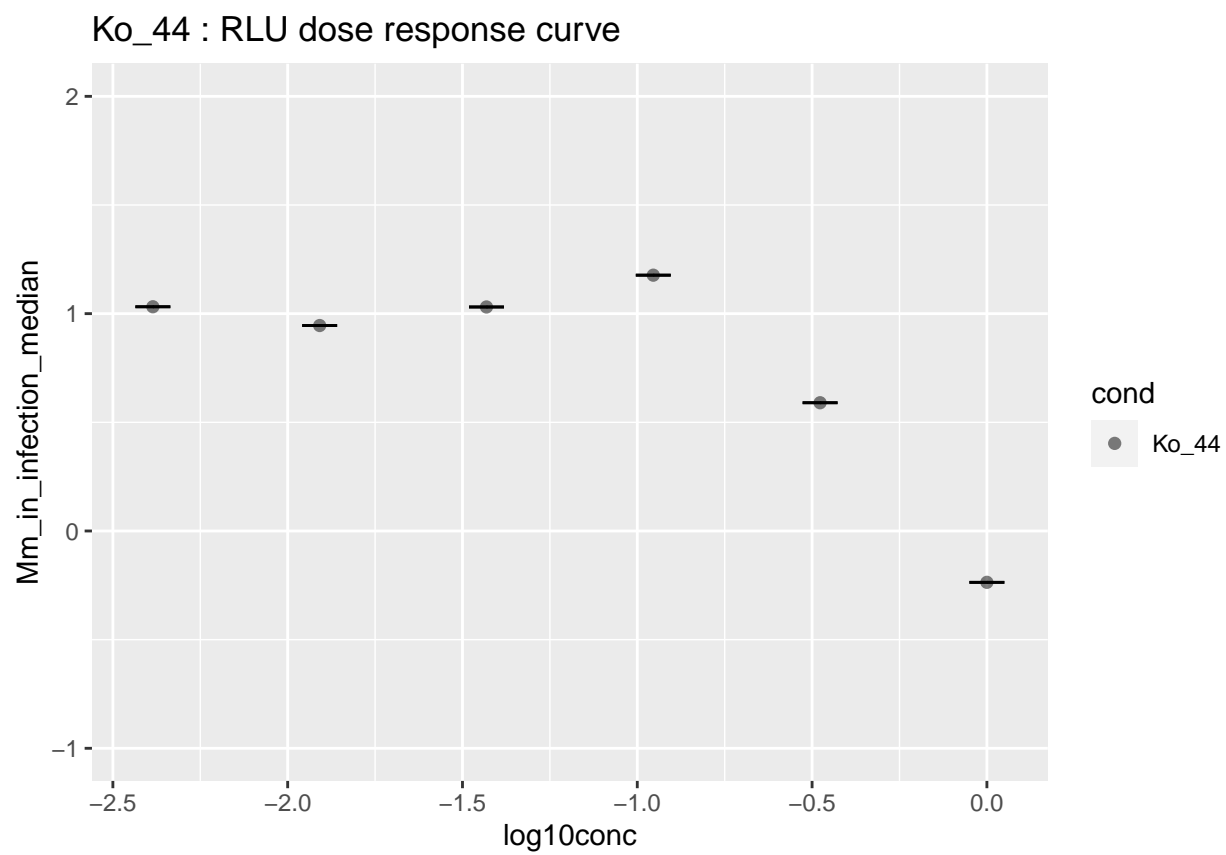


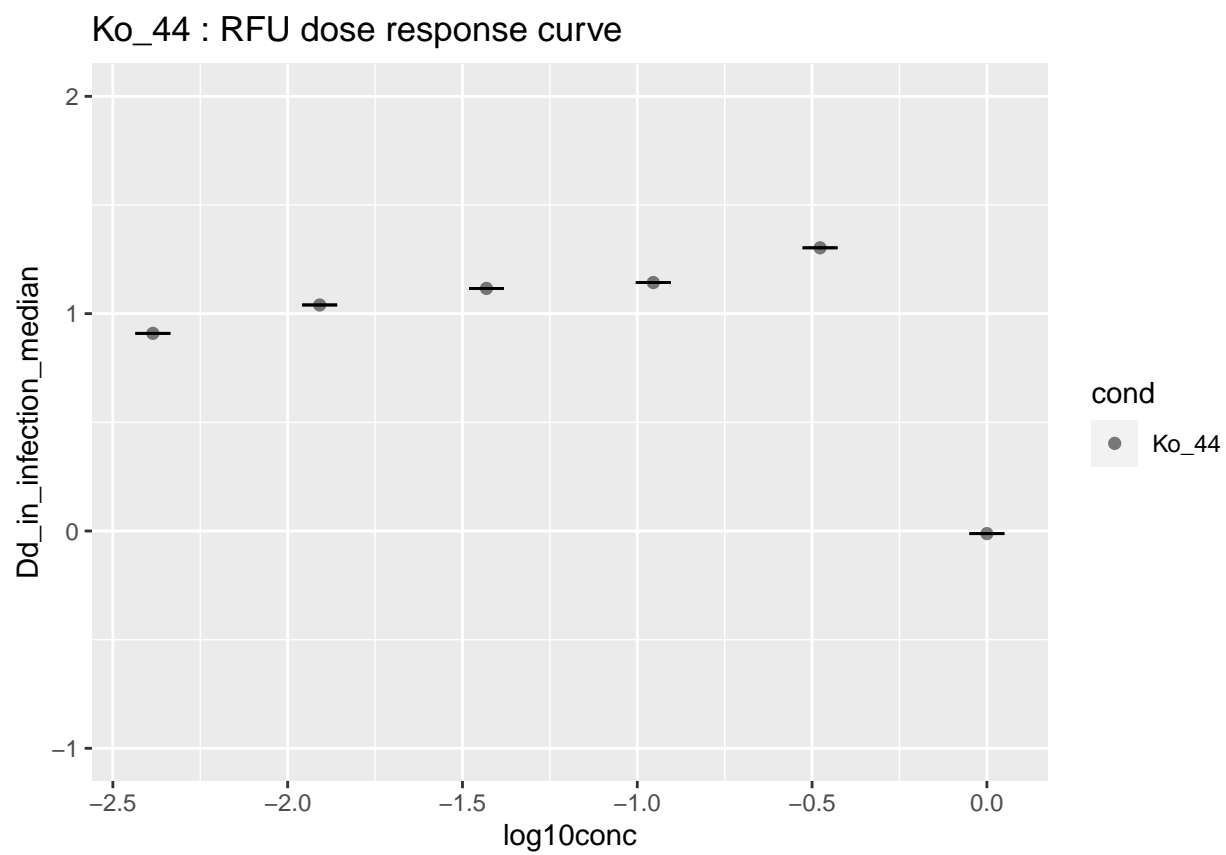


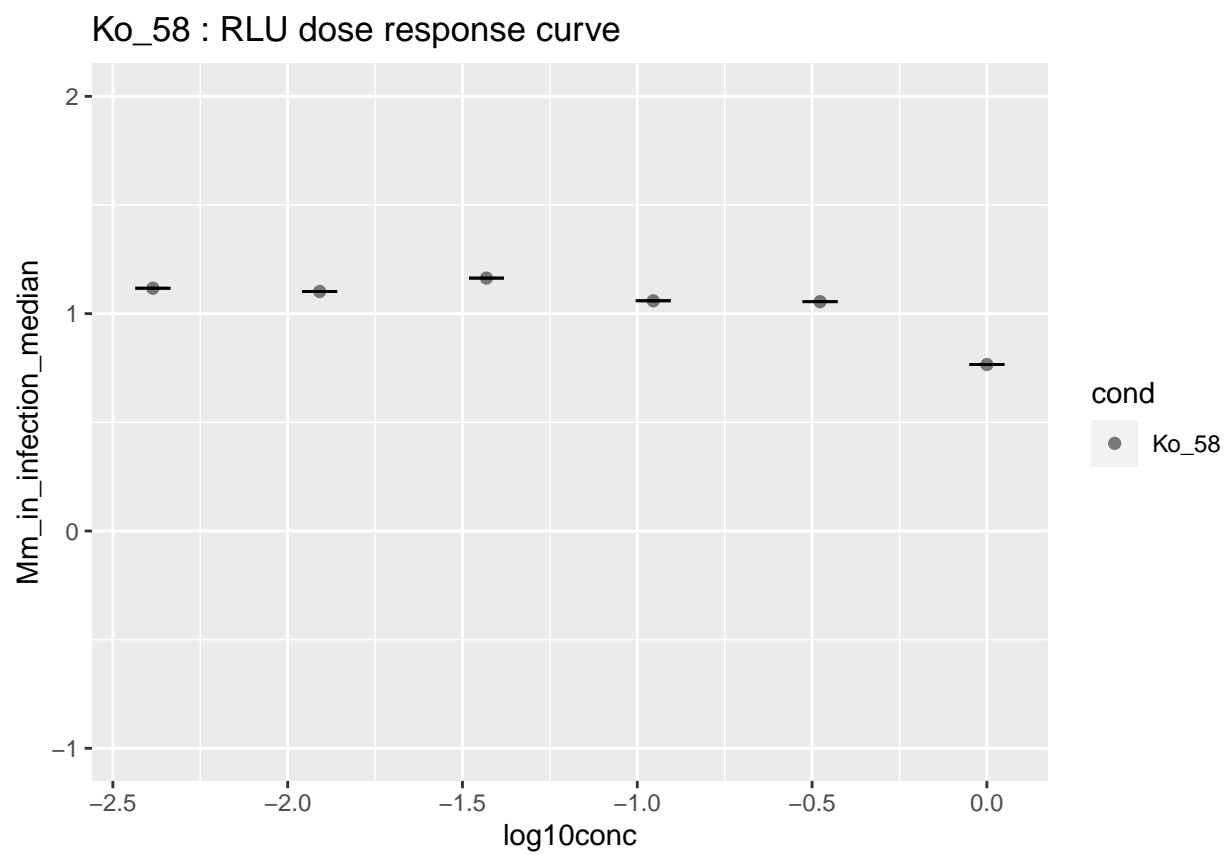


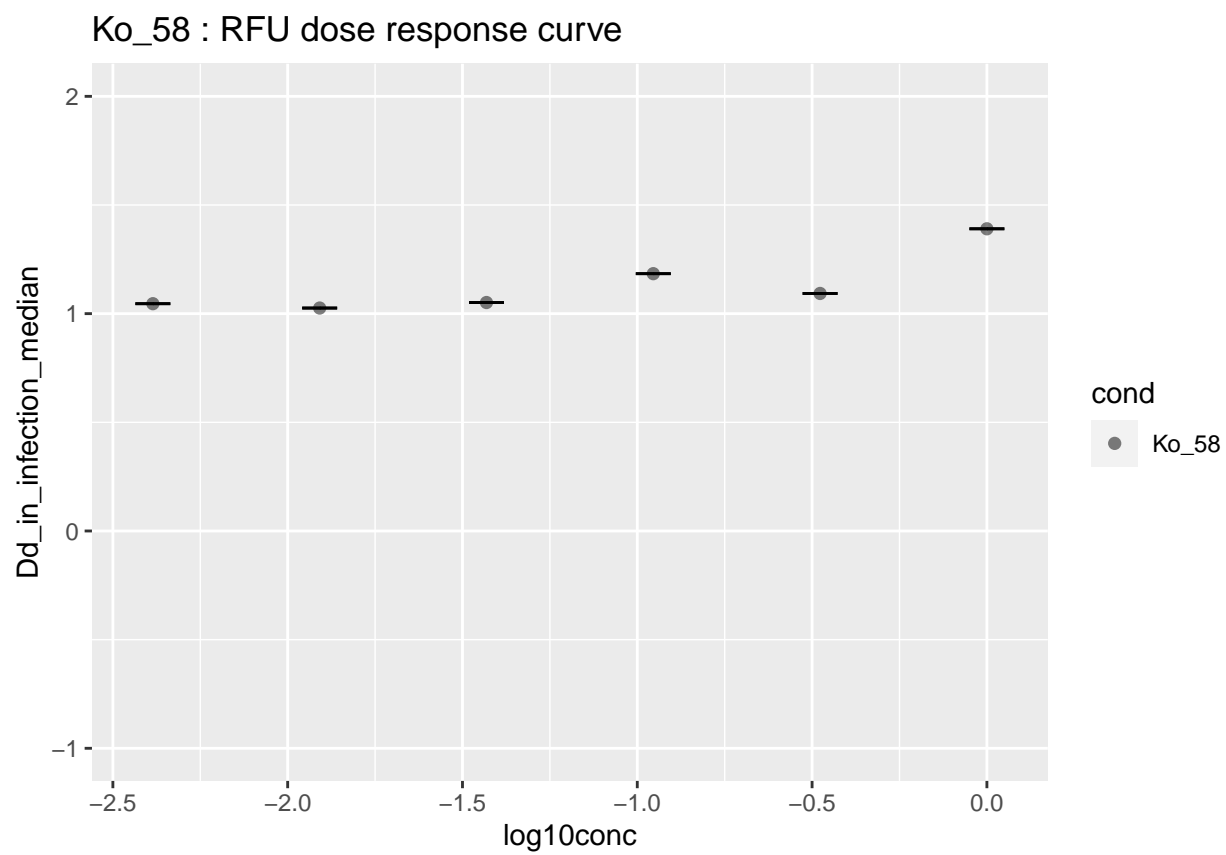


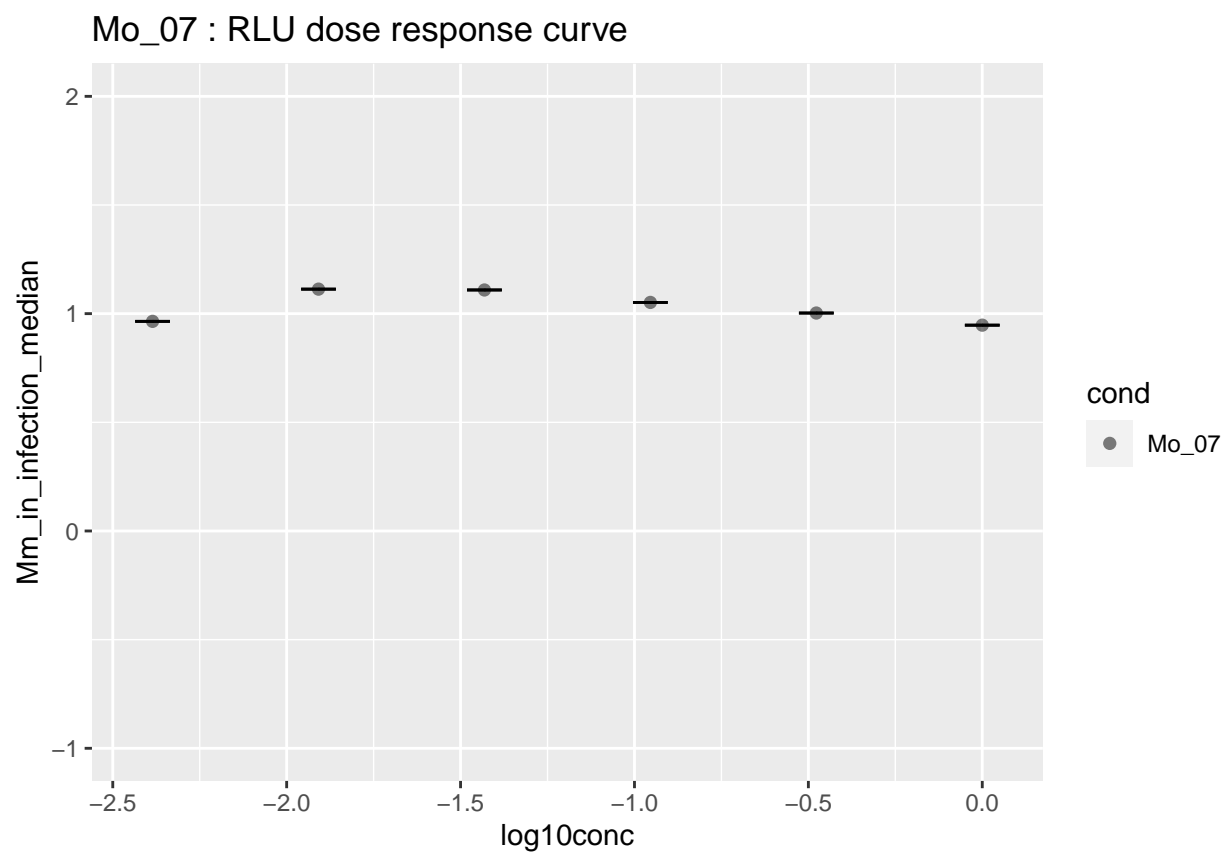


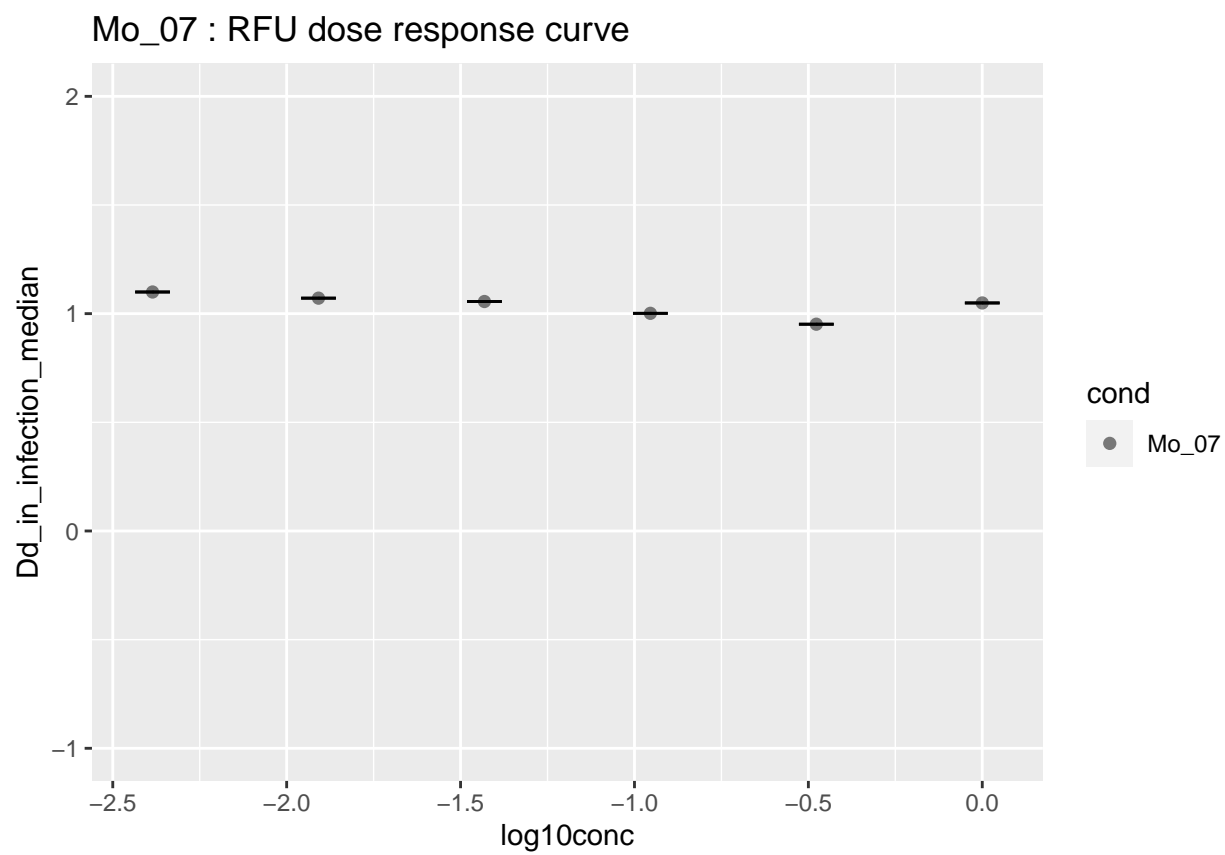


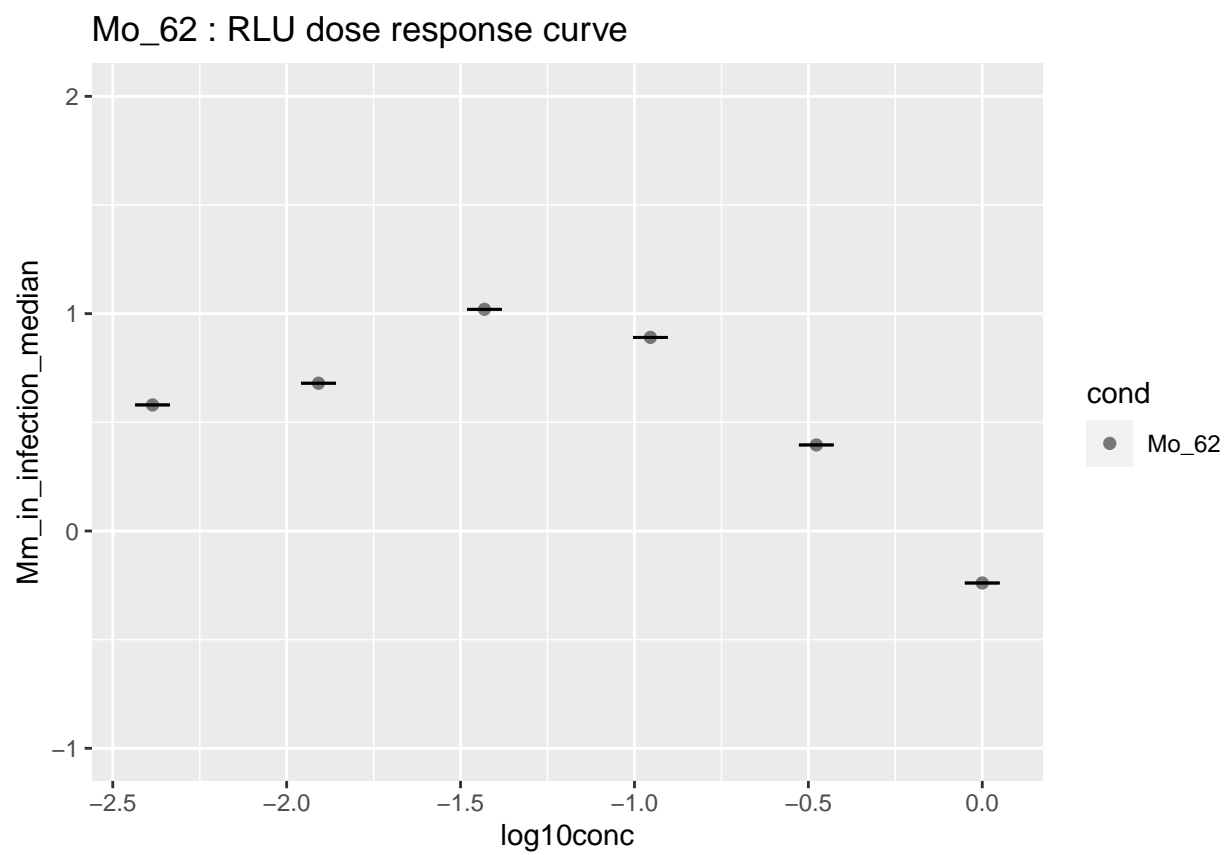


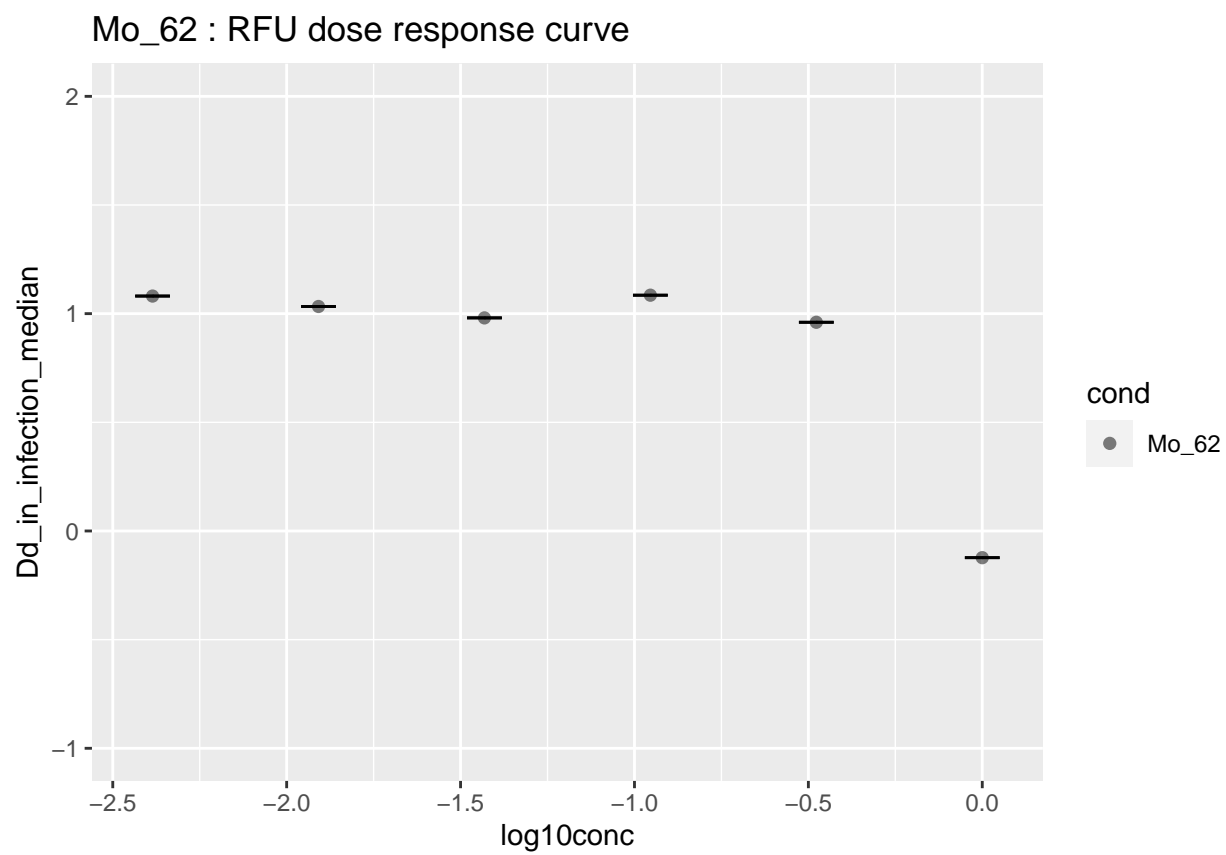


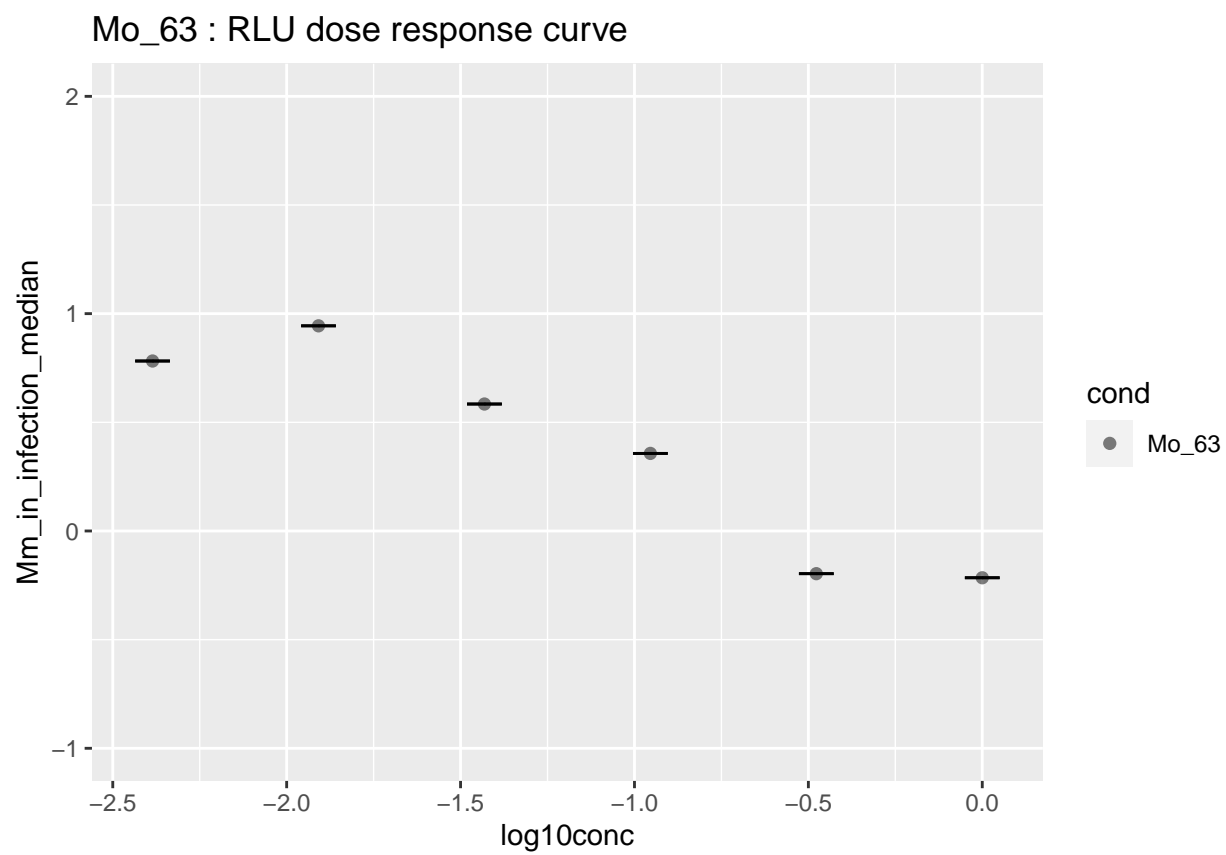


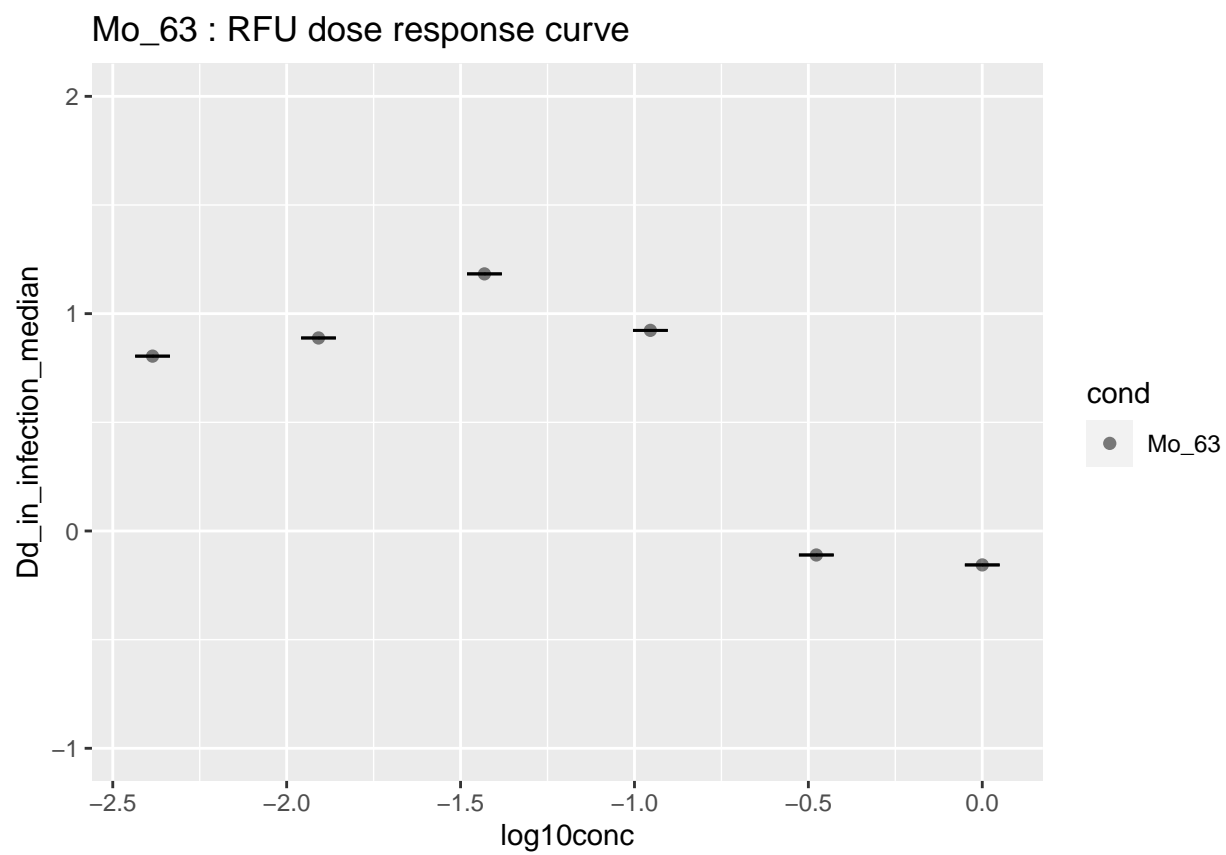


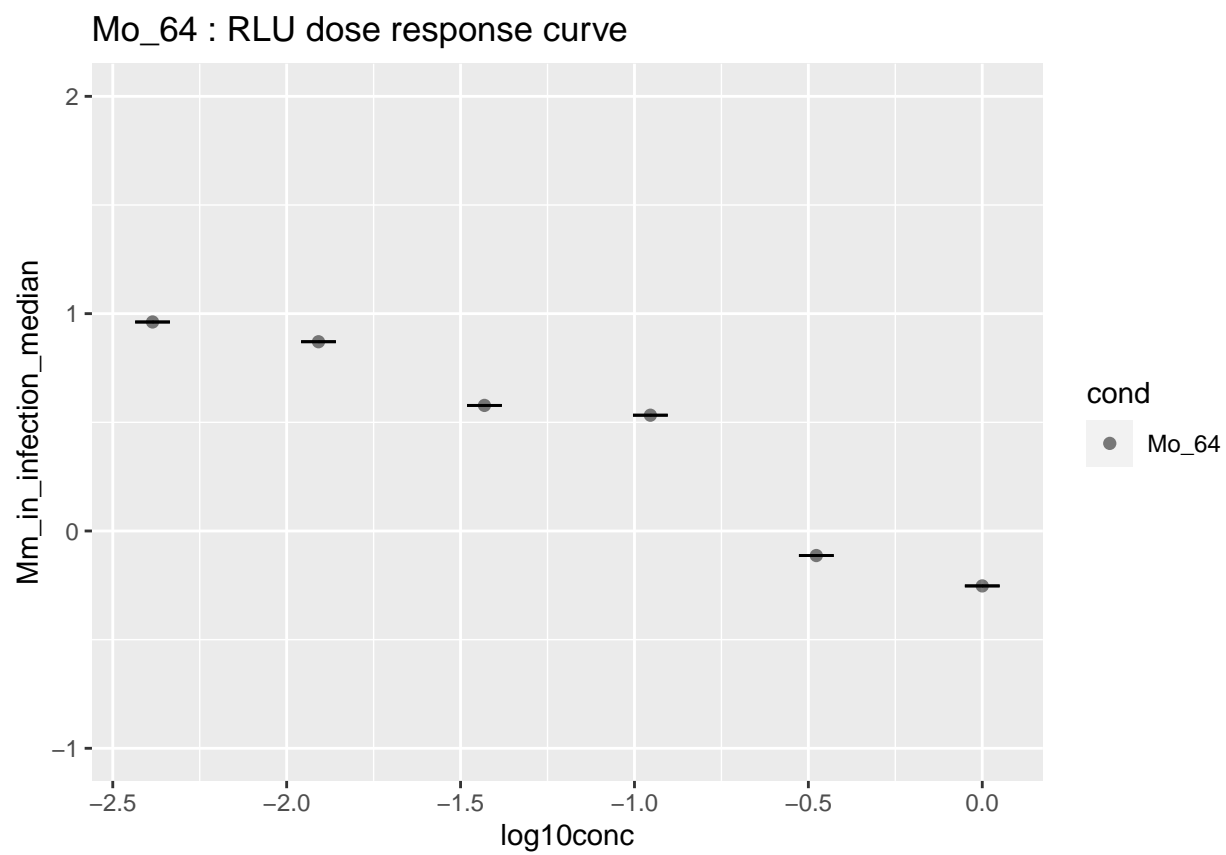


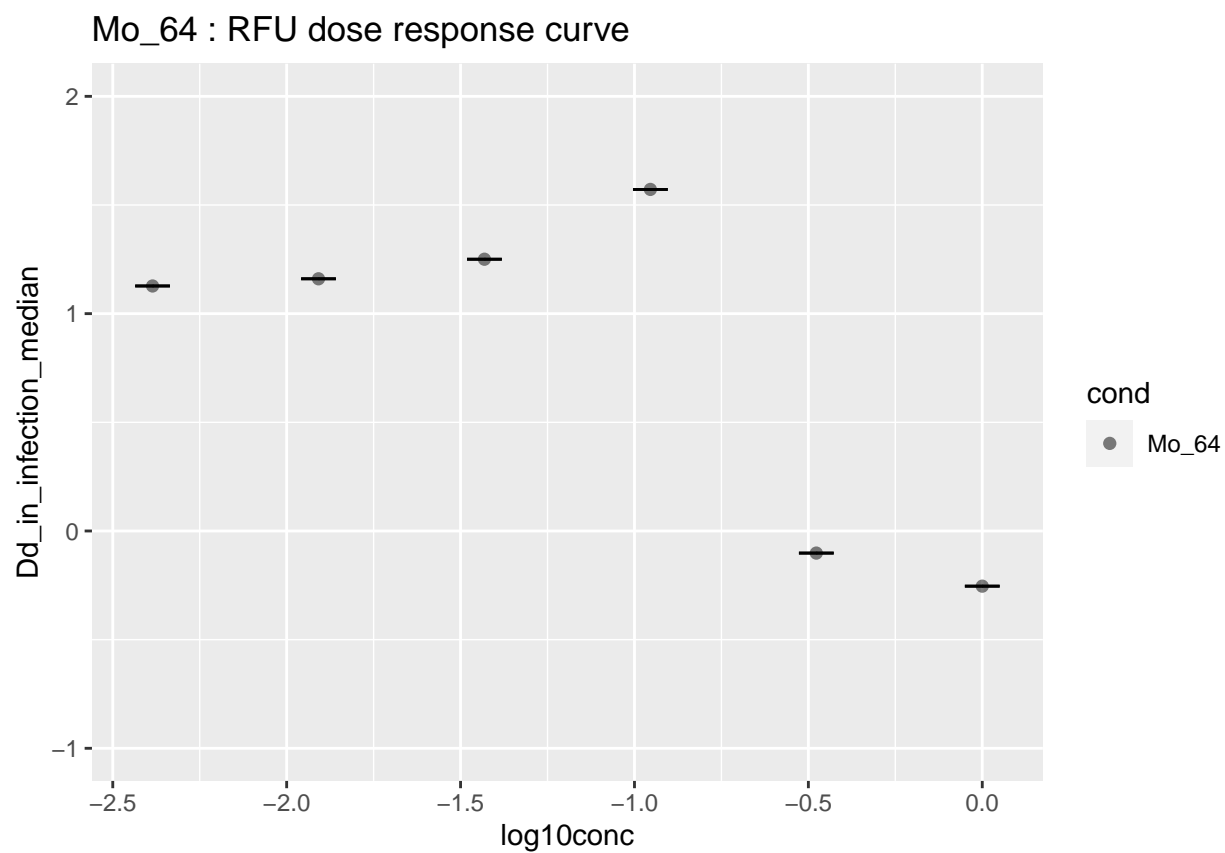


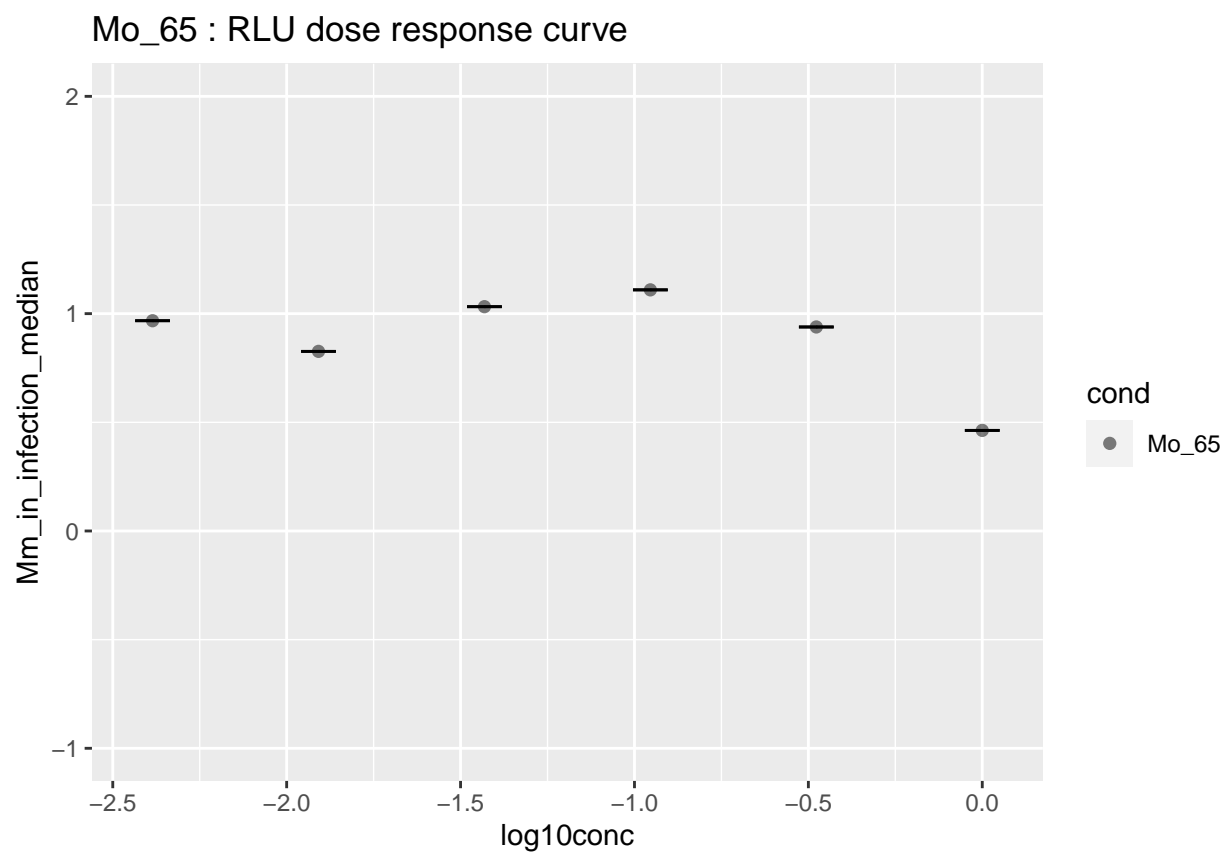


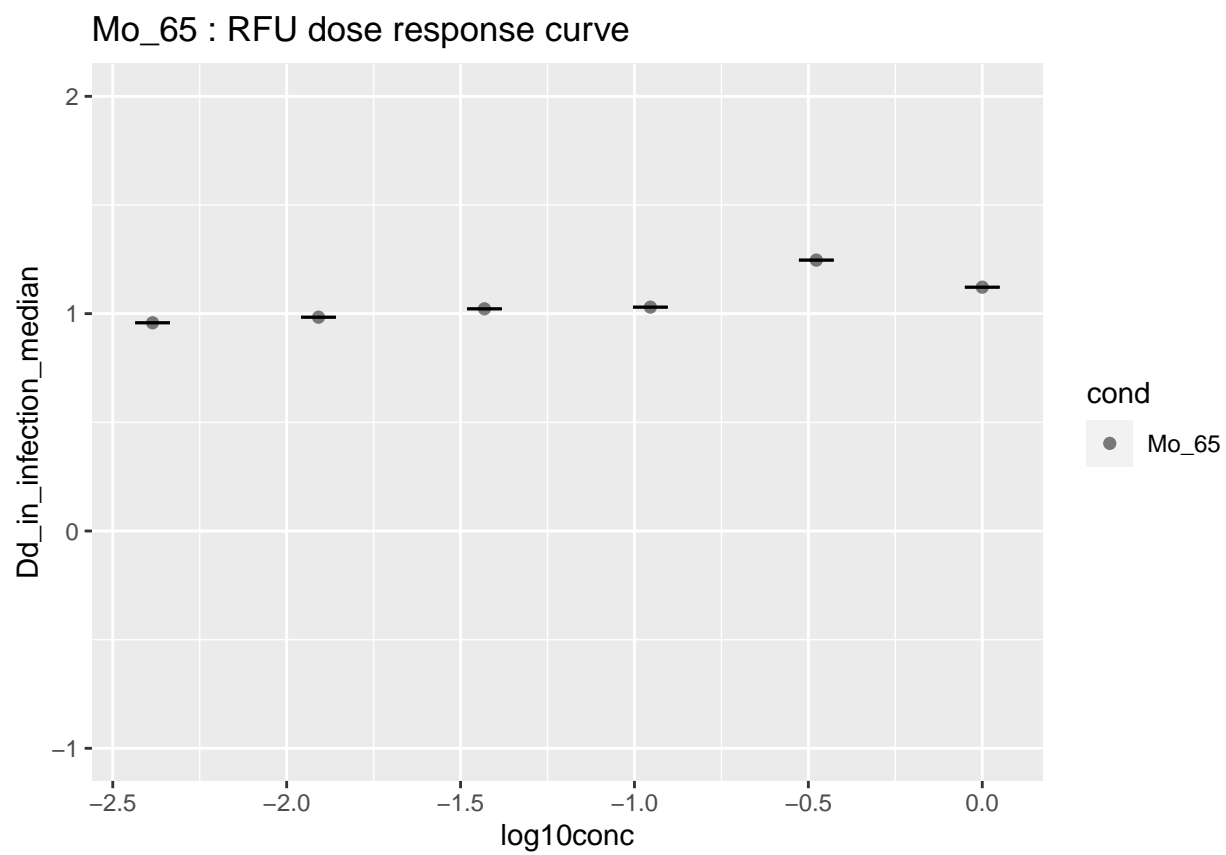


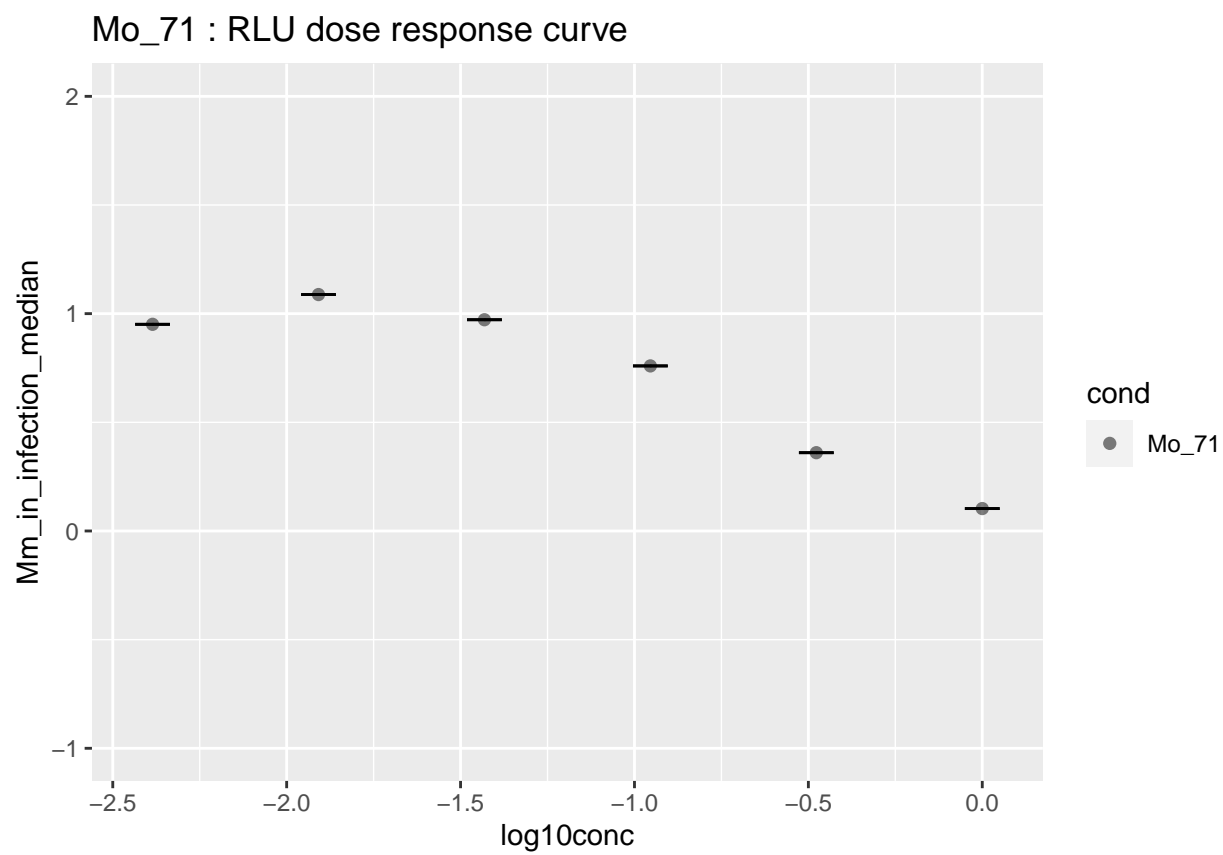


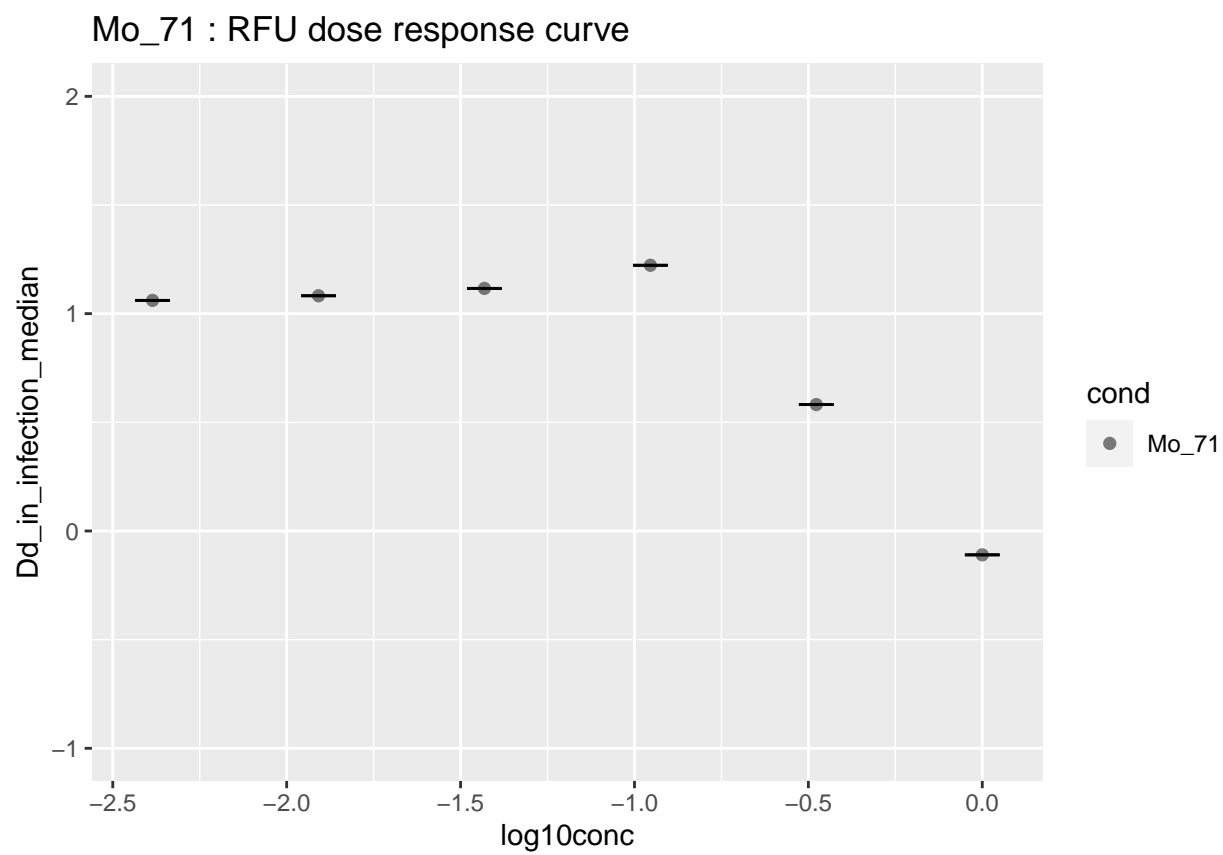


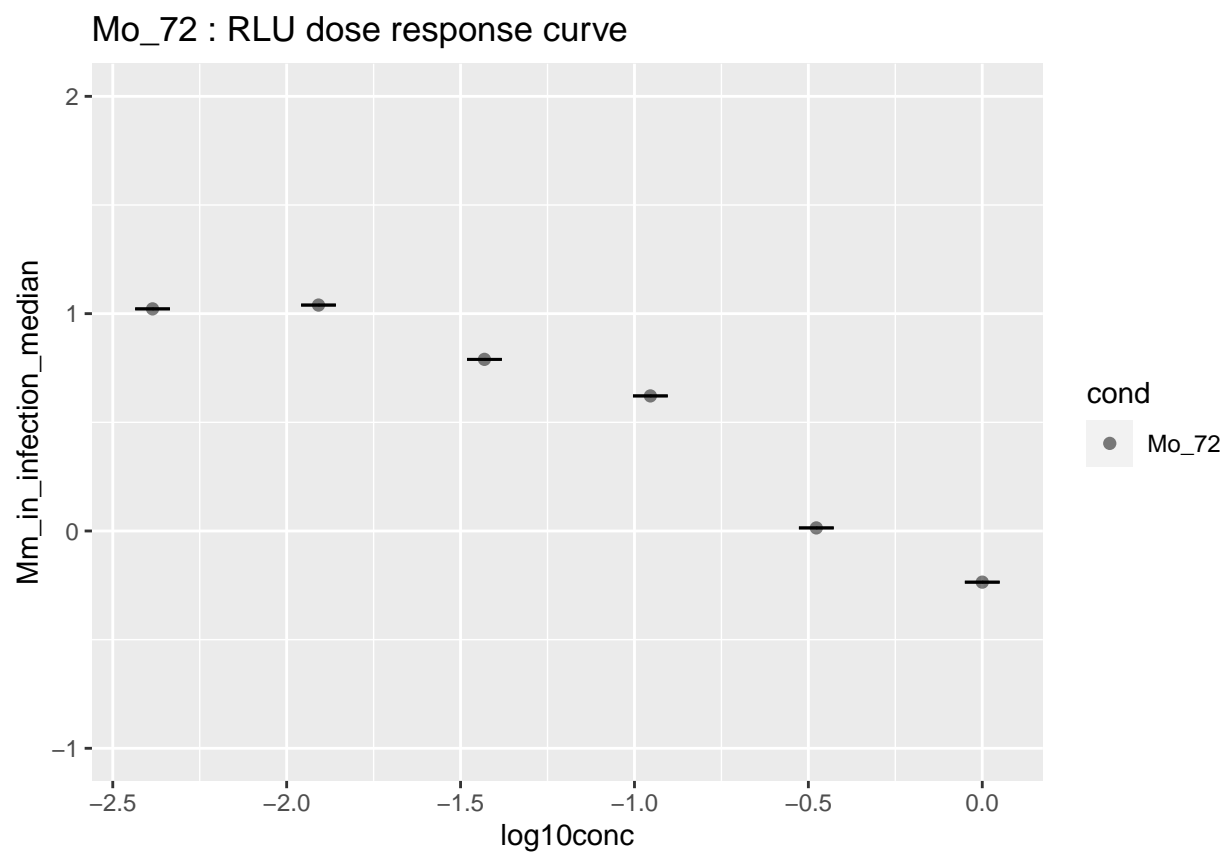


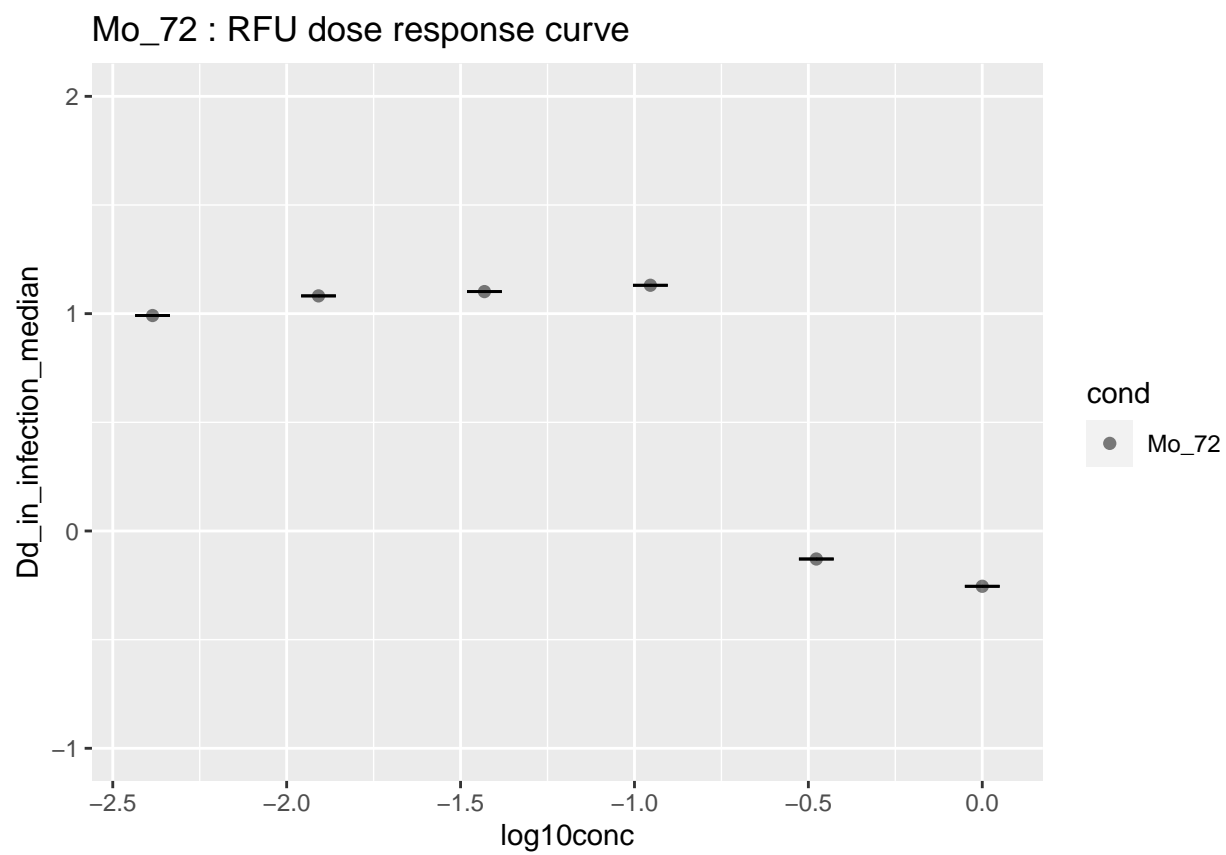


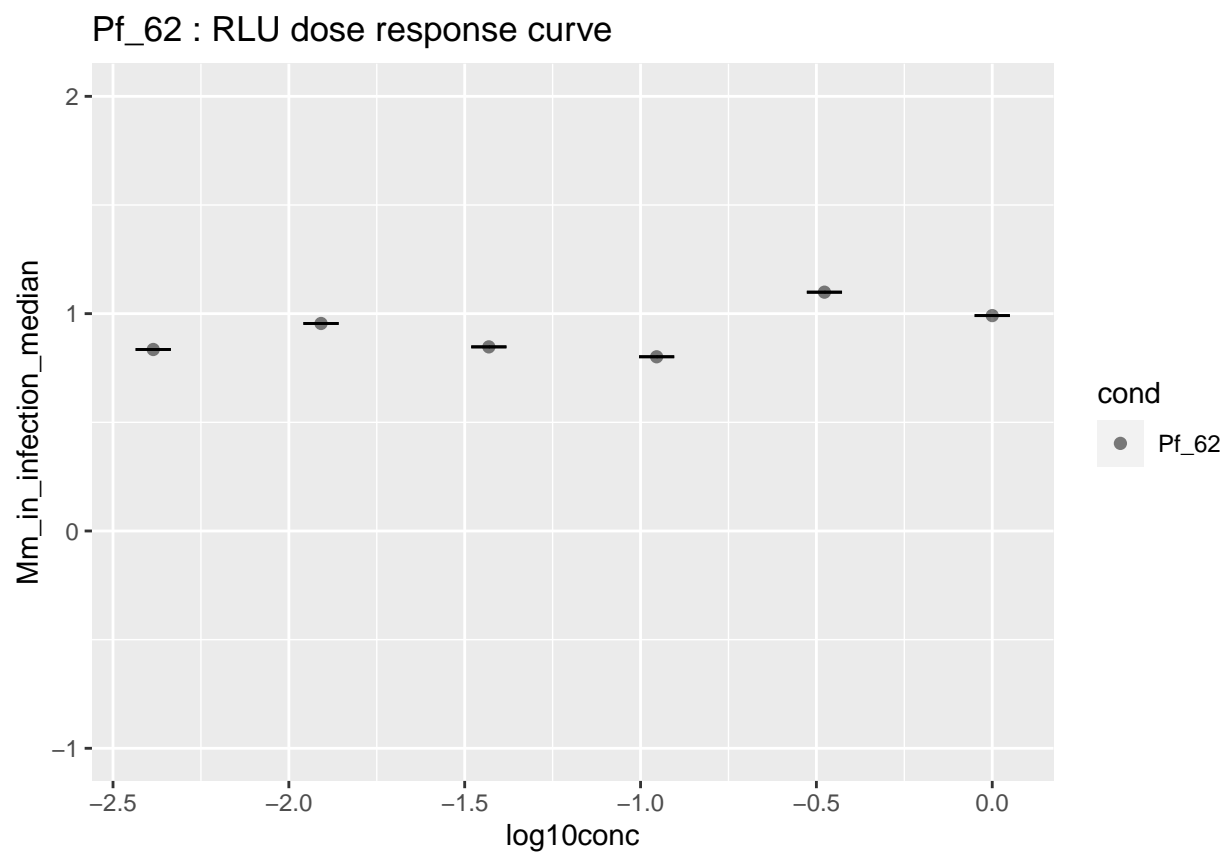


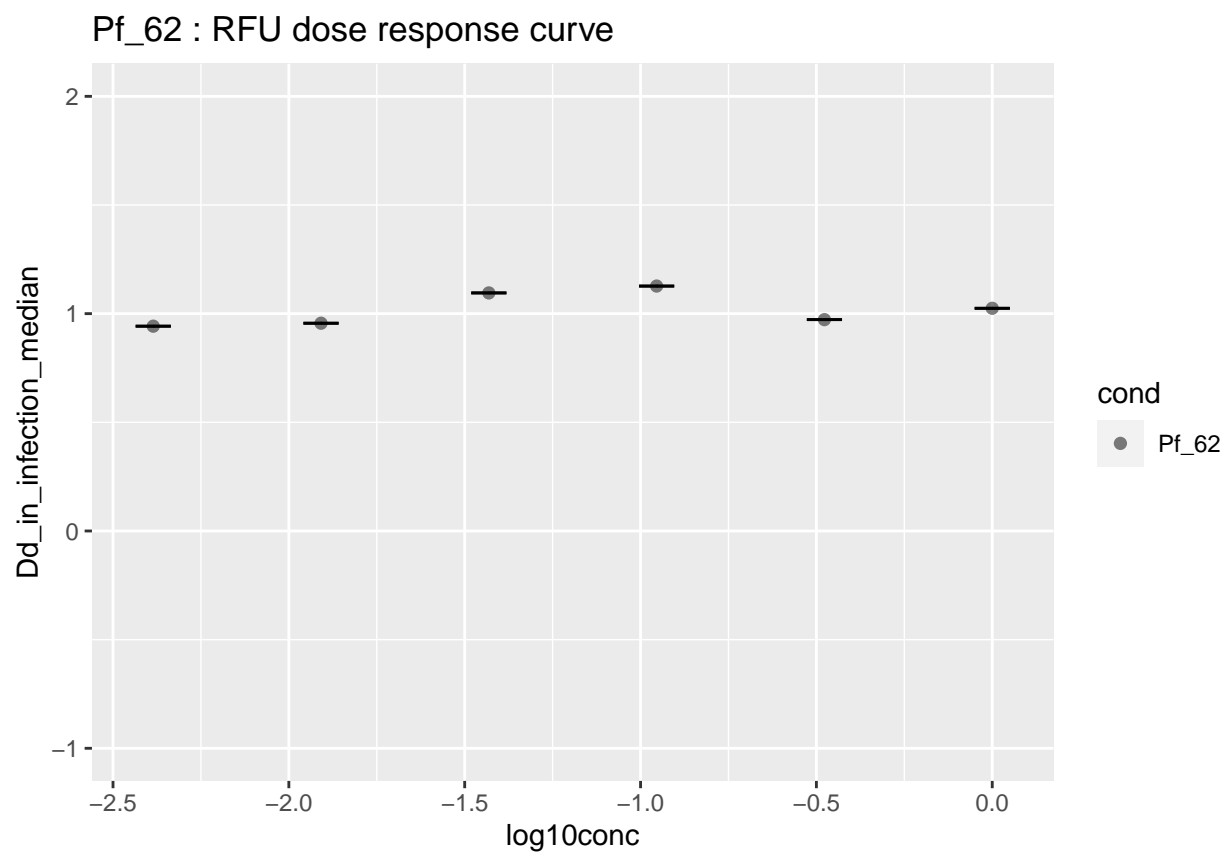




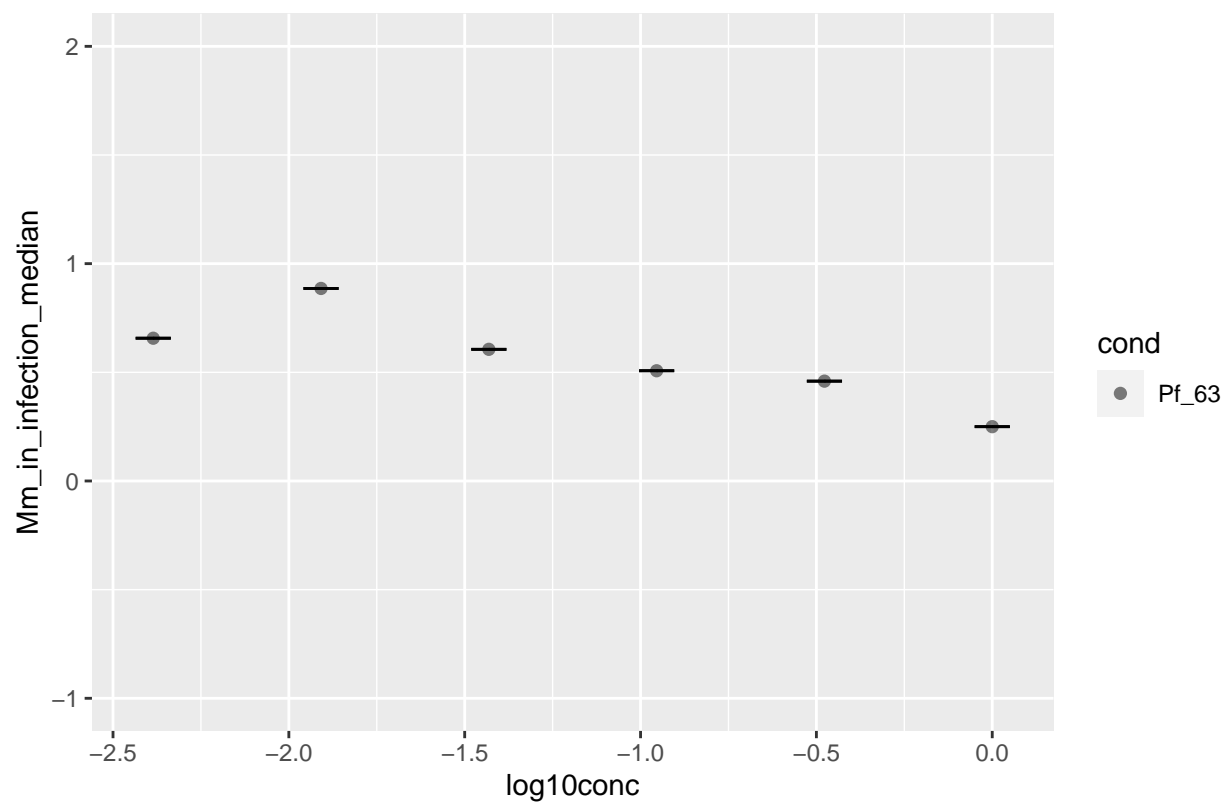


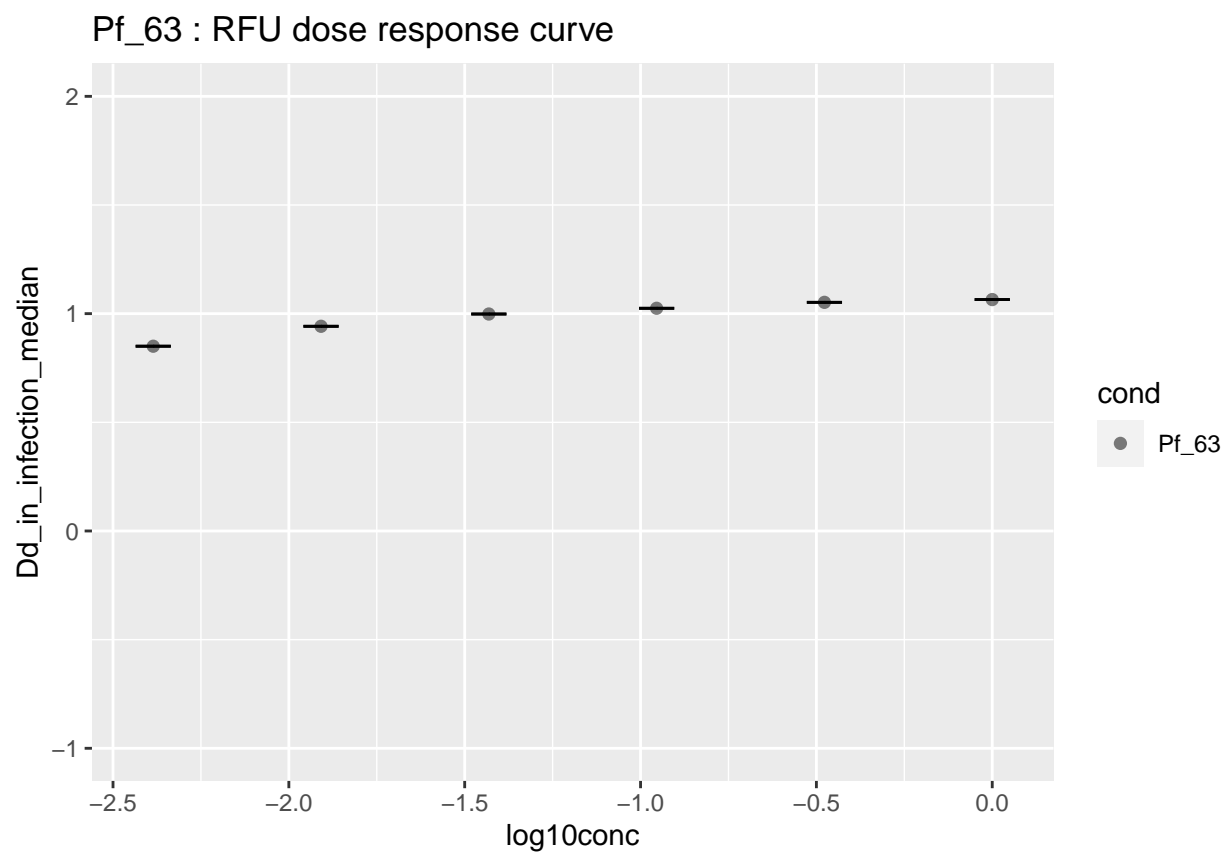


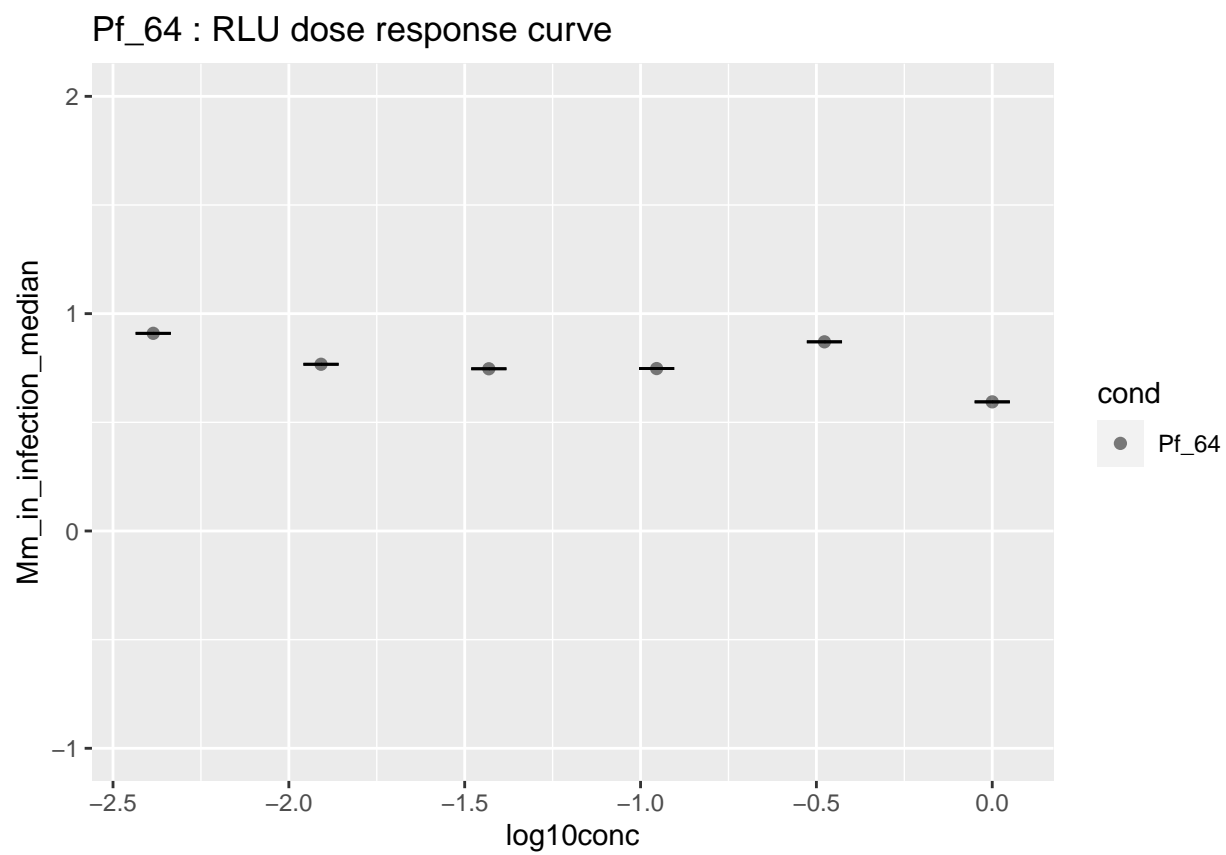


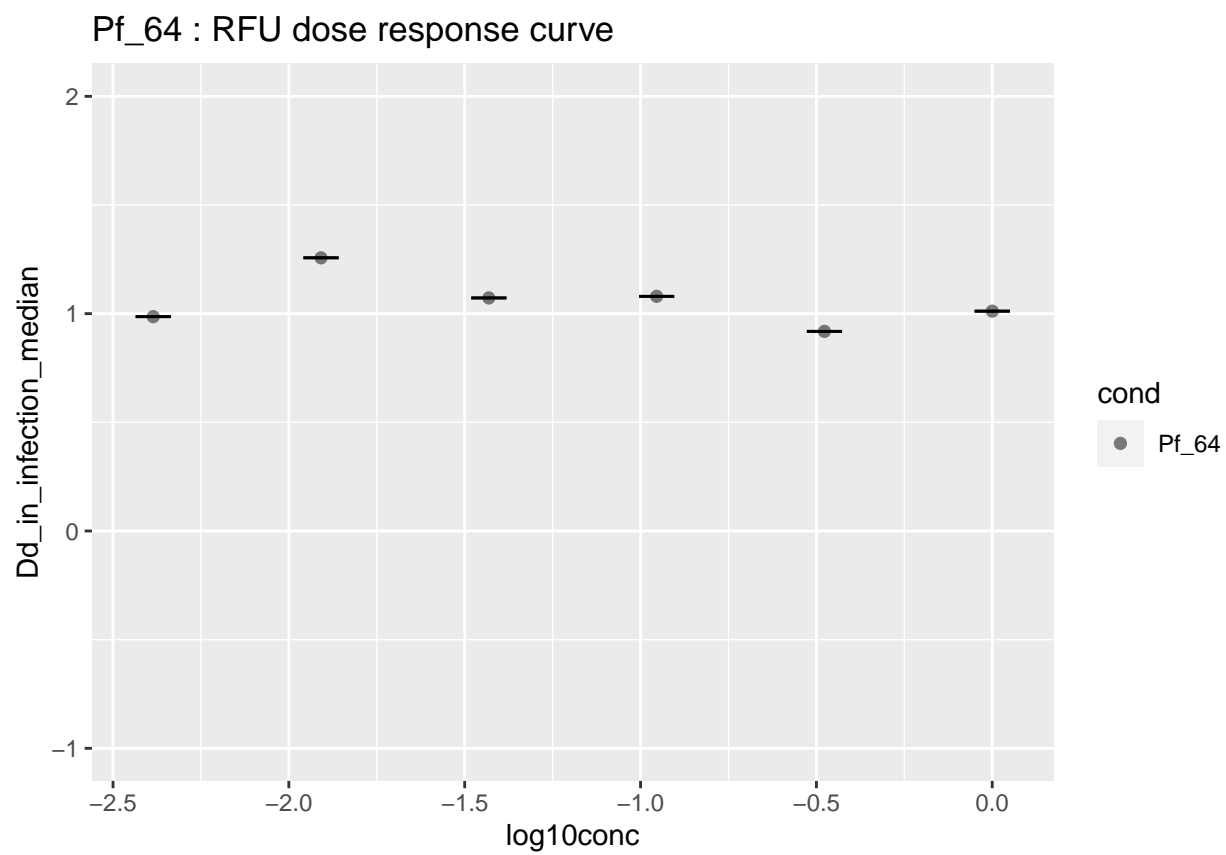


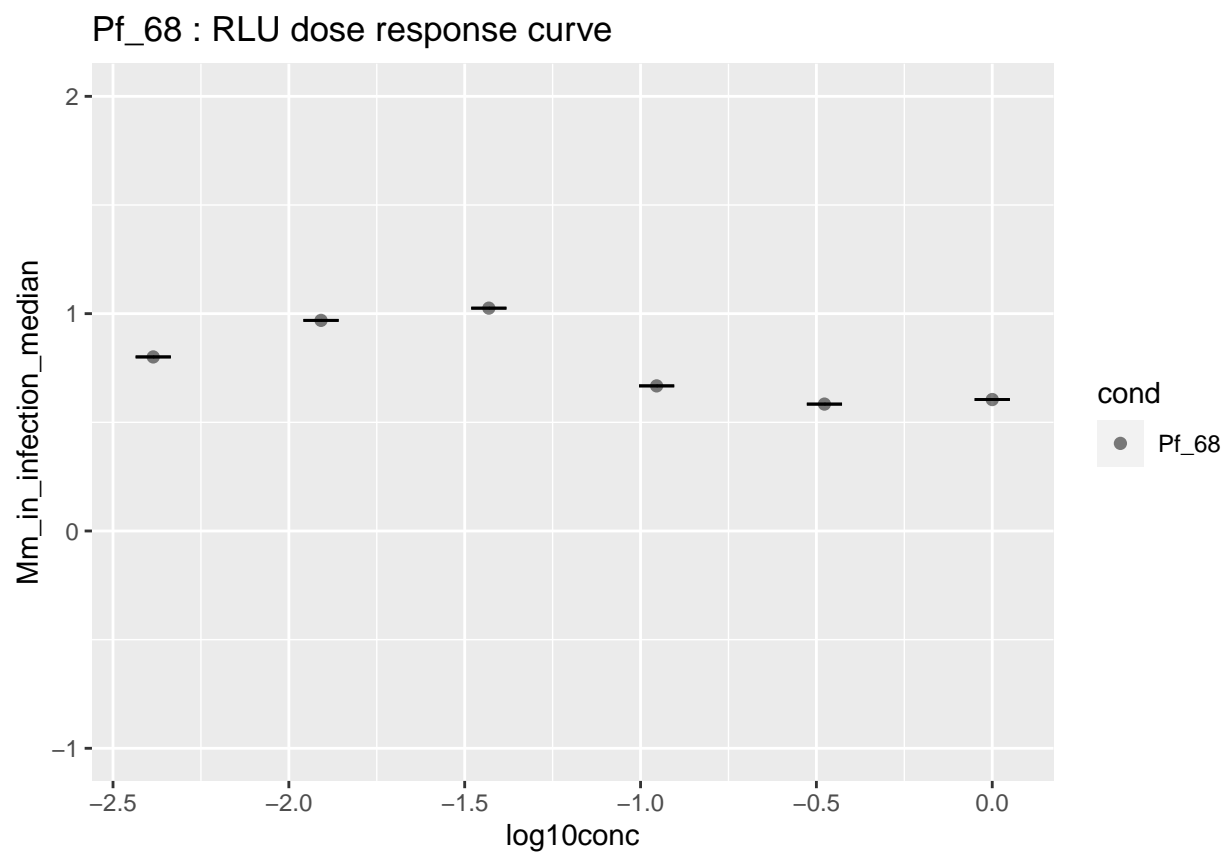
Pf_63 : RLU dose response curve



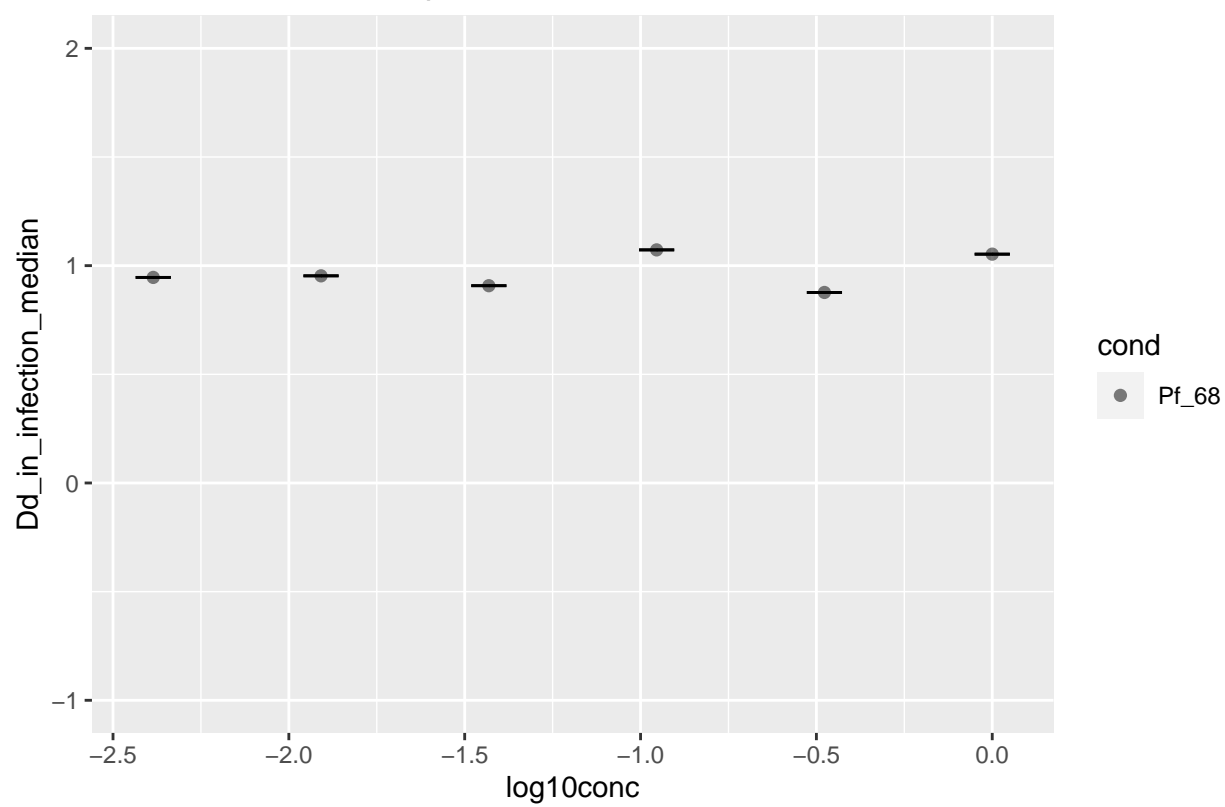




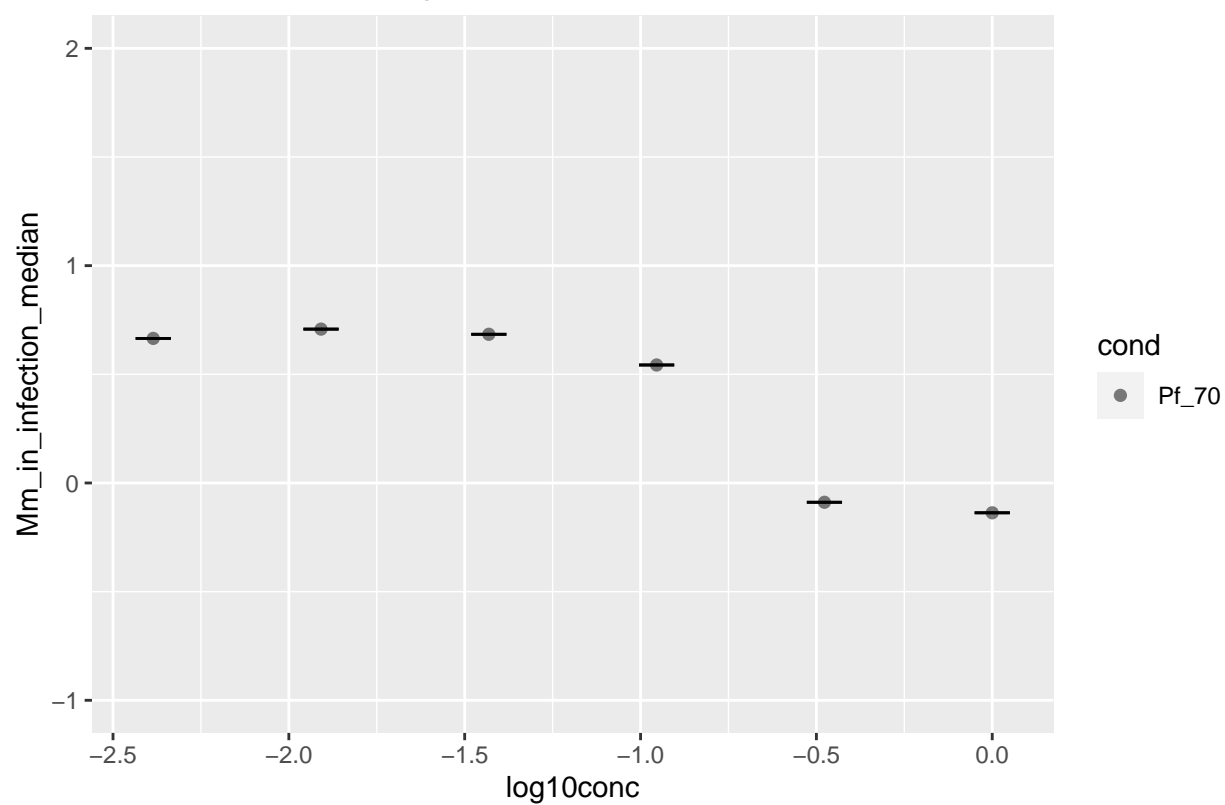


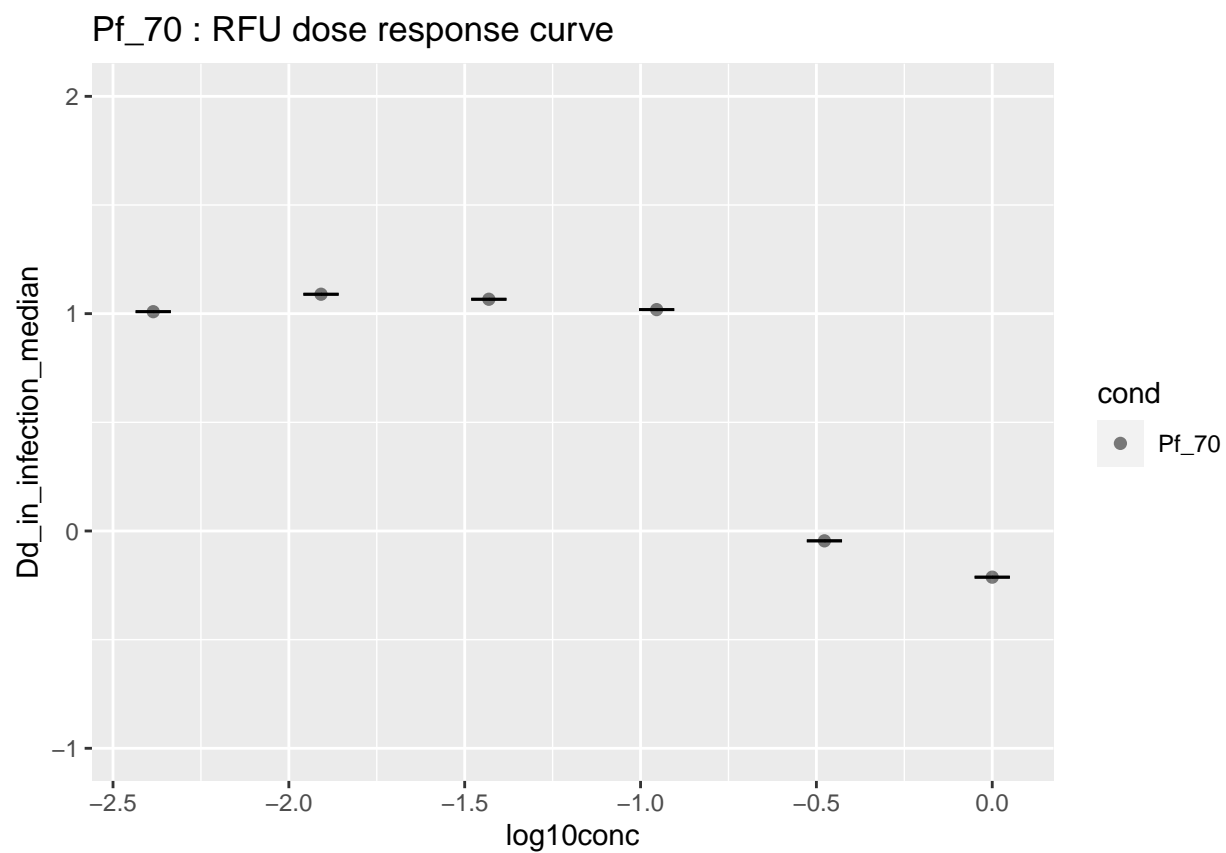


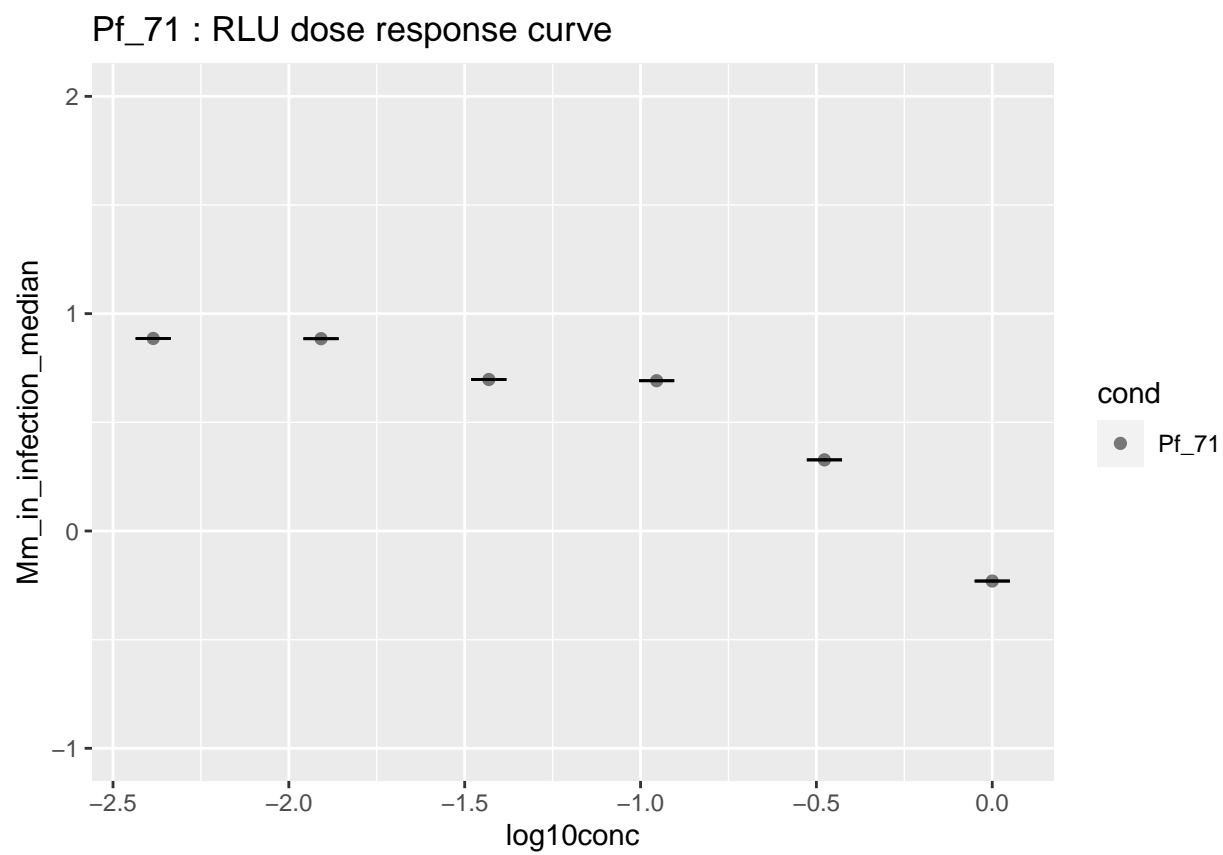
Pf_68 : RFU dose response curve

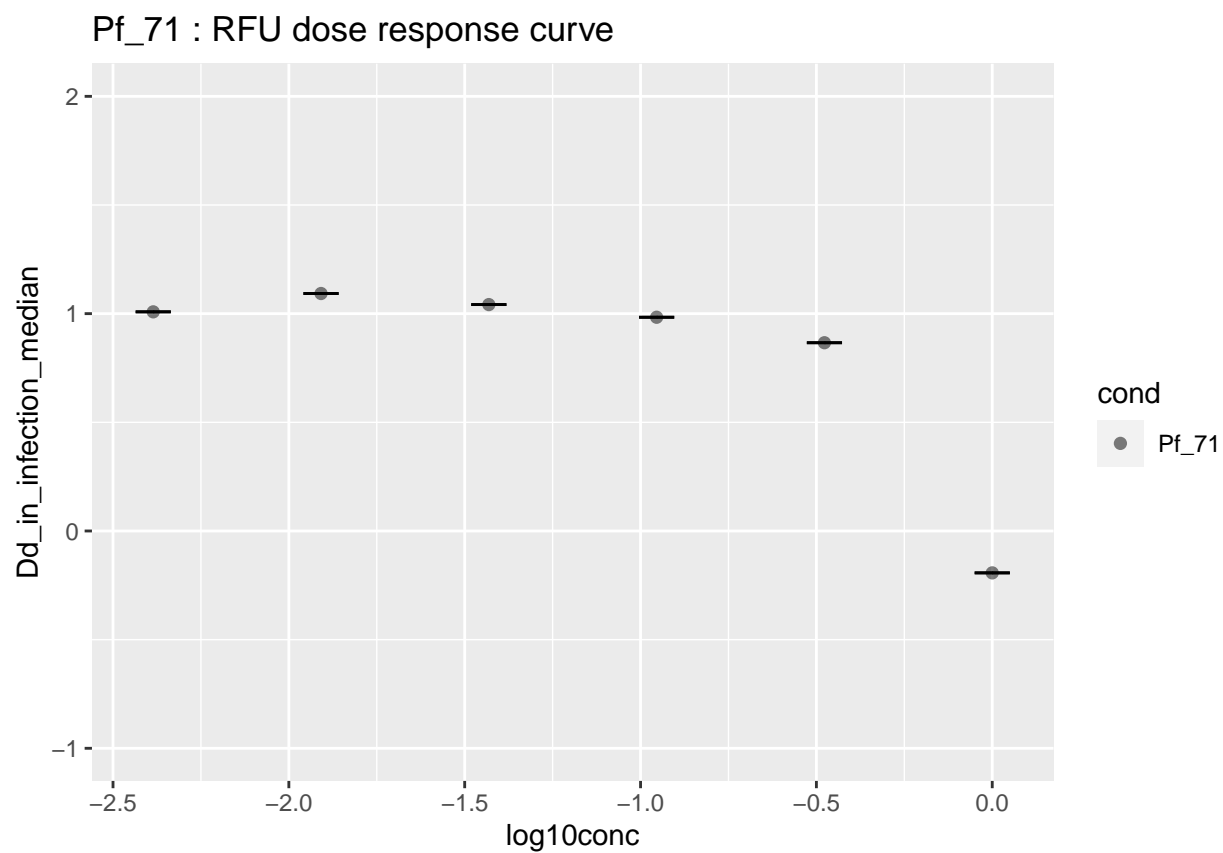


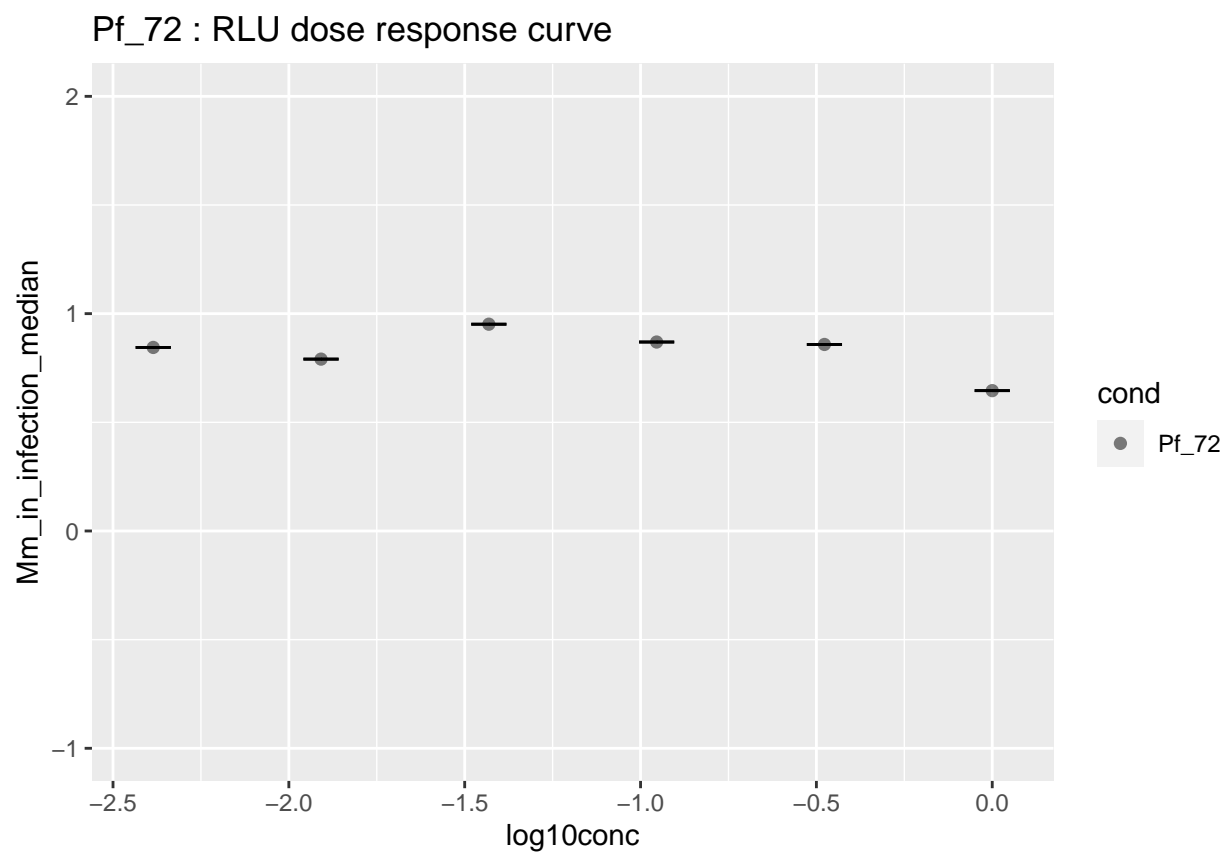
Pf_70 : RLU dose response curve

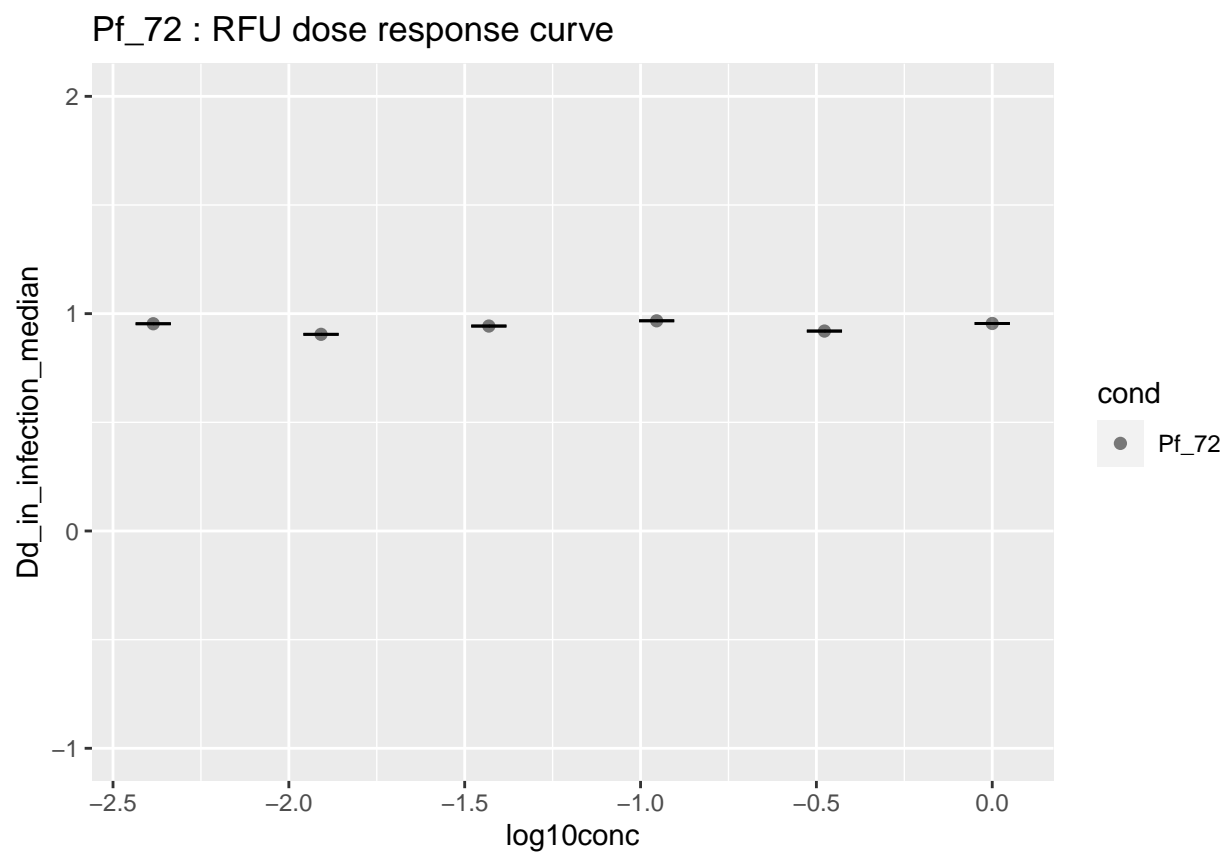


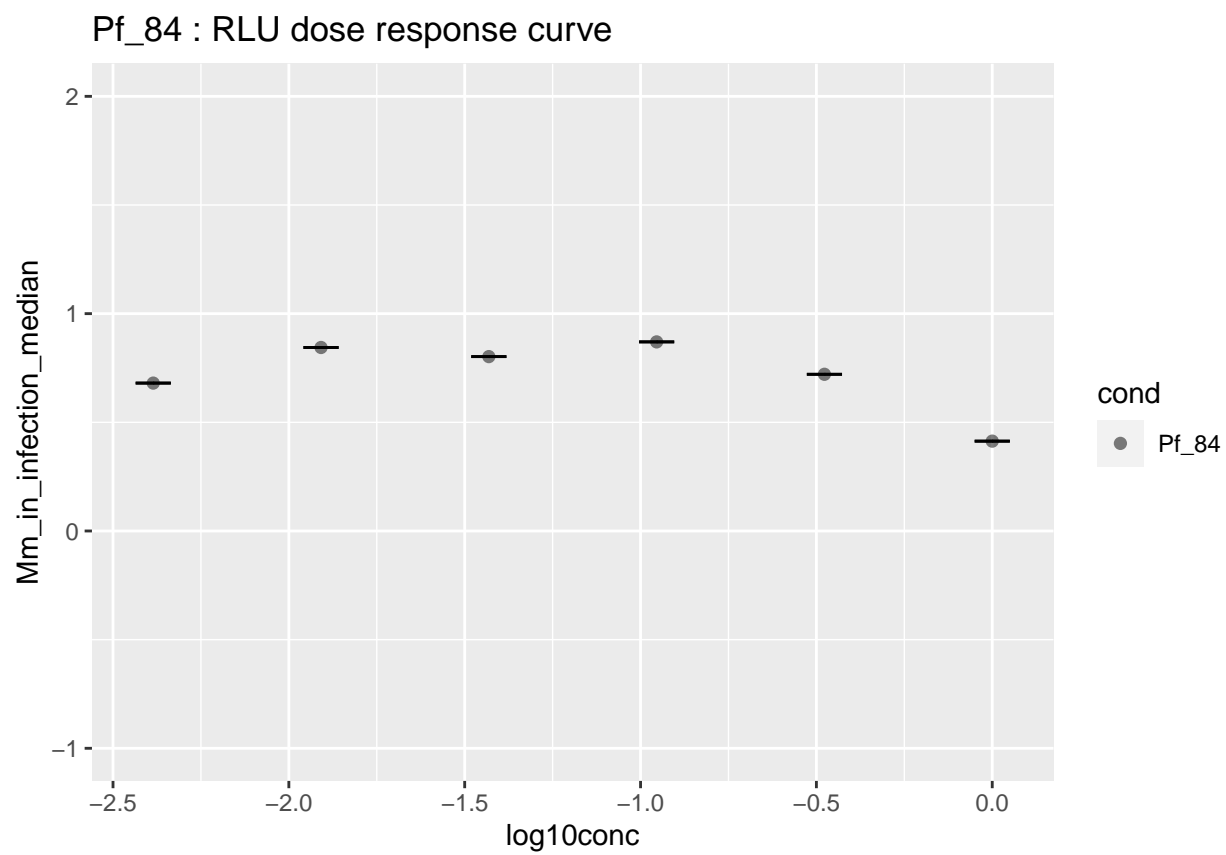


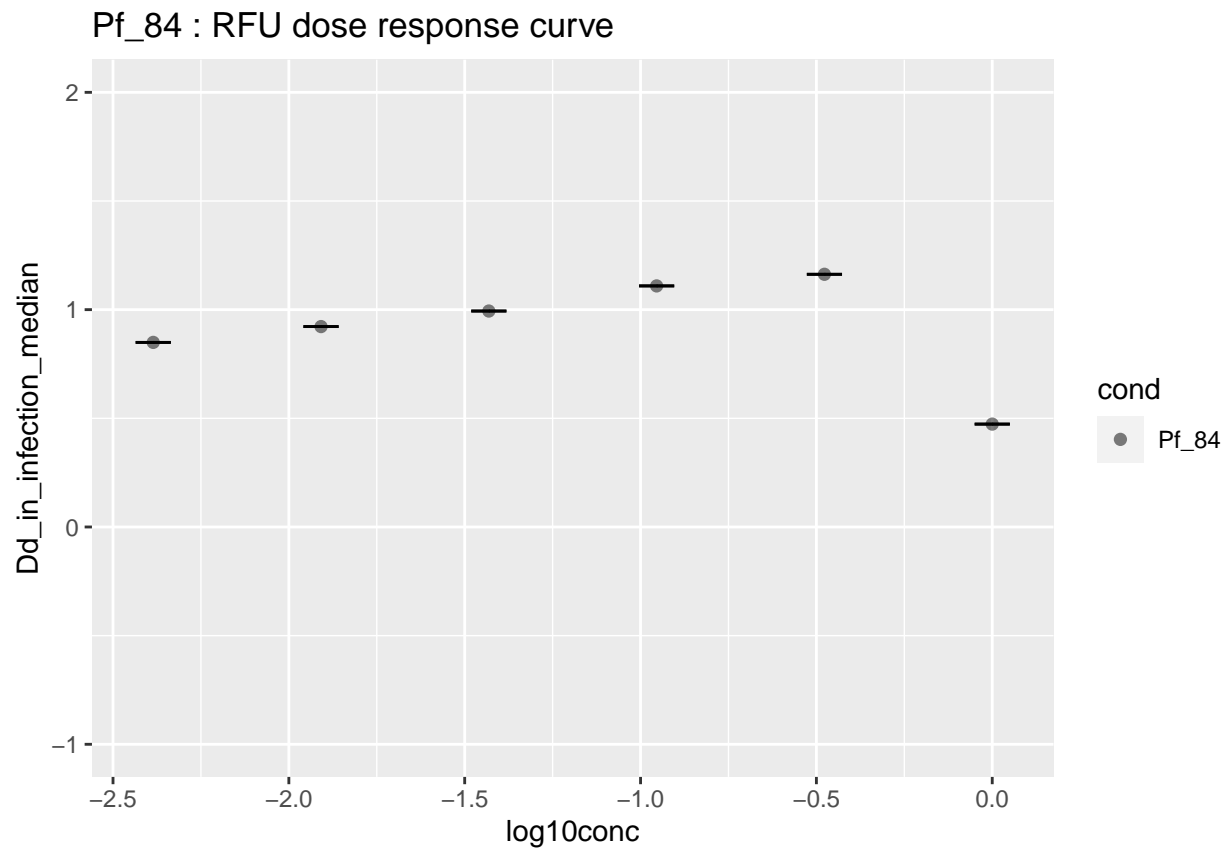


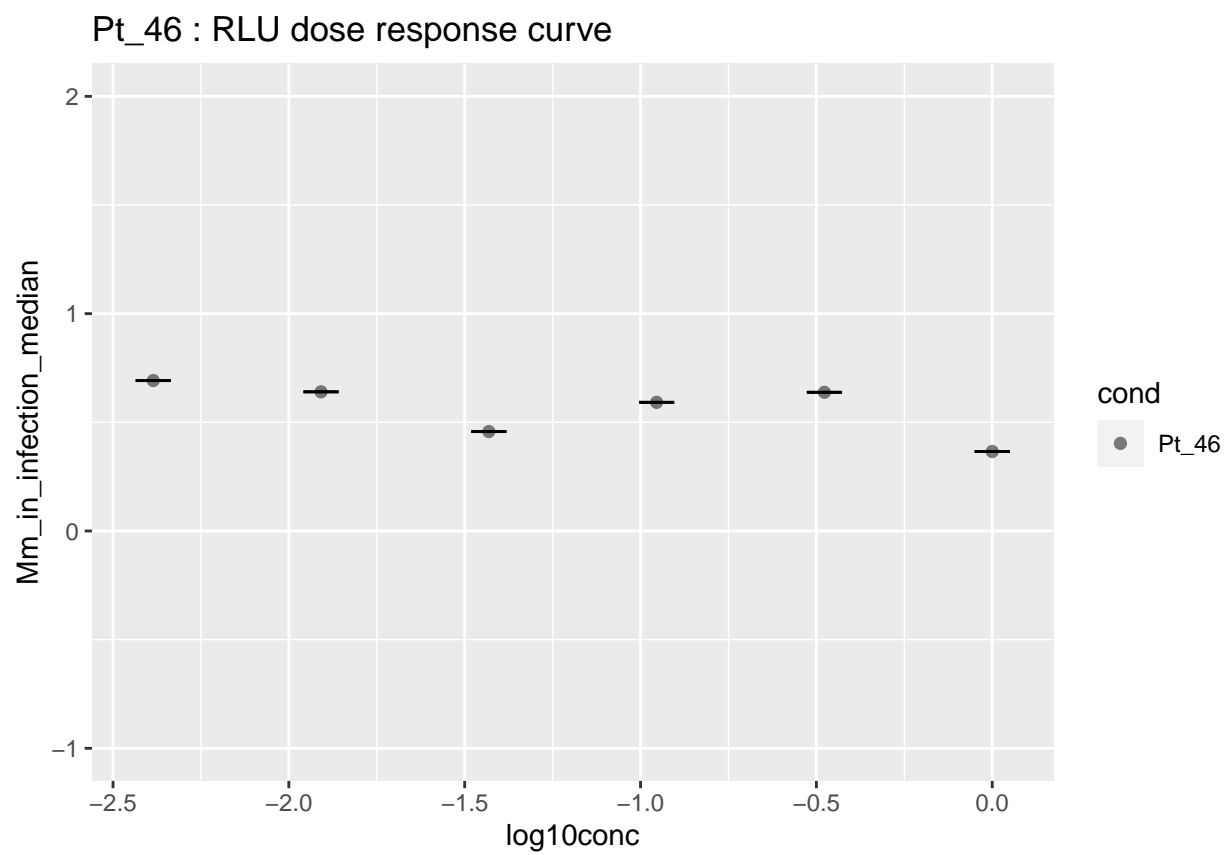


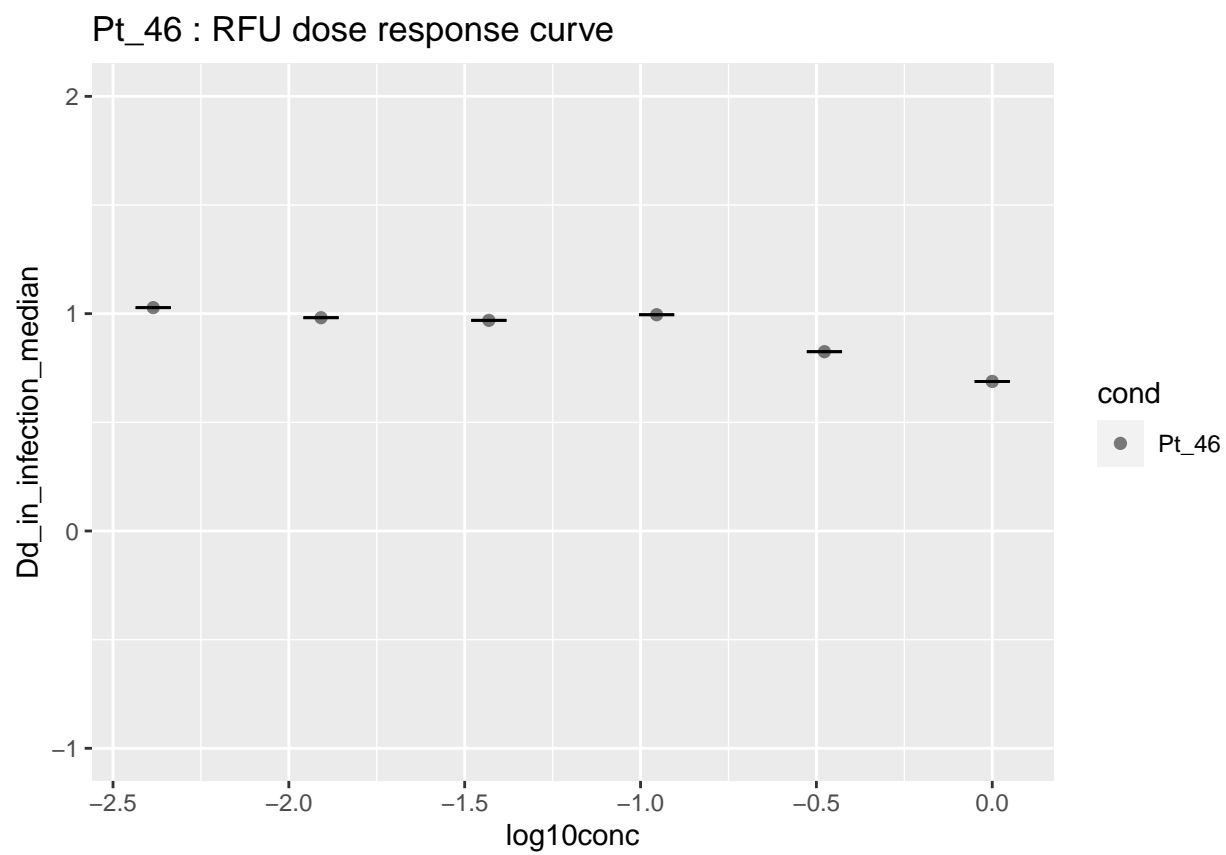




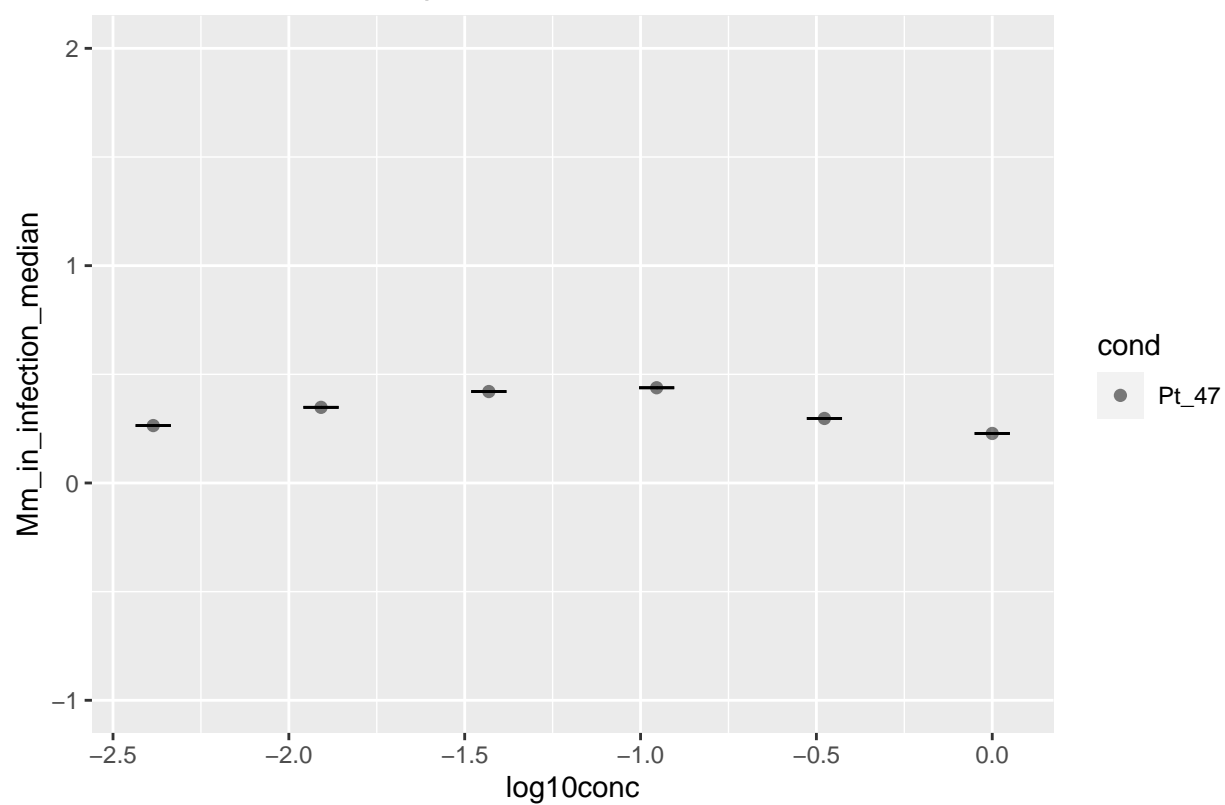


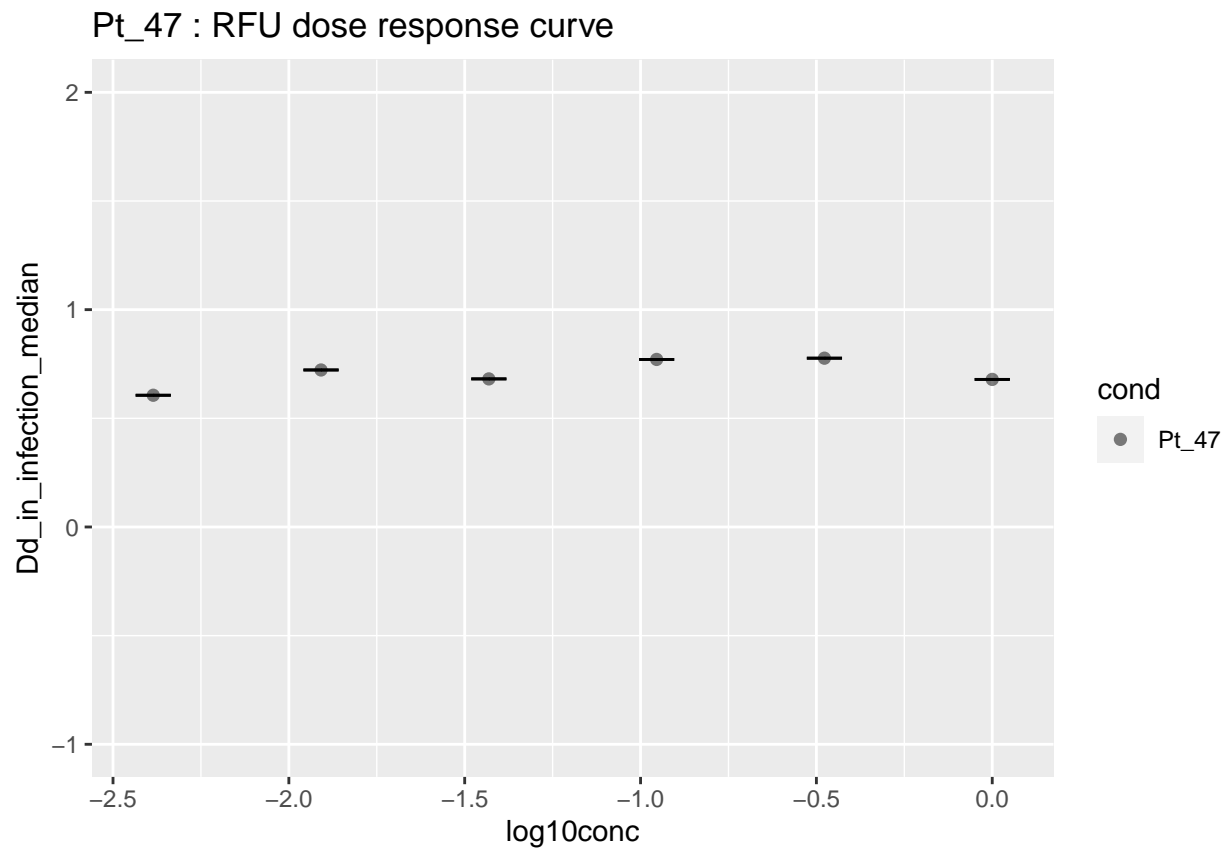




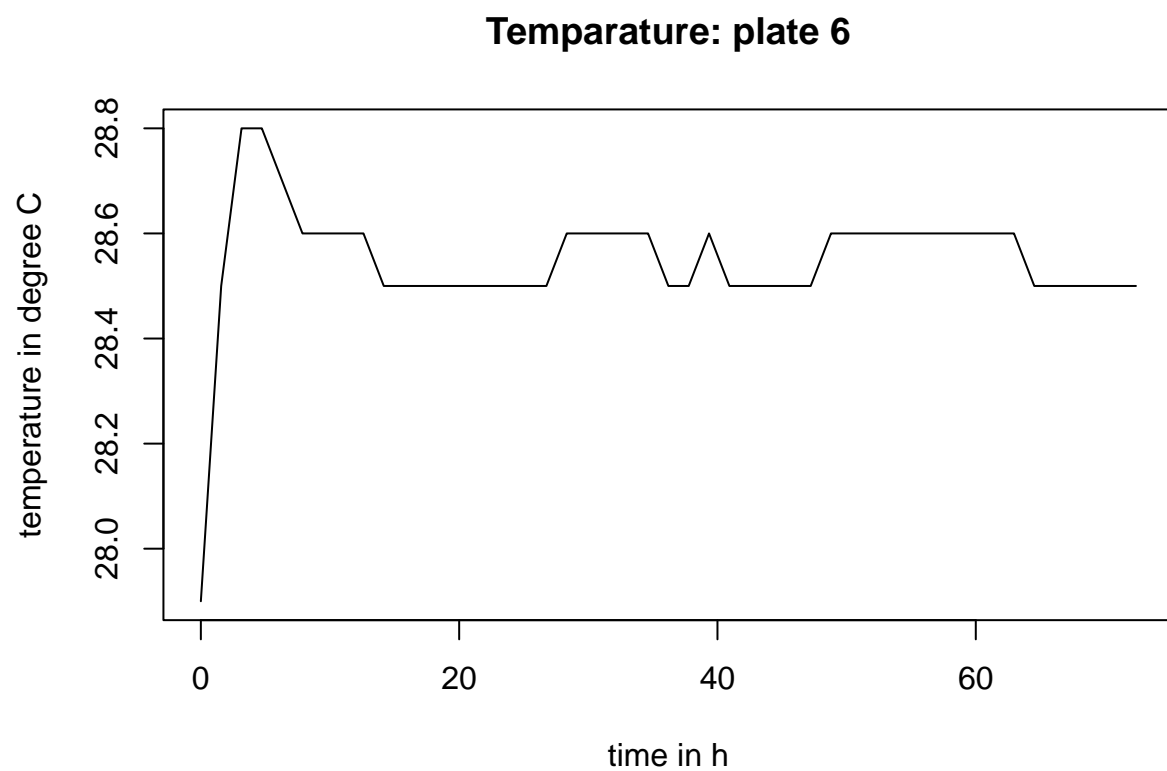


Pt_47 : RLU dose response curve

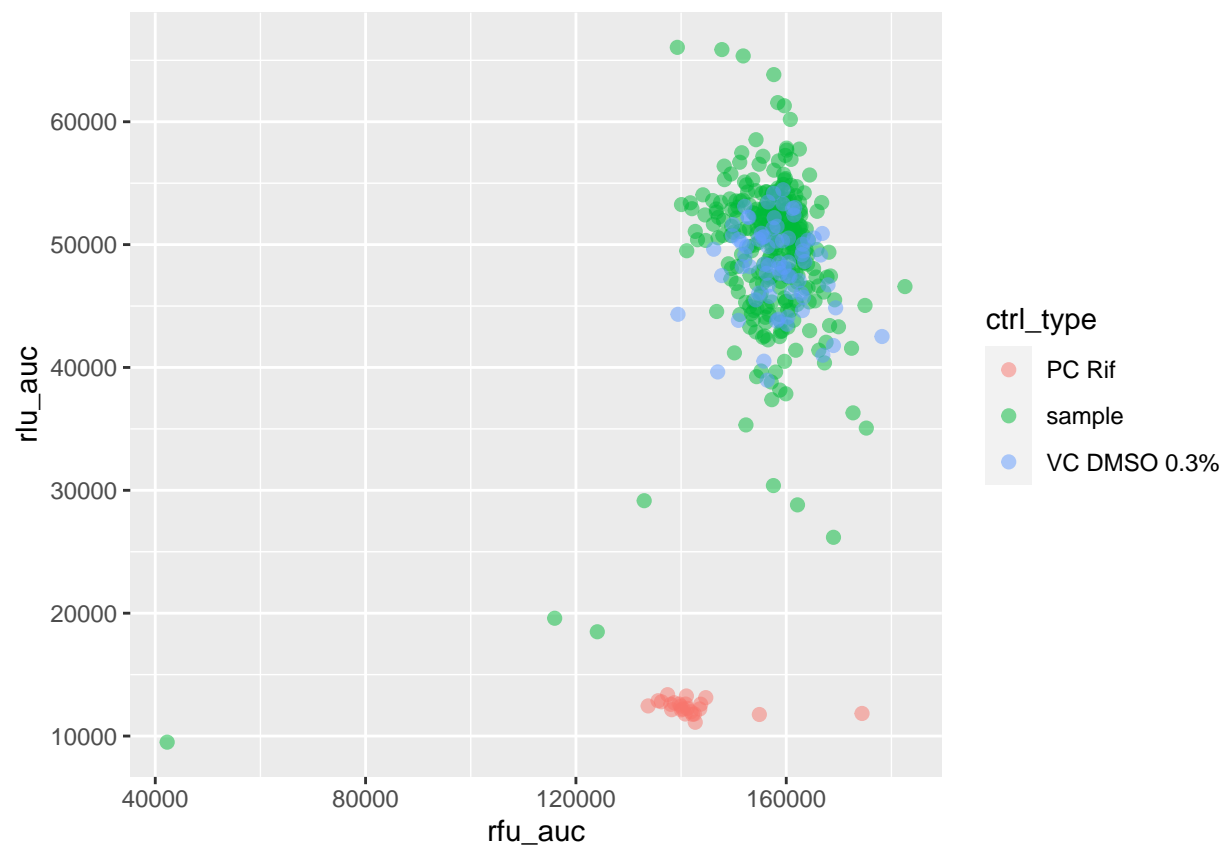


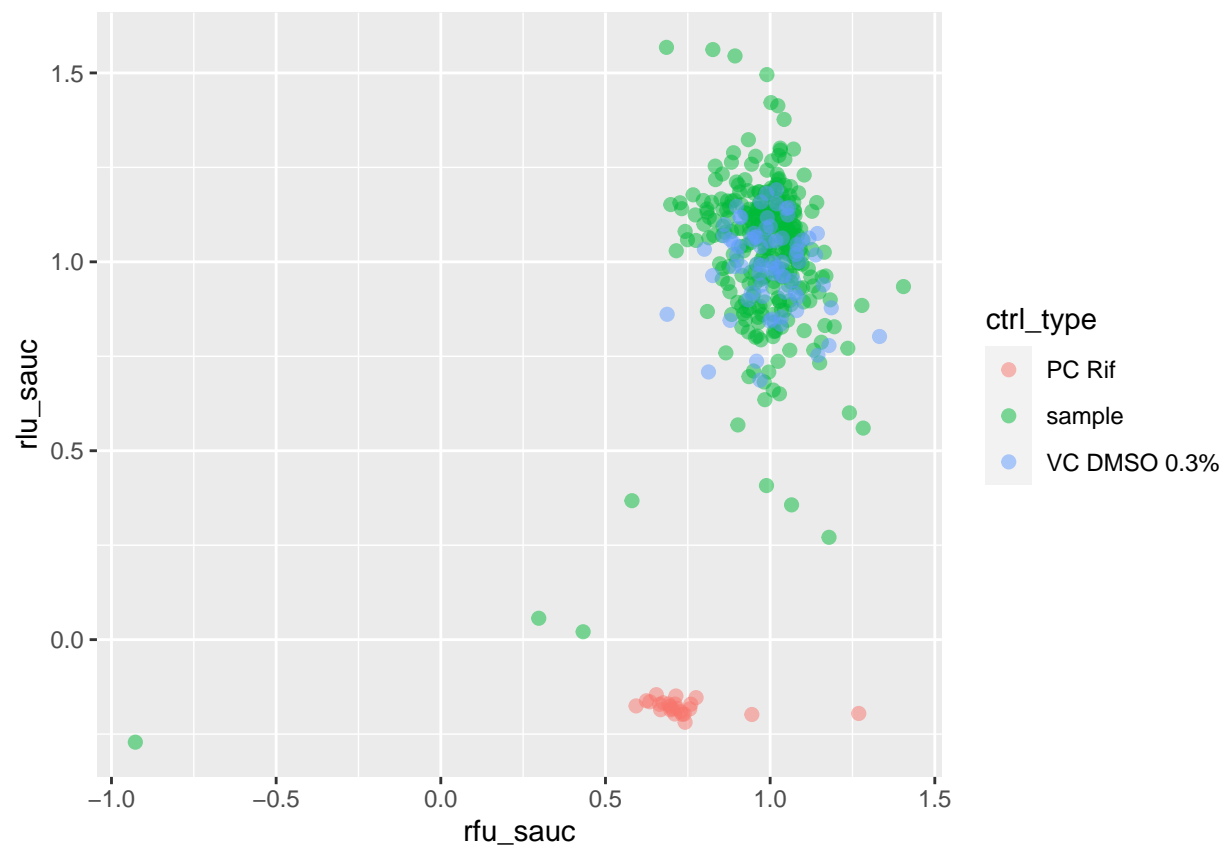


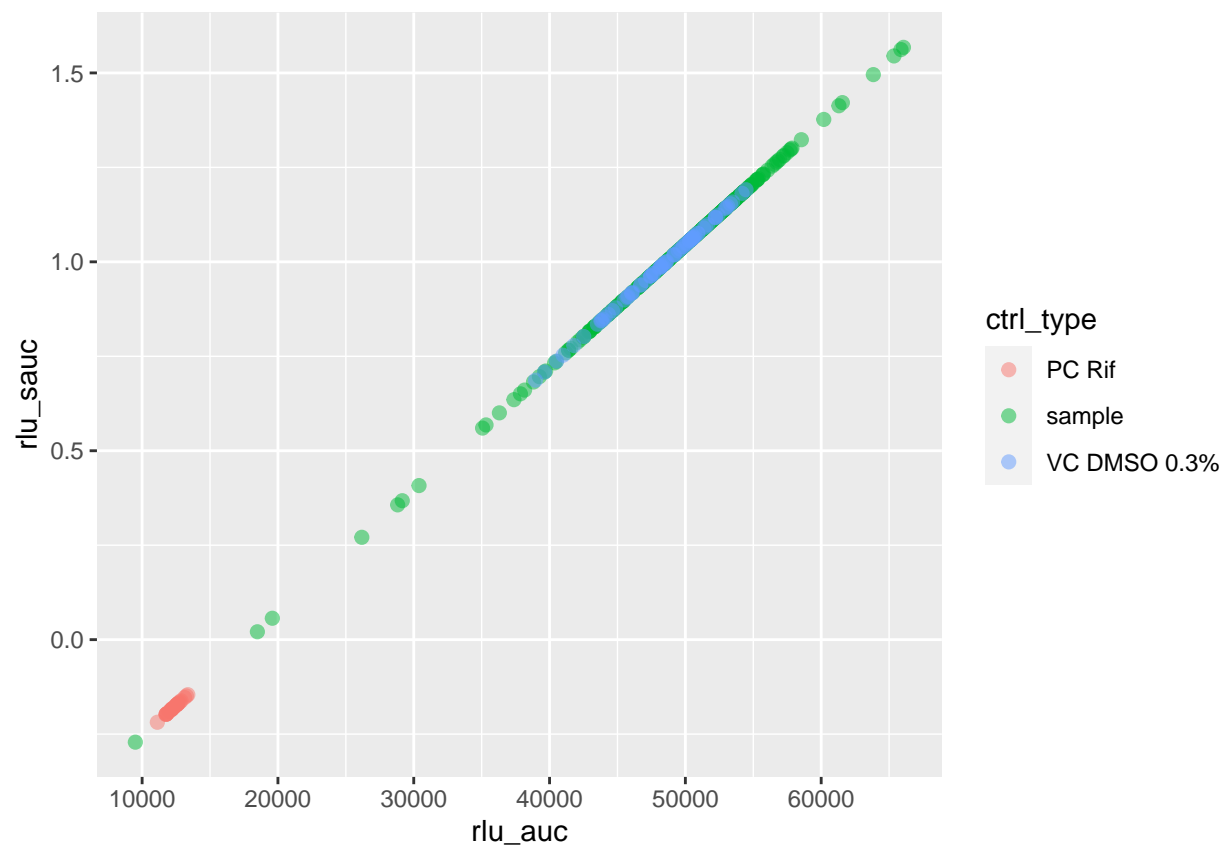
```
## [1] "analysis for plate07.xlsx"
```

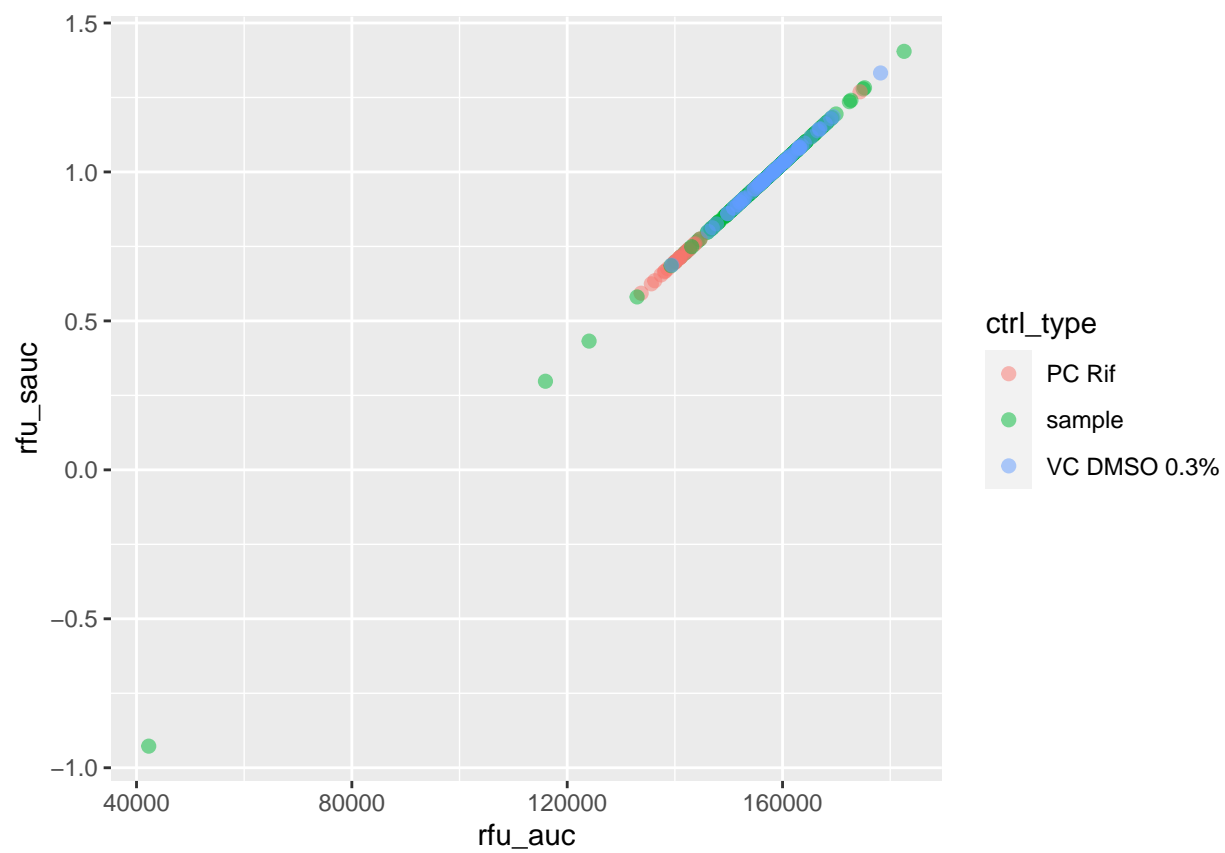


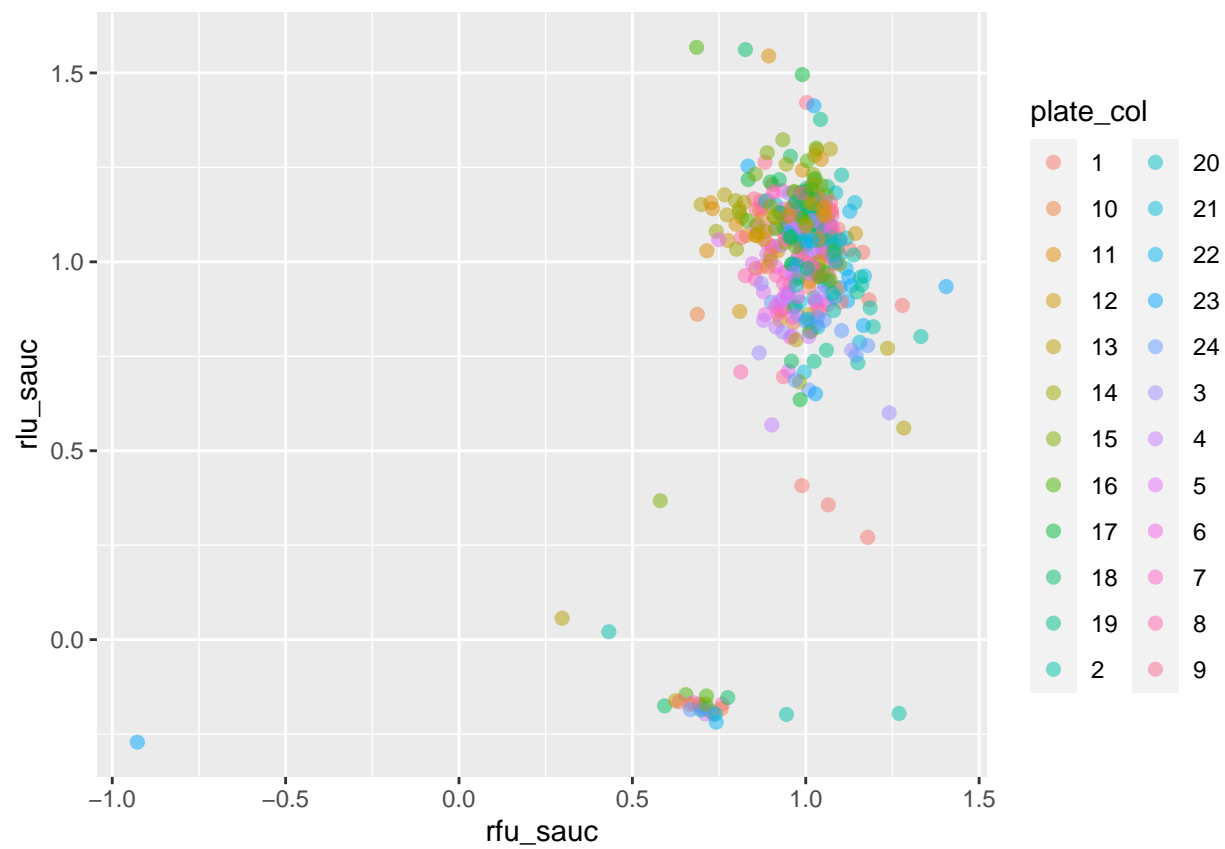
```
## [1] "VC DMSO 0.3% Robust z'-factor of rlu for plate plate07.xlsx, biorep 3 : "  
## [1] 0.68
```

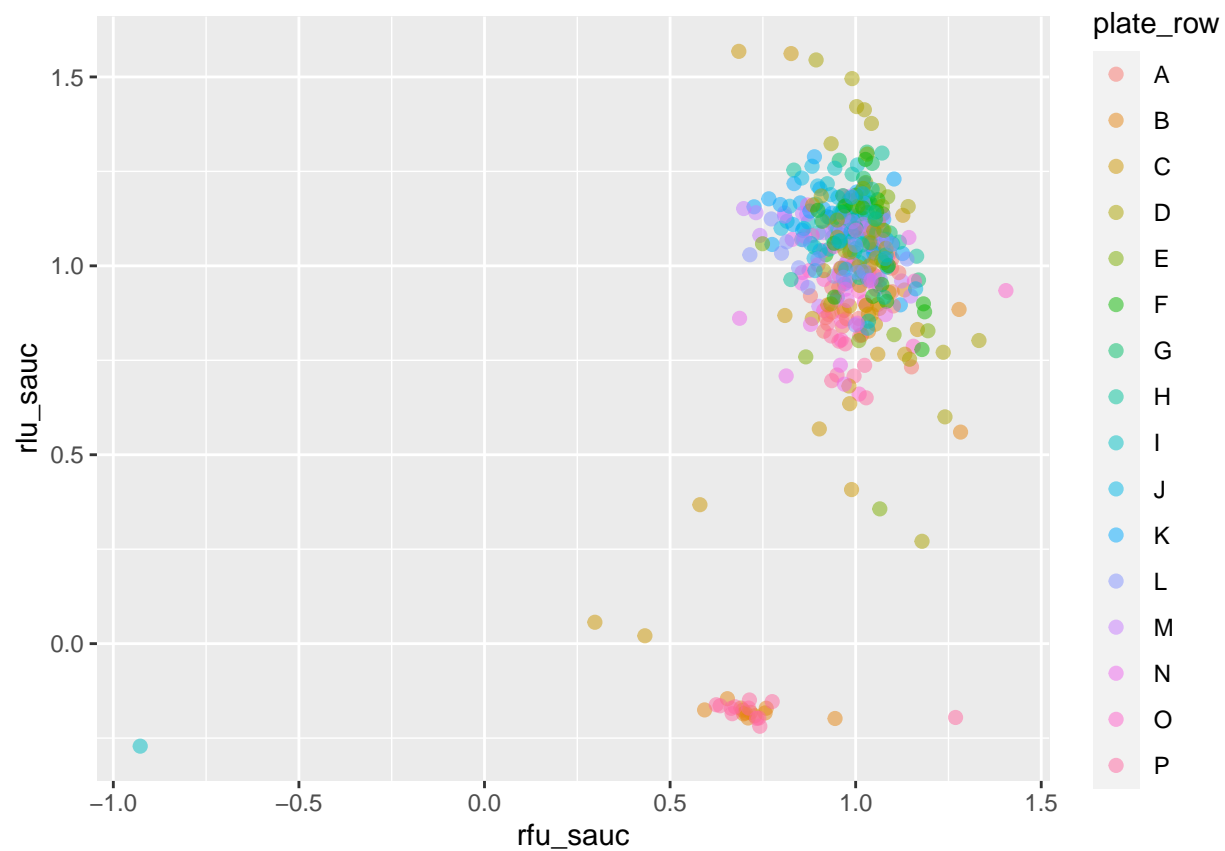





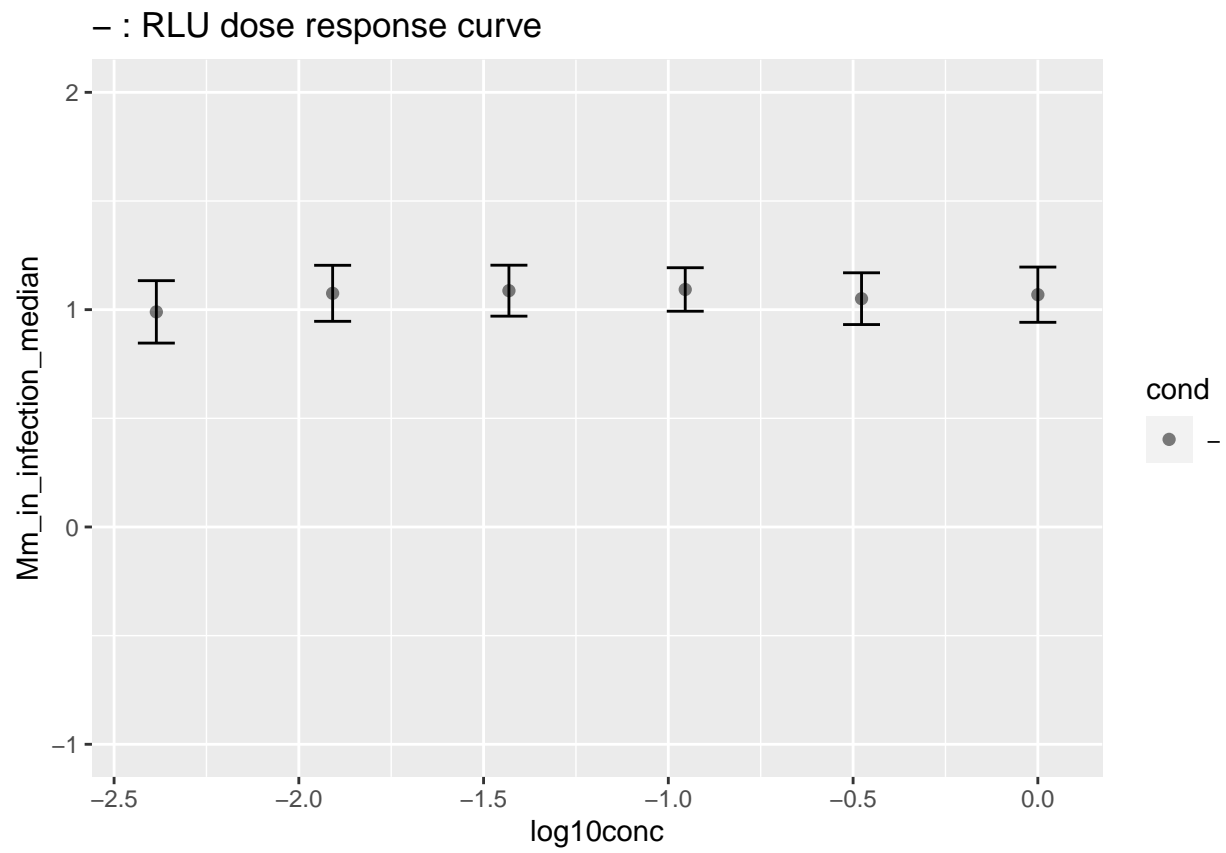


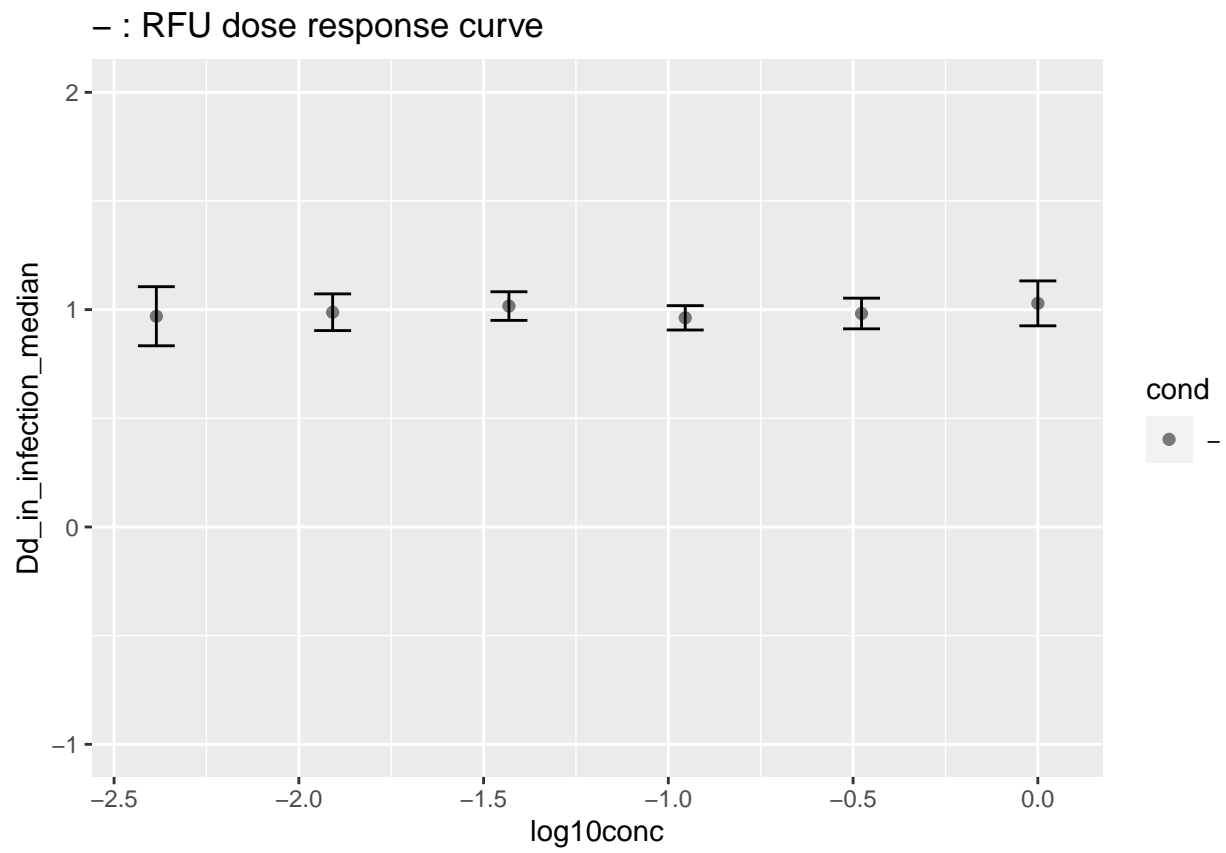


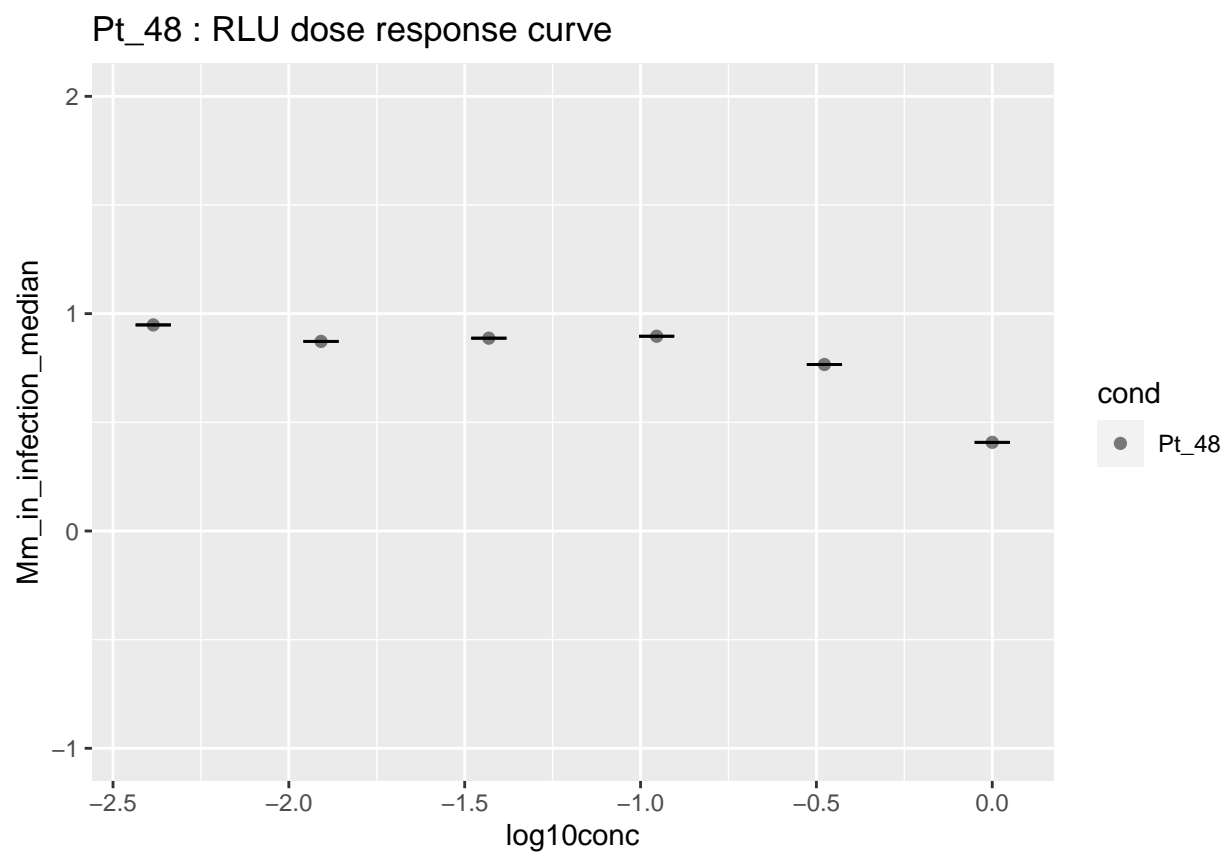


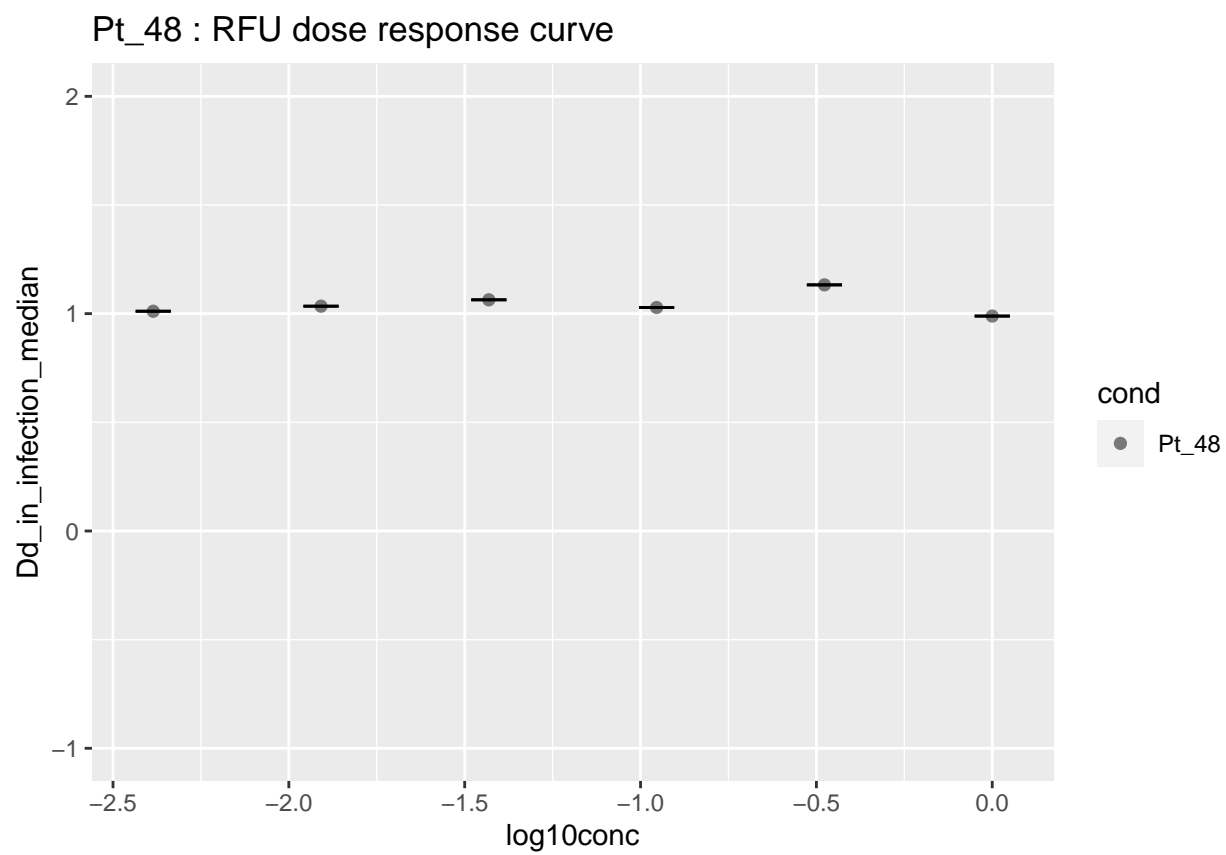


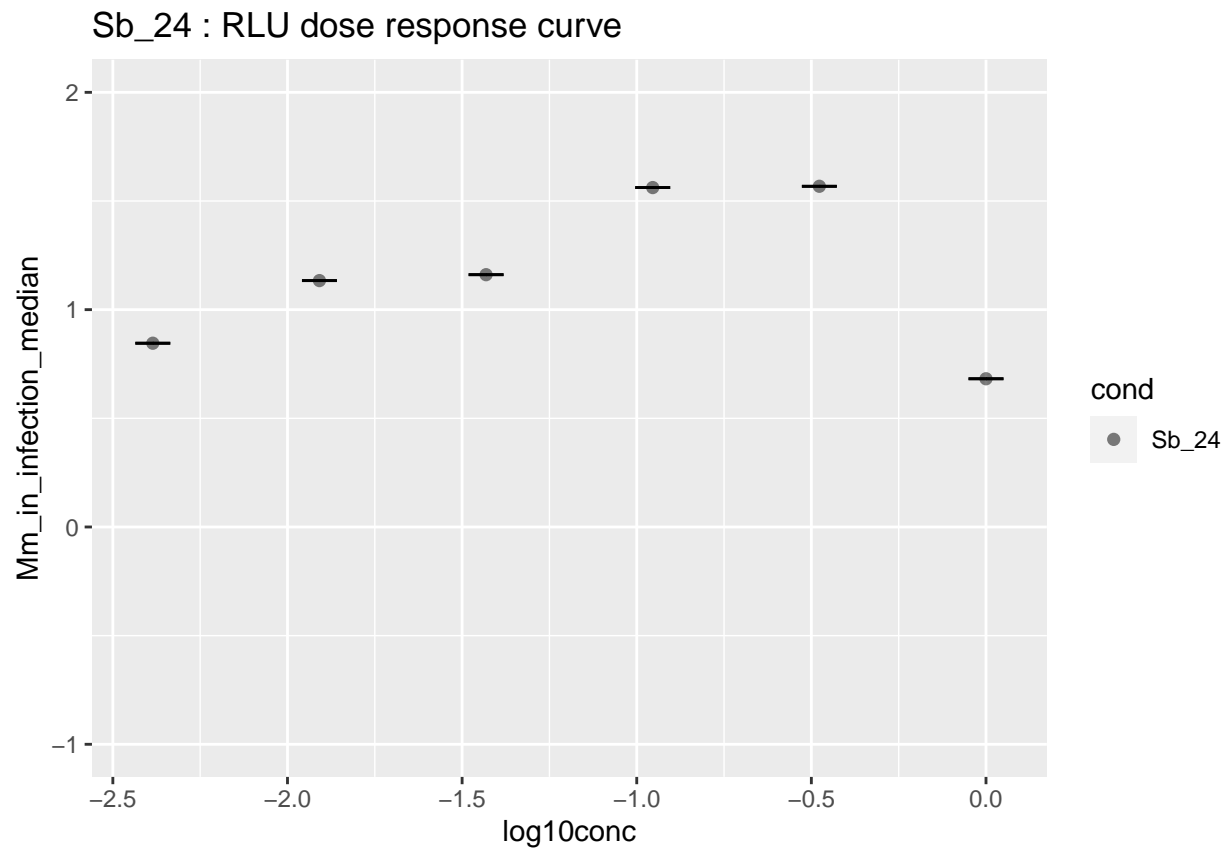
```
## [1] "Dose response curves over all bioreps within this plate"
```

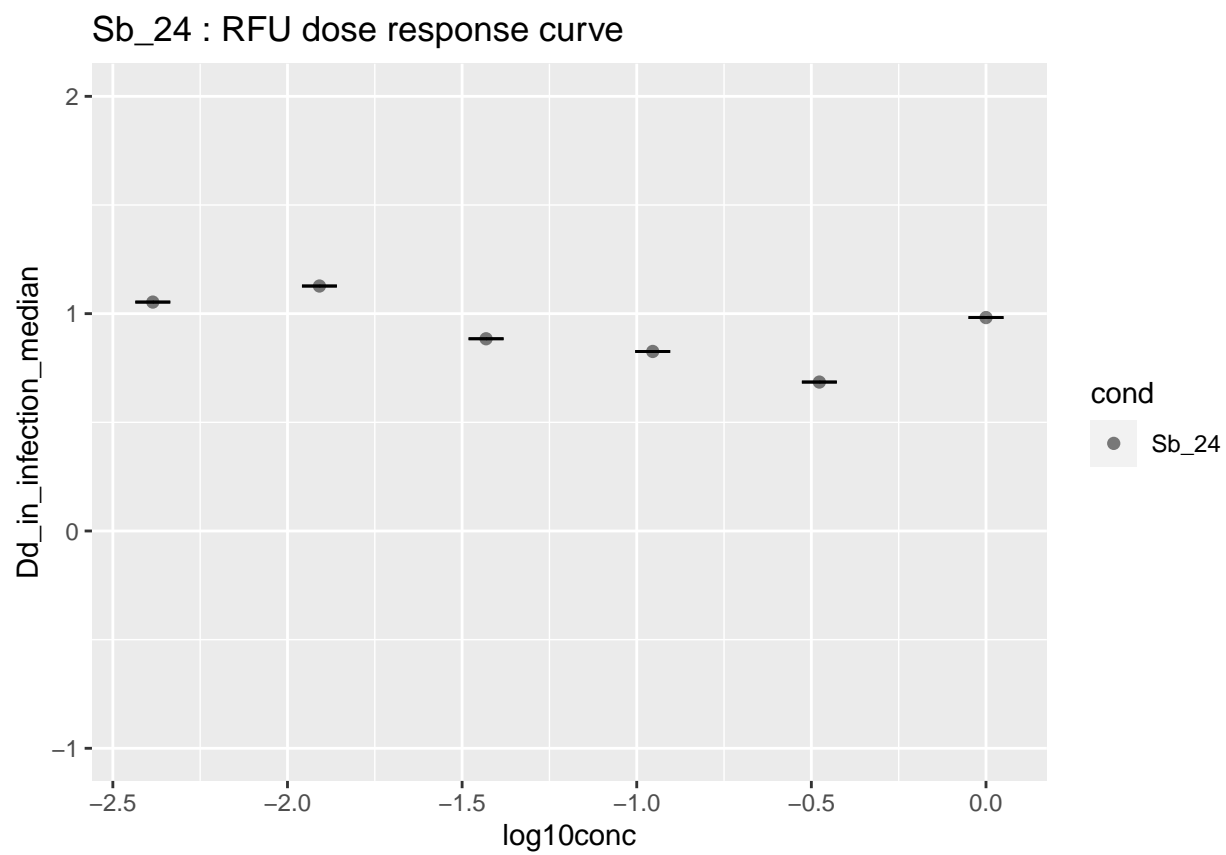


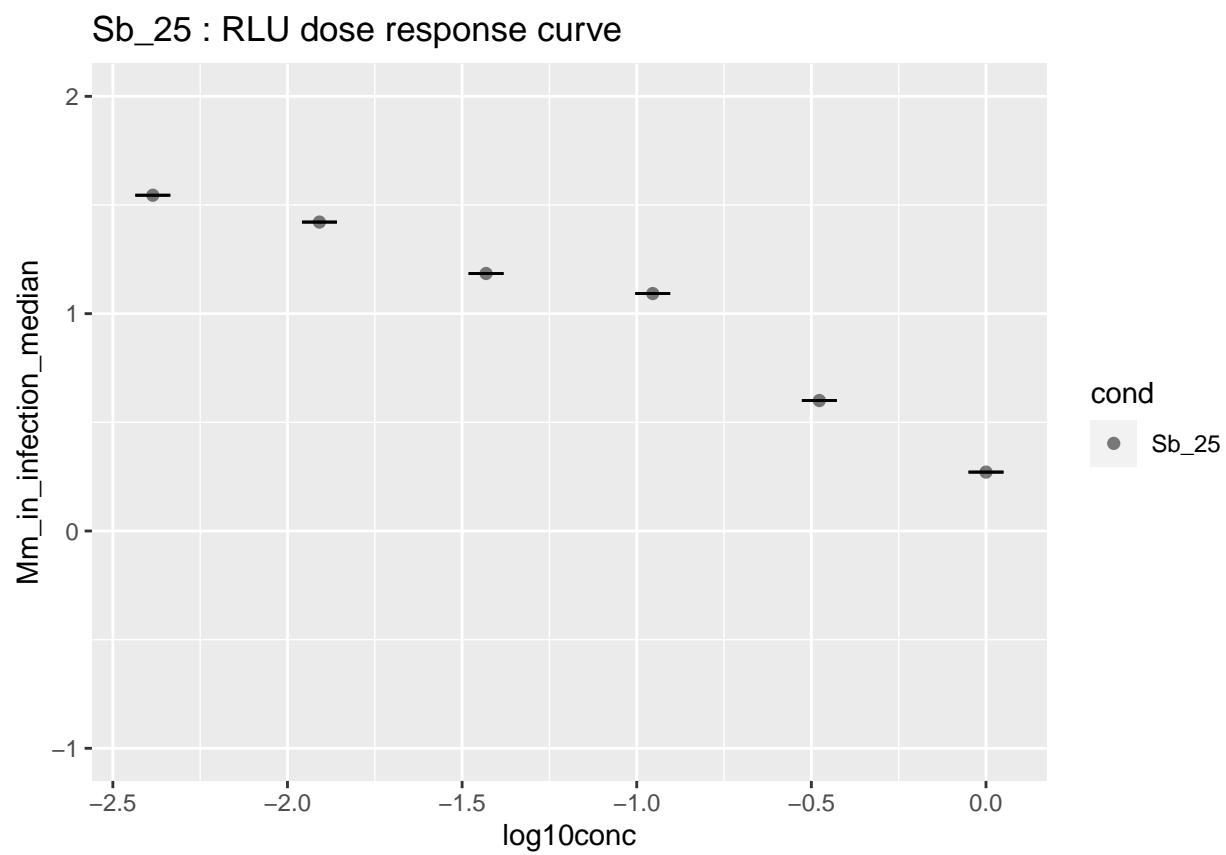


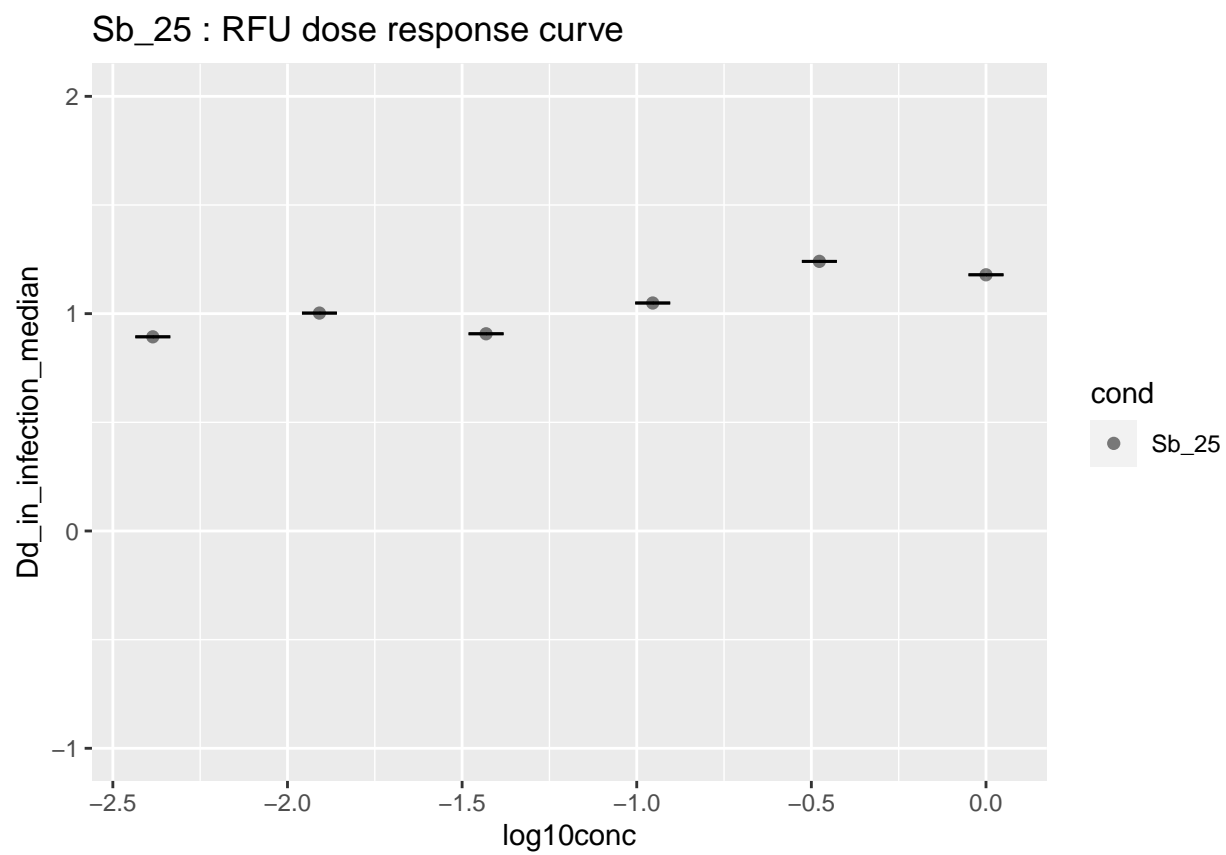


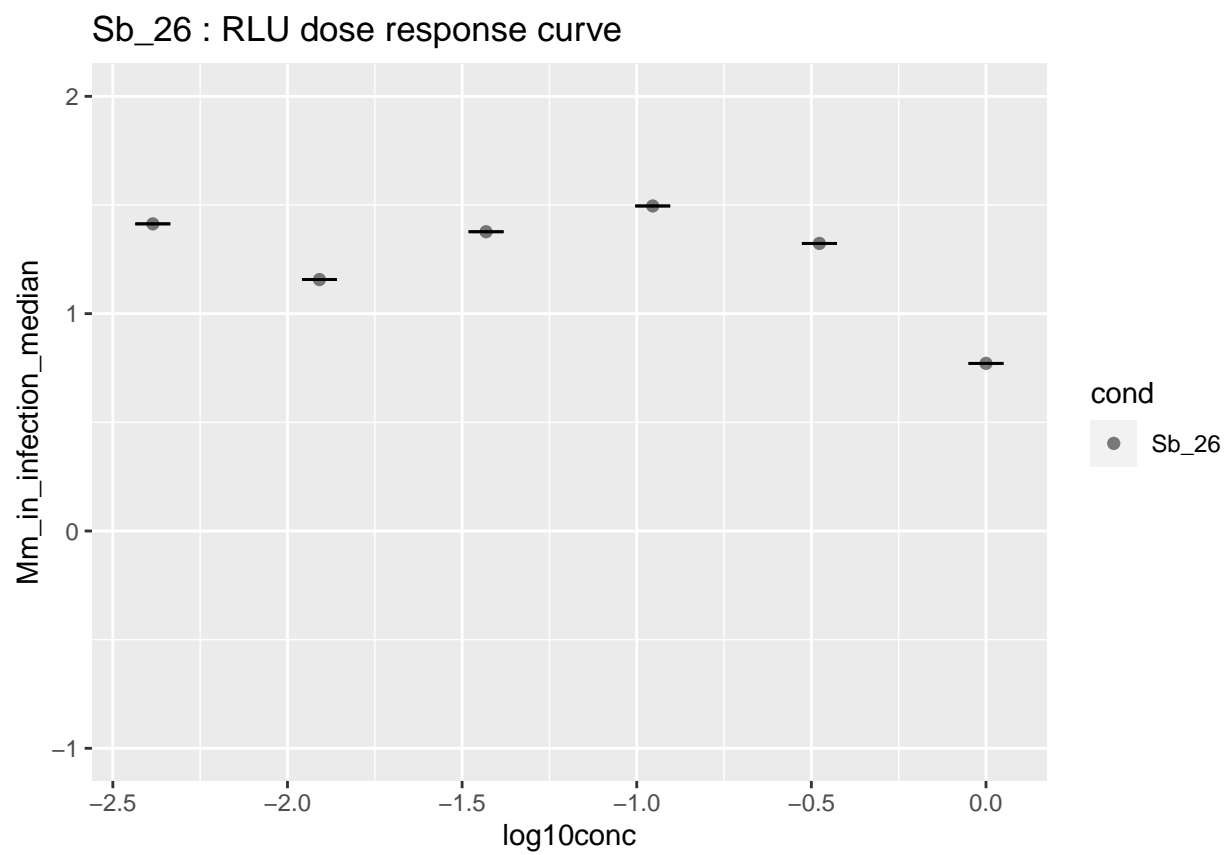


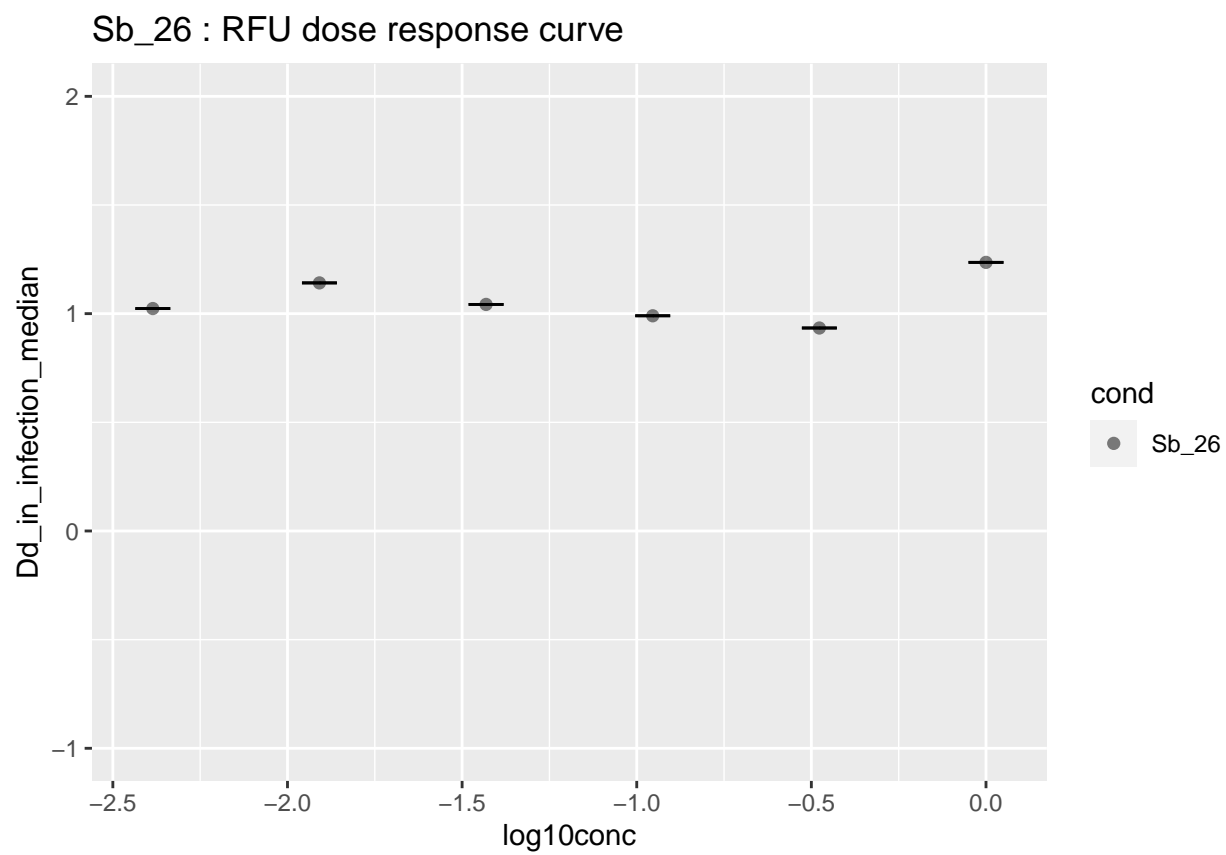


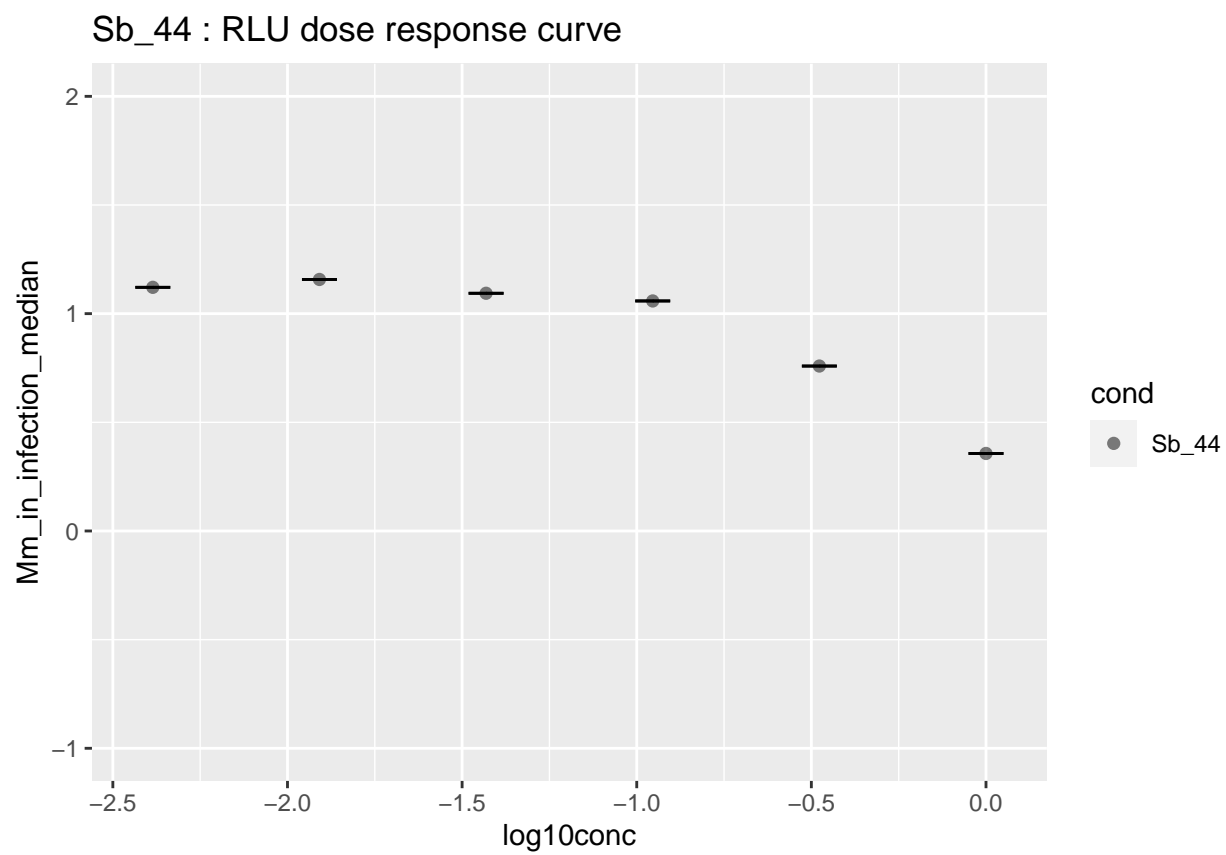


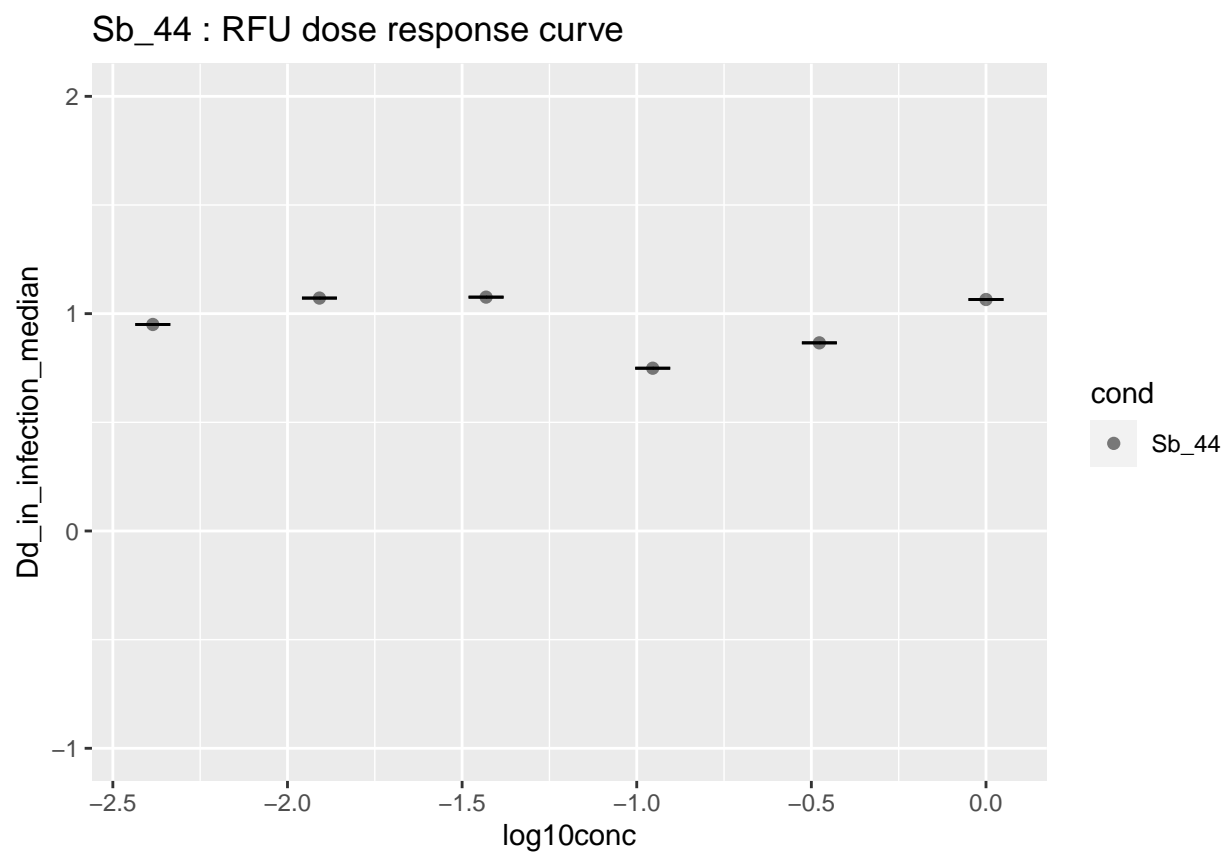


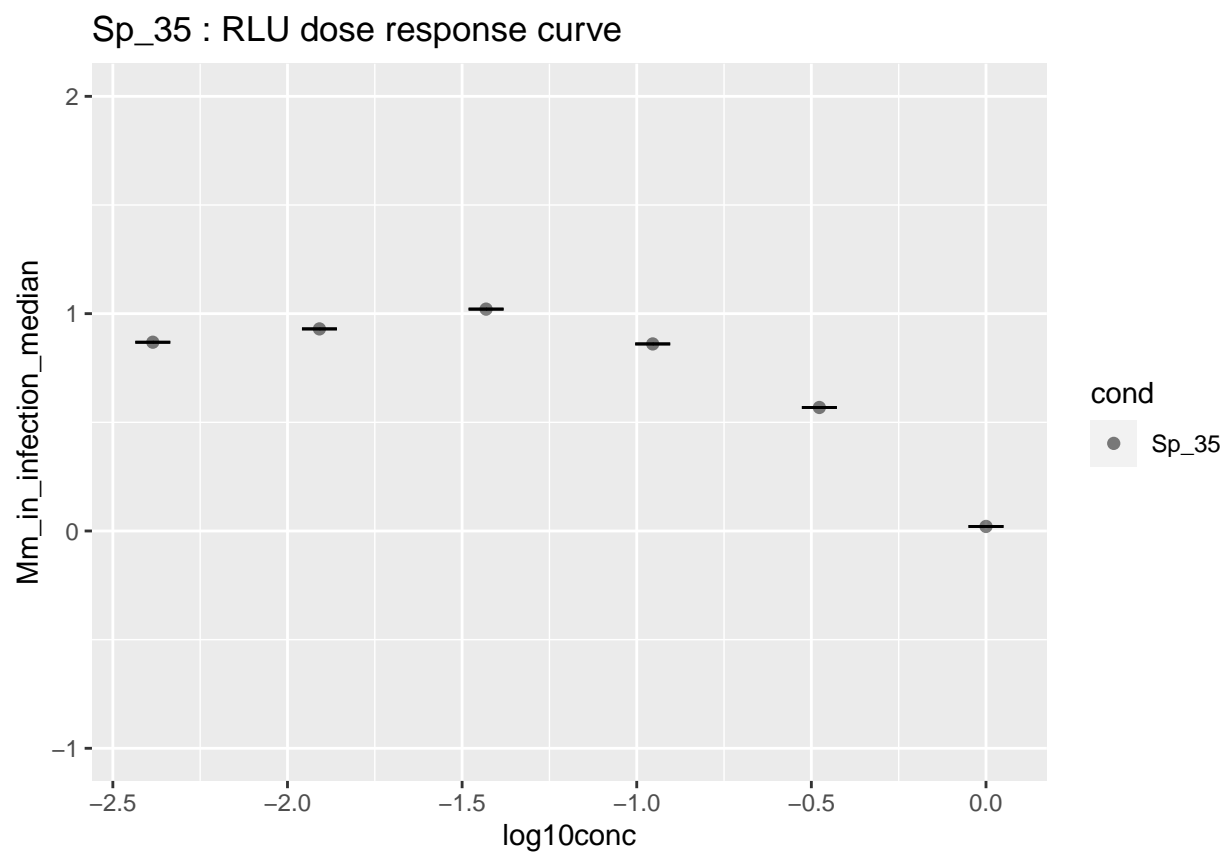


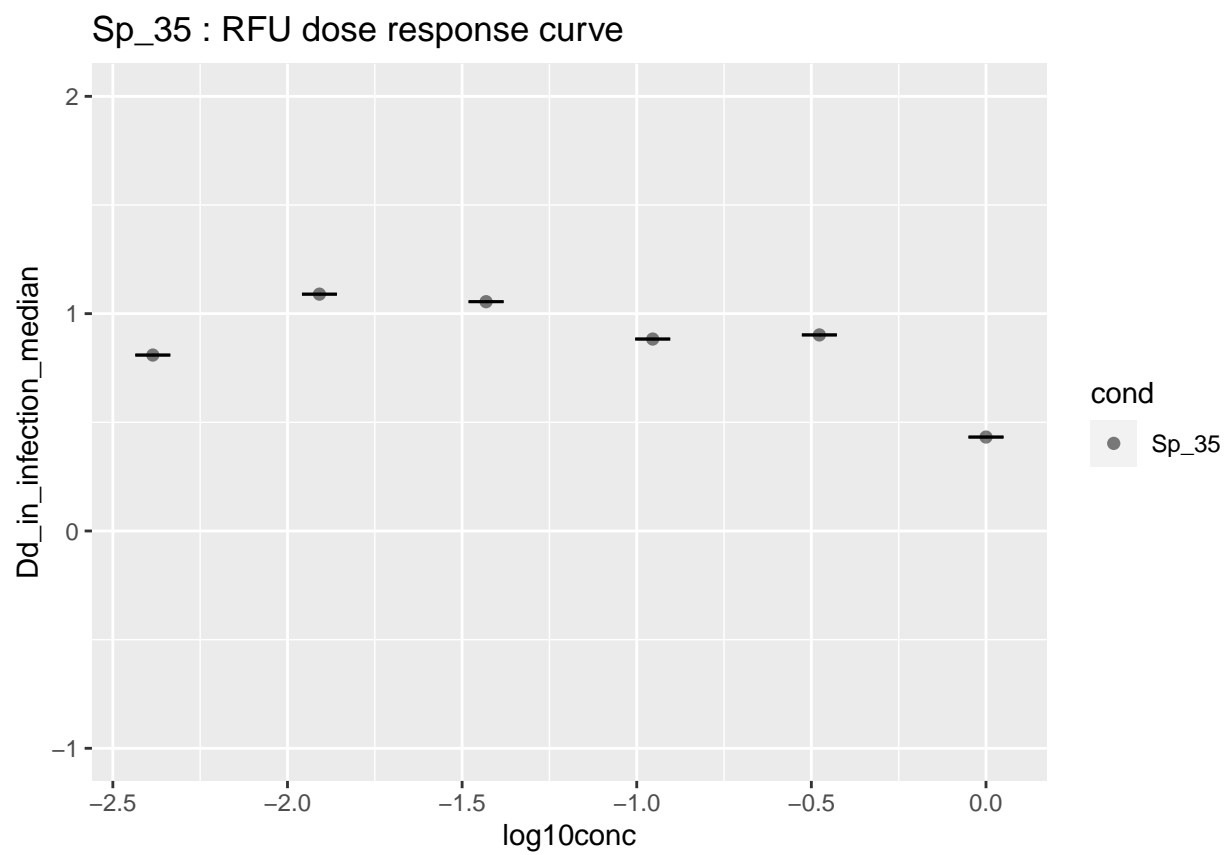


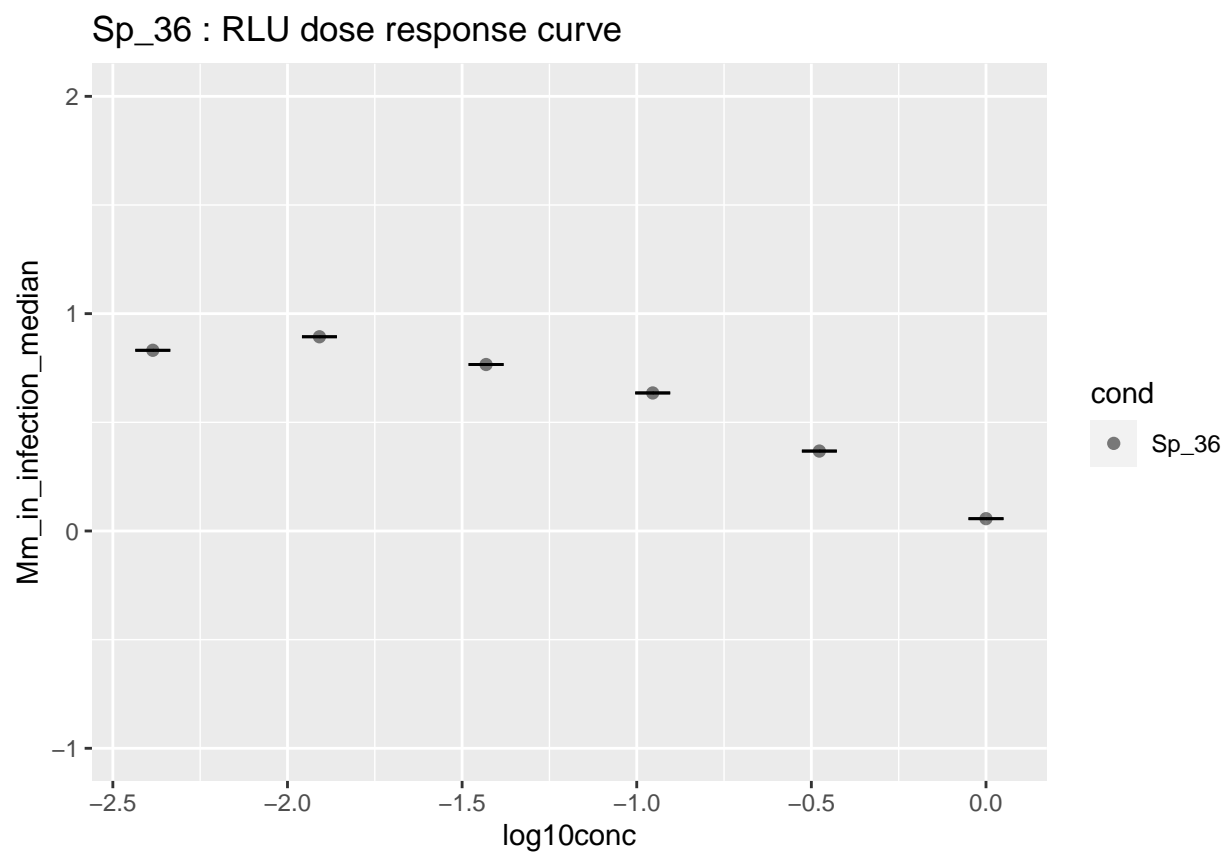


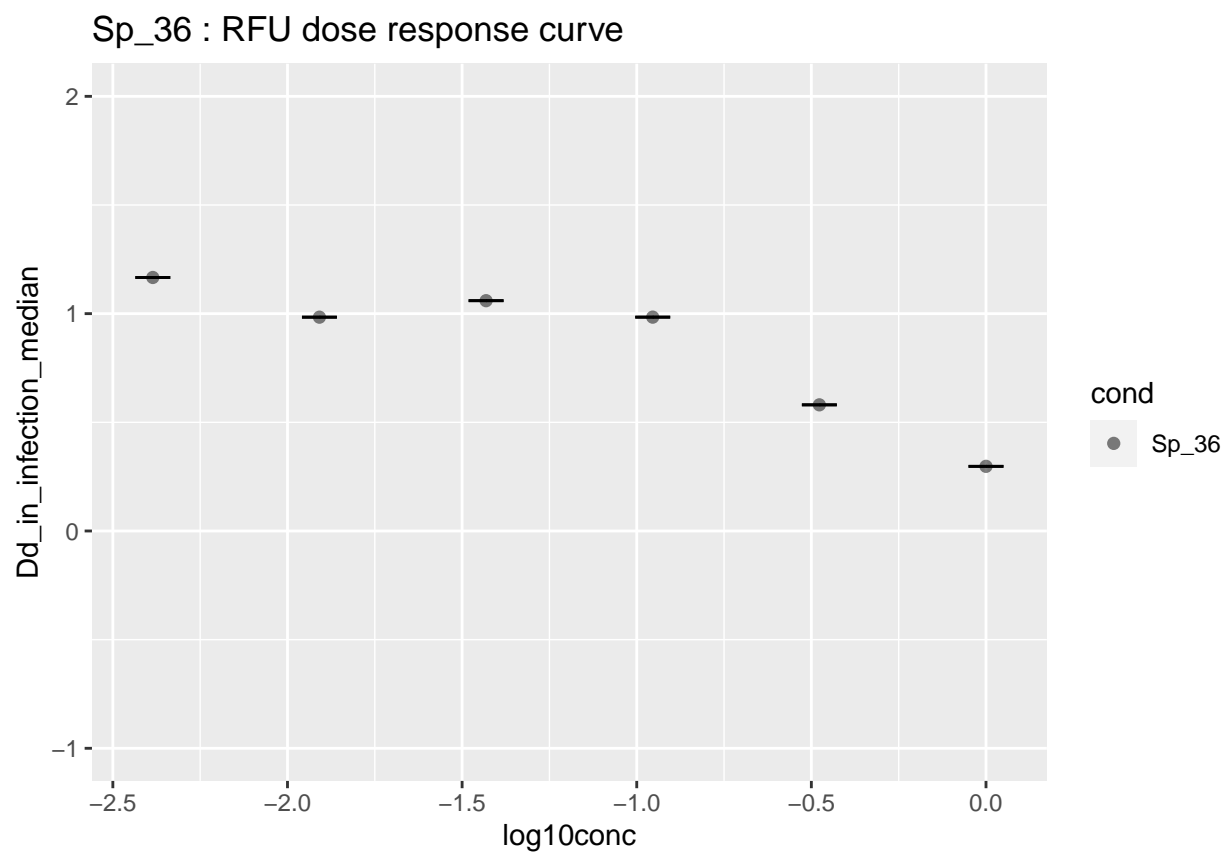


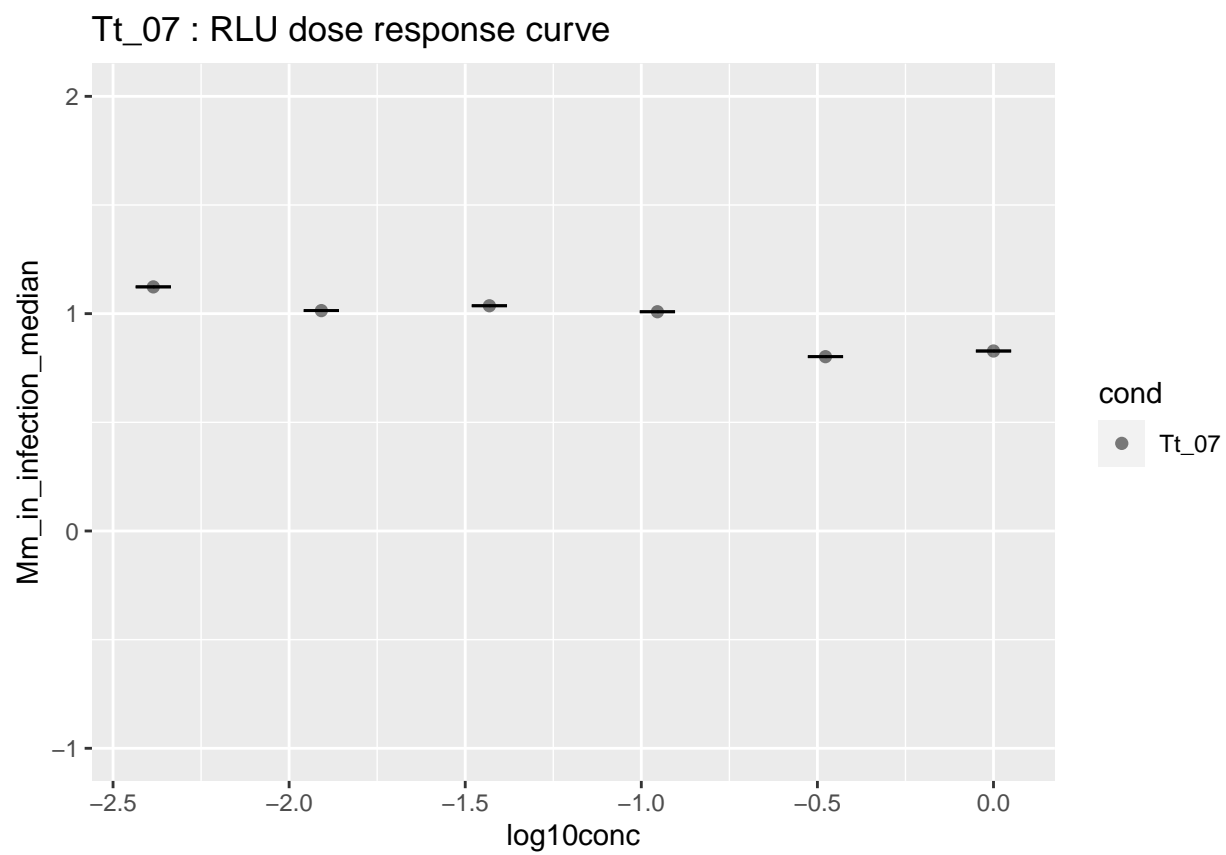


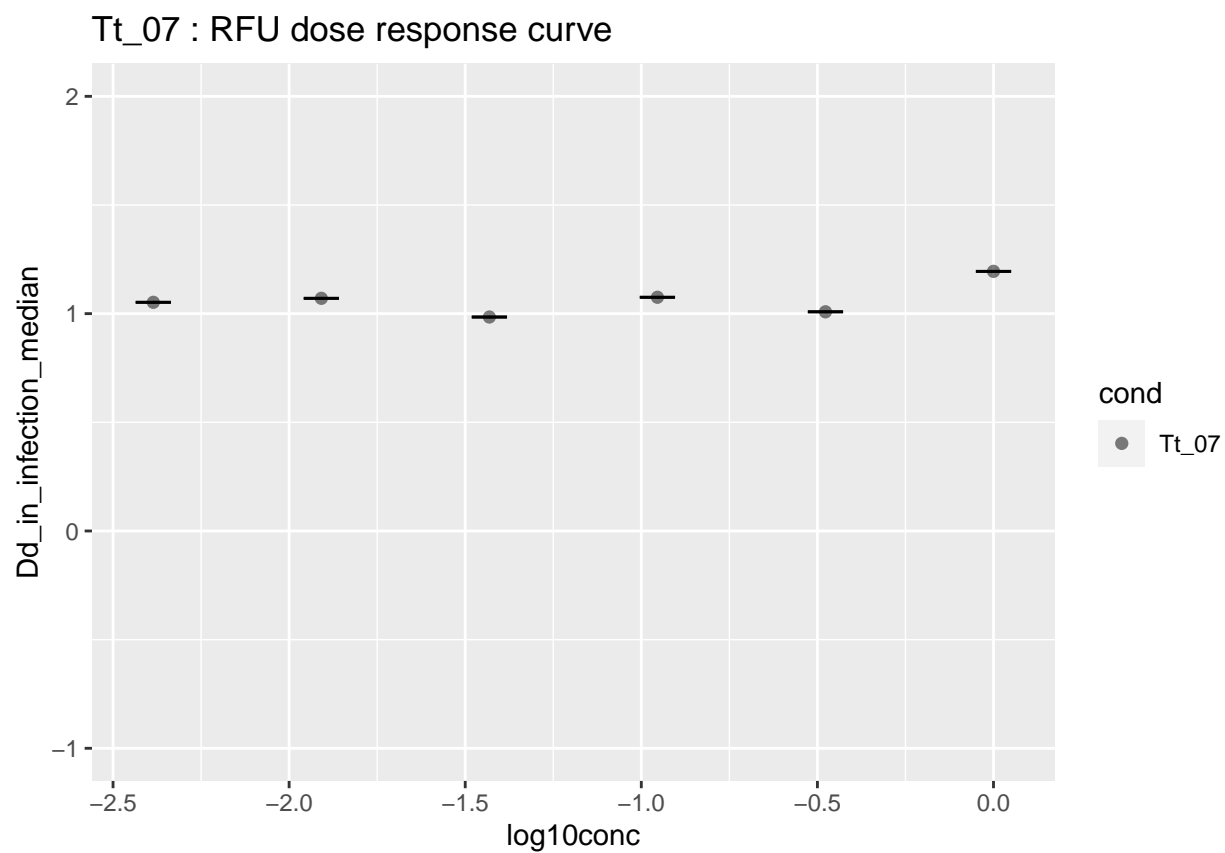


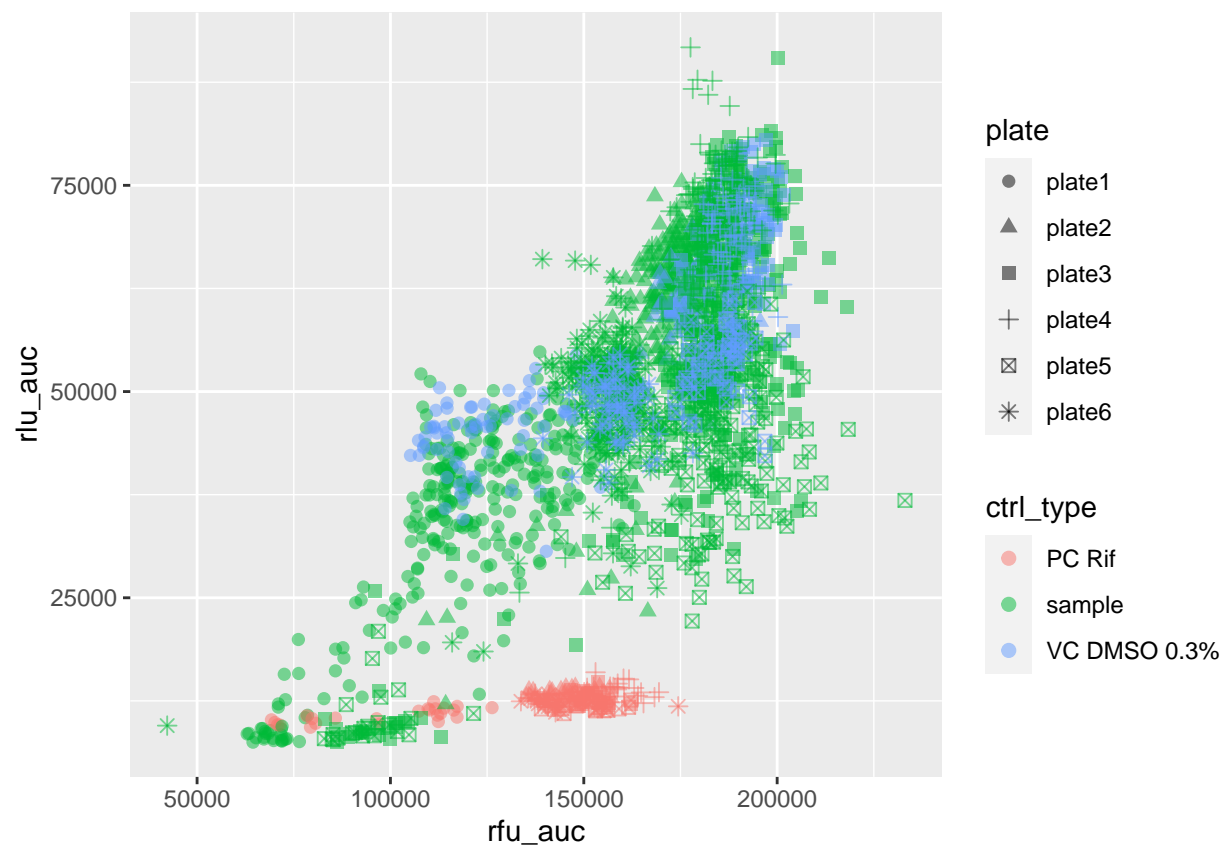


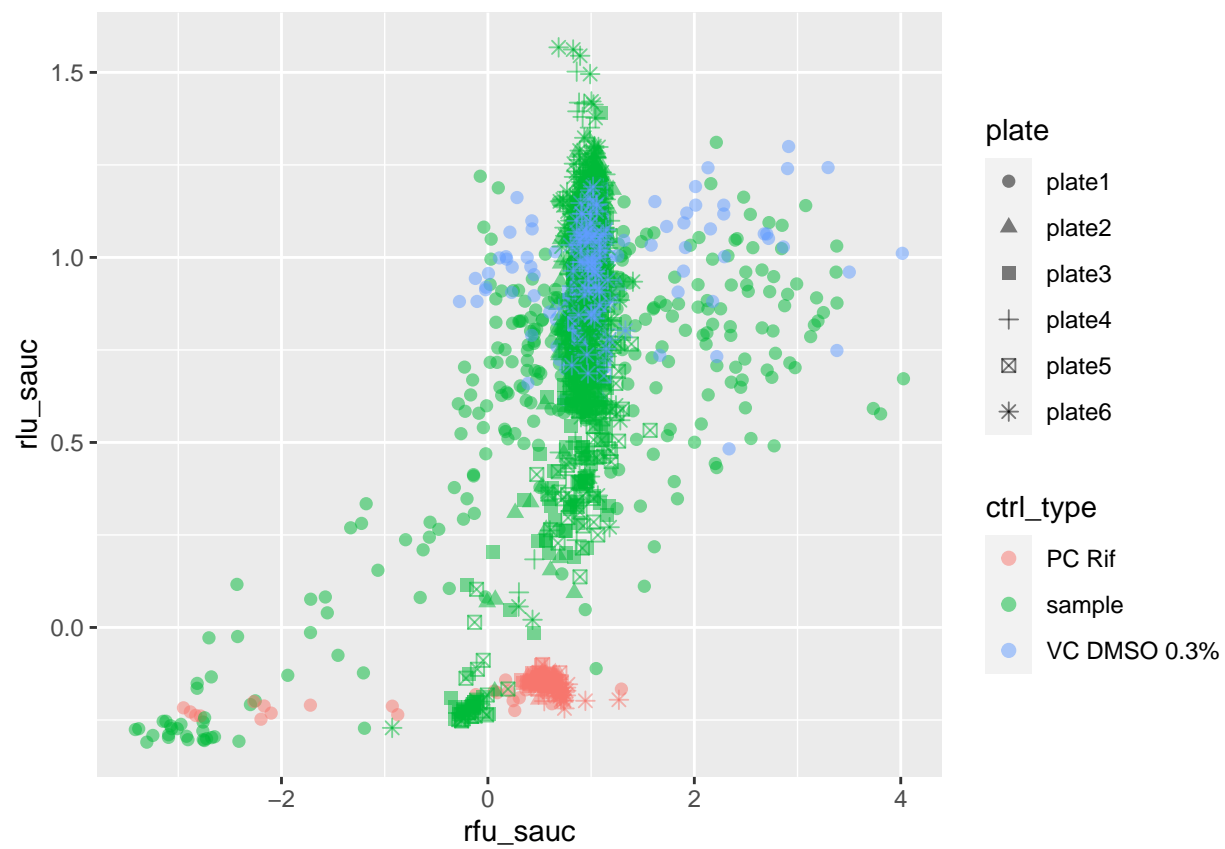


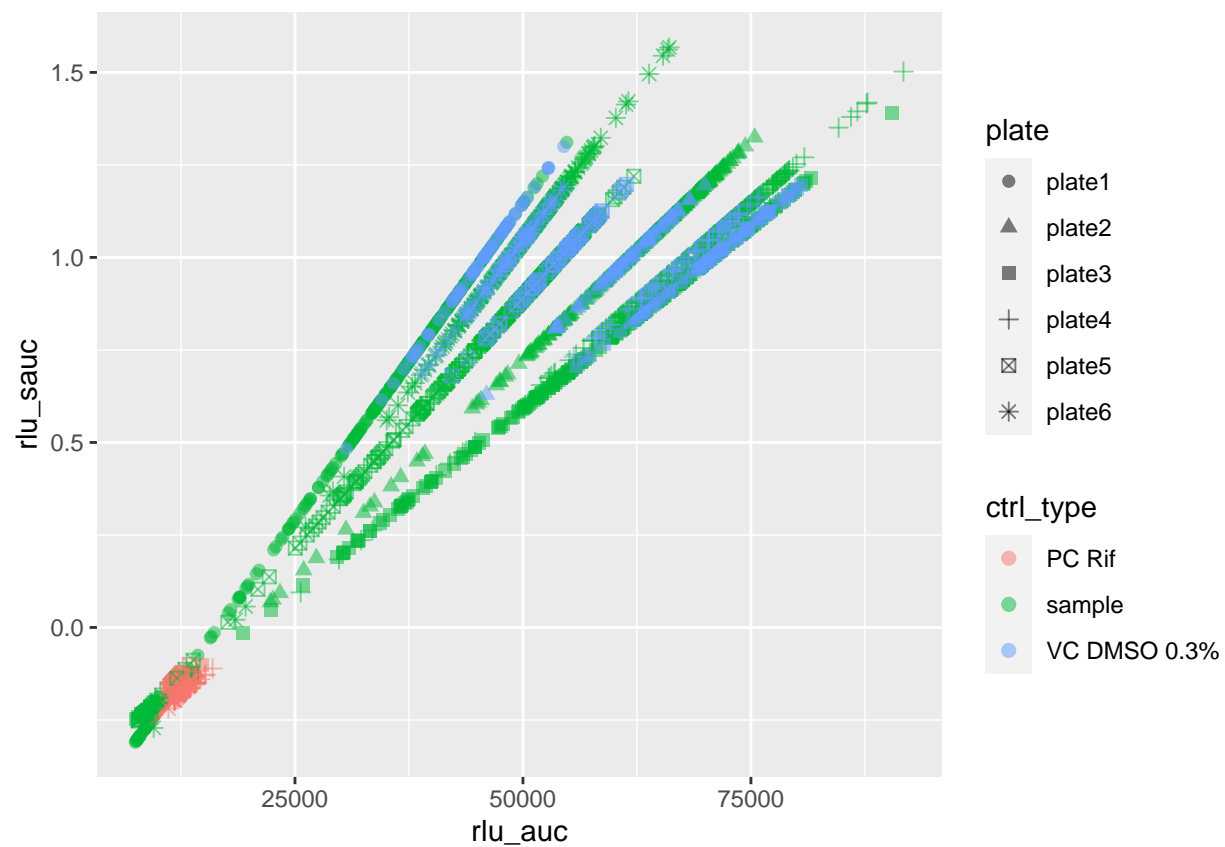


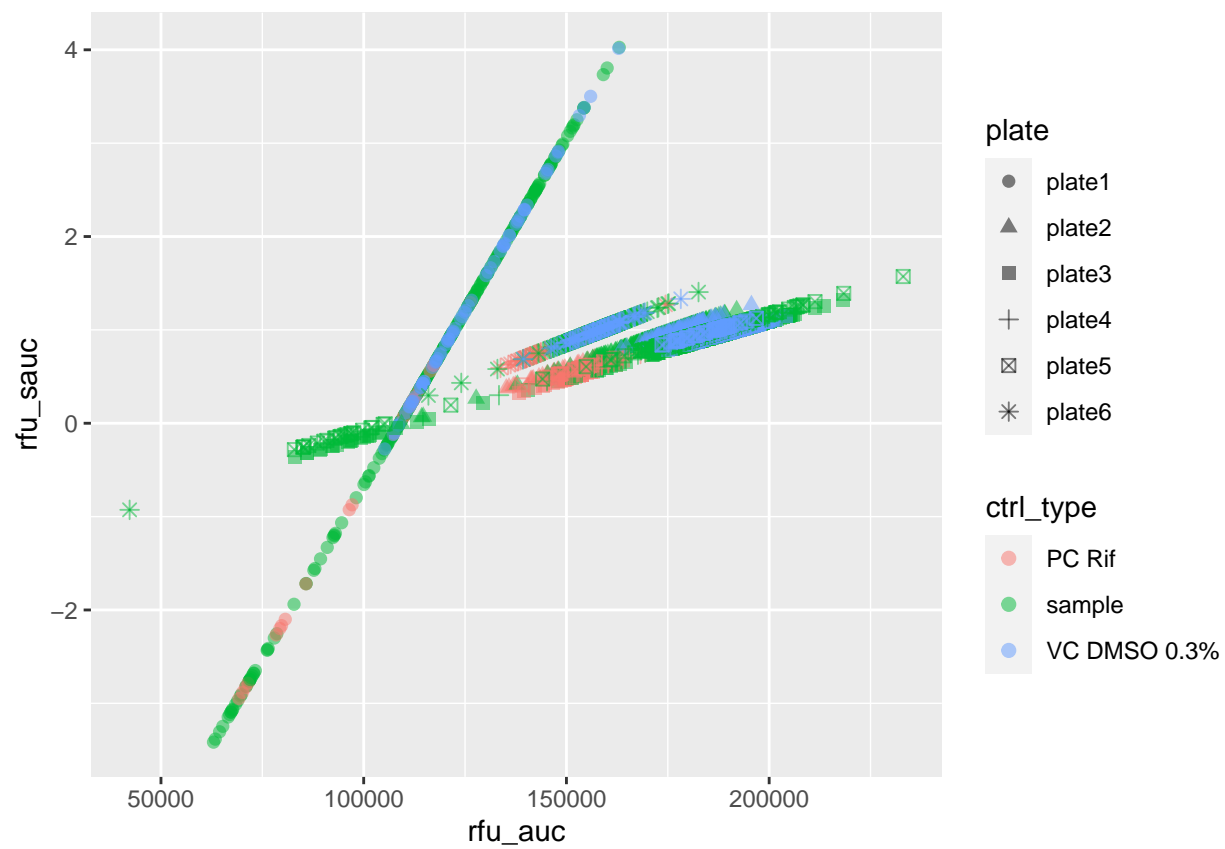


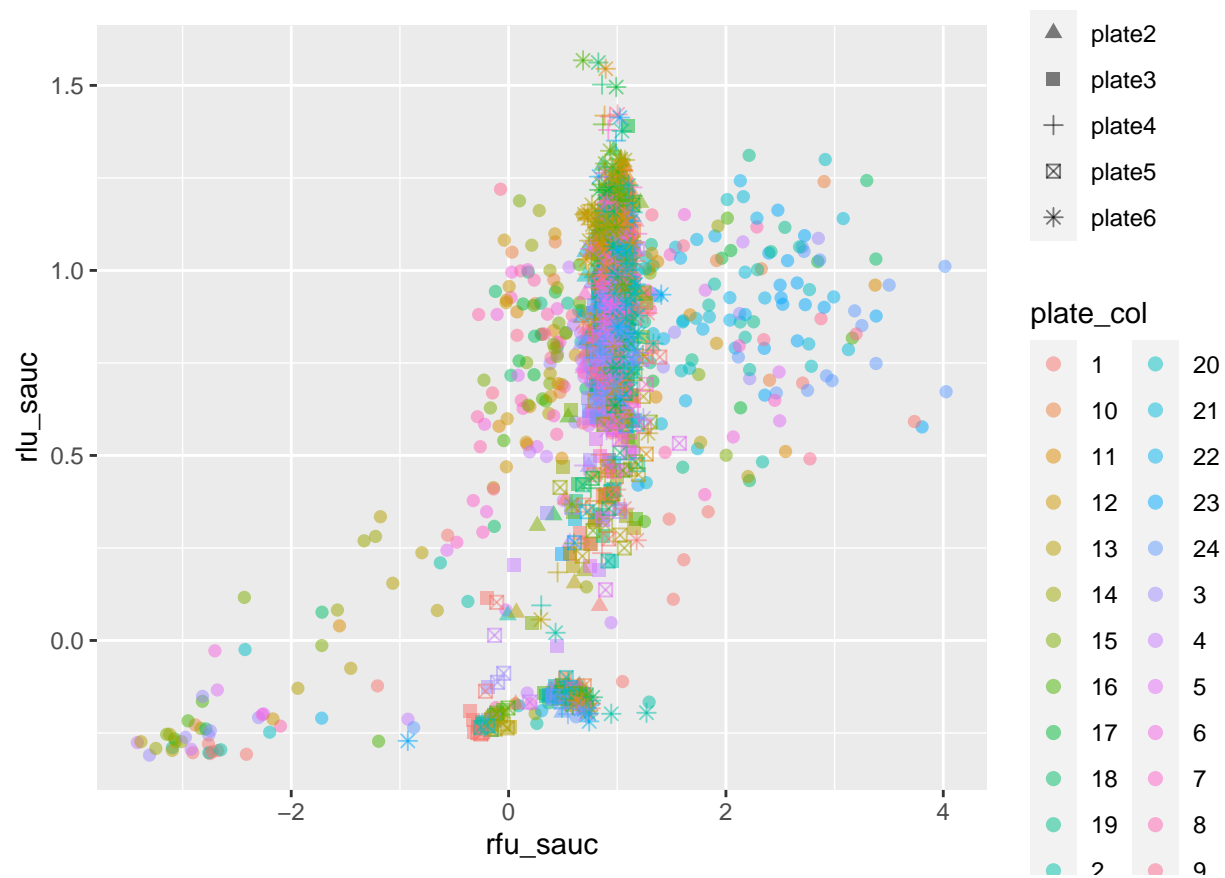


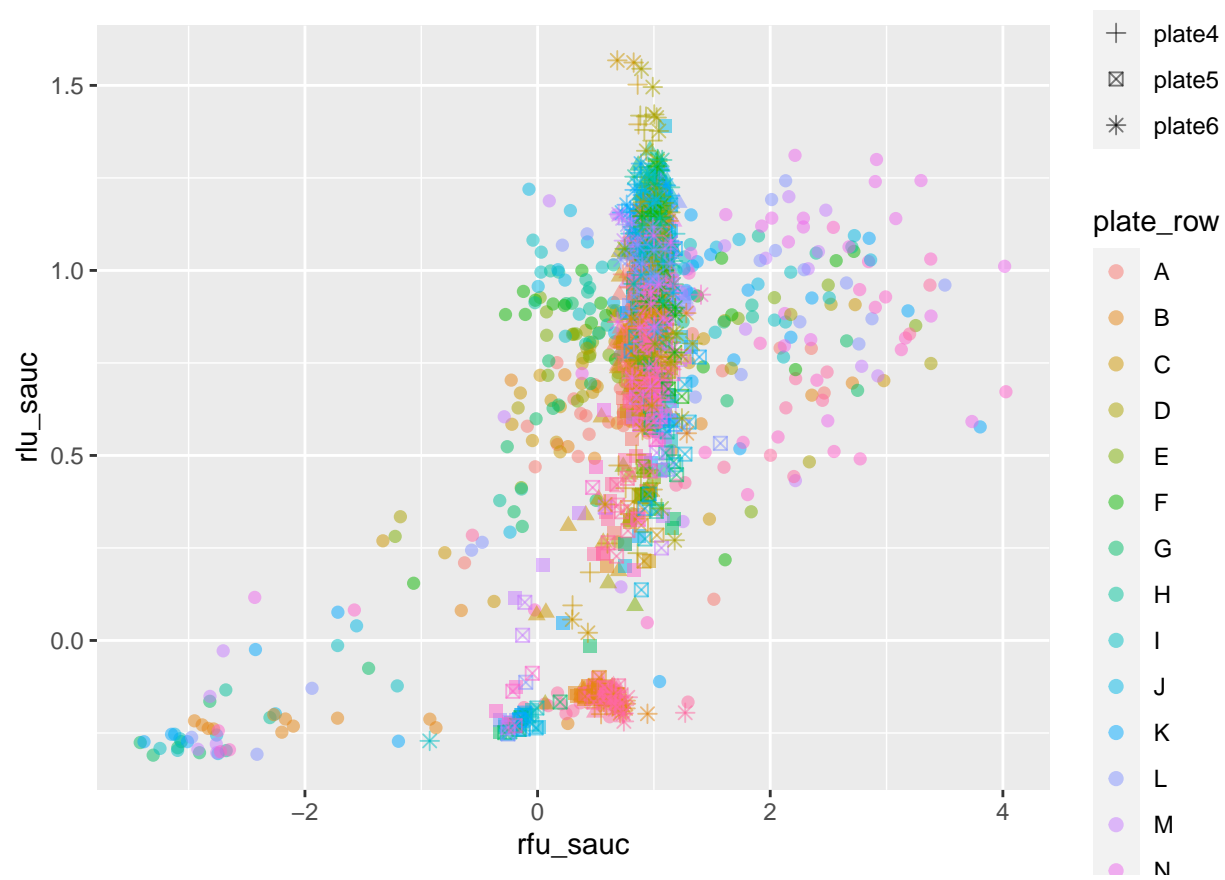


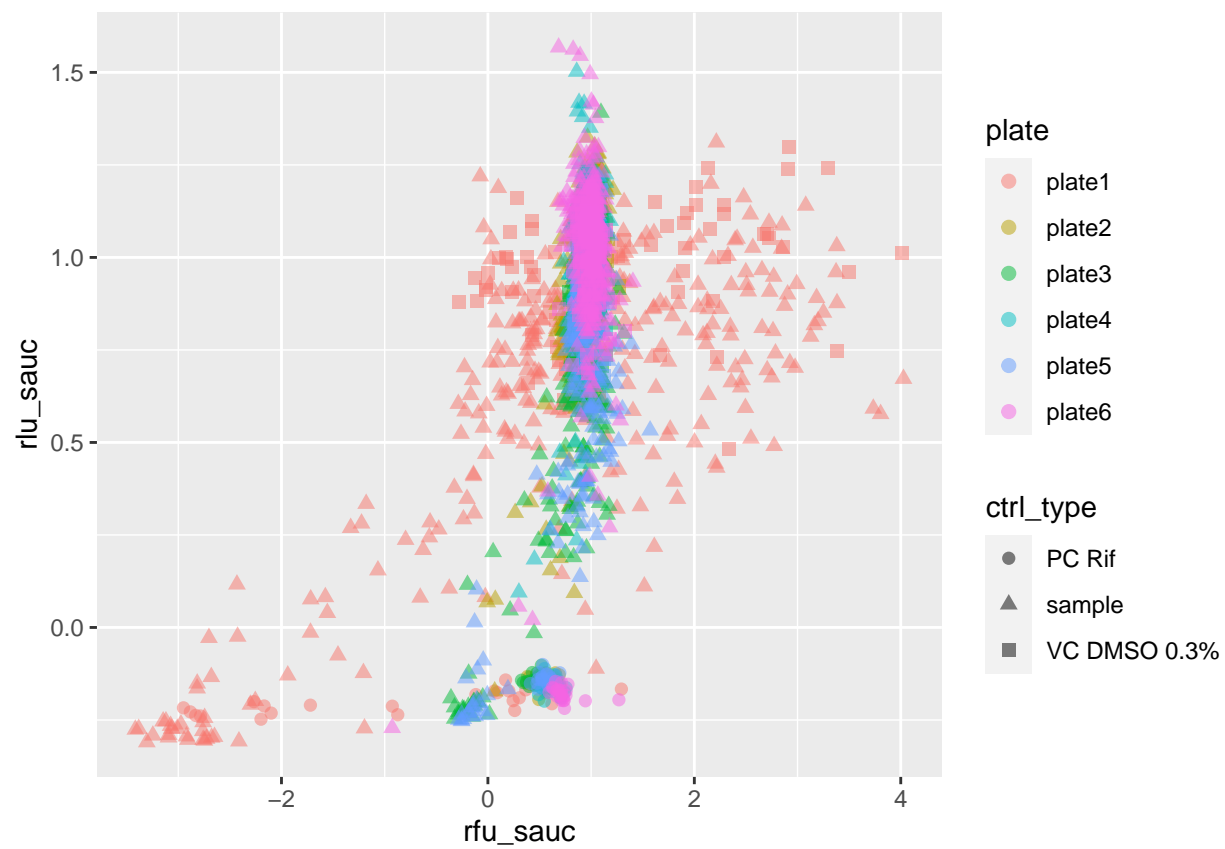


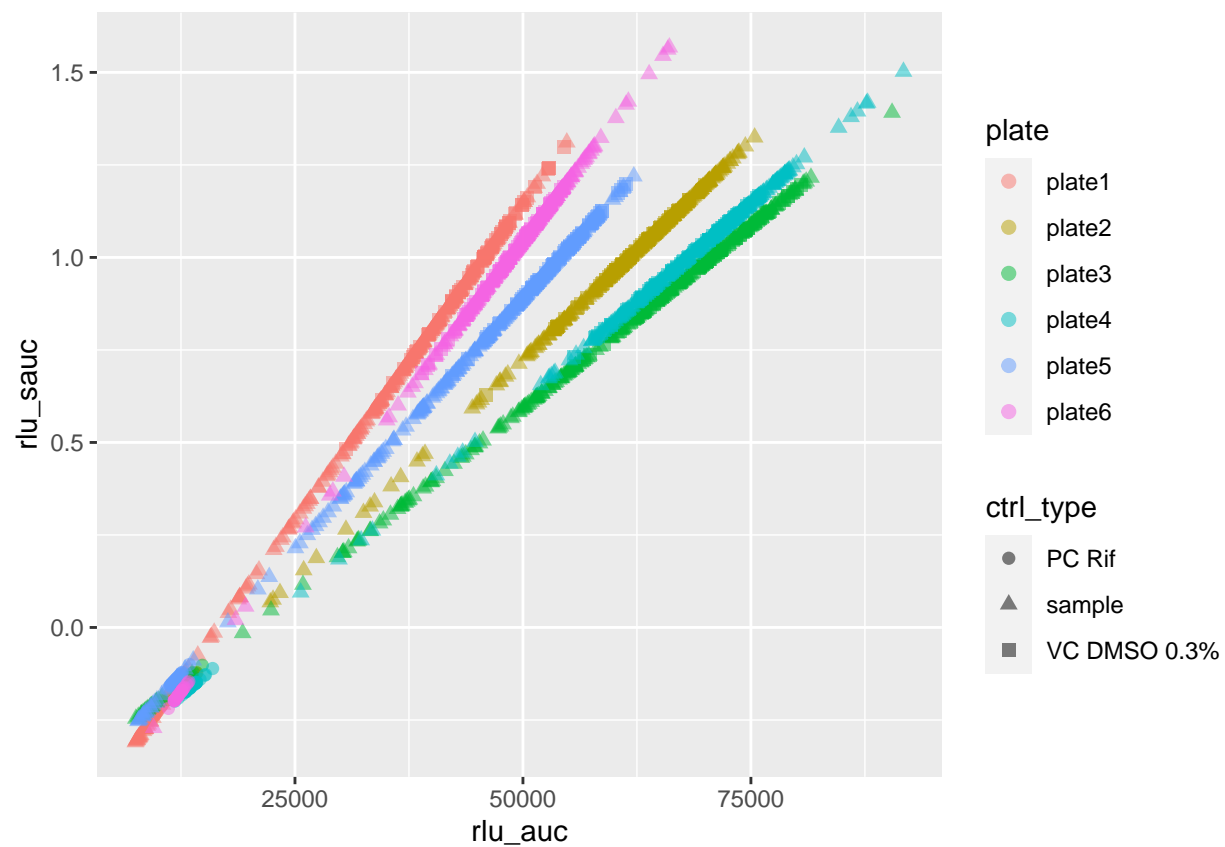


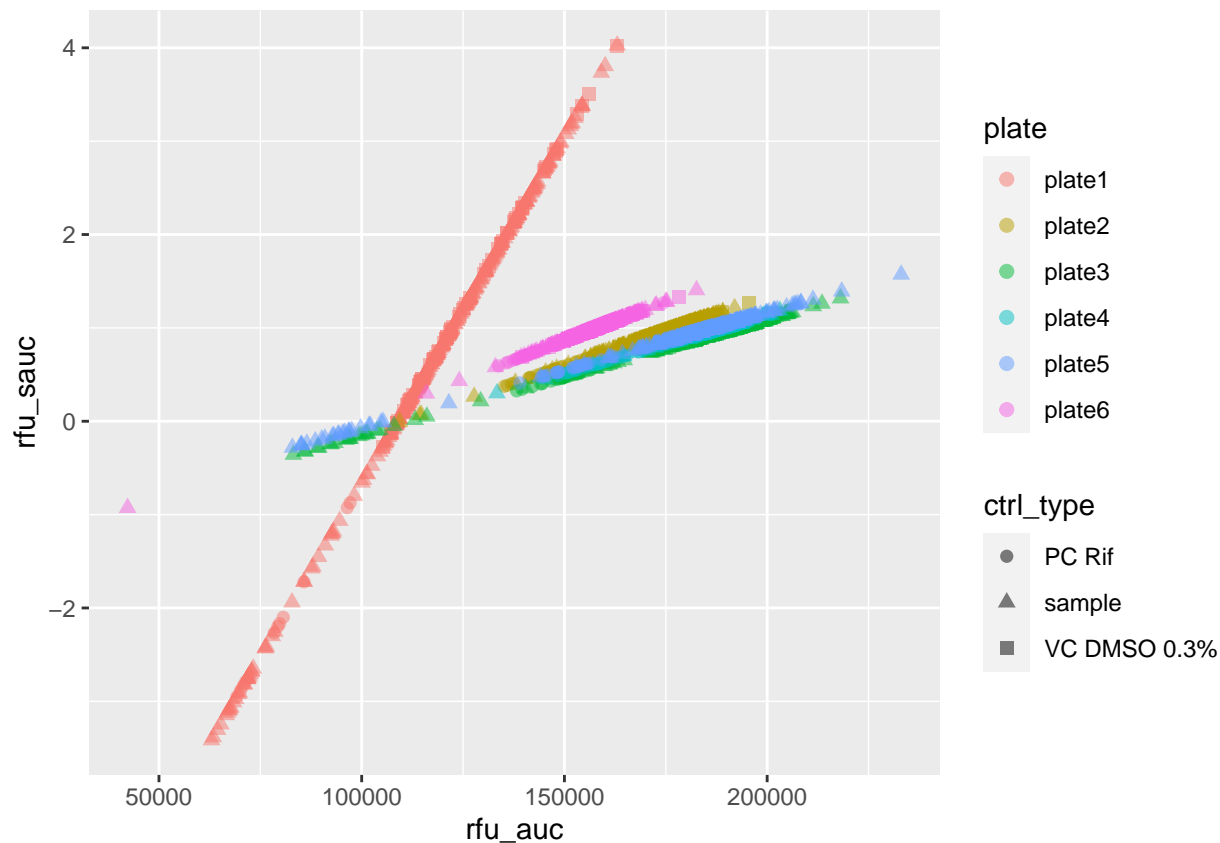












```
## R version 4.3.1 (2023-06-16 ucrt)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19045)
##
## Matrix products: default
##
##
## locale:
## [1] LC_COLLATE=English_Switzerland.utf8  LC_CTYPE=English_Switzerland.utf8
## [3] LC_MONETARY=English_Switzerland.utf8 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] tidyr_1.3.0  dplyr_1.1.3  ggplot2_3.4.4 gplots_3.1.3 readxl_1.4.3
## [6] xlsx_0.6.5
##
## loaded via a namespace (and not attached):
## [1] gtable_0.3.4      compiler_4.3.1    gtools_3.9.4      tidyselect_1.2.0
## [5] bitops_1.0-7      scales_1.2.1      yaml_2.3.7         fastmap_1.1.1
## [9] R6_2.5.1          labeling_0.4.3    generics_0.1.3     knitr_1.44
```

## [13]	tibble_3.2.1	munSELL_0.5.0	pillar_1.9.0	rlang_1.1.1
## [17]	utf8_1.2.3	xfun_0.40	caTools_1.18.2	cli_3.6.1
## [21]	withr_2.5.1	magrittr_2.0.3	digest_0.6.33	grid_4.3.1
## [25]	rstudioapi_0.15.0	rJava_1.0-6	lifecycle_1.0.3	vctrs_0.6.3
## [29]	KernSmooth_2.23-21	evaluate_0.22	glue_1.6.2	farver_2.1.1
## [33]	cellranger_1.1.0	xlsxjars_0.6.1	fansi_1.0.5	colorspace_2.1-0
## [37]	rmarkdown_2.25	purrr_1.0.2	tools_4.3.1	pkgconfig_2.0.3
## [41]	htmltools_0.5.6.1			