

Testing package with devtools

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02/02/2021

Notes before release

- package
 - **TODO** automatically load libraries when loading *npde* ()
 - ggplot2 not loaded, gridExtra, mclust not loaded
 - check: mclust
 - compilation: compiled package as npde 3.0
 - * version number 3.0 because reference profiles included
 - * lots of error messages for “no visible binding for global variable” in variables related to ggplot2 (Undefined global functions or variables) => hope this does not block us for CRAN
- TODO *npde website* (bookdown, Eco)
- code **TODO** for 3.1
 - add an option to remove all output when required and remove warnings if requested
- **TODO** simplify output for *show()* function (v3.0 or v3.1 ?)

Setup, loading libraries

Install package in development mode

```
dev_mode() # development mode

## Dev mode: ON
install.packages(pkgs=file.path(workDir, "npde_3.0.tar.gz"), repos=NULL)

## Installing package into 'C:/Users/Romai/OneDrive/Documents/R-dev'
## (as 'lib' is unspecified)
library(npde)
```

Running examples from documentation

Checking help files work

OK

```
## starting httpd help server ... done
```

Computing npde for theophylline data

Checking that the results are the same when we use data frames versus files on disk: OK

Missing data (lines with MDV=1) removed from the data for plots (also removed from simulated data for approx.pi=TRUE, as per documentation the simulated and observed data must have compatible dimensions).

```
## -----
```

```

## Distribution of npde :
##   nb of obs: 120
##   mean= 0.0668  (SE= 0.095 )
##   variance= 1.074  (SE= 0.14 )
##   skewness= 0.511
##   kurtosis= 0.2912
## -----
## 
## Statistical tests
##   t-test : 0.481
##   Fisher variance test : 0.55
##   SW test of normality : 0.00273 **
##   Global adjusted p-value : 0.00819 **
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## 

## Object of class NpdeObject
## -----
## ---- Component data ----
## 
## Object of class NpdeData
##   Structured data: Conc ~ Time | ID
## This object has the following components:
##   data: data
##   with 12 subjects
##   120 observations
## The data has the following components
##   X: Time (hr)
##   Y: Conc (mg/L)
##   missing data: mdv (1=missing)
## 
## ---- Component results ----
## 
## Object of class NpdeRes
##   containing the following elements:
##   predictions (ypred)
##   prediction discrepancies (pd)
##   normalised prediction distribution errors (npde)
##   completed responses (ycomp) for censored data
##   decorrelated responses (ydobs)
##   the dataframe has 120 non-missing observations and 132 lines.
## First 10 lines of results, removing missing observations:
##   ypred ycomp pd ydobs npde tnpde
## 2 2.923864 2.84 0.550 -0.05124648 0.1256613 5.403817
## 3 4.682299 6.57 0.850 1.96398150 2.0537489 11.302178
## 4 6.264357 10.50 0.990 2.56602650 2.3263479 12.136107
## 5 6.986255 9.66 0.980 0.41616411 0.5244005 6.623631
## 6 6.511039 8.58 0.930 0.28430866 0.2533471 5.794430
## 7 5.895675 8.36 0.960 0.54879386 0.6744898 7.082780
## 8 5.064736 7.47 0.970 1.79335938 1.6448536 10.051295
## 9 4.302909 6.89 0.990 0.80506269 0.7721932 7.381673
## 10 3.294020 5.94 0.995 1.91537662 1.7506861 10.375055
## 11 1.168743 3.28 0.995 3.25535923 2.5758293 12.899315

```

```

## -----
## Distribution of npde :
##     nb of obs: 120
##         mean= 0.0668   (SE= 0.095 )
##     variance= 1.074   (SE= 0.14 )
##     skewness= 0.511
##     kurtosis= 0.2912
## -----
## 
## Statistical tests
##     t-test                 : 0.481
##     Fisher variance test   : 0.55
##     SW test of normality   : 0.00273 **
##     Global adjusted p-value : 0.00819 **
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----
##      ypred  ycomp    pd      ydobs      npde      tnpde
## 1     NaN    NaN  NaN        NA       NA       NA
## 2 2.923864 2.84 0.55 -0.05124648 0.1256613 5.403817
## 3 4.682299 6.57 0.85  1.96398150 2.0537489 11.302178
## 4 6.264357 10.50 0.99  2.56602650 2.3263479 12.136107
## 5 6.986255 9.66 0.98  0.41616411 0.5244005  6.623631
## 6 6.511039 8.58 0.93  0.28430866 0.2533471  5.794430
## Same results with data frame and with data on disk: TRUE

```

Show and print function :

- modified so that print(theofit) shows the results of gof.test applied to the object
- show displays the first 10 lines of the results datafram
- **TODO** simplify output for *show()* function (v3.0 or v3.1 ?)

```

## Object of class NpdeObject
## -----
## ----- Component data -----
## 
## Object of class NpdeData
##     Structured data: Conc ~ Time | ID
## This object has the following components:
##     data: data
##     with 12 subjects
##     120 observations
## The data has the following components
##     X: Time (hr)
##     Y: Conc (mg/L)
##     missing data: mdv (1=missing)
## -----
## ----- Component results -----
## 
## Object of class NpdeRes
##     containing the following elements:
##     predictions (ypred)
##     prediction discrepancies (pd)
##     normalised prediction distribution errors (npde)

```

```

##      completed responses (ycomp) for censored data
##      decorrelated responses (ydobs)
##      the dataframe has 120 non-missing observations and 132 lines.
## First 10 lines of results, removing missing observations:
##      ypred ycomp    pd      ydobs    npde    tnpde
## 2  2.923864  2.84 0.550 -0.05124648 0.1256613 5.403817
## 3  4.682299  6.57 0.850  1.96398150 2.0537489 11.302178
## 4  6.264357 10.50 0.990  2.56602650 2.3263479 12.136107
## 5  6.986255  9.66 0.980  0.41616411 0.5244005 6.623631
## 6  6.511039  8.58 0.930  0.28430866 0.2533471 5.794430
## 7  5.895675  8.36 0.960  0.54879386 0.67444898 7.082780
## 8  5.064736  7.47 0.970  1.79335938 1.6448536 10.051295
## 9  4.302909  6.89 0.990  0.80506269 0.7721932 7.381673
## 10 3.294020  5.94 0.995  1.91537662 1.7506861 10.375055
## 11 1.168743  3.28 0.995  3.25535923 2.5758293 12.899315

## Object of class NpdeObject
## -----
## ----- Data -----
## -----
## Object of class NpdeData
##   longitudinal data
##   Structured data: Conc ~ Time | ID
##   predictor: Time (hr)
## Dataset characteristics:
##   number of subjects:      12
##   number of non-missing observations: 120
##   average/min/max nb obs: 10.00 / 10 / 10
## First 10 lines of data:
##   index ID  Time  Conc mdv
## 1     1  1  0.00   NA   1
## 2     1  1  0.25  2.84   0
## 3     1  1  0.57  6.57   0
## 4     1  1  1.12 10.50   0
## 5     1  1  2.02  9.66   0
## 6     1  1  3.82  8.58   0
## 7     1  1  5.10  8.36   0
## 8     1  1  7.03  7.47   0
## 9     1  1  9.05  6.89   0
## 10    1  1 12.12  5.94   0
##
## Summary of original data:
##   vector of predictor Time
##   Min. 1st Qu. Median Mean 3rd Qu. Max.
##   0.000  0.595  3.530  5.895  9.000 24.650
##   vector of response Conc
##   Min. 1st Qu. Median Mean 3rd Qu. Max. NA's
##   0.850  3.513  5.665  5.447  7.325 11.400      12
## -----
## ----- Key options -----
## -----
## Methods
##   compute prediction discrepancies (pd): yes
##   compute normalised prediction distribution errors (npde): yes

```

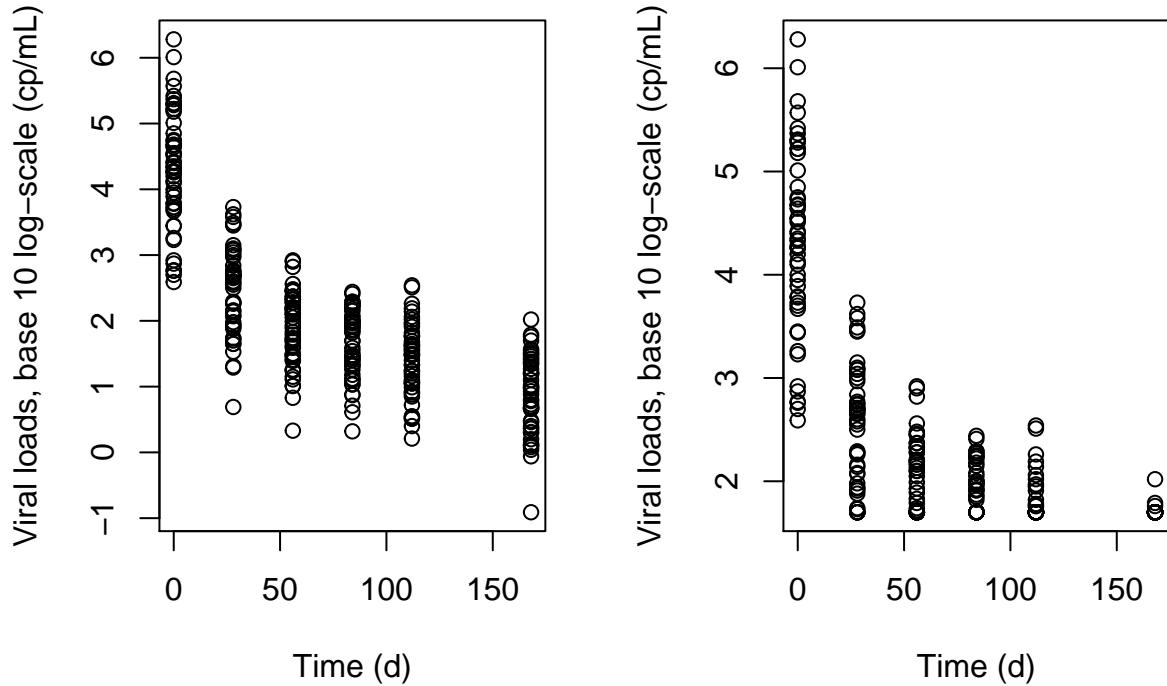
```

##      method for decorrelation: Cholesky decomposition (upper triangular)
##      method to treat censored data: Impute pd* and compute y* as F-1(pd*)
## Input/output
##      verbose (prints a message for each new subject): FALSE
##      save the results to a file, save graphs: FALSE
## -----
## ----- Results -----
## -----
## Object of class NpdeRes
##      resulting from a call to npde or autopnde
##      containing the following elements:
##      predictions (ypred)
##      Min. 1st Qu. Median Mean 3rd Qu. Max.
##      0.8897 4.0194 5.5705 5.5208 7.1204 10.0162
##      prediction discrepancies (pd)
##      Min. 1st Qu. Median Mean 3rd Qu. Max.
##      0.0500 0.2600 0.4300 0.4807 0.7100 0.9950
##      normalised prediction distribution errors (npde)
##      Min. 1st Qu. Median Mean 3rd Qu. Max.
##      -2.3263 -0.5102 0.0000 0.0668 0.5244 2.5758
##      completed responses (ycomp) for censored data
##      decorrelated responses (ydobs)
##      the dataframe has 120 non-missing observations and 132 lines.
## -----
## Distribution of npde :
##      nb of obs: 120
##      mean= 0.0668 (SE= 0.095 )
##      variance= 1.074 (SE= 0.14 )
##      skewness= 0.511
##      kurtosis= 0.2912
## -----
## 
## Statistical tests
##      t-test : 0.481
##      Fisher variance test : 0.55
##      SW test of normality : 0.00273 **
##      Global adjusted p-value : 0.00819 **
## -----
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----

```

Computing npde for viral load data

- For censored viral load data (20 or 50 cp/mL), we compute the npde using the different methods implemented in the package. We expect:
 - NS for yvir1, yvir20, yvir50 (complete datasets, and censored datasets with imputed BQL)
 - significant for ppred method
 - NS for omit method because we also omit BQL data in the simulations (so OK)
 - NS for ipred method



```

## -----
## Distribution of npde :
##   nb of obs: 300
##   mean= 0.03821  (SE= 0.053 )
##   variance= 0.8327  (SE= 0.068 )
##   skewness= -0.04464
##   kurtosis= -0.2207
## -----
## 
## Statistical tests
##   t-test : 0.469
##   Fisher variance test : 0.032 *
##   SW test of normality : 0.845
##   Global adjusted p-value : 0.0959 .
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## 

## -----
## Distribution of npde :
##   nb of obs: 300
##   mean= -0.0343  (SE= 0.06 )
##   variance= 1.072  (SE= 0.088 )
##   skewness= 0.009137
##   kurtosis= 0.1297
## -----
## 
```

```

## Statistical tests
##   t-test : 0.566
##   Fisher variance test : 0.375
##   SW test of normality : 0.898
## Global adjusted p-value : 1
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----
## 
## Distribution of npde :
##   nb of obs: 300
##   mean= -0.0518 (SE= 0.052 )
##   variance= 0.8019 (SE= 0.066 )
##   skewness= -0.1503
##   kurtosis= 0.02162
## -----
## 
## Statistical tests
##   t-test : 0.317
##   Fisher variance test : 0.00998 **
##   SW test of normality : 0.377
## Global adjusted p-value : 0.0299 *
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----
## 
## Distribution of npde :
##   nb of obs: 300
##   mean= -0.02917 (SE= 0.053 )
##   variance= 0.8509 (SE= 0.07 )
##   skewness= -0.1542
##   kurtosis= 0.03102
## -----
## 
## Statistical tests
##   t-test : 0.584
##   Fisher variance test : 0.0582 .
##   SW test of normality : 0.687
## Global adjusted p-value : 0.175
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----
## 
## Distribution of npde :
##   nb of obs: 169
##   mean= 0.1433 (SE= 0.07 )
##   variance= 0.8186 (SE= 0.089 )
##   skewness= -0.03812
##   kurtosis= -0.3733
## -----
## 
## Statistical tests

```

```

##   t-test          : 0.041 *
##   Fisher variance test    : 0.0822 .
##   SW test of normality     : 0.687
## Global adjusted p-value    : 0.123
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----
## -----
## Distribution of npde :
##   nb of obs: 300
##   mean= 0.03101  (SE= 0.057 )
##   variance= 0.9715  (SE= 0.079 )
##   skewness= -0.006498
##   kurtosis= 0.8122
## -----
## 
## Statistical tests
##   t-test          : 0.586
##   Fisher variance test    : 0.746
##   SW test of normality     : 0.00121 **
## Global adjusted p-value    : 0.00364 **
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----
## -----
## Distribution of npde :
##   nb of obs: 300
##   mean= 0.03058  (SE= 0.062 )
##   variance= 1.164  (SE= 0.095 )
##   skewness= 0.04433
##   kurtosis= -0.05092
## -----
## 
## Statistical tests
##   t-test          : 0.624
##   Fisher variance test    : 0.0539 .
##   SW test of normality     : 0.973
## Global adjusted p-value    : 0.162
## ---
## Signif. codes: '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1
## -----

```

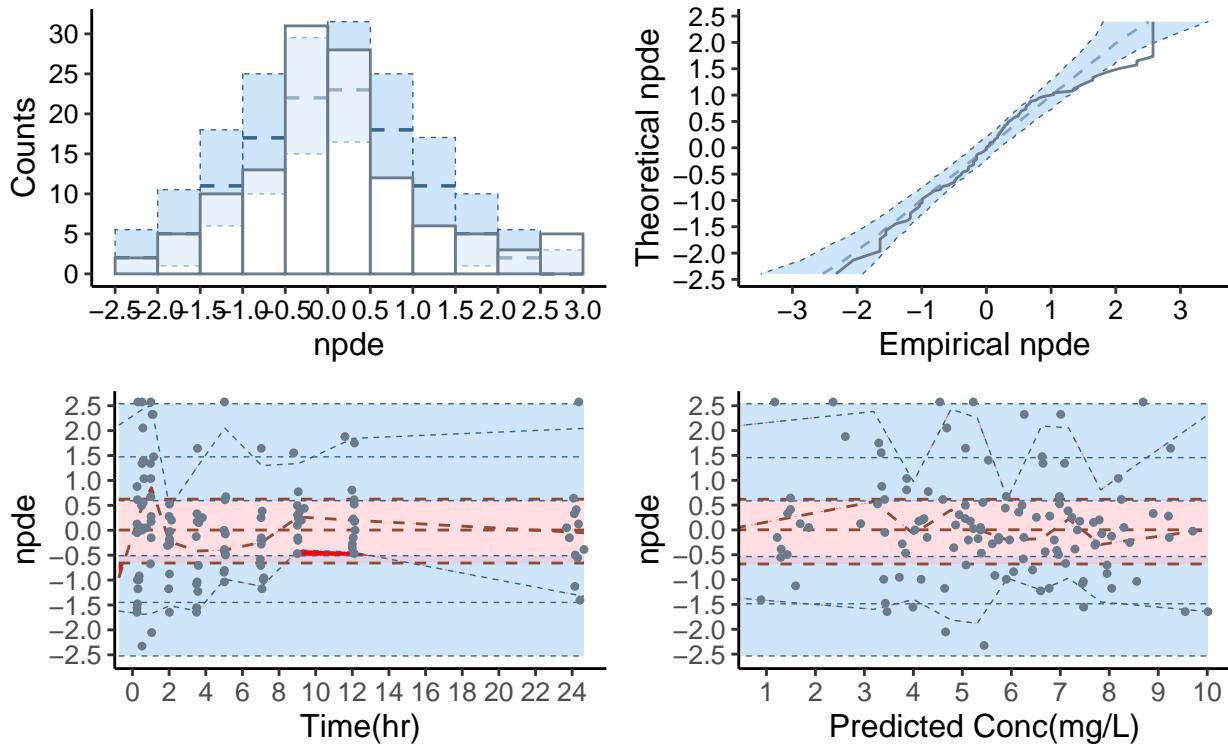
Test plots - default options

Default plots for theophylline data

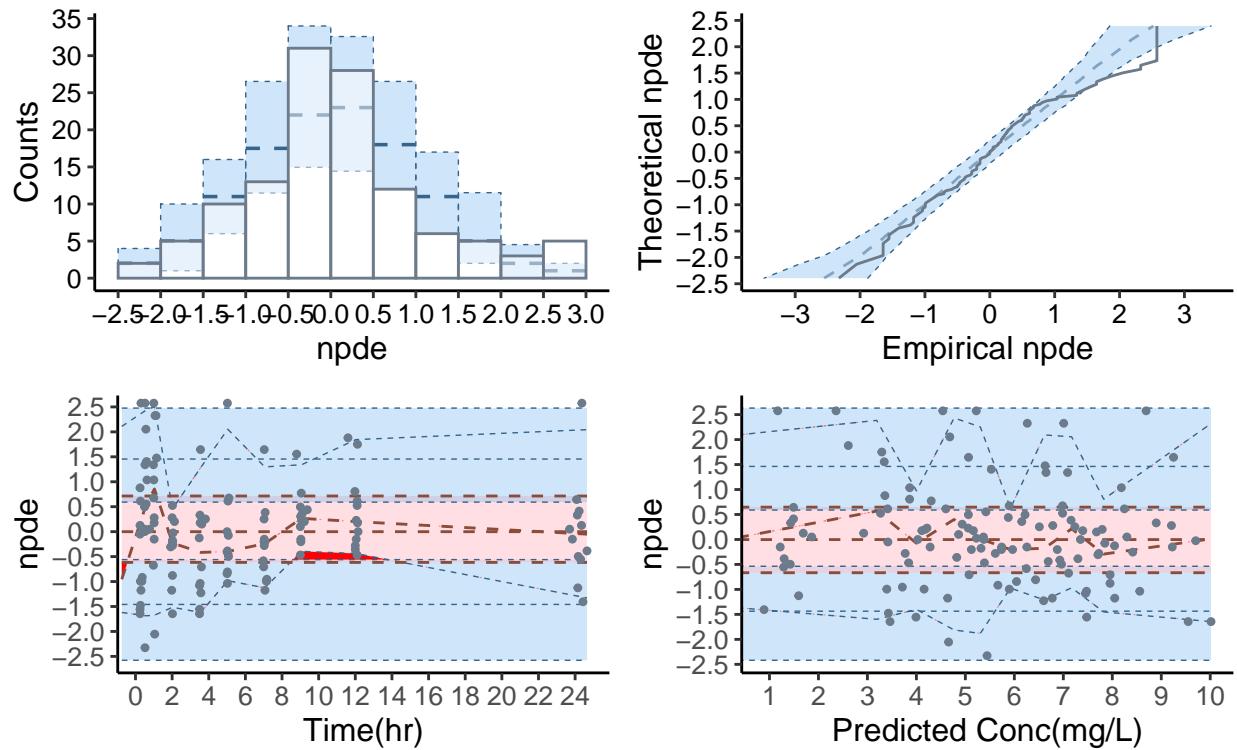
- **BUGS Romain TODOO**

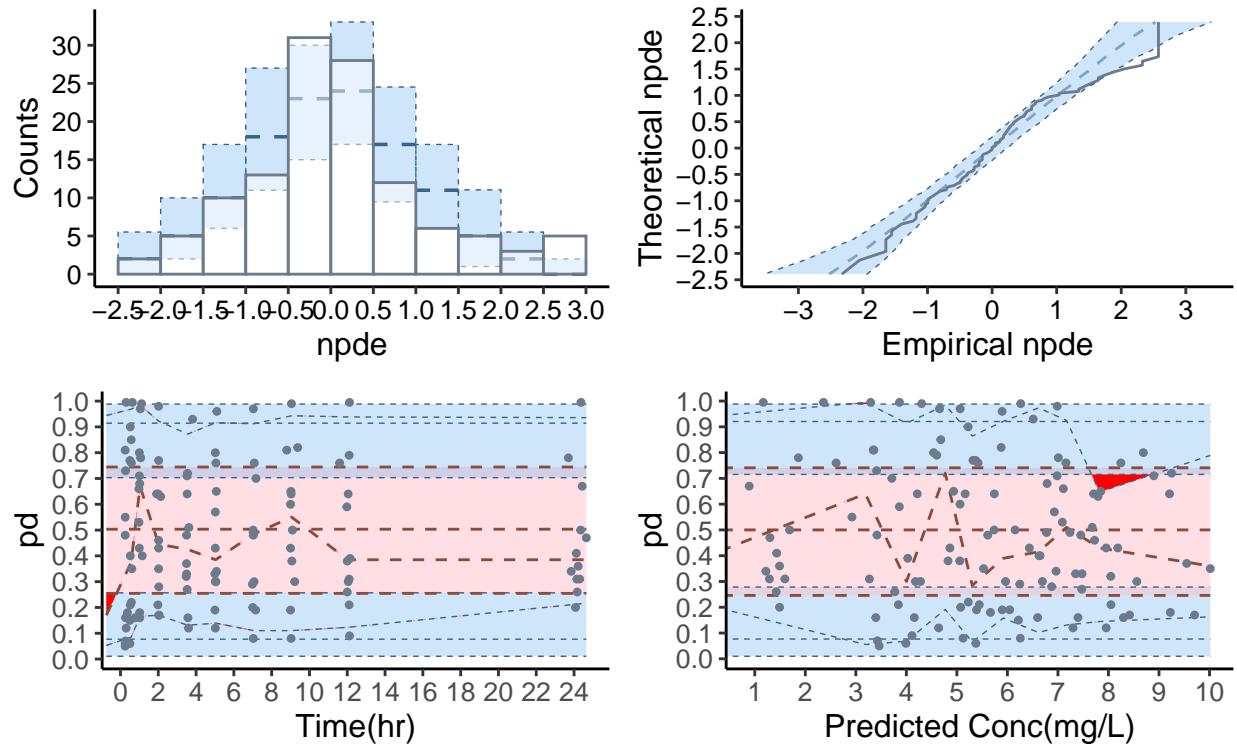
- **solved** title: in the waffle plot, only one title should appear, above all the 4 graphs (not one title per graph)
- ajout de la condition plot.opt\$plot.default==FALSE) pour l'affichage des titres dans les plot
- **solved** no axes in the histogram plot => add
- **solved X-range on PI** extend prediction band to min/max X (it seems to stop at the center of the bin) => option ???

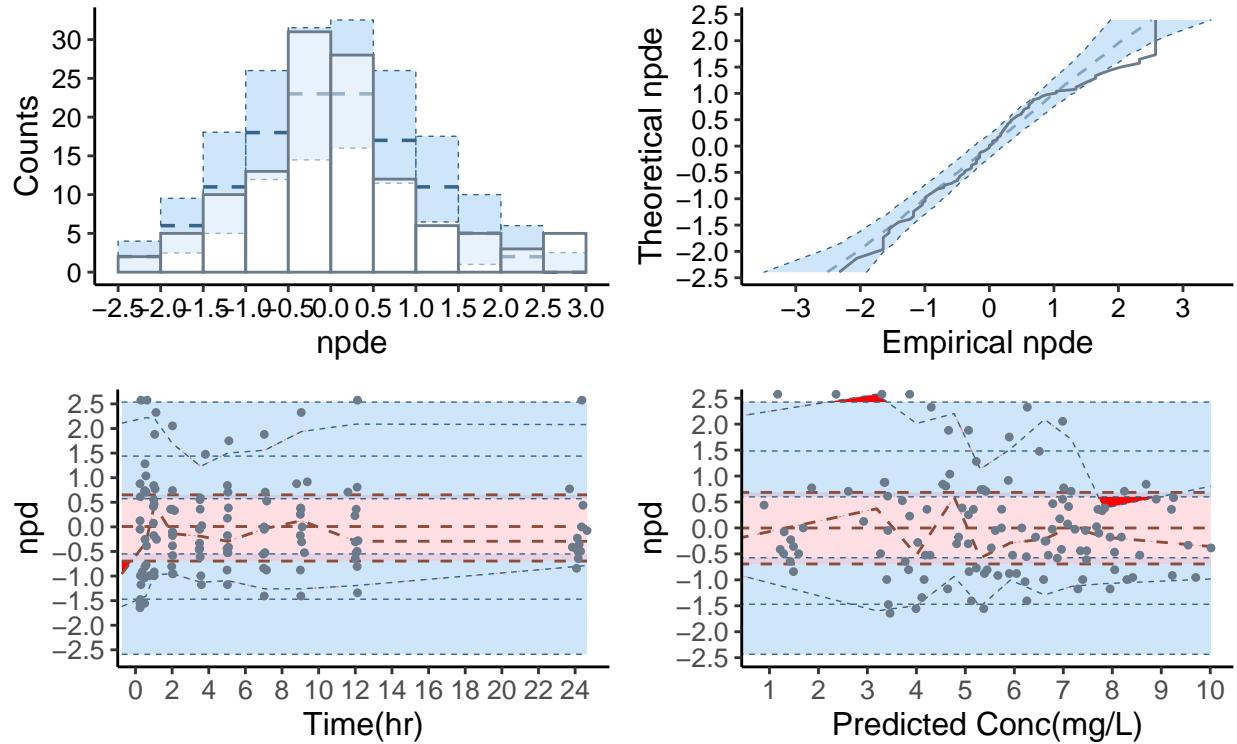
- dans plotNpde-auxScatterPlot: 1er bin avec valeur minimal de la borne et dernier bin avec valeur maximal de la borne. 1er et dernier le plot de s'arrête pas au milieu.
- **global options** not passing on to the functions
 - * **solved** size
 - **solved** normalement si l'utilisateur spécifie "size=2" sans spécifier size.pobs, size.pobs devrait passer à 1.5, et là ça ne semble rien changer
 - **solved** ni dans le graphe par défaut (ie les 4 graphes), ni individuellement
 - * **solved** colours
 - **solved** when only col is specified, it should affect other elements (but not sure which we had decided), here only affects histogram and col.lobs for qqplot
 - **solved** col => col.pobs, col.lobs, col.pcens ?
 - **solved** col.lobs ne change rien, devrait changer les lignes correspondant aux percentiles et les lignes dans l'histogramme
- **WARNING** attention aux effets de bord !!!
 - dans **set.plotoptions.NpdeObject**, xlab et ylab étaient fixés par défaut or on veut pouvoir les fixer automatiquement dans les fonctions graphiques tout en laissant l'utilisateur les passer en argument => on ne peut pas les mettre par défaut (xlab et ylab dépendent ici des variables qu'on veut tracer) => réfléchir avant de faire un truc comme ça (ou bien documenter qu'on puisse voir pourquoi ça plante, sinon on perd du temps à trouver pourquoi ça ne marche pas)
- **Note** : dans la dernière version, la fonction NpdeControl a toujours dans ses arguments xlab="" et ylab="" + which et xaxis Dabs plotNpde-distributionPlot 1.156-1.168 il y a différents tests qui restent en conséquence. Pour le moment ça n'empêche pas à l'utilisateur/utilisatrice de changer les labels à sa guise.



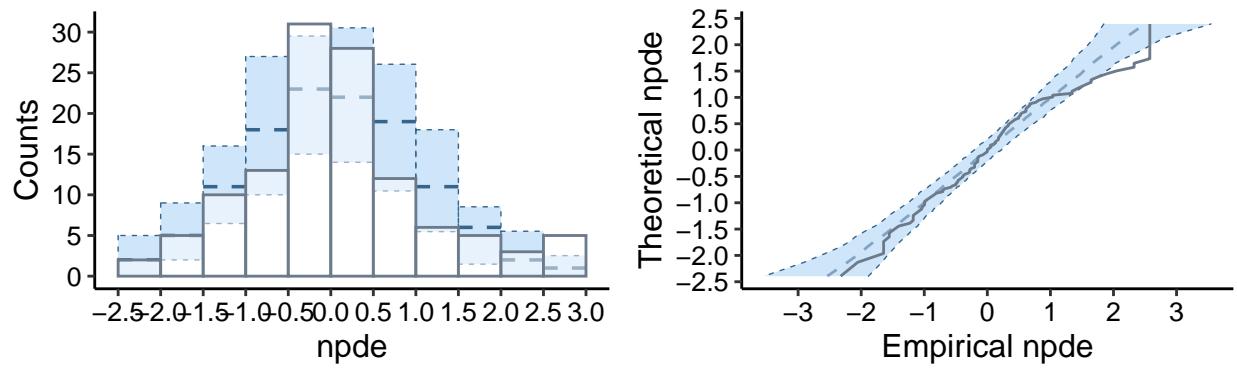
Theophylline data

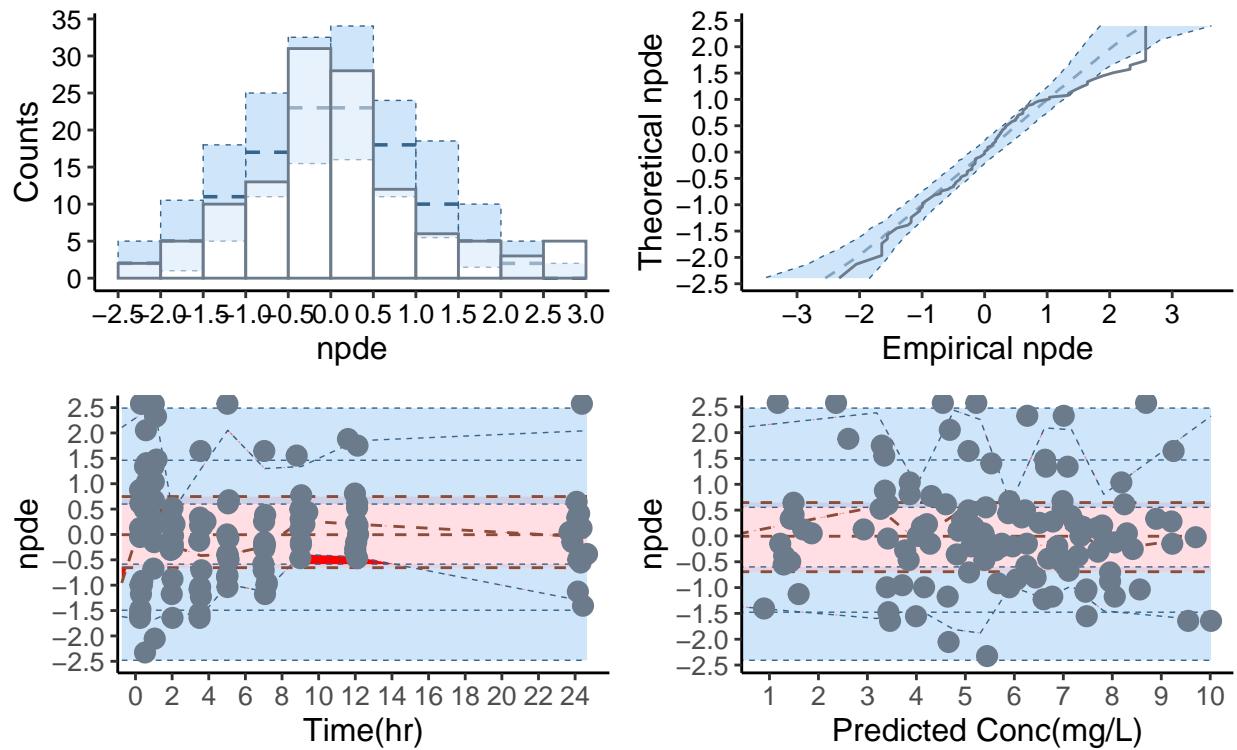


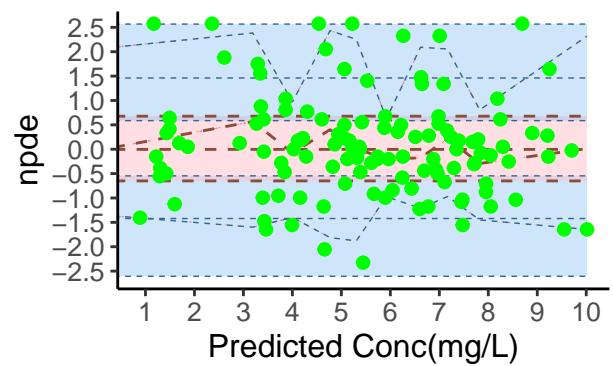
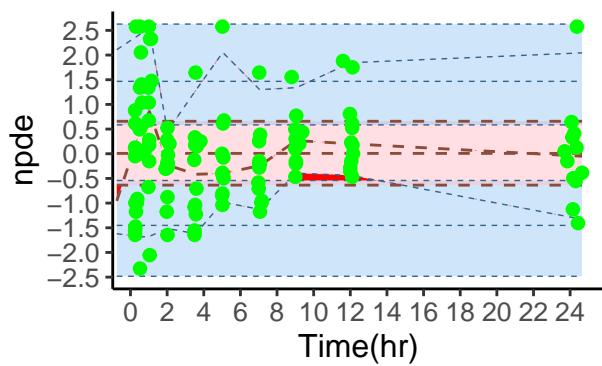
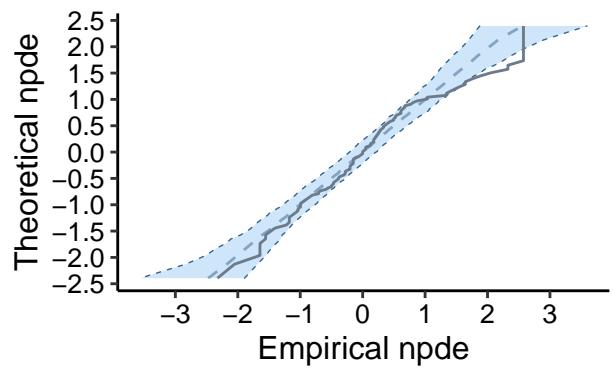
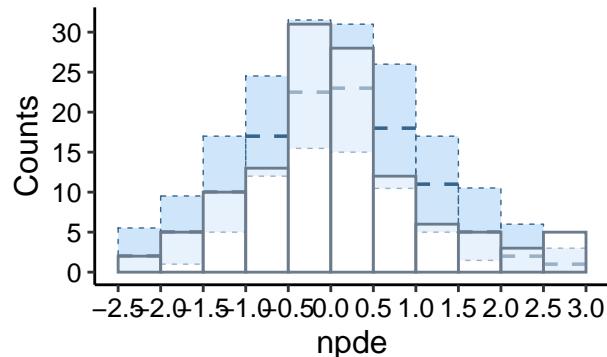


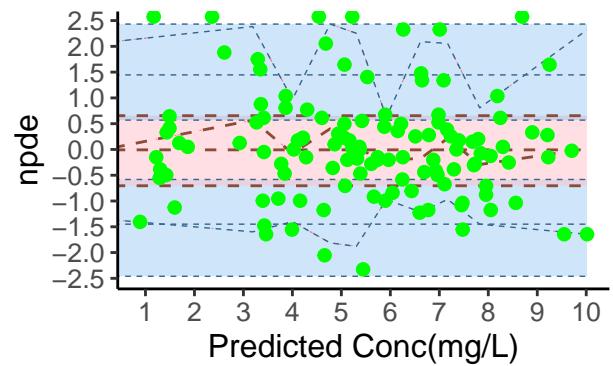
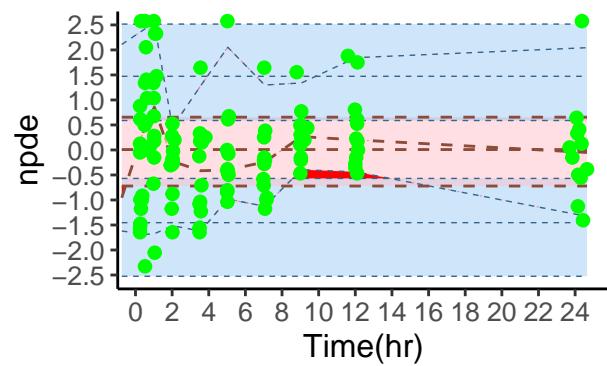
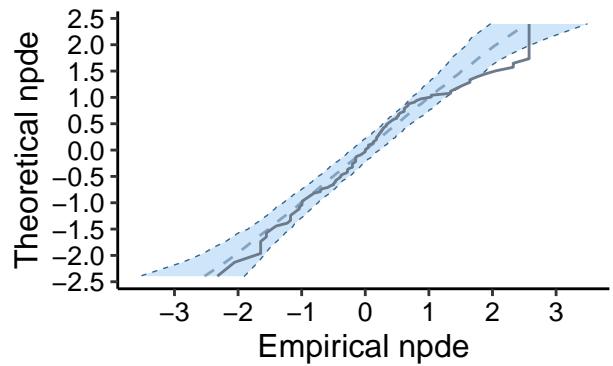
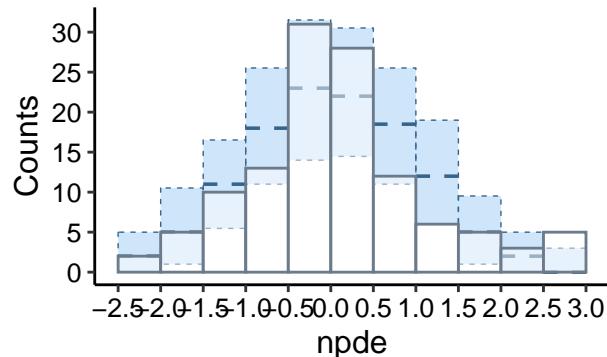


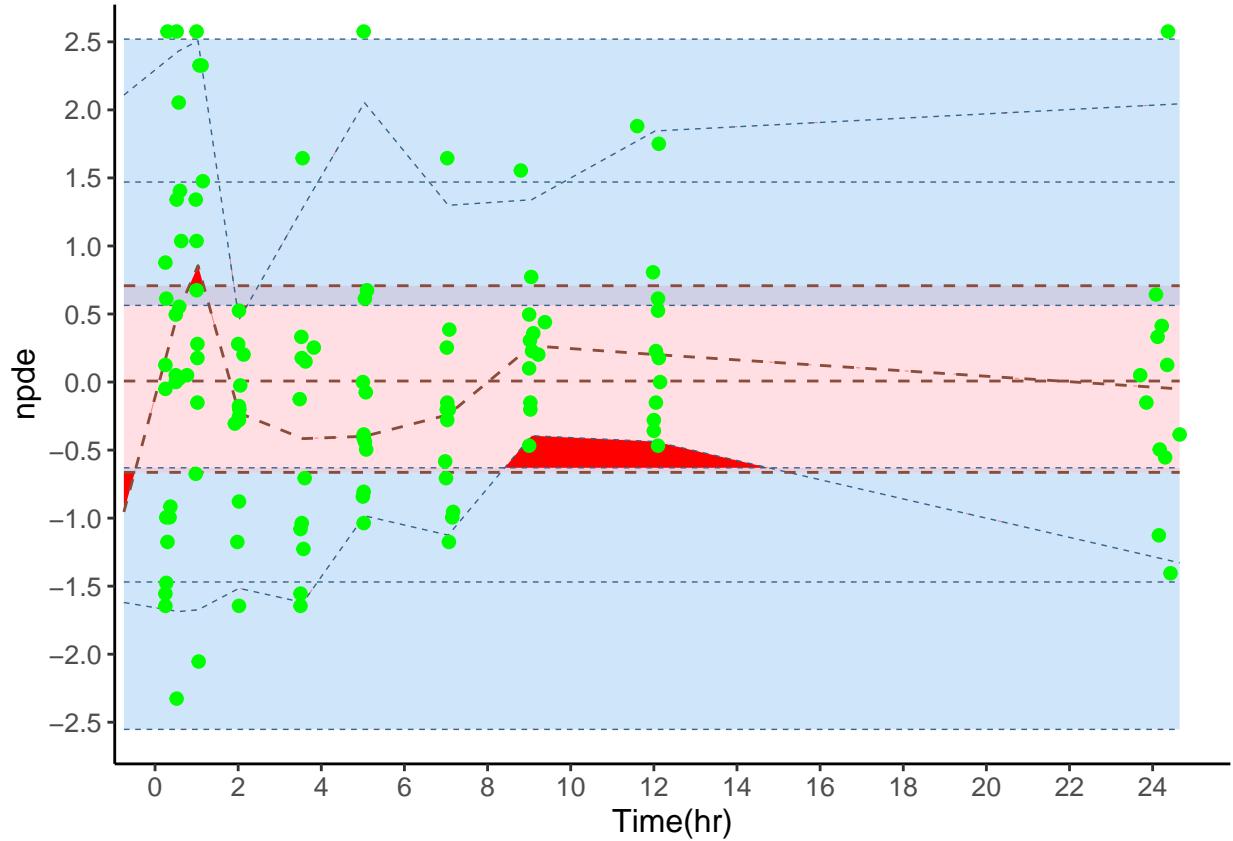
```
## Option which.y= faux not recognised
## Option which.y= faux not recognised
```

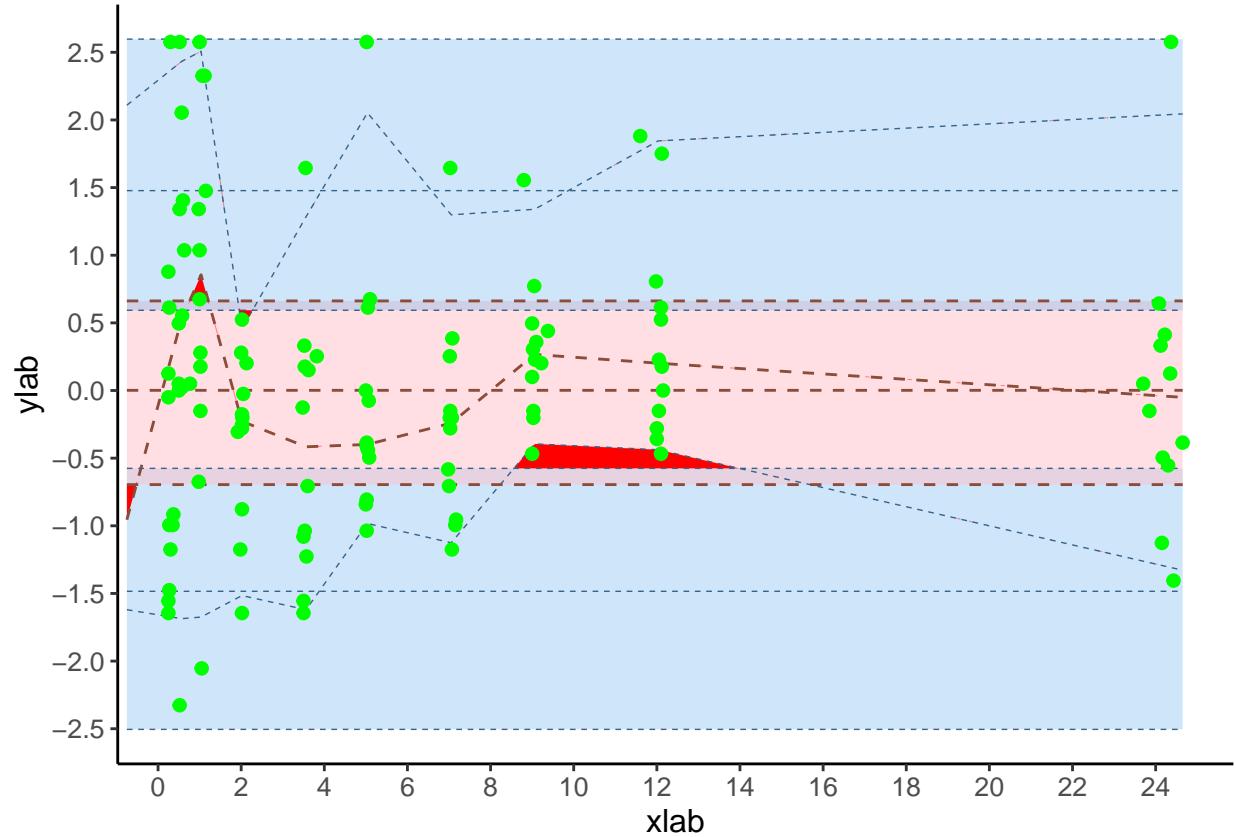








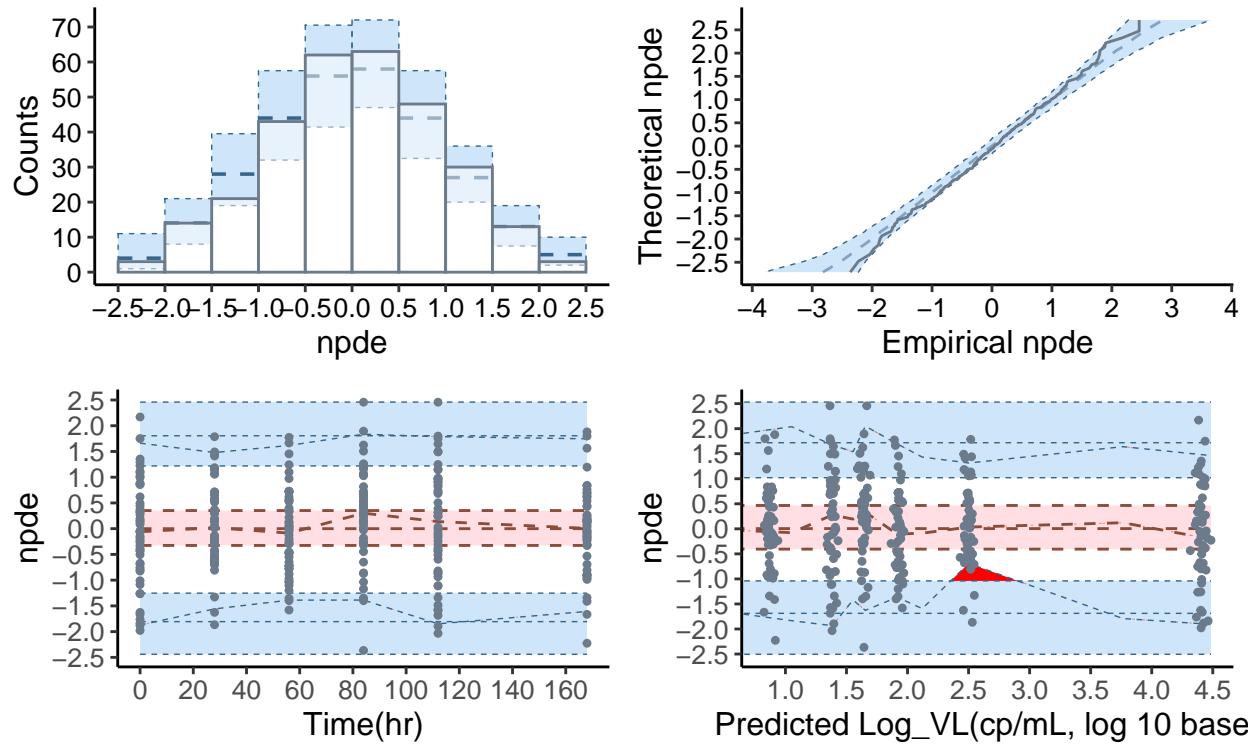




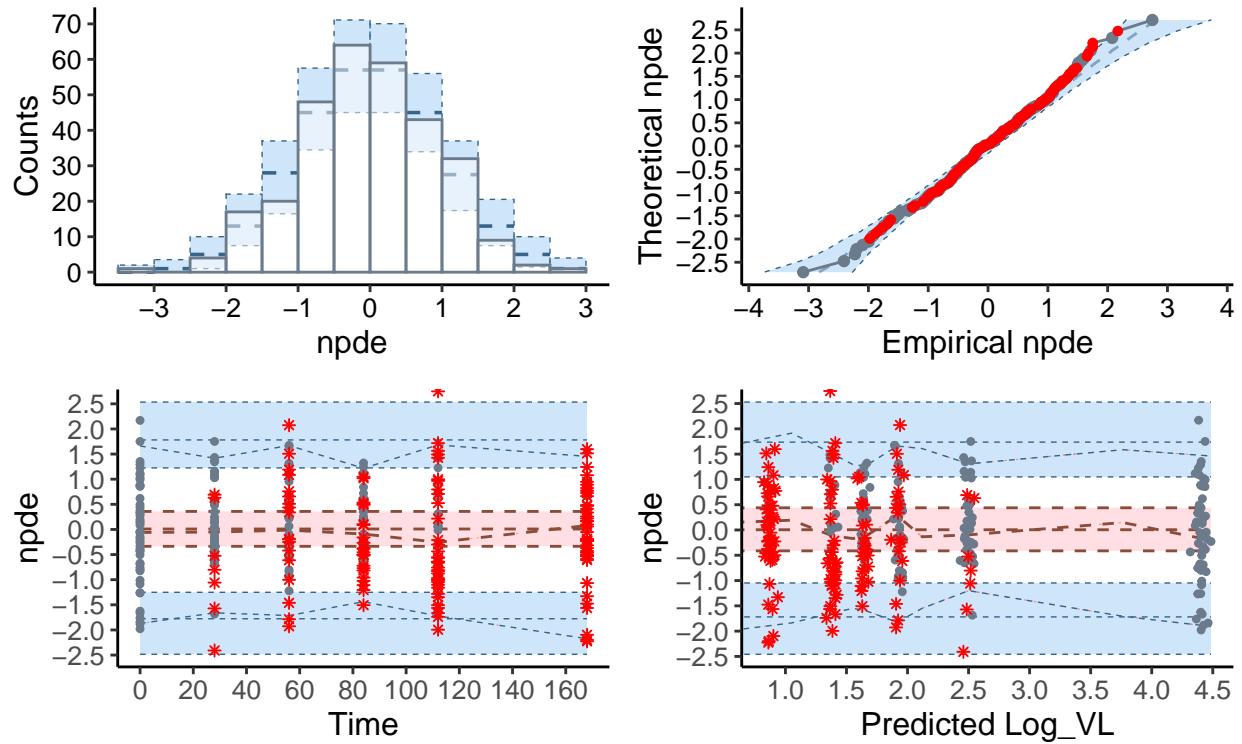
Default plots for viral load data

- computation of the PI: seems appropriate in the different cases
- **Problems (Eco)**
 - PI for omit aren't consistent with the VPC later on
 - * **documentation** explain how PI are computed in the omit case

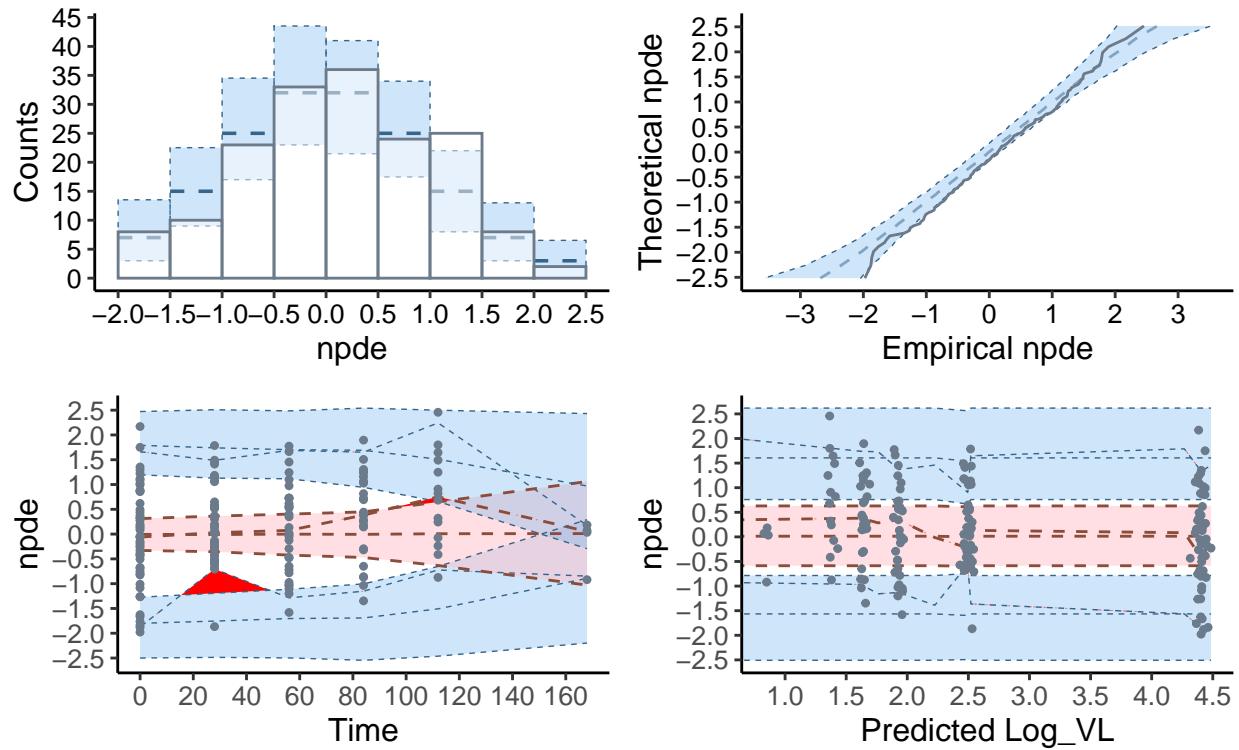
Complete dataset



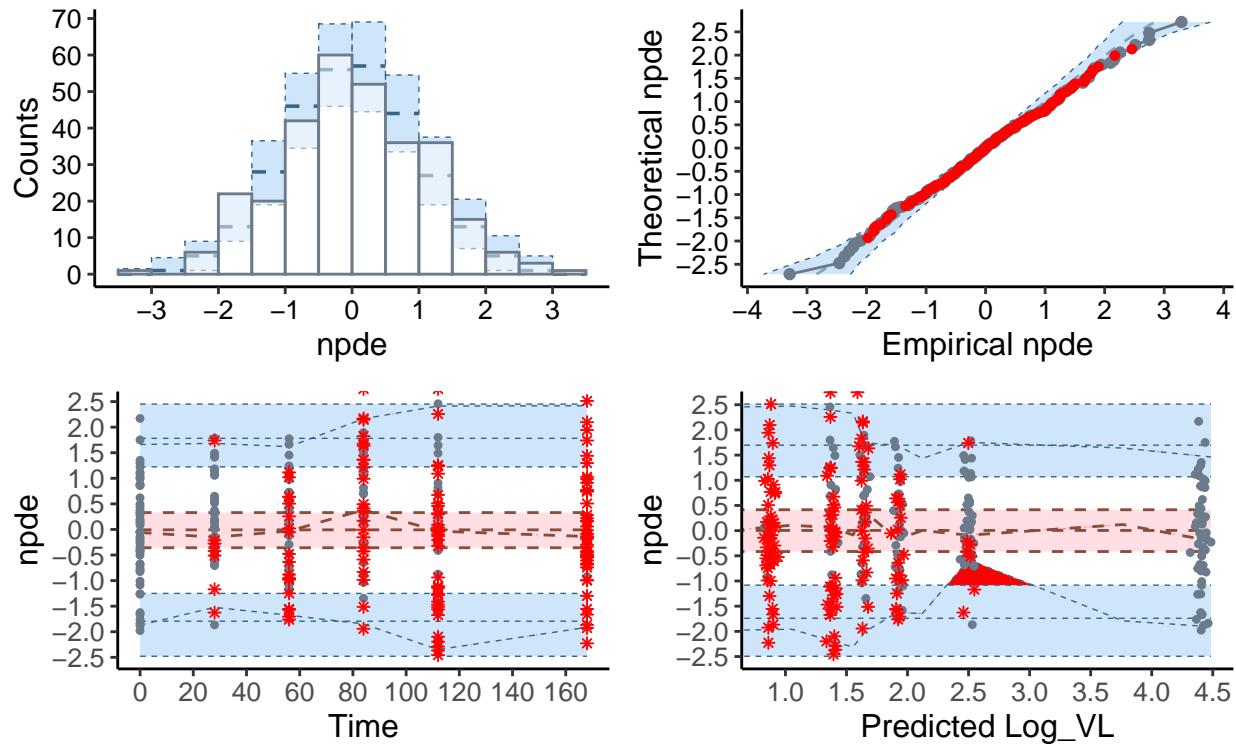
Dataset censored 50 cp/mL



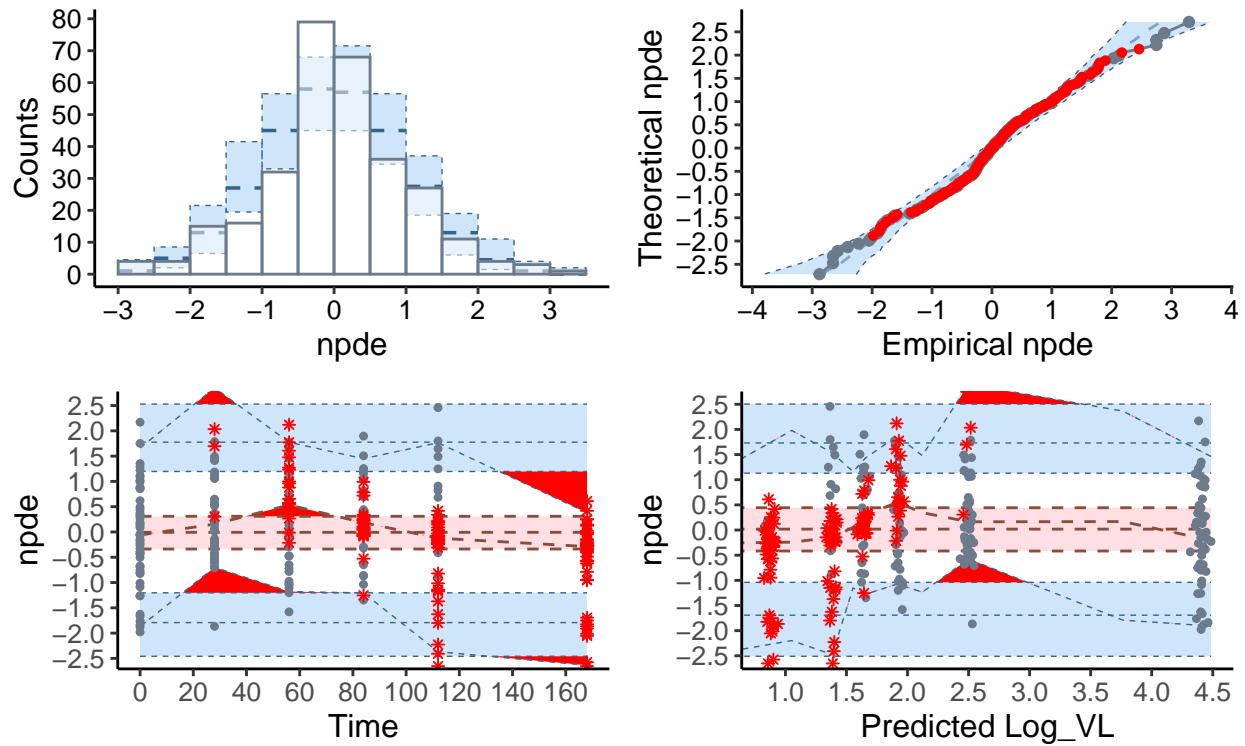
Censored 50, censoring=omit

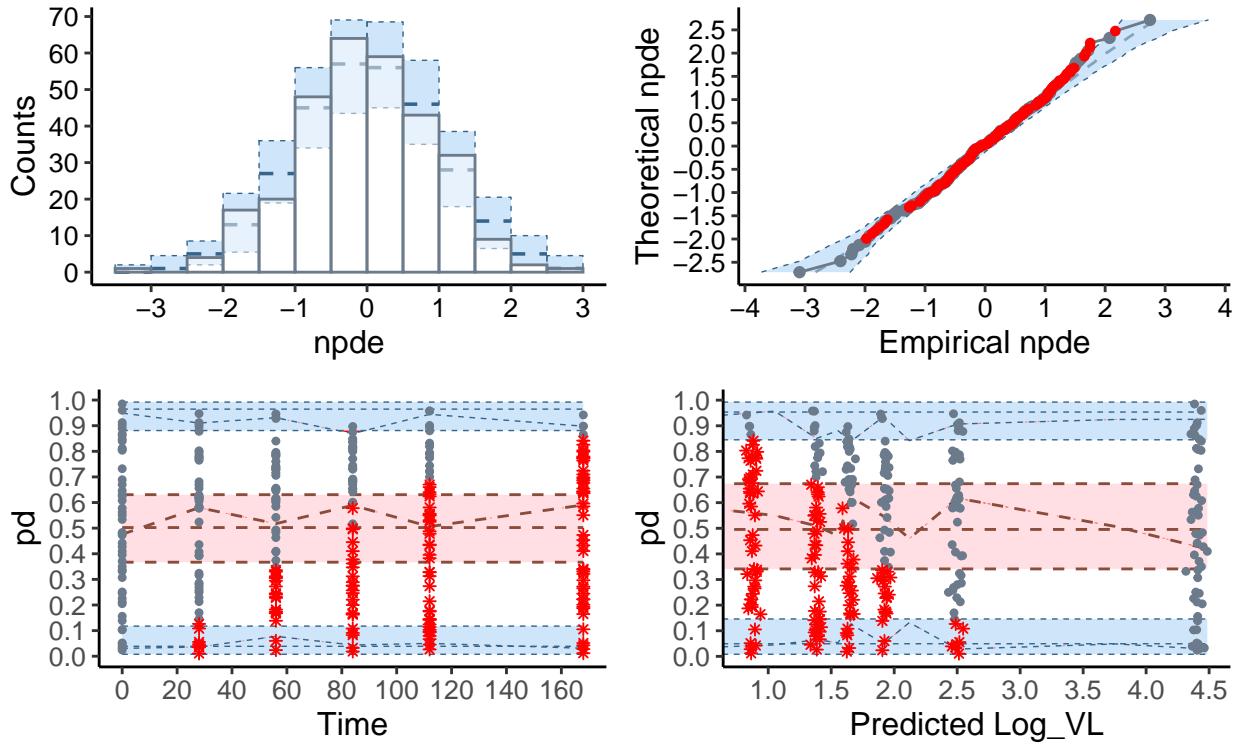


Censored 50, censoring=ipred



Censored 50, censoring=ppred

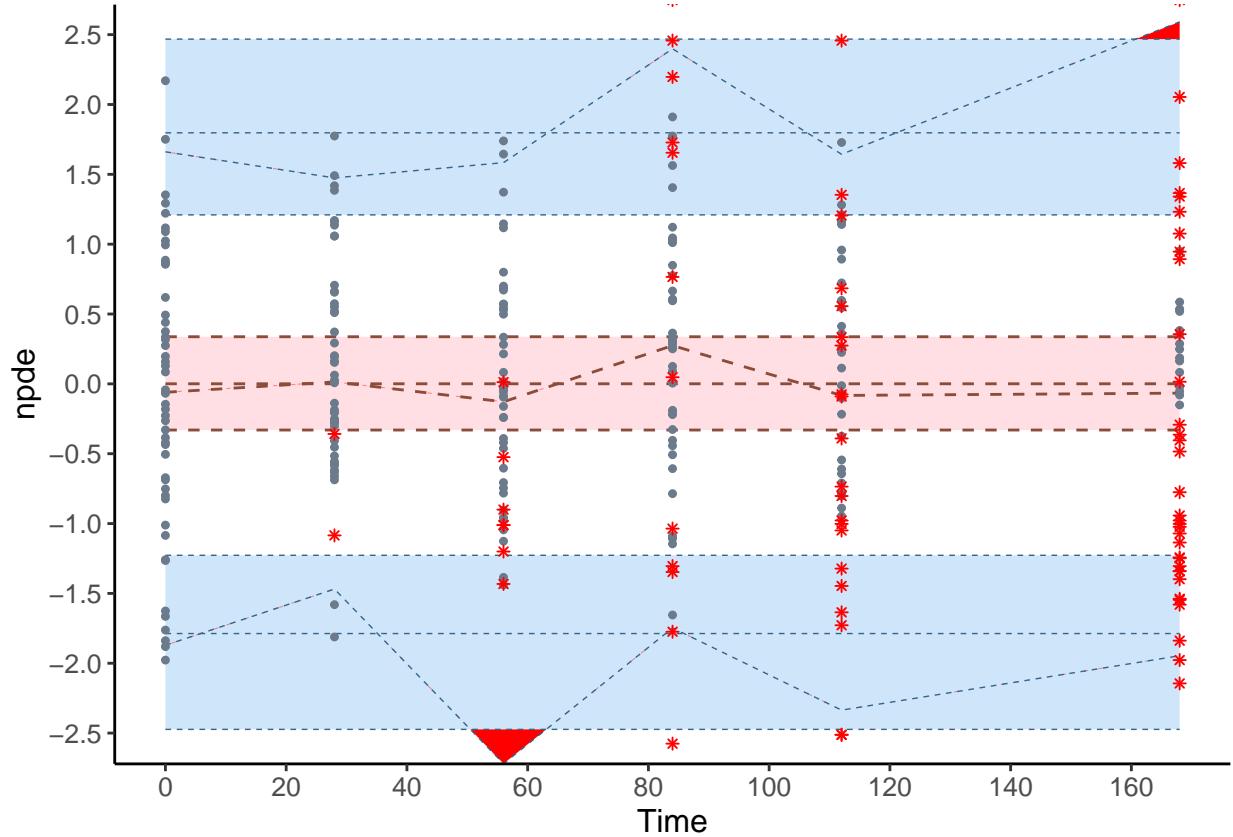


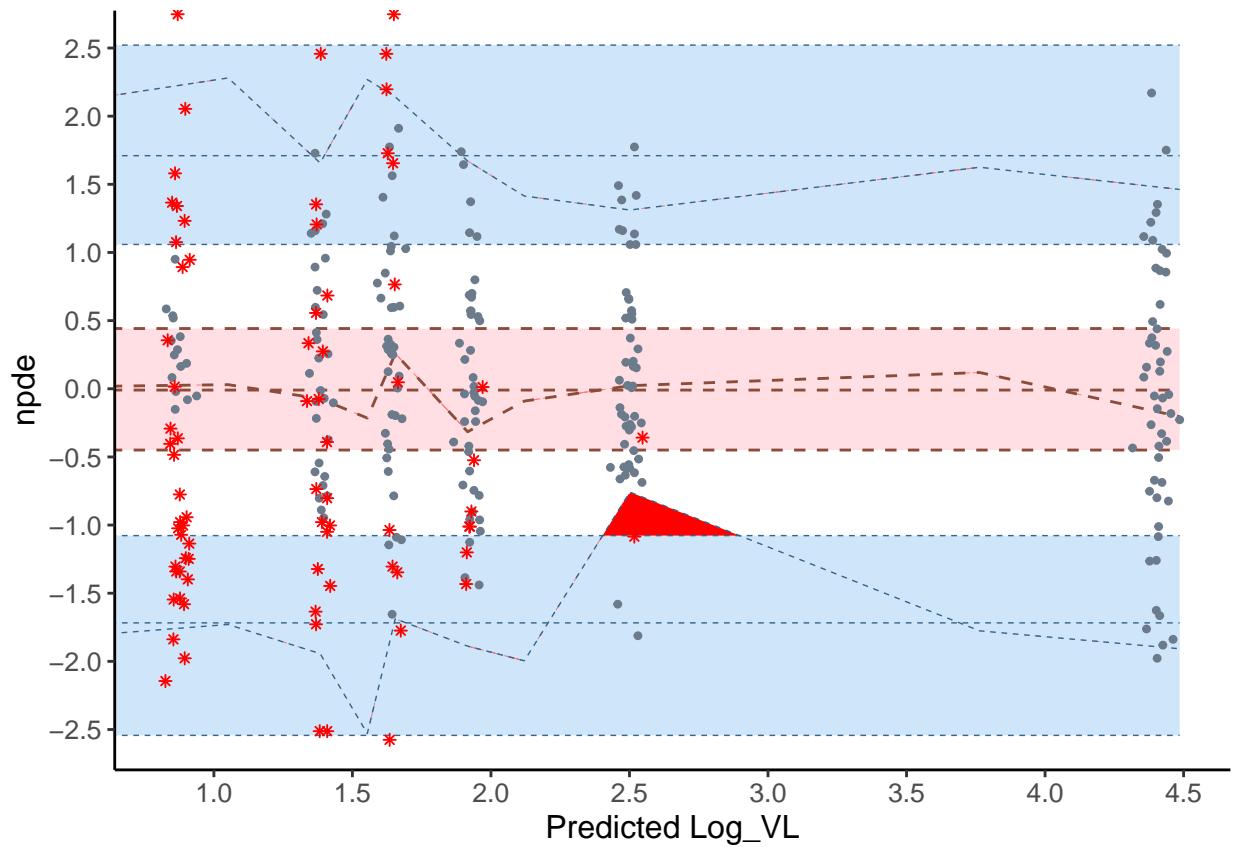


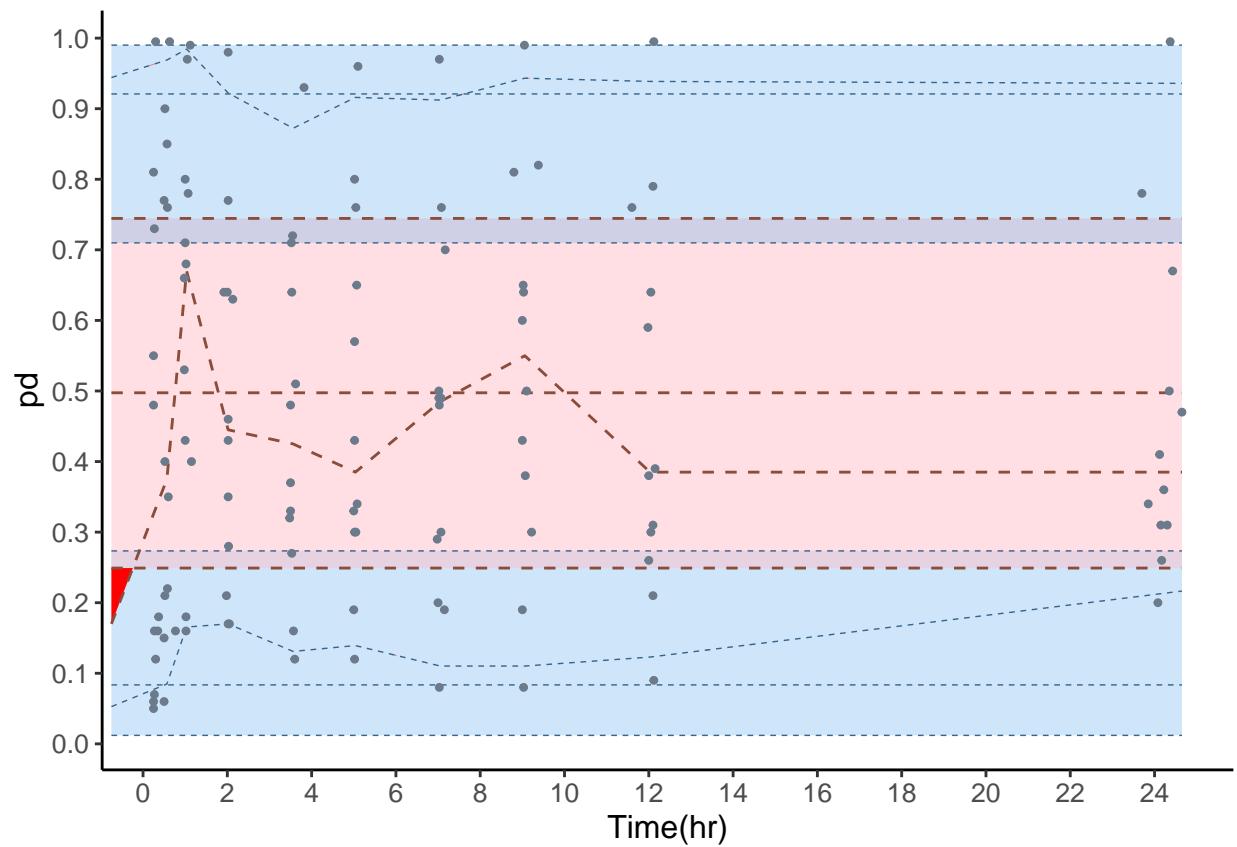
Plots with default options

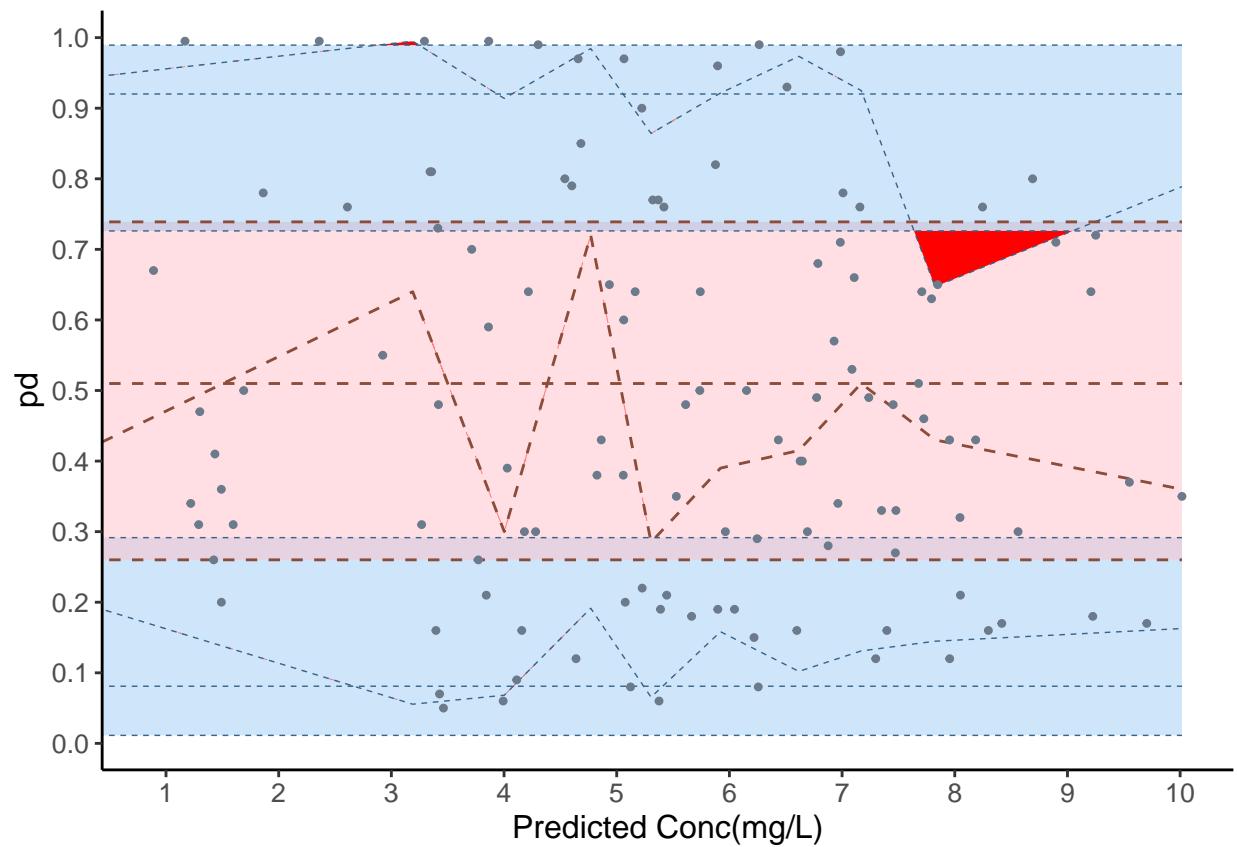
Scatterplots

- **Problems (à résoudre Romain)**
 - solved ylab for scatterplot doesn't appear (but works in waffle plot, wtf ?)
 - solved size ne passe pas (devrait changer à la fois size.pobs et size.pcens)
 - * solved même pb que dans le graphe par défaut (ie les 4 graphes)
 - solved calling plot with plot.type="x.scatter" and which="pd" (eg) triggers the message "Option which.y= x.scatter not recognised" => debug call to intermediate functions with arguments in **plotNpde-methods.R** (dispatching arguments given in ...)
 - solved plot.box has a weird aspect with seemingly empty boxes for the first 2 bins ?
- **Check**
 - solved size of boxes when plot.box=TRUE

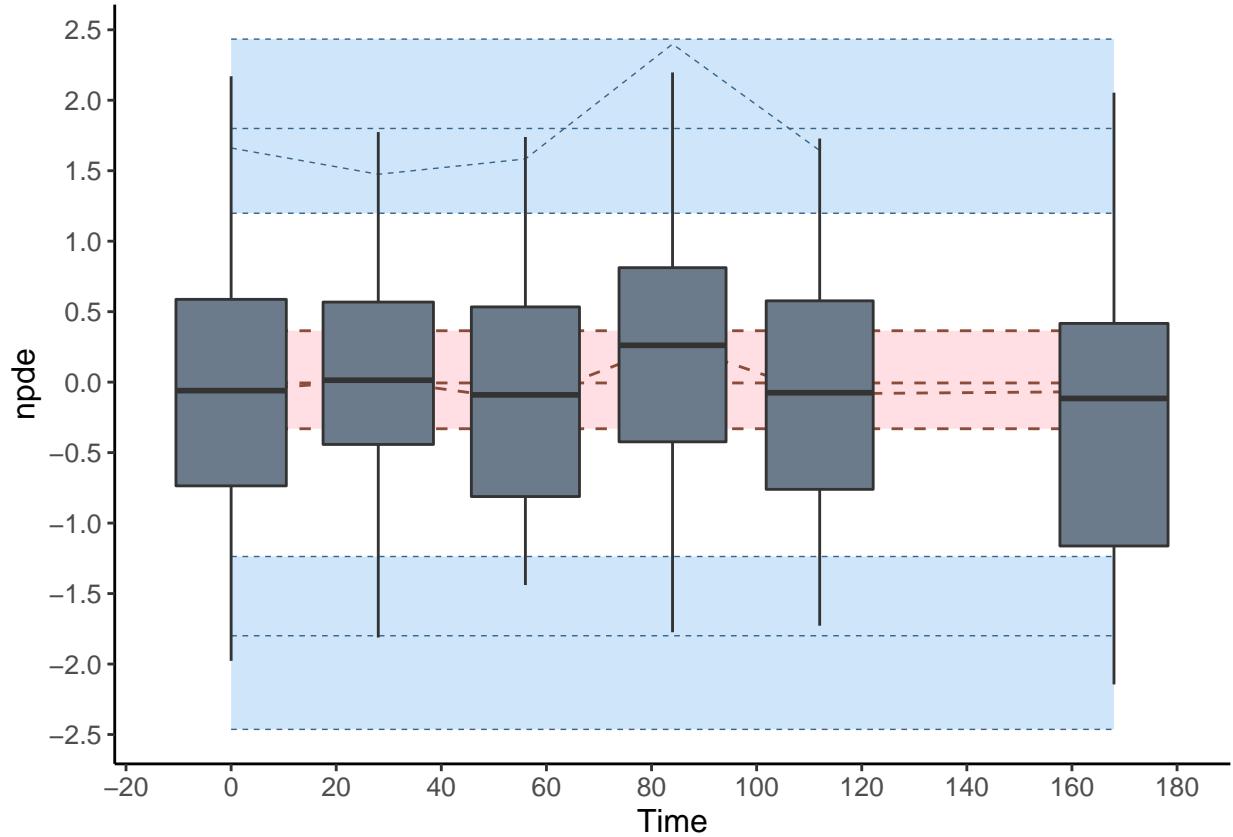




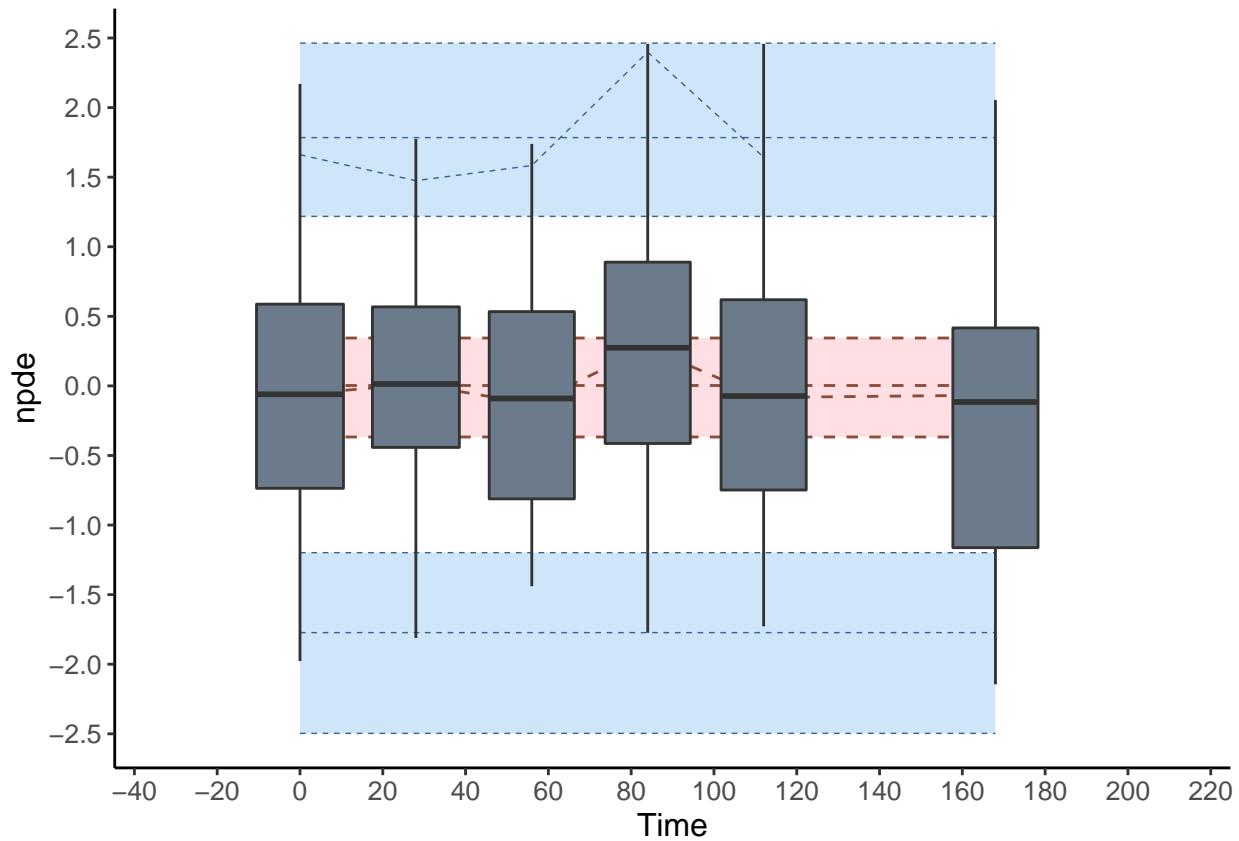


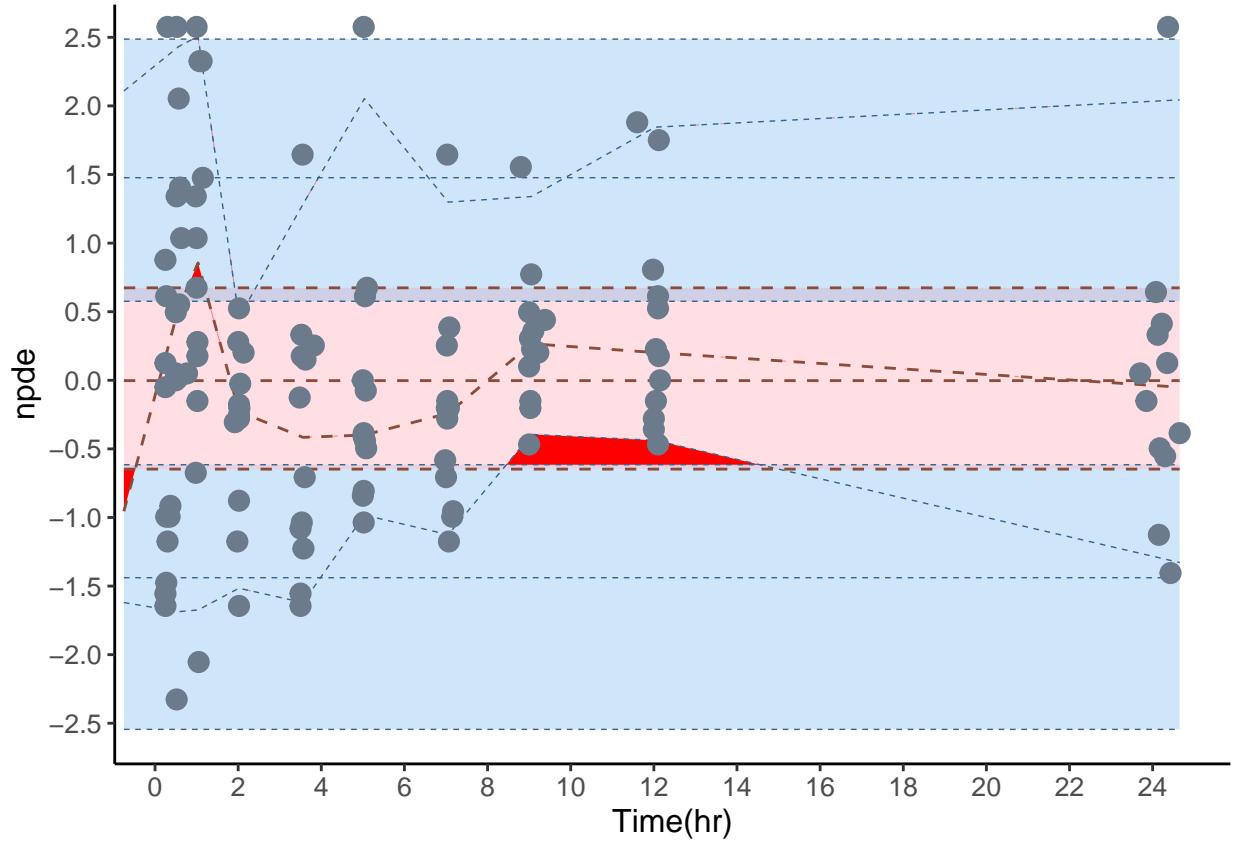


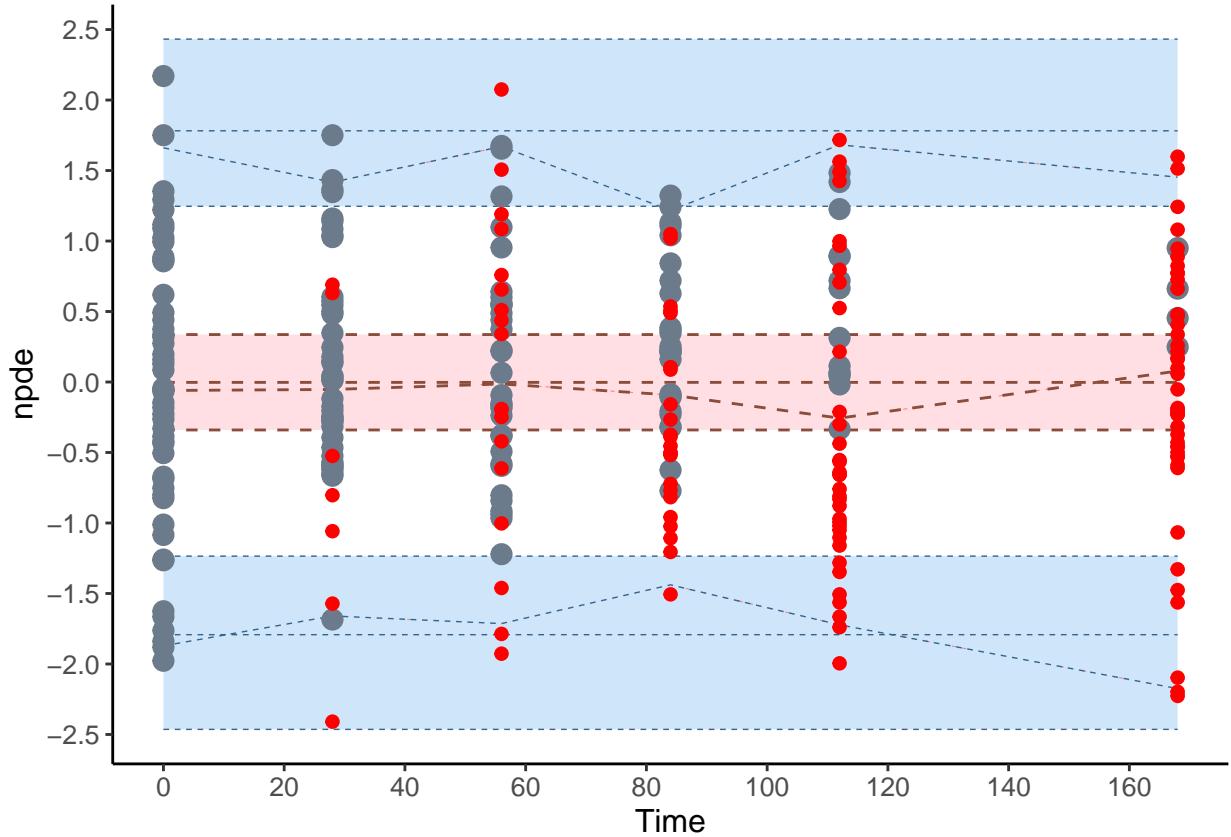
```
## Warning: Removed 10 rows containing non-finite values (stat_boxplot).
## Warning: Removed 1 row(s) containing missing values (geom_path).
```



```
## Warning: Removed 8 rows containing non-finite values (stat_boxplot).  
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

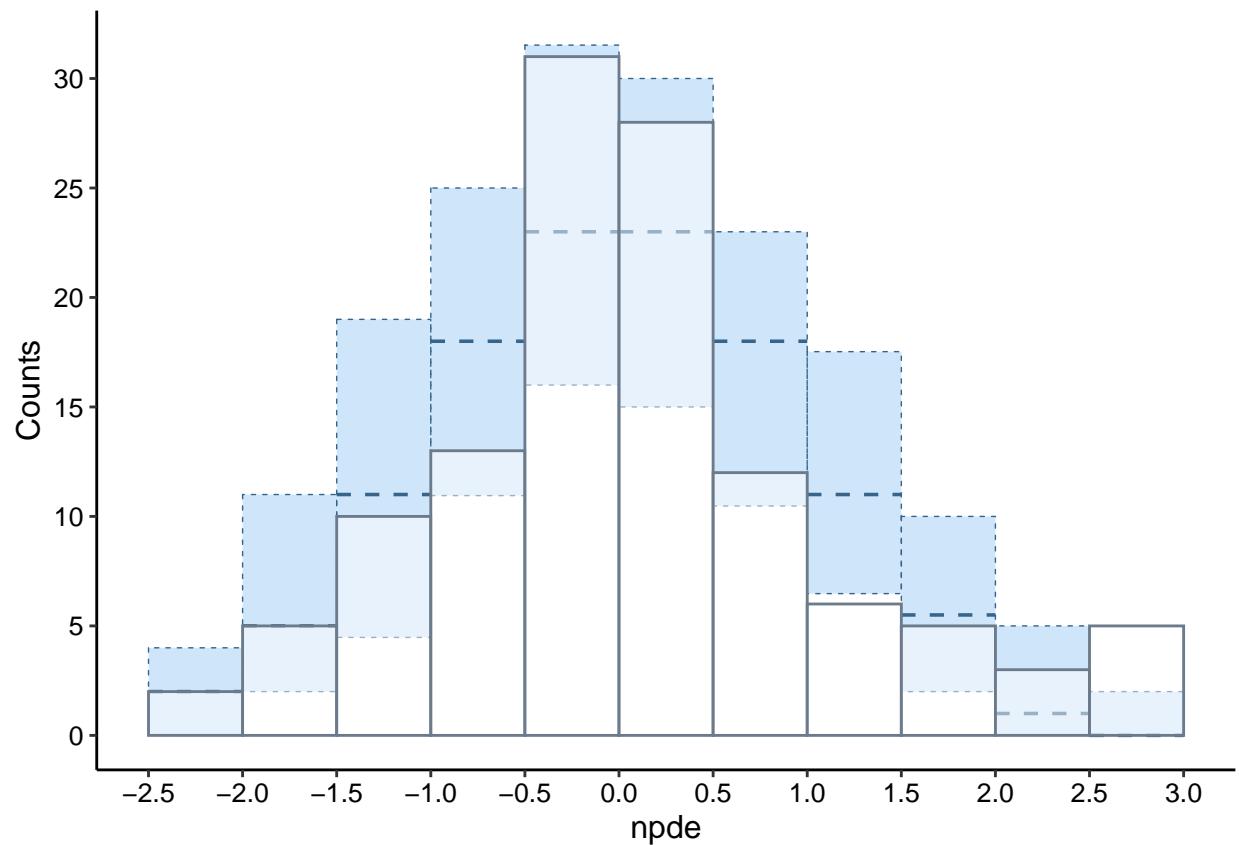


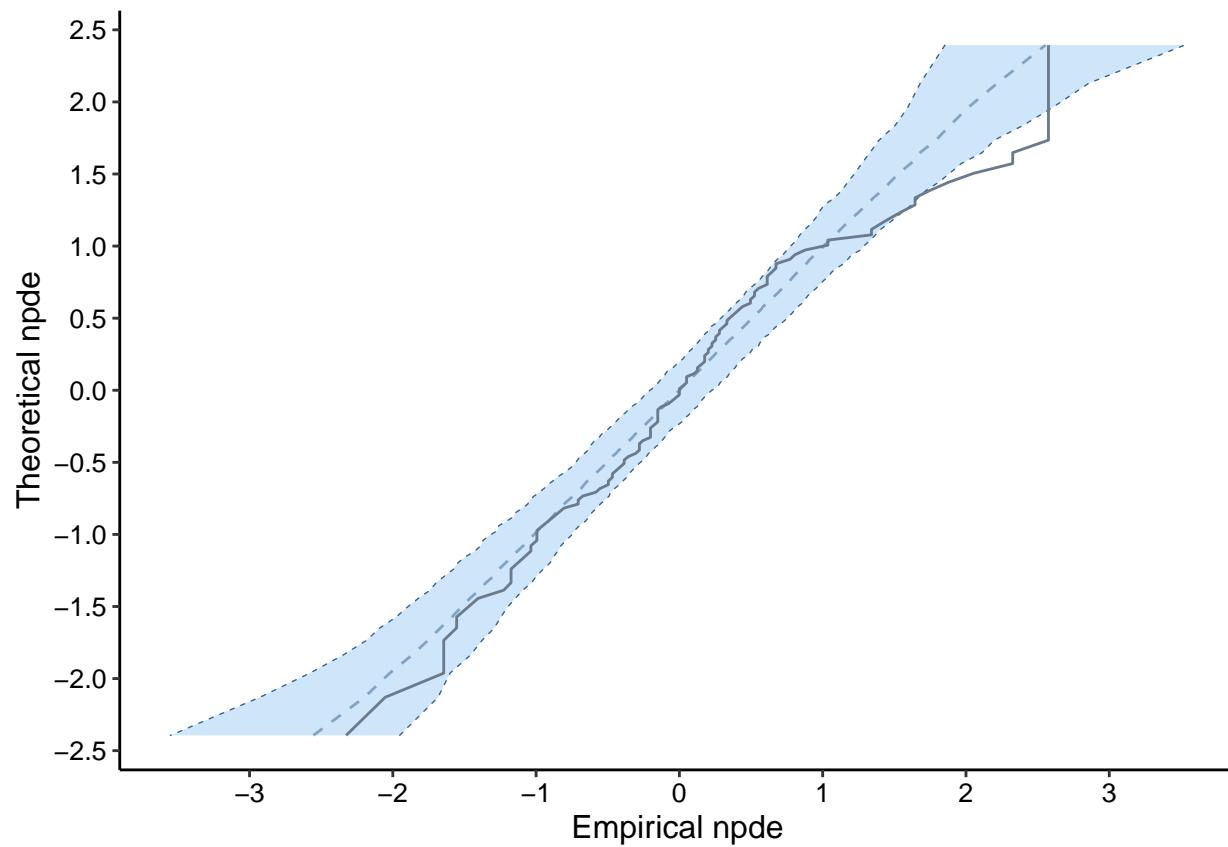


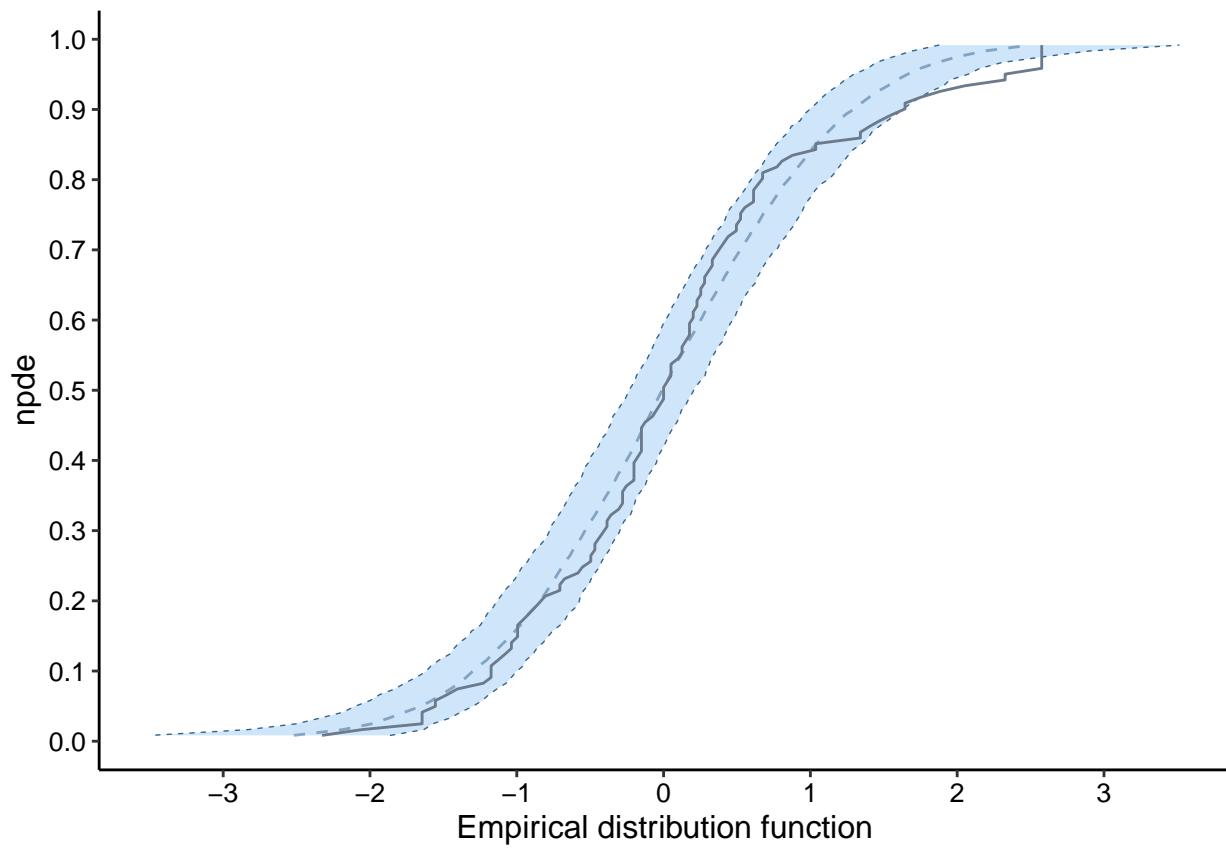


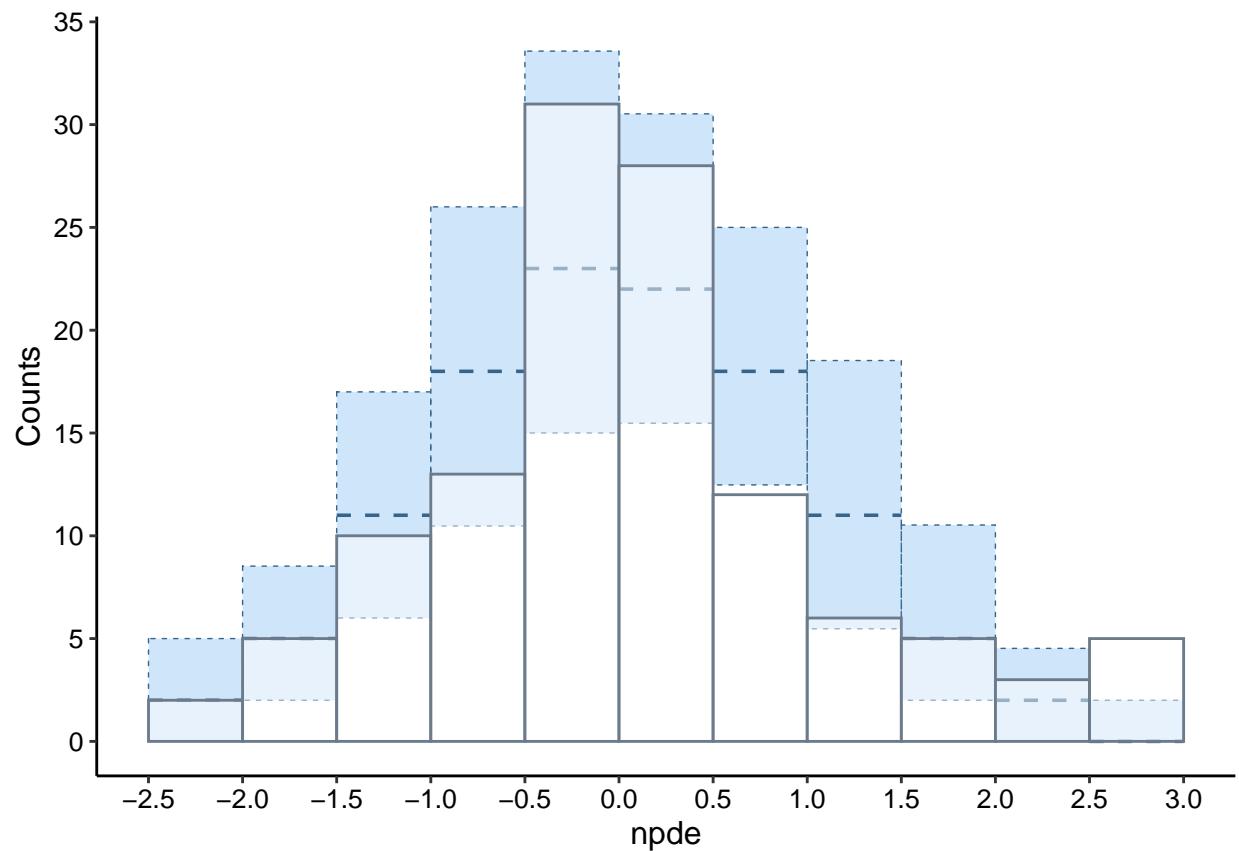
Distribution plots

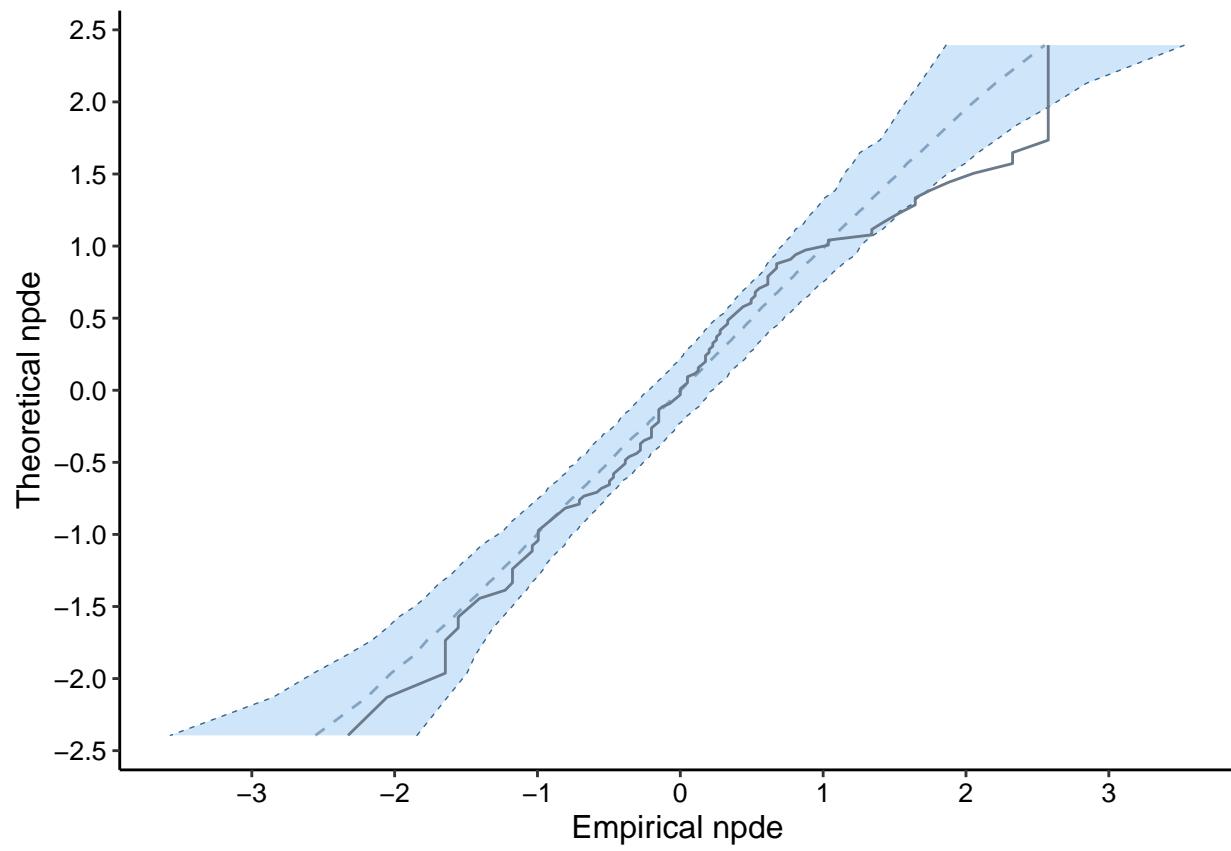
- **Problems (à résoudre Romain)**
 - solved pb with `size` (comme plus haut, size ne passe pas, size.pobs/size.pcens passent)
- **Problems (à résoudre Eco)**
 - **axe title** for ecdf plot (should be: ‘npde’ on X-axis (/pd/npd) and ‘Empirical distribution function’ on Y-axis) [done, check it works]

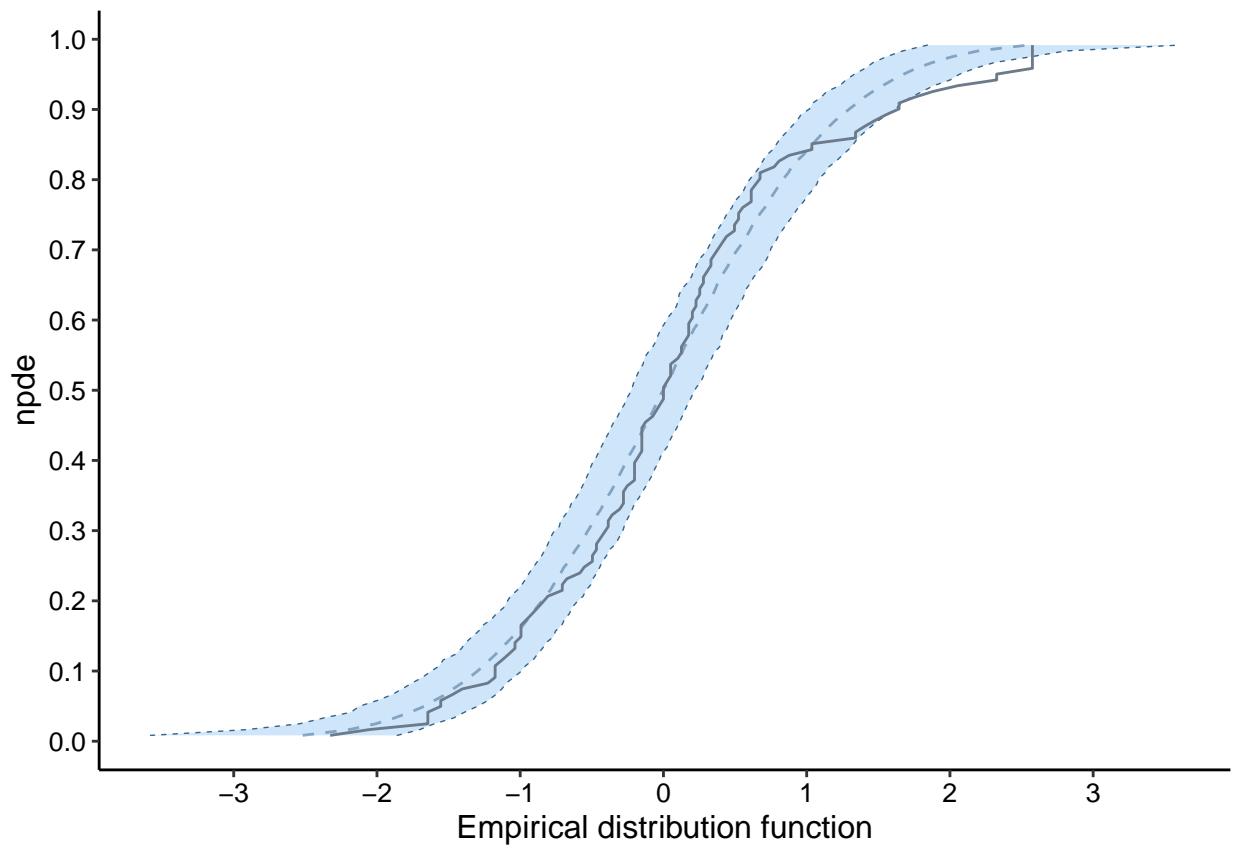


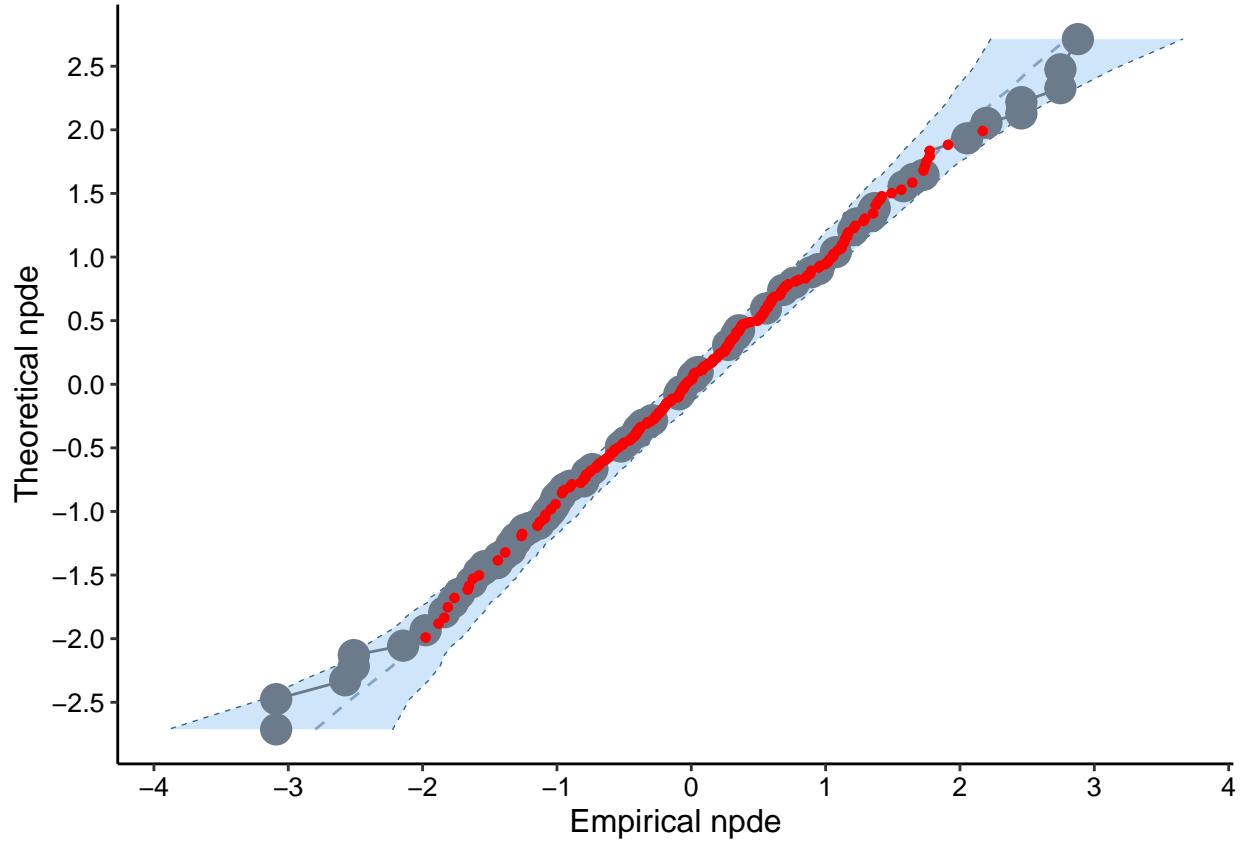


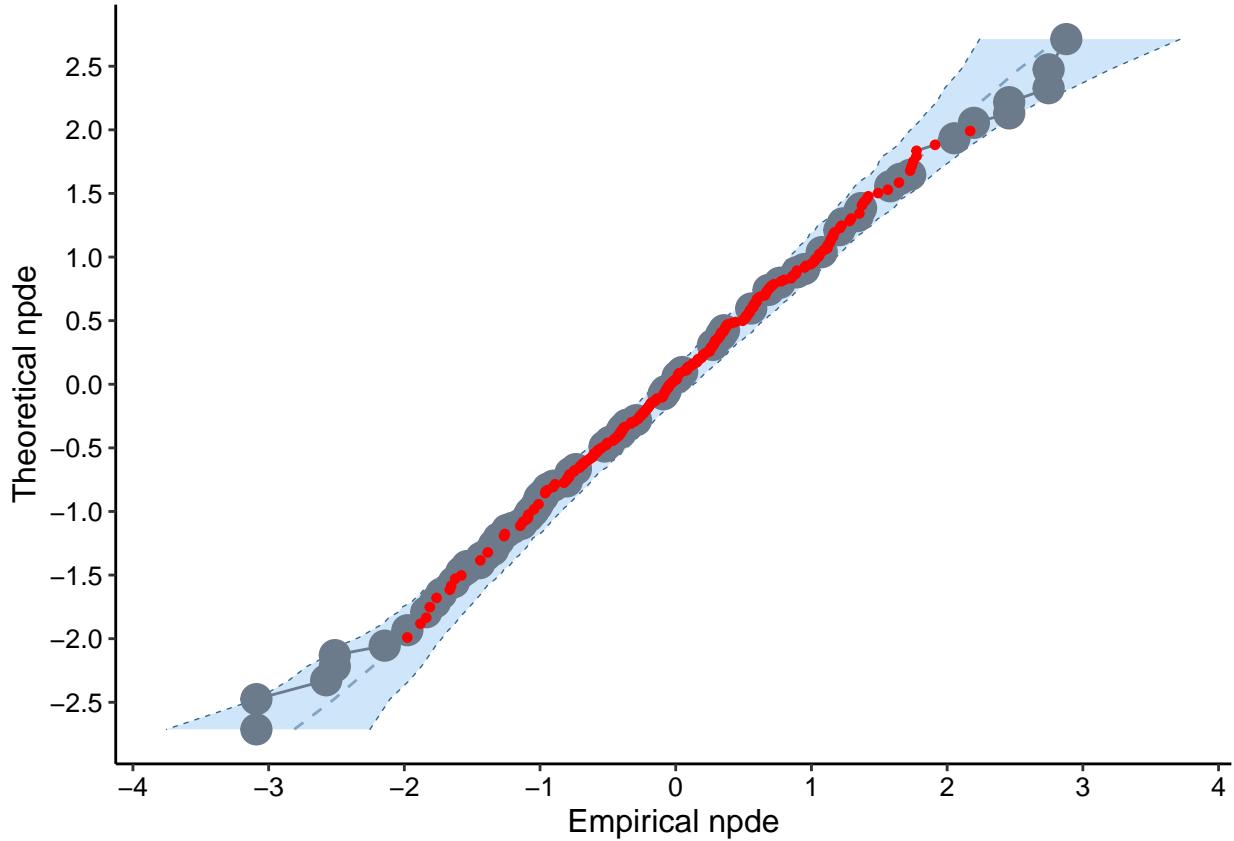








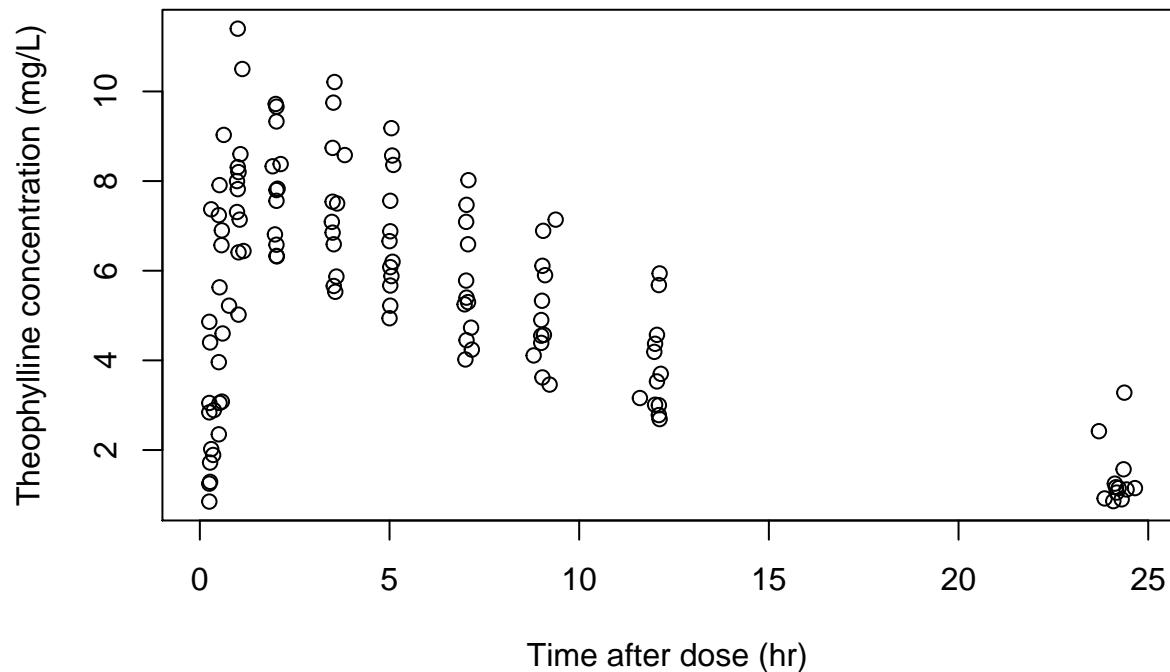




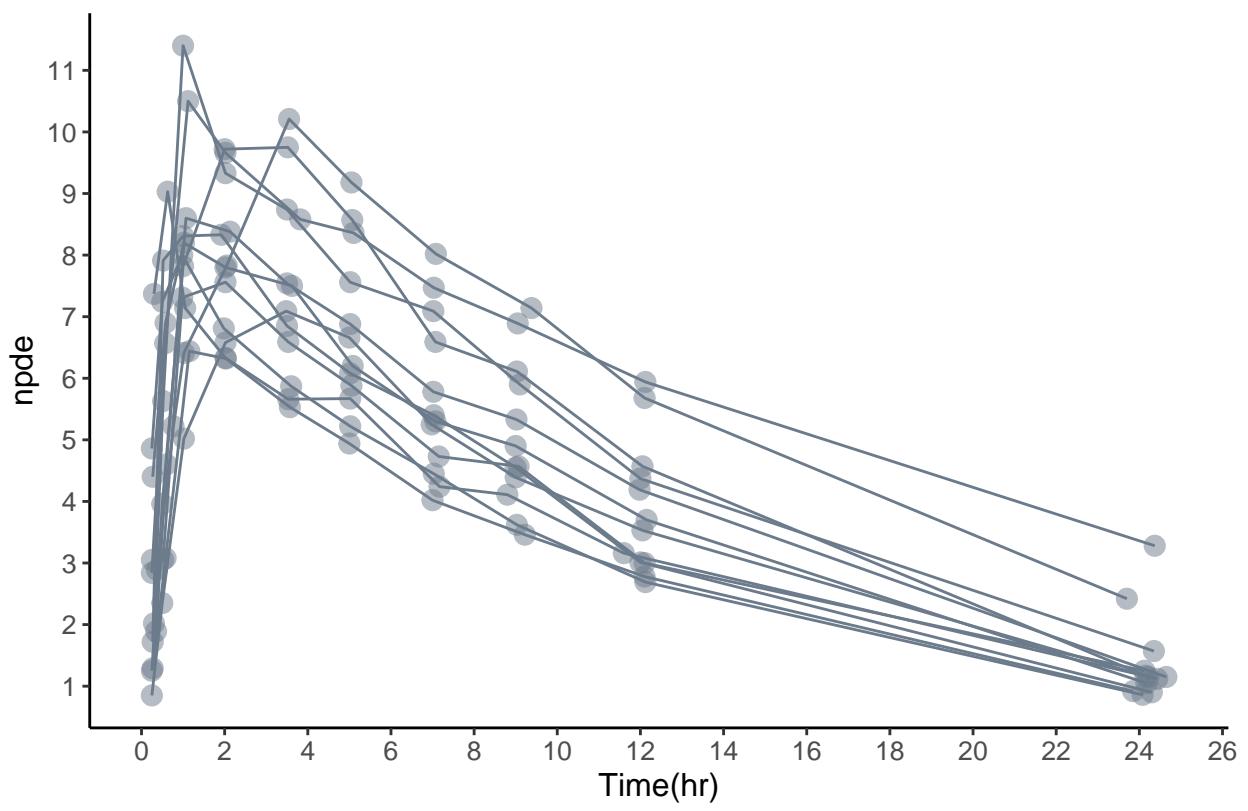
Data plots

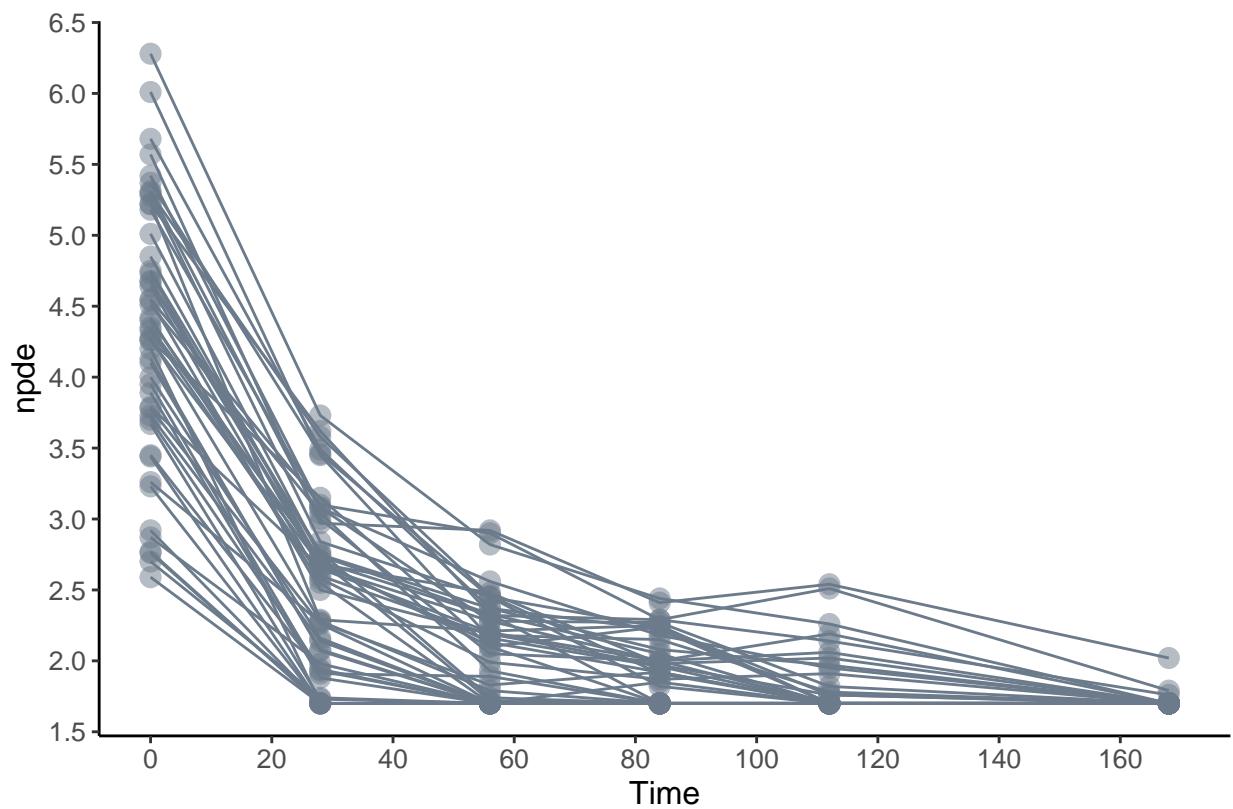
- **Problems (à résoudre Romain)**
- **Note** pour cette version ça fait beaucoup à modifier, à commencer par `plotNpde-methods` puis `npde.plot.data` où le slot `data` est à modifier. Il faudrait distinguer le plot par défaut et `plot(yivr50 @data)` sans arguments supplémentaires.
 - plot d'un **objet npdeData** ne marche pas
 - * par contre `plot.type="data"` de l'objet `npdeObject` fonctionne => bizarre, vérifier [normalement `plot.type="data"` devrait appeler `plot(objet@data)` mais là ça semble court-circuiter ceci => à rectifier, on veut plutôt l'inverse]
 - **pour cette version trop de chose à modifier pour ça**
 - problem with the data in the presence of **LOQ data**, the data being plotted is the LOQ
 - * if applied directly to an objet `npdeData` (eg `plot(yivr50 @data)`) => should plot `y` or the LOQ (ie censoring value)
 - * if applied to the **npdeObject** resulting from a run => data plotted should depend on the censoring method (same as VPC)
 - for `cdf`, plot imputed `y`
 - for `omit`, don't plot anything
 - for `ipred`, plot `ipred`
 - for `ppred`, plot `ppred`
 - * all other options should be the same [axis titles, axes, grid, default colours, etc...]
 - **solved lines for axes** don't appear
 - **solved size**
 - * size should be controlled by `size.pobs` and `size.pcens`, currently too small (not the same defaults as for the other plots ?)

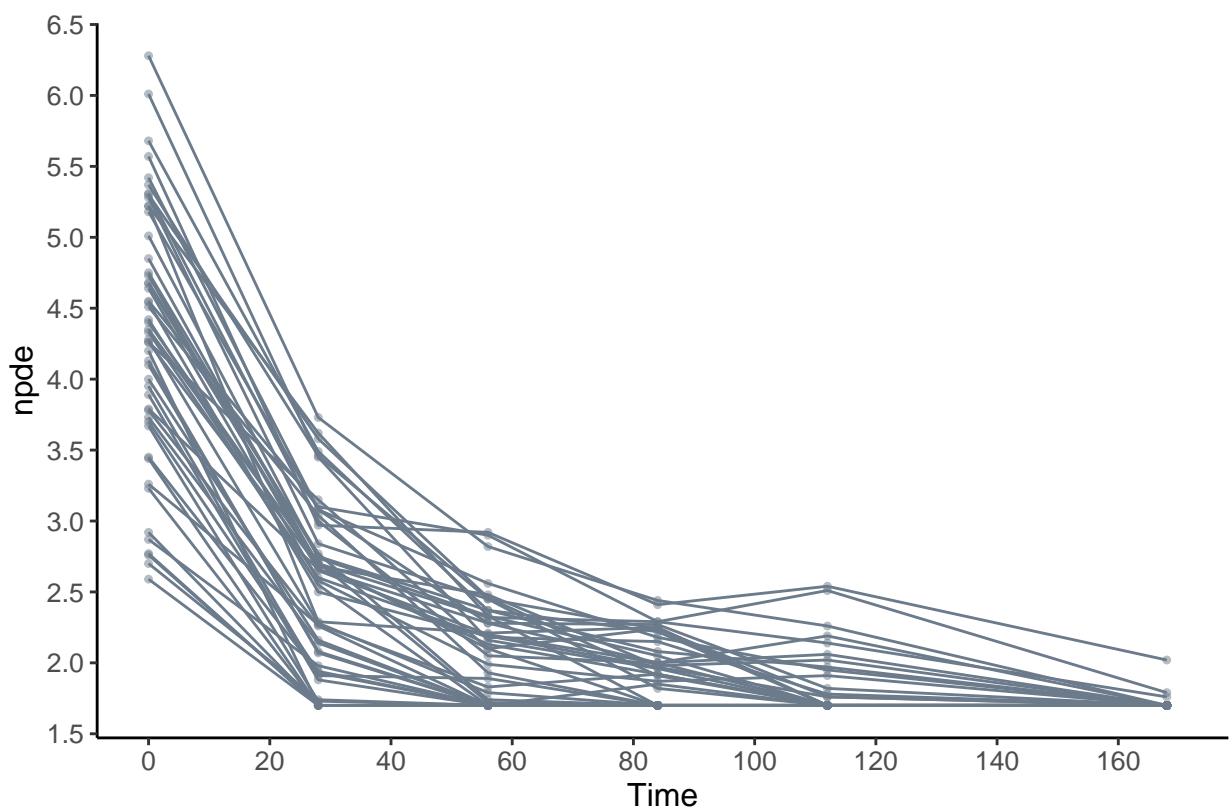
- * size doesn't change size.pobs and size.lobs as it should
- * size.lobs doesn't work
- **solved axis titles**
 - * missing for theofit, why ? (present for yvir50, so odd)
- **solved line.loq** doesn't work (no line appears at the LOQ)

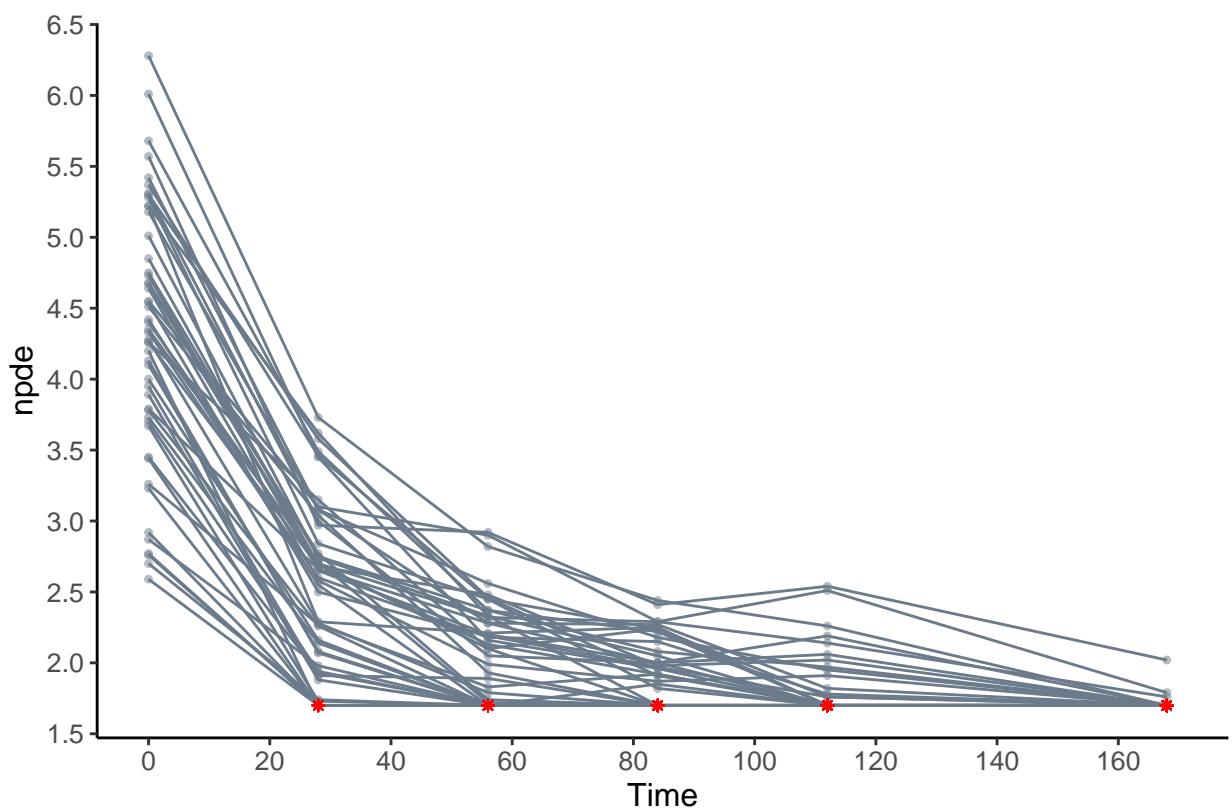


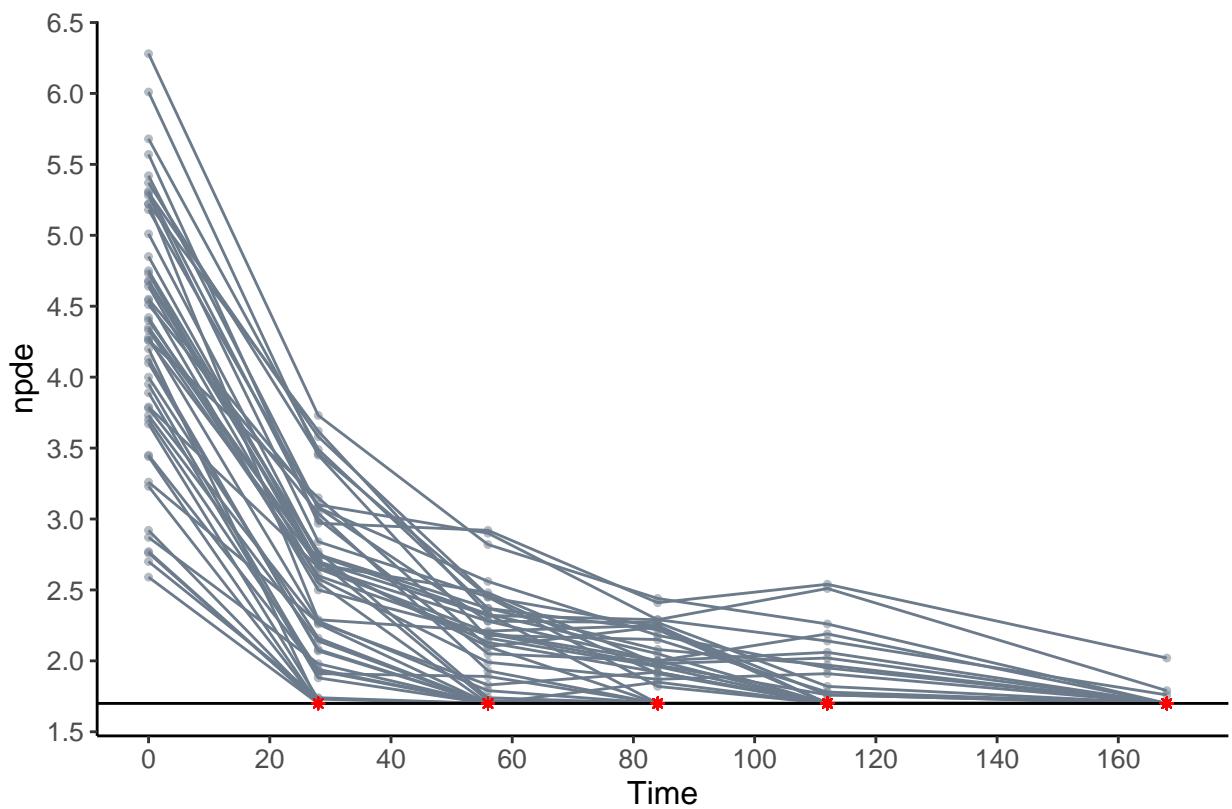
```
## Failed test, no output
## Warning: Removed 12 rows containing missing values (geom_point).
## Warning: Removed 12 row(s) containing missing values (geom_path).
```

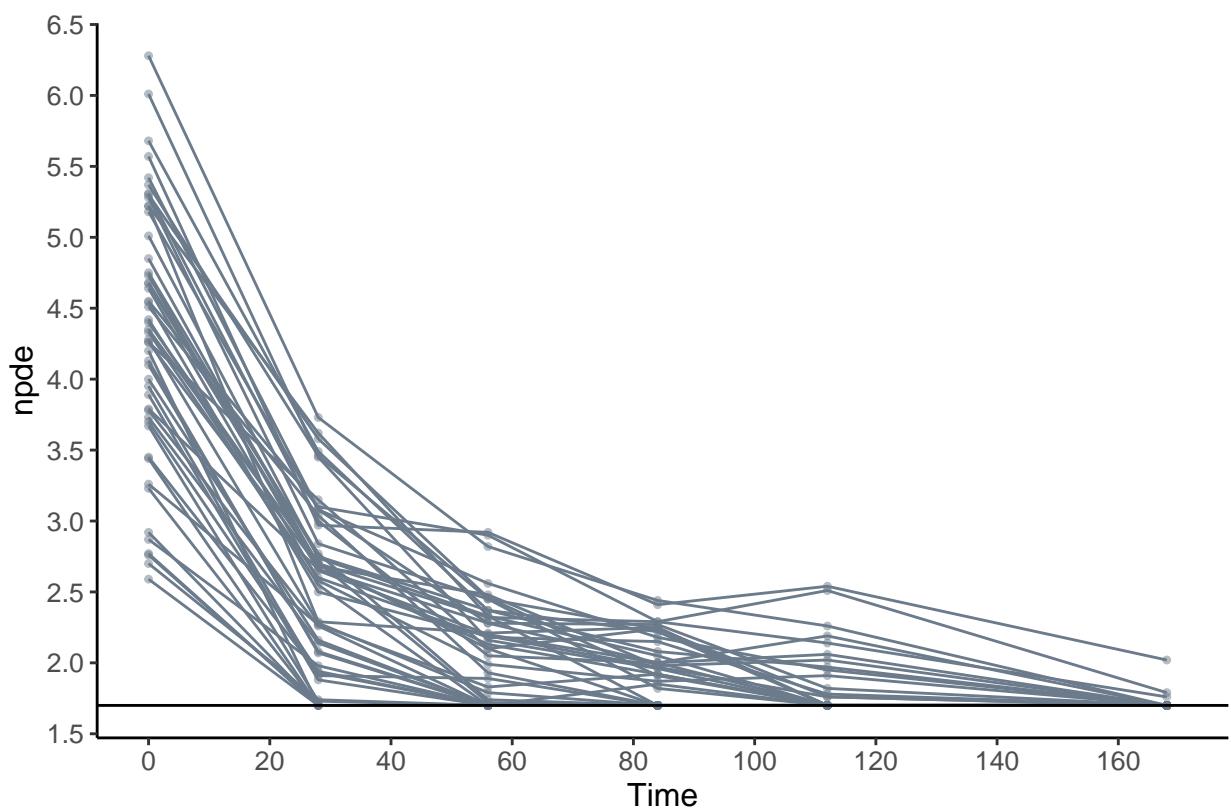


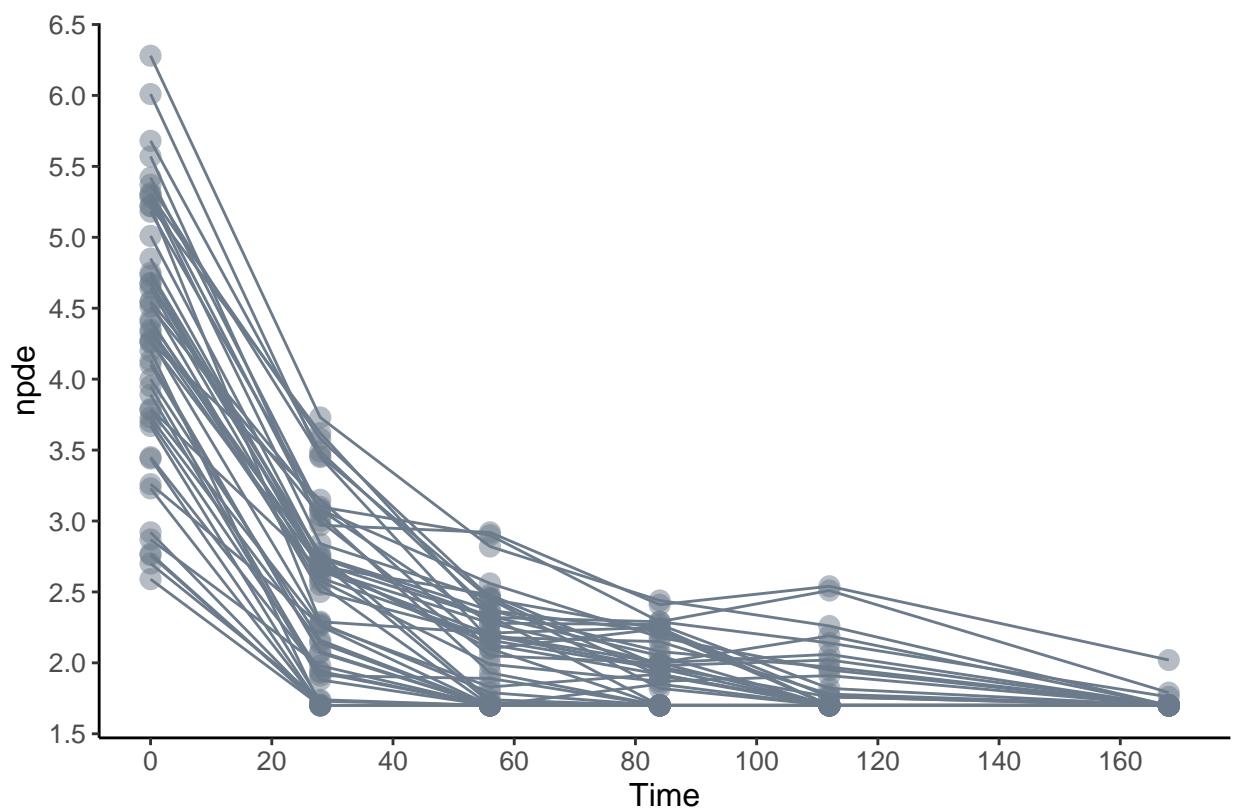


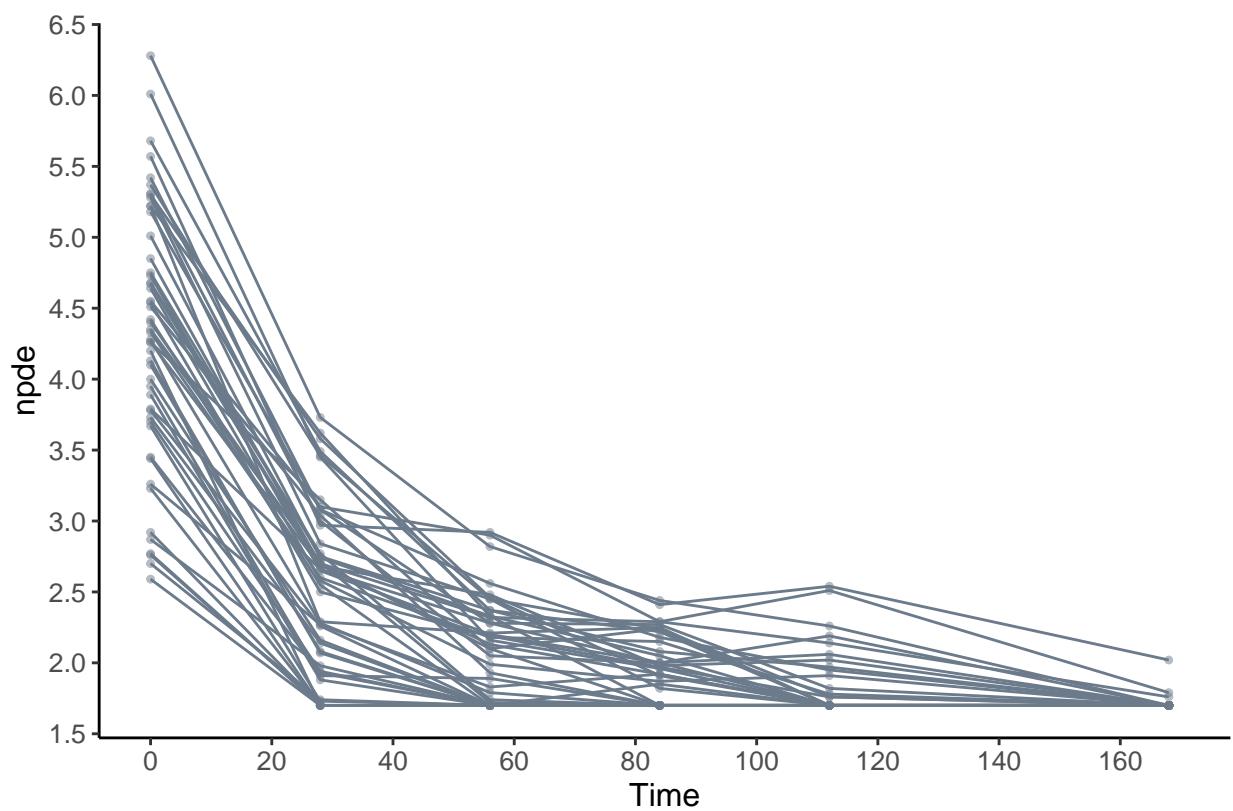


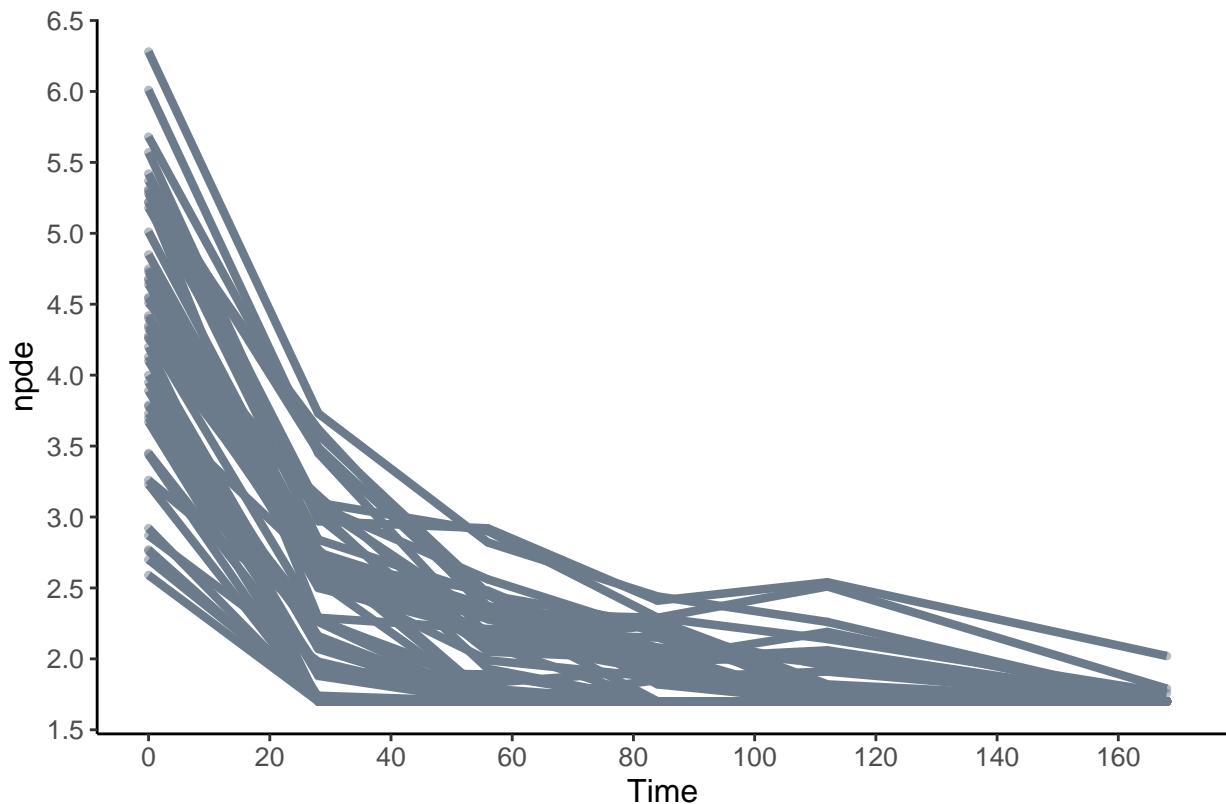






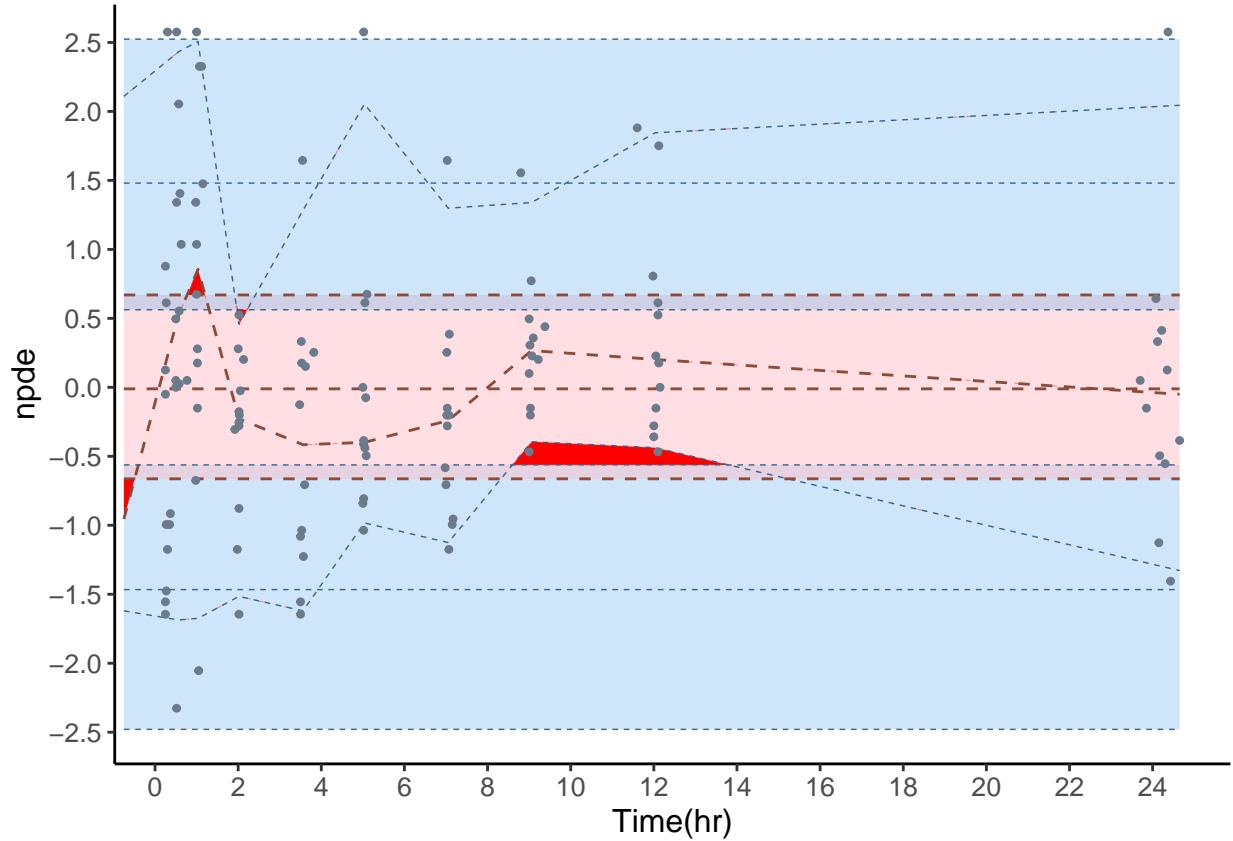




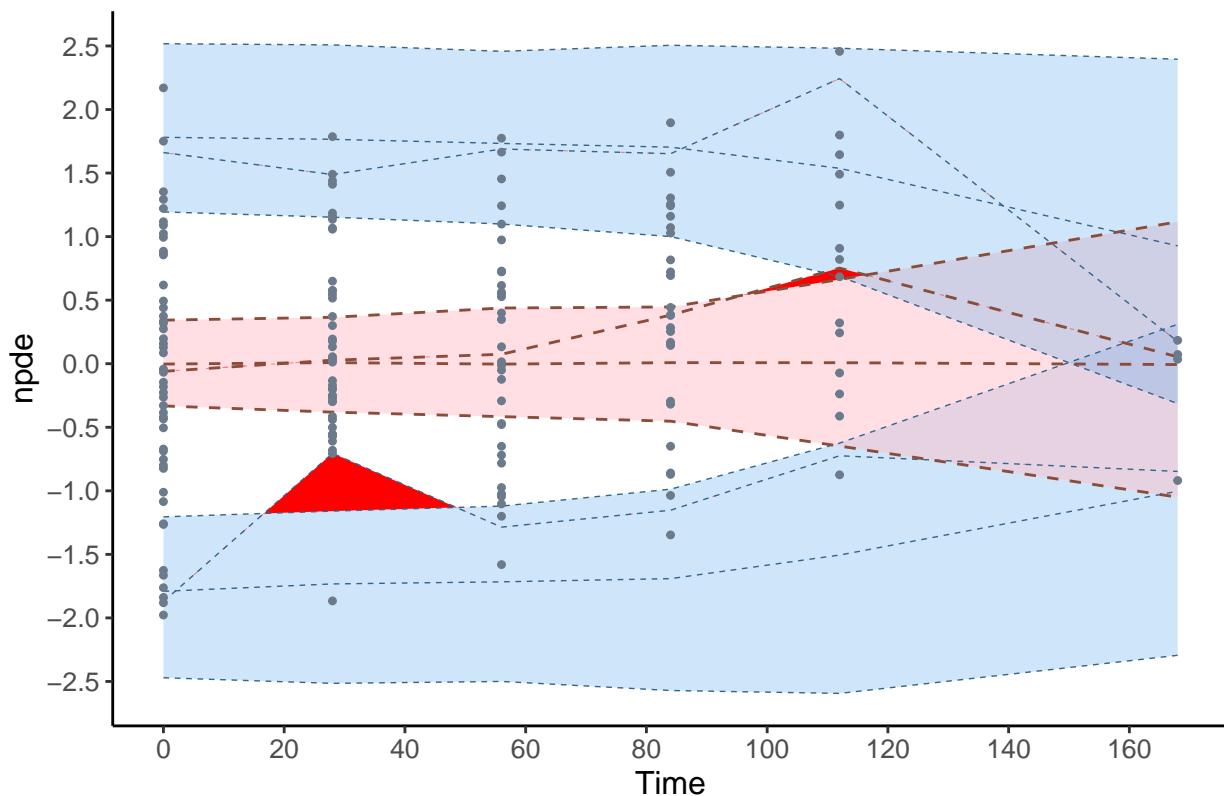


VPC

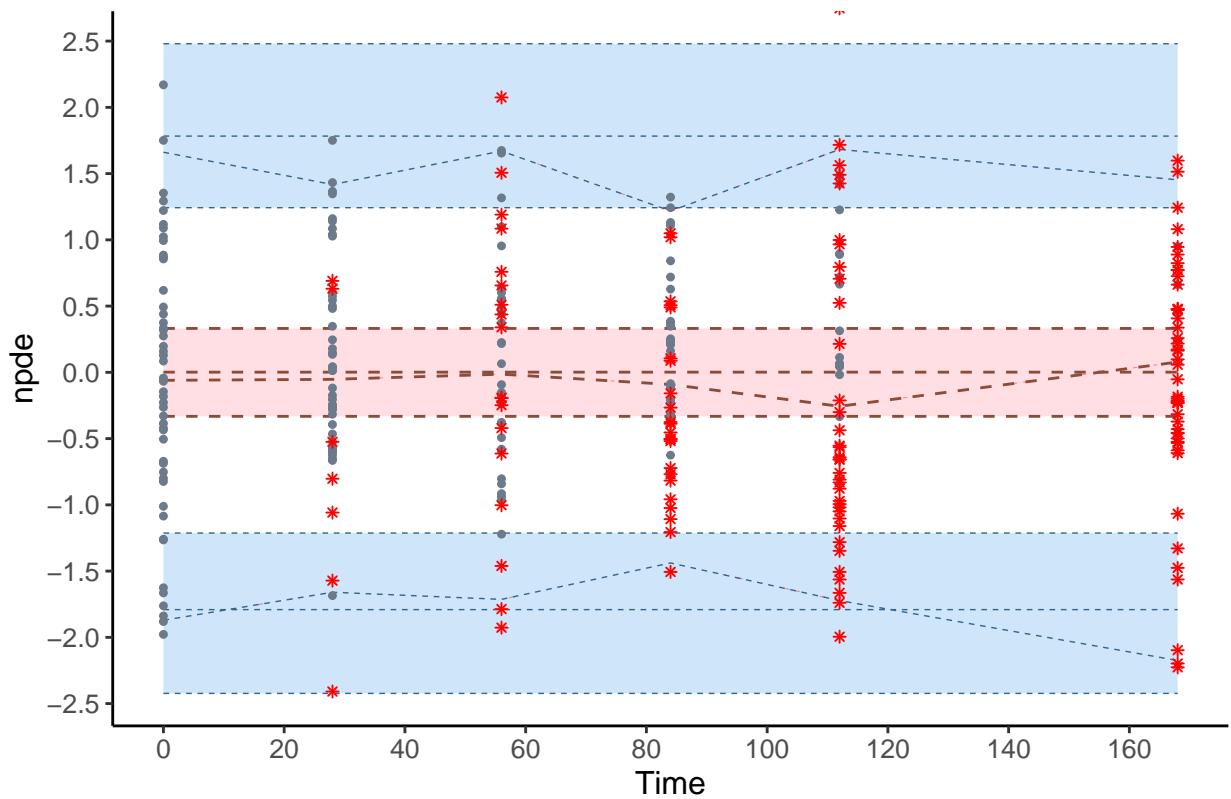
- **Problems (à résoudre Romain)**
 - Y-axis missing
 - problem with the PI not being interpolated properly (??? thought that worked :-/)
- **Notes to check Eco**
 - computation of the PI for **omit** consistent with the PI computed for **npde** ?
 - * compute the PI in this case by omitting the data>LOQ for the simulated data as well, to be consistent with the way the PI are computed for **npde**
 - * check also that the PI are computed consistently for **ppred**
 - * **ipred** ??? in this case can't be consistent (we have **isim** in the dataset but not **ipred**) so keep the simulations as is
 - **but** VPC with LOQ data done as here => **Eco TODO:** expliquer dans la doc les méthodes de censoring et la différence entre les graphes **npde** et **VPC** dans le cas de la méthode **omit** (message=the discrepancy seen in **VPC** comes from not treating observed and simulated data in the same way, “corrected” in **npde** but at the price of a loss of power)



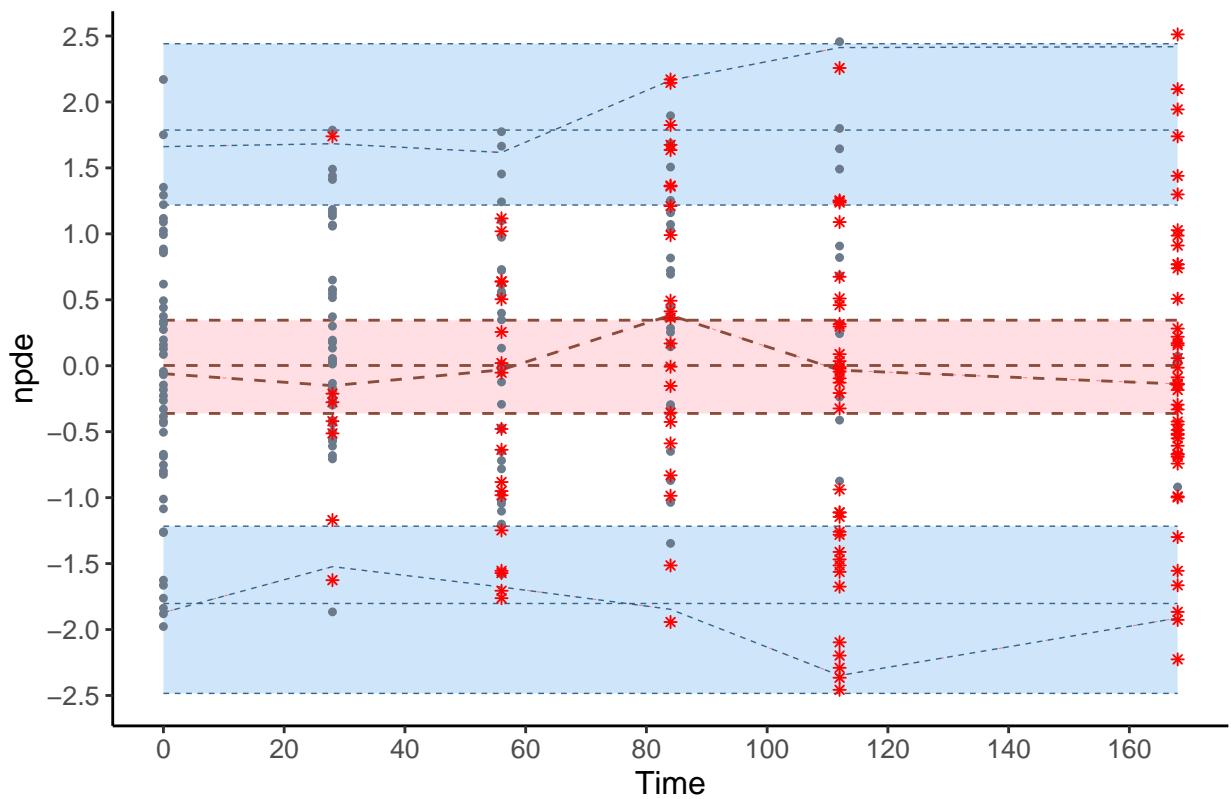
Viral load, method=omit



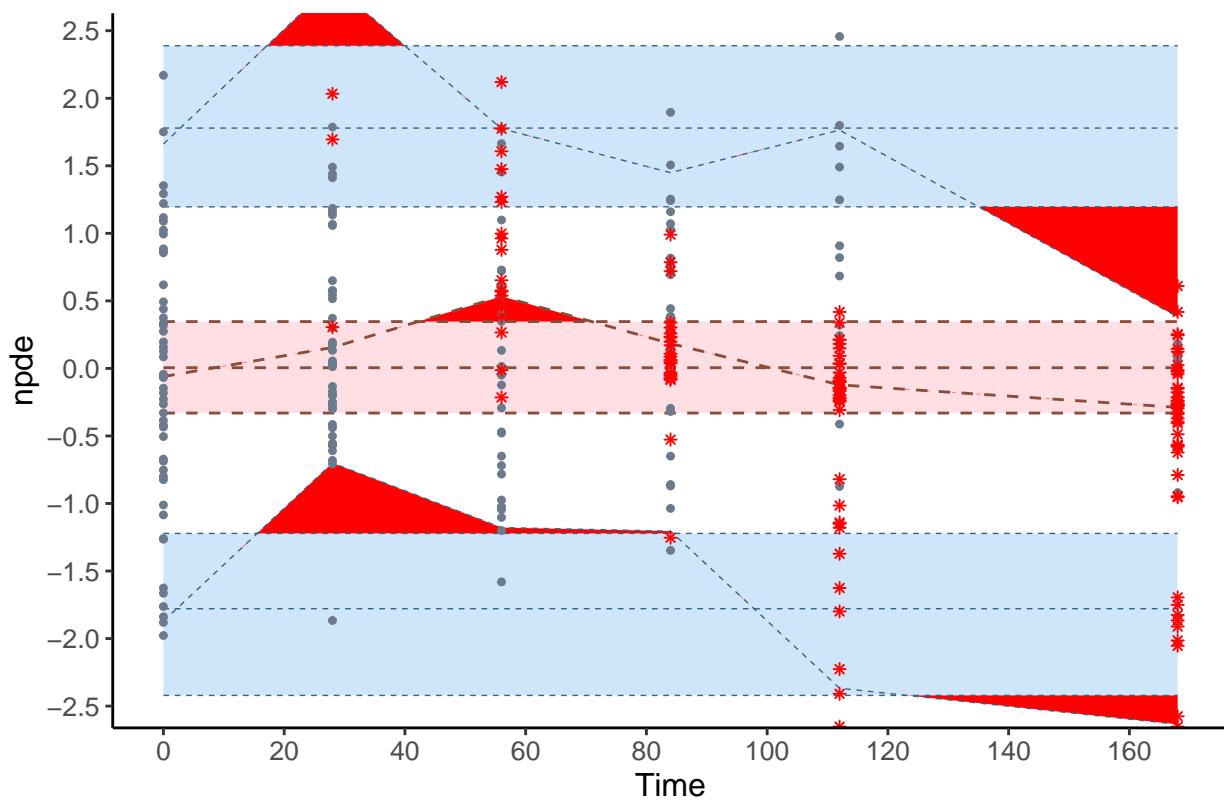
Viral load, method=cdf



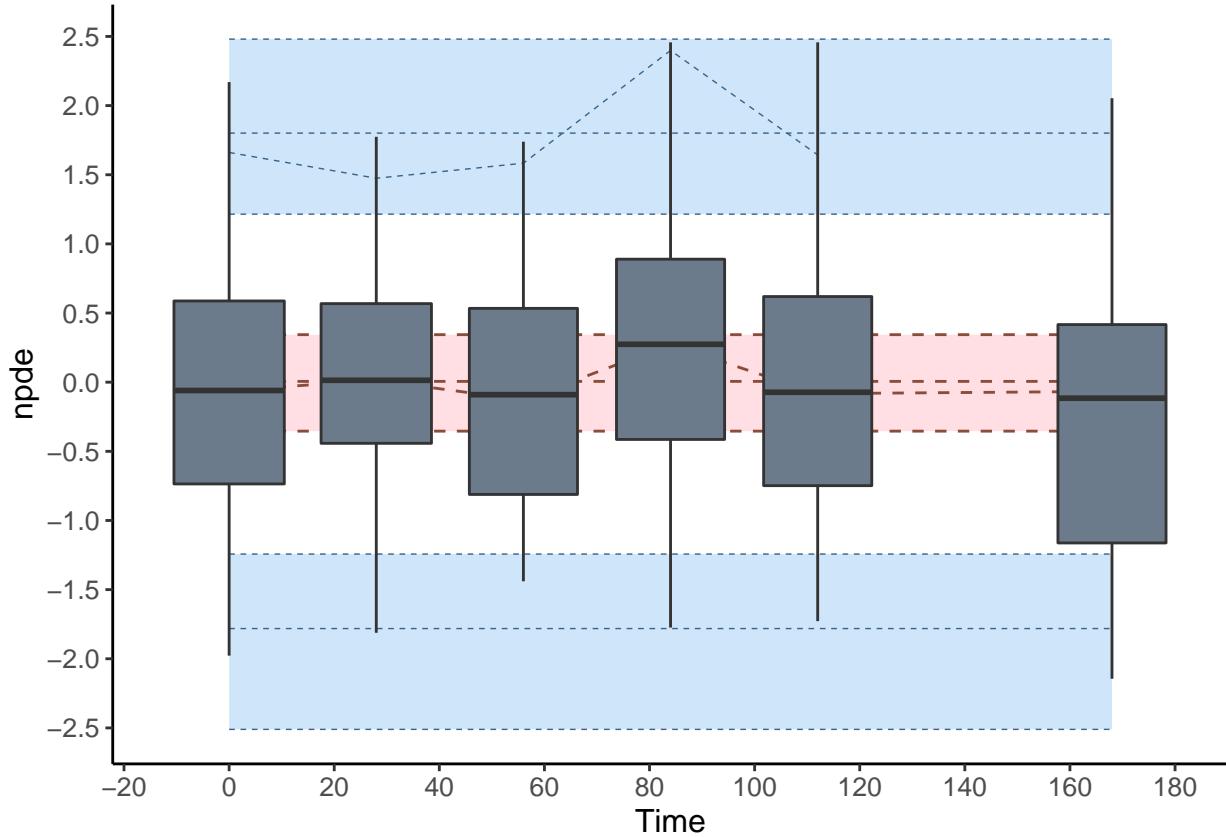
Viral load, method=ipred



Viral load, method=ppred

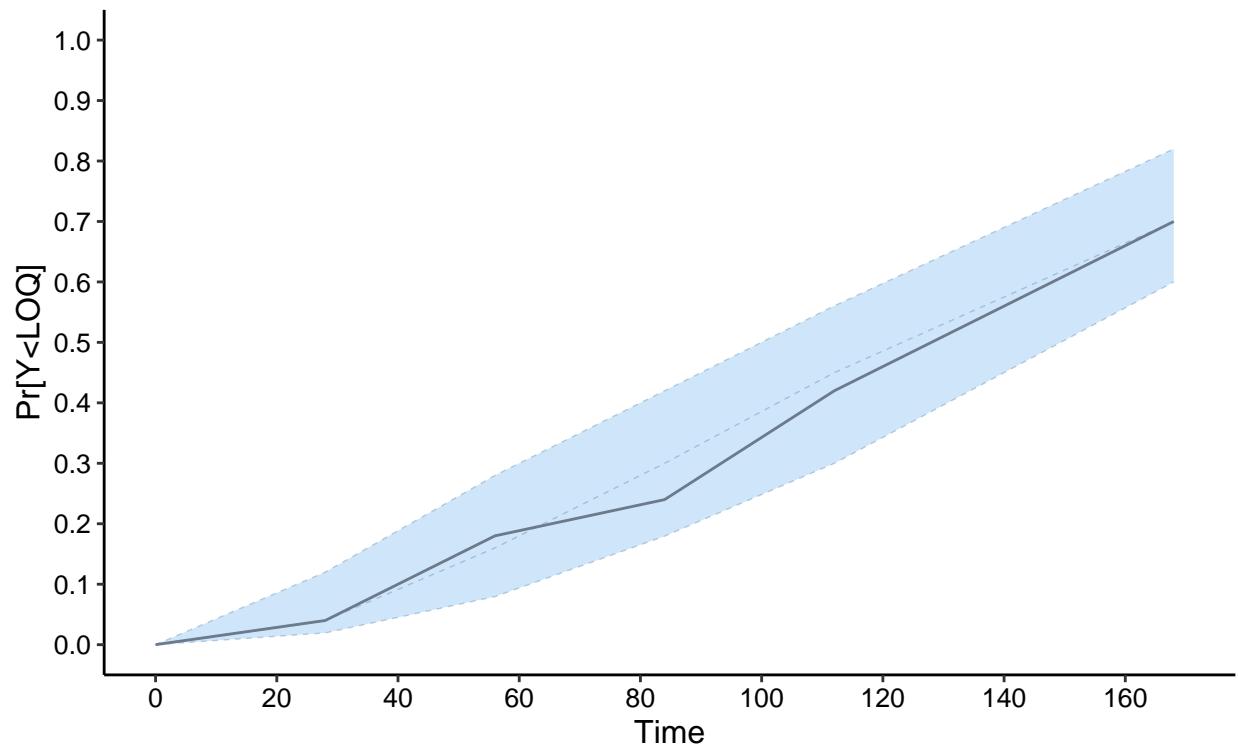


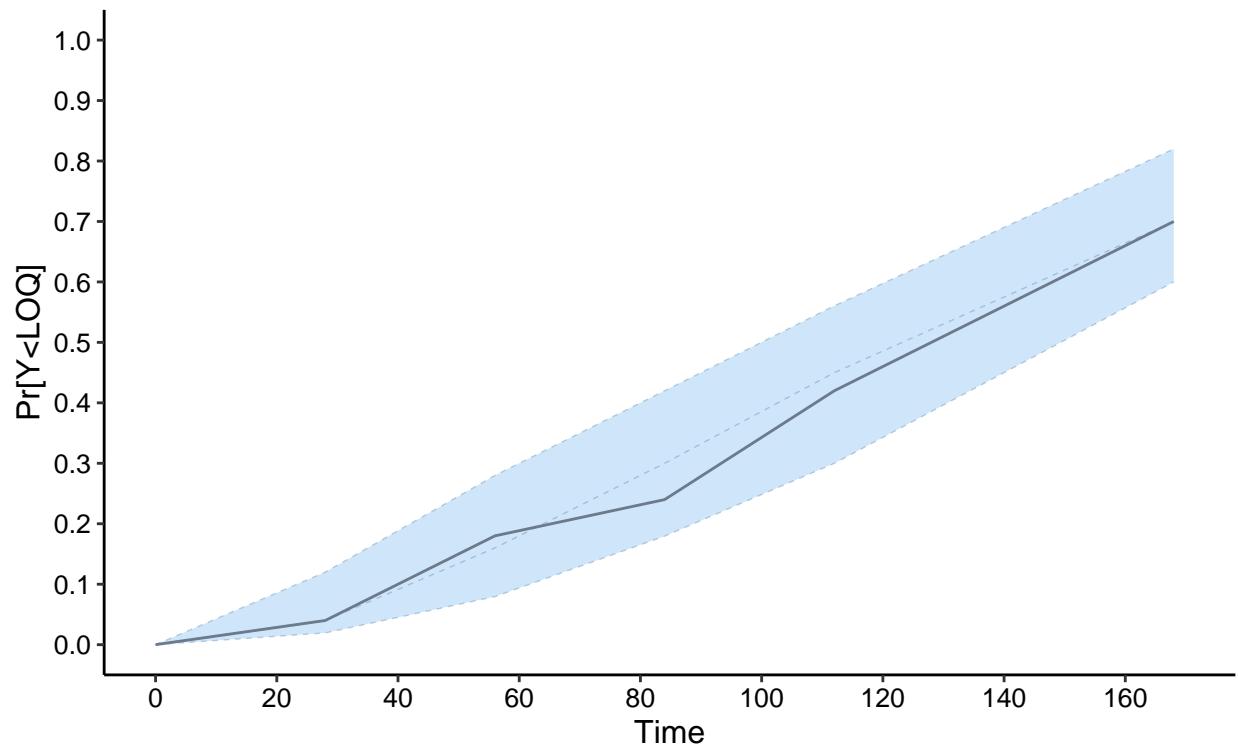
```
## Warning: Removed 8 rows containing non-finite values (stat_boxplot).  
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

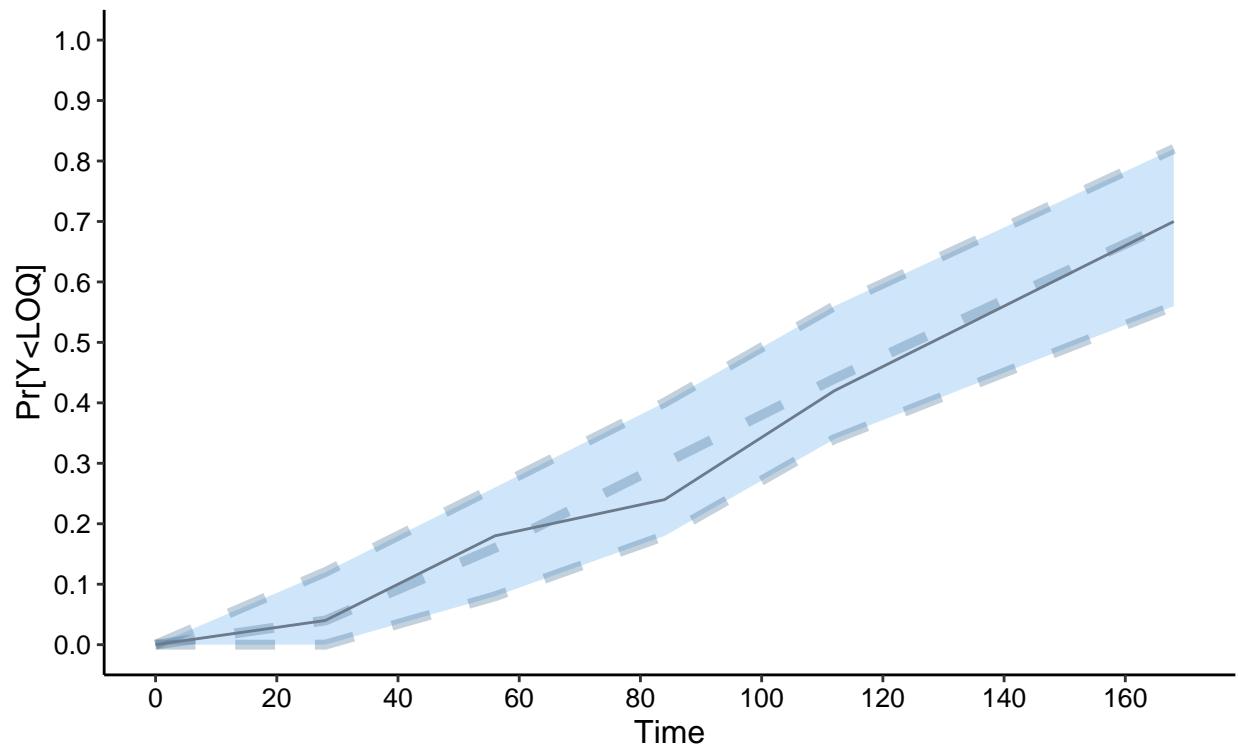


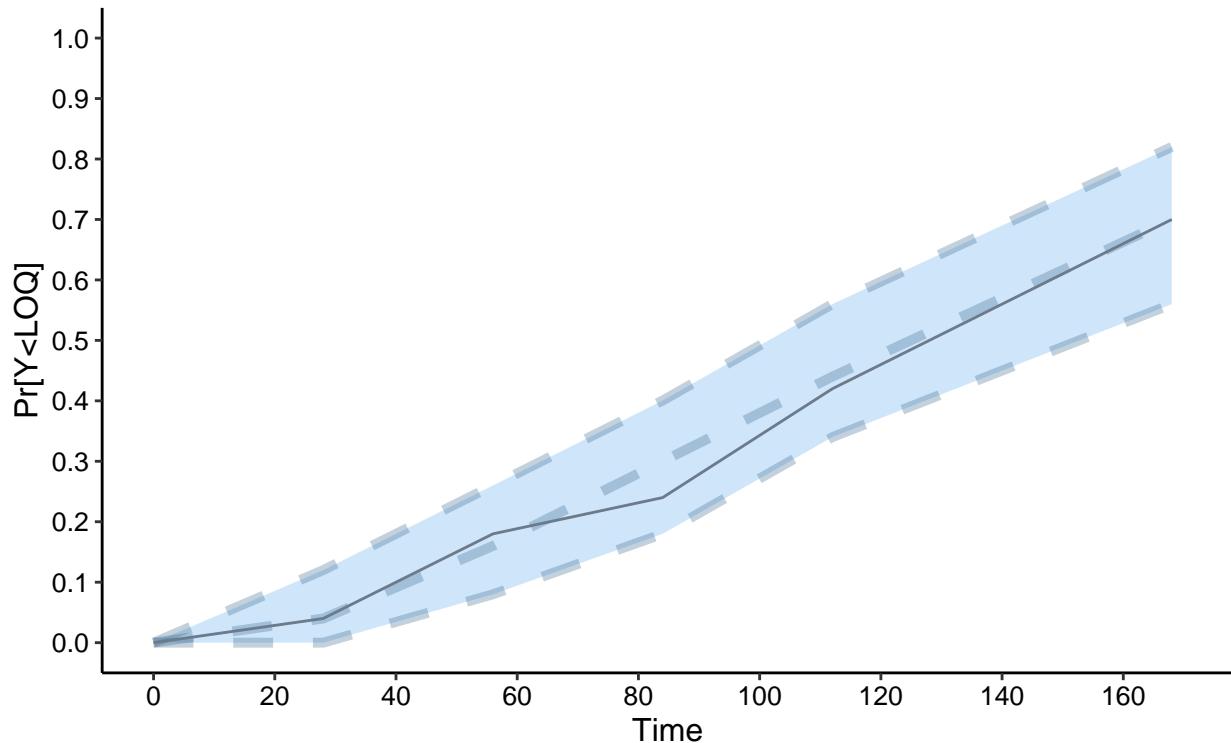
P(Y<LOQ)

- **Problems** (à résoudre Romain)
- **solved** fails
- avant, vérifier: - **solved PI** on ne voit pas les bornes extérieures du PI (ça devrait être contrôlé par lty.bands et col.bands aussi) - **solved axes** missing lines for axes - **solved lwd.bands** should probably be increased for this plot :-/ (compared to the others), but never mind => show an example in Beautiful graphs







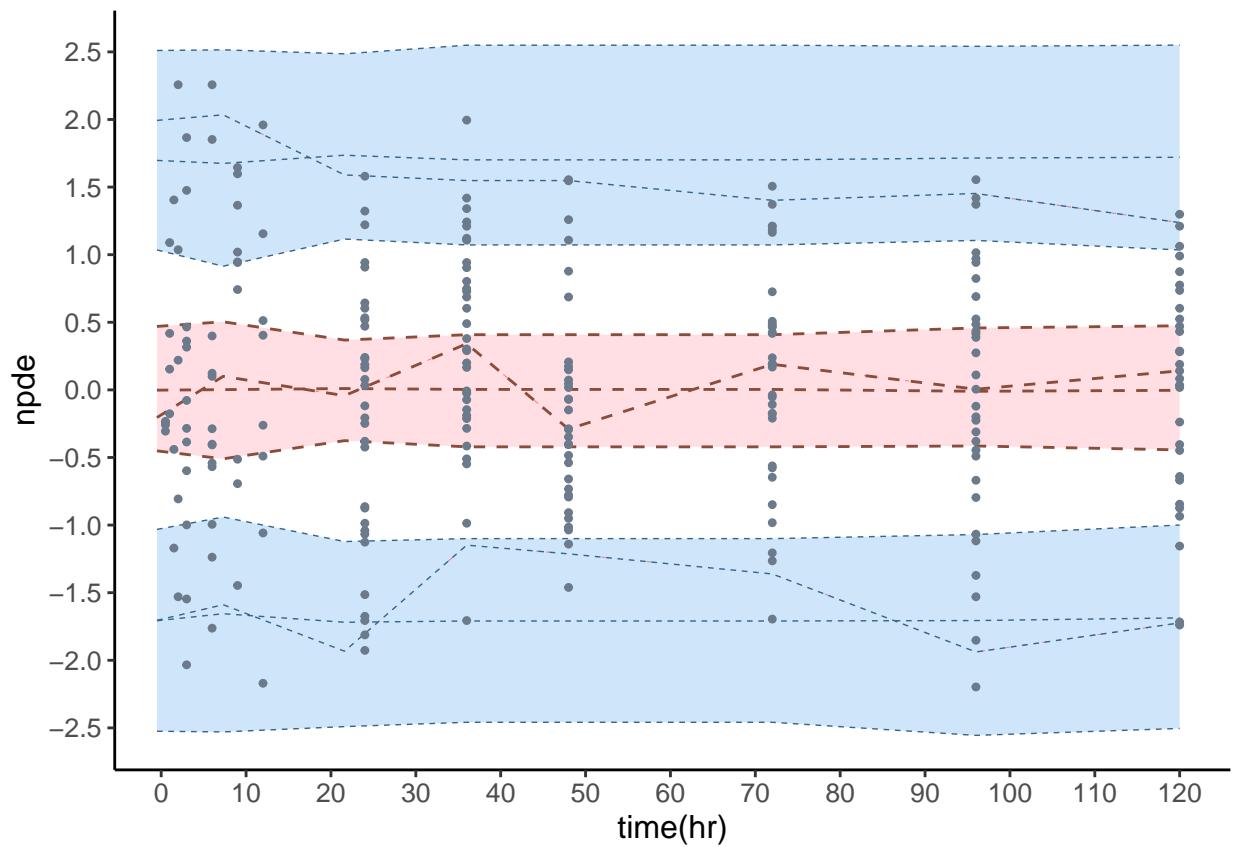


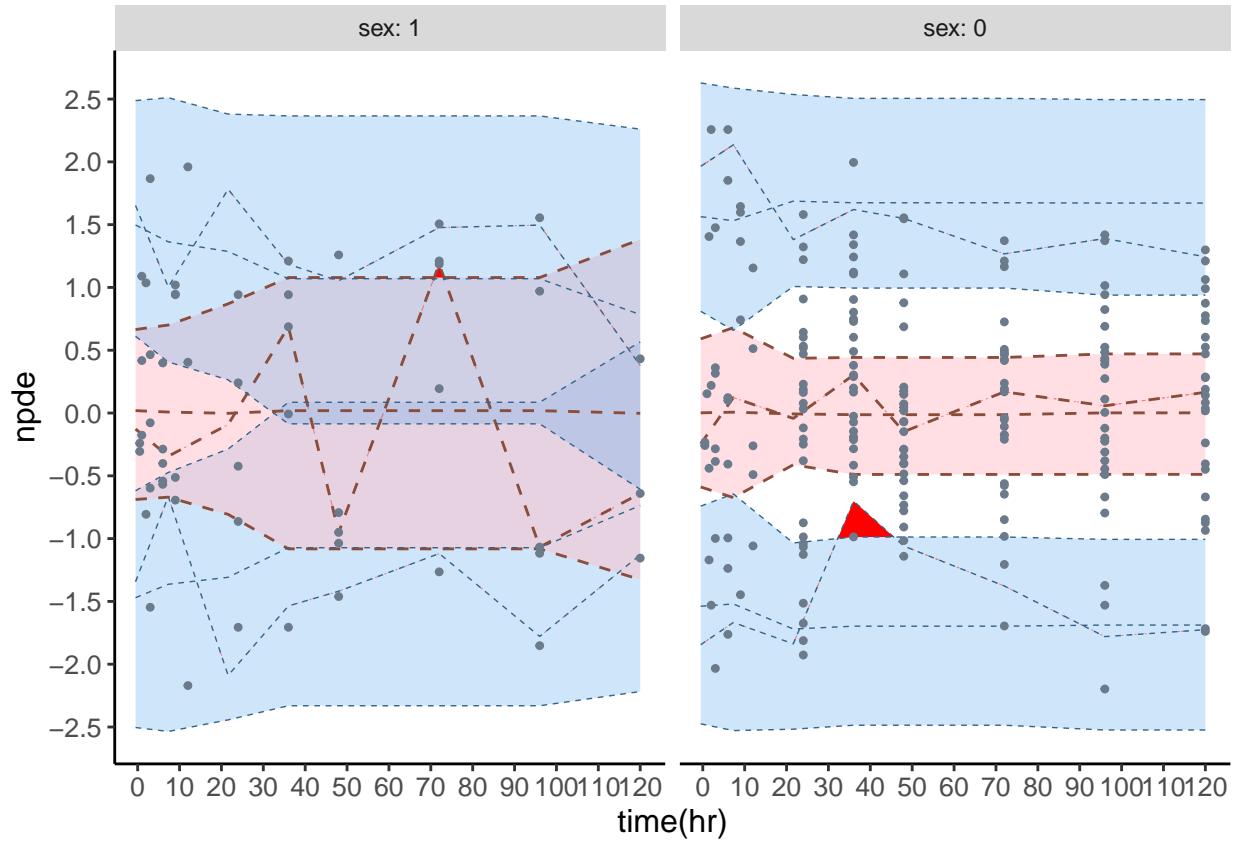
Covariate plots

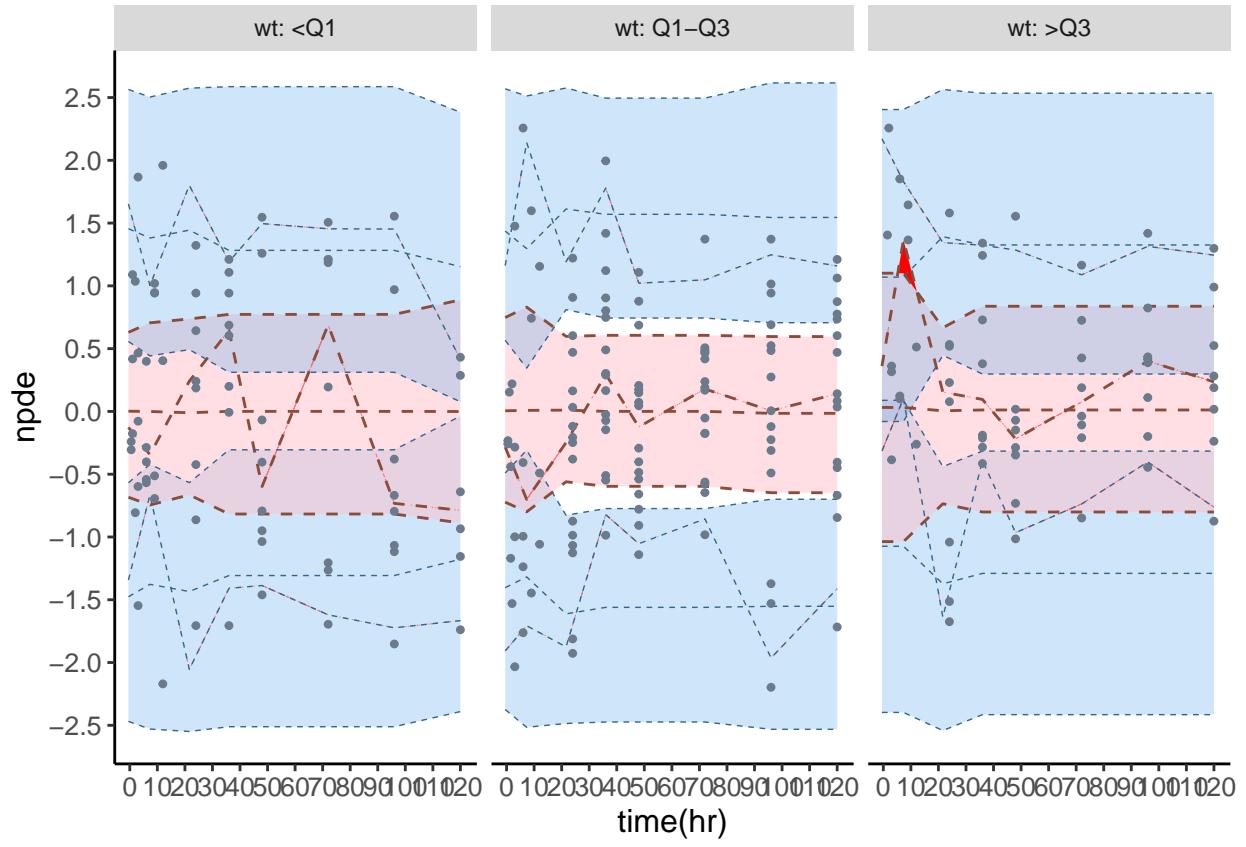
Covariates

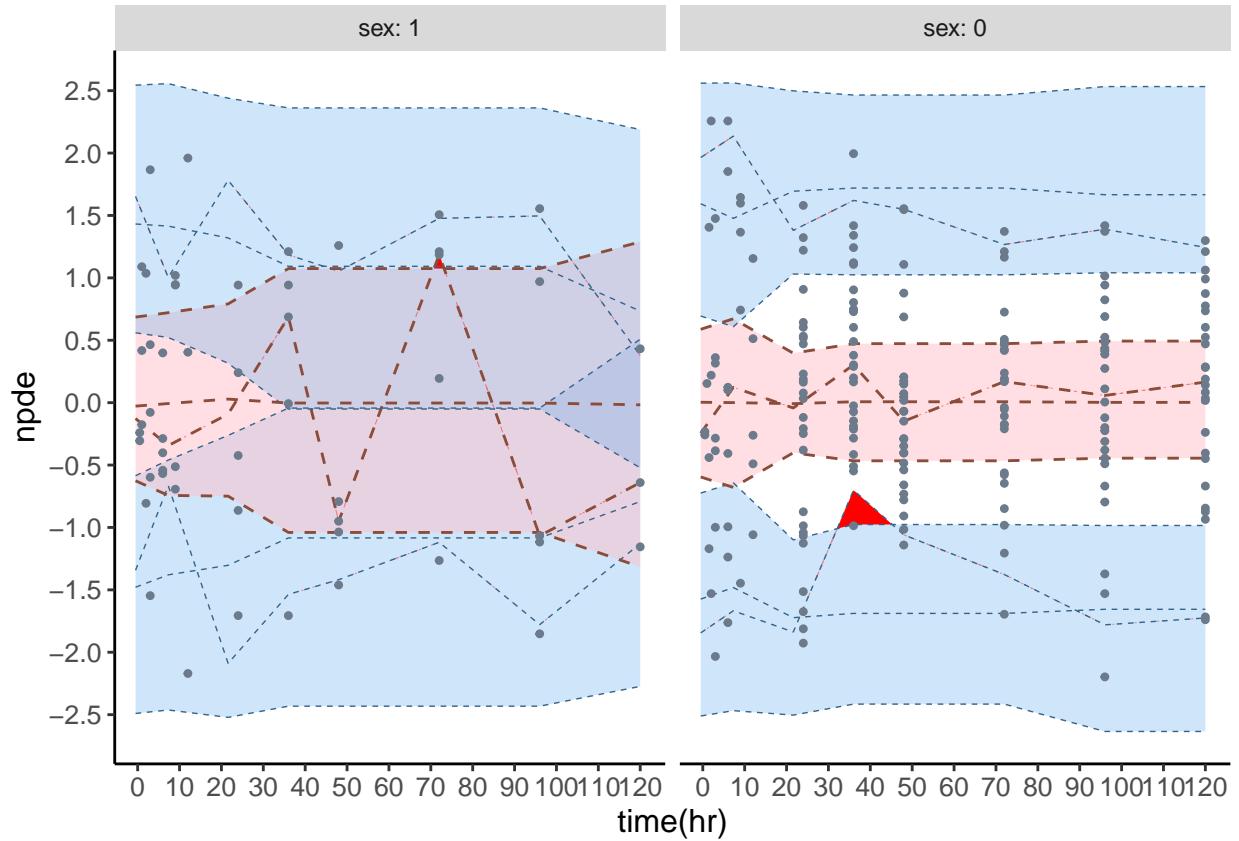
Splitting plots according to covariates

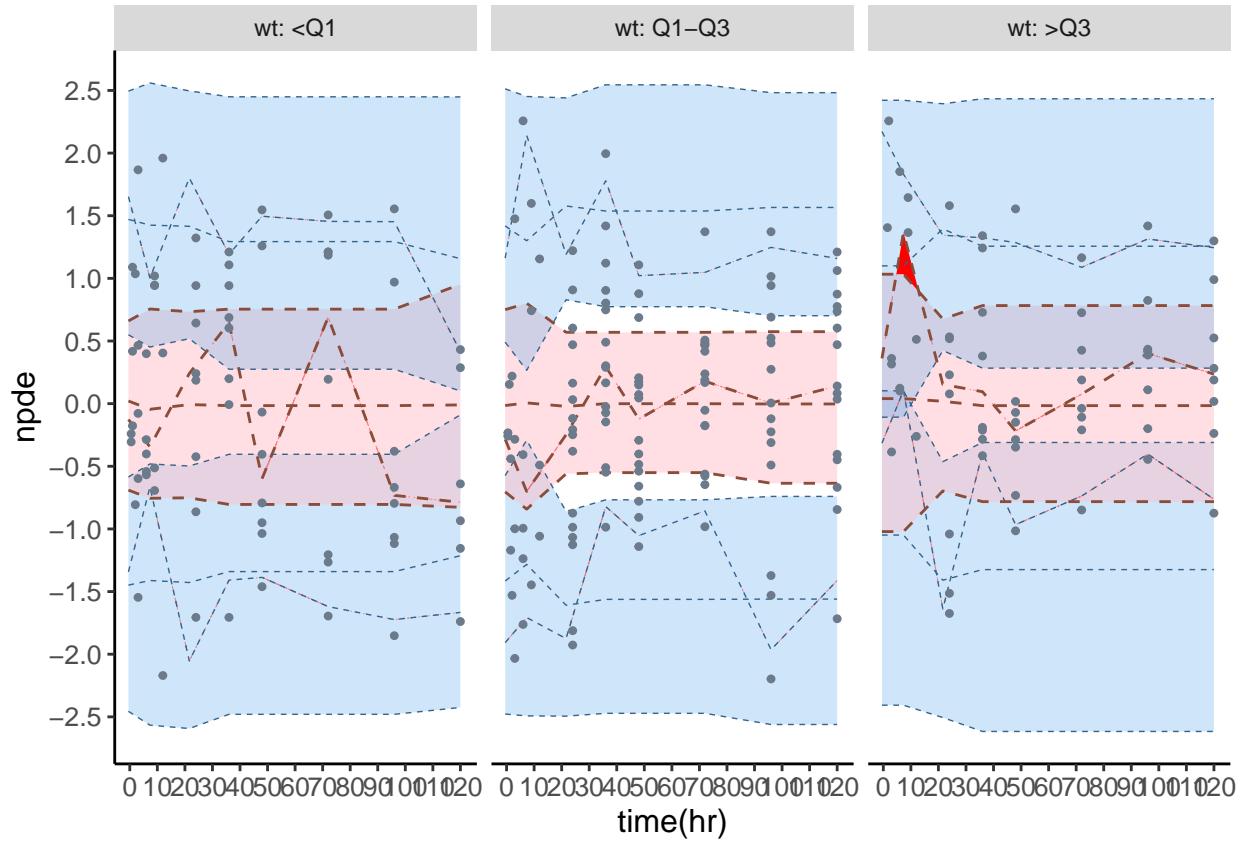
- **Problems (à résoudre Romain)**
 - **solved** yes here we really need to extend PI to cover the whole X-range for x.scatter and pred.scatter (not VPC)
 - * just the PI (not the observed percentiles or outlier bands)
 - **covsplit**
 - * **solved** doesn't work with the default plots => behaviour TBD
 - with one covariate, gives default plot (not split by the covariate)
 - with several covariates, message `arrangeGrob(...)` ...: `nrow ncol >= n` is not TRUE
 - we should get default plots for each category of the covariate but maybe only for one covariate ??? (too crowded otherwise)
- **Problems (à résoudre Eco)**
 - ordering categorical variable (?)
 - why is there a blank space in the pred.scatter plot for sex=1 ?
- **other checks to do**
 - check if PI extend on the X-range when less homogenous data

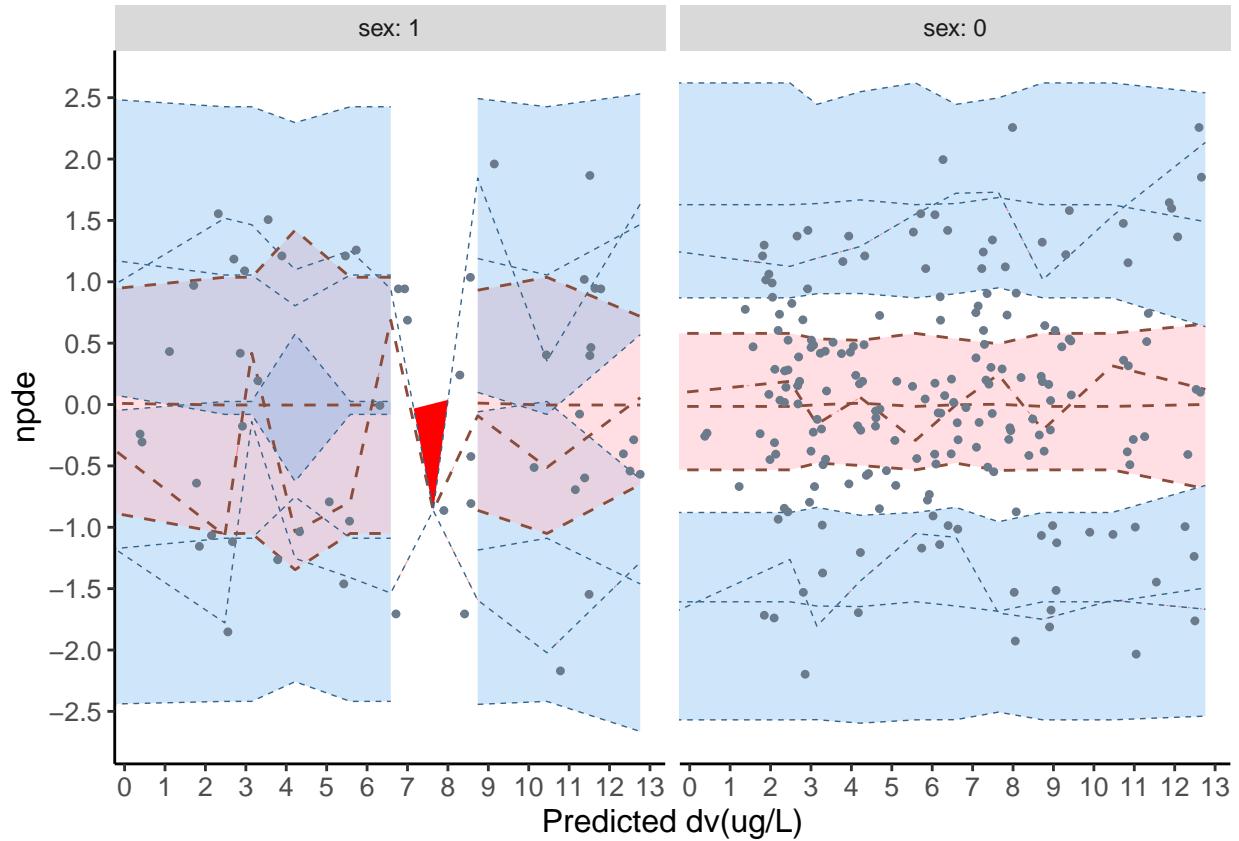


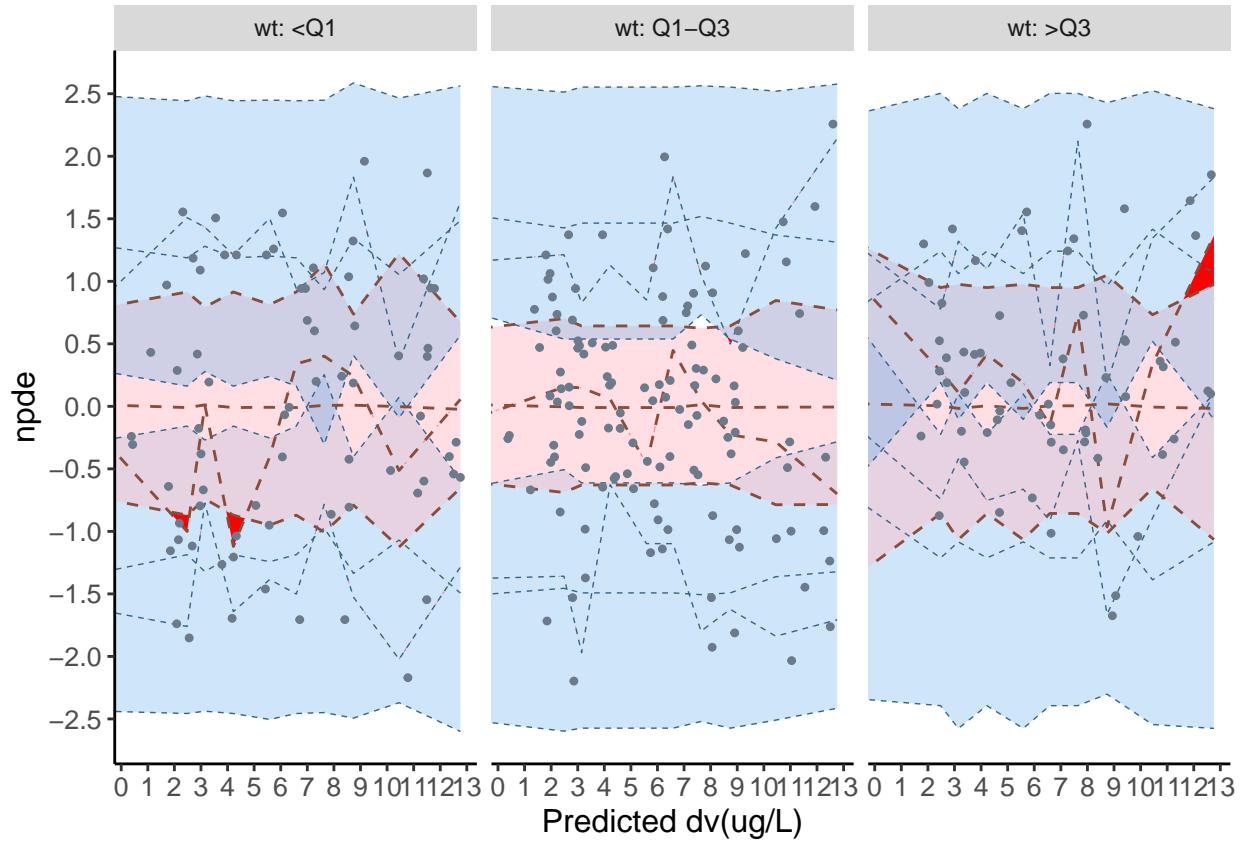


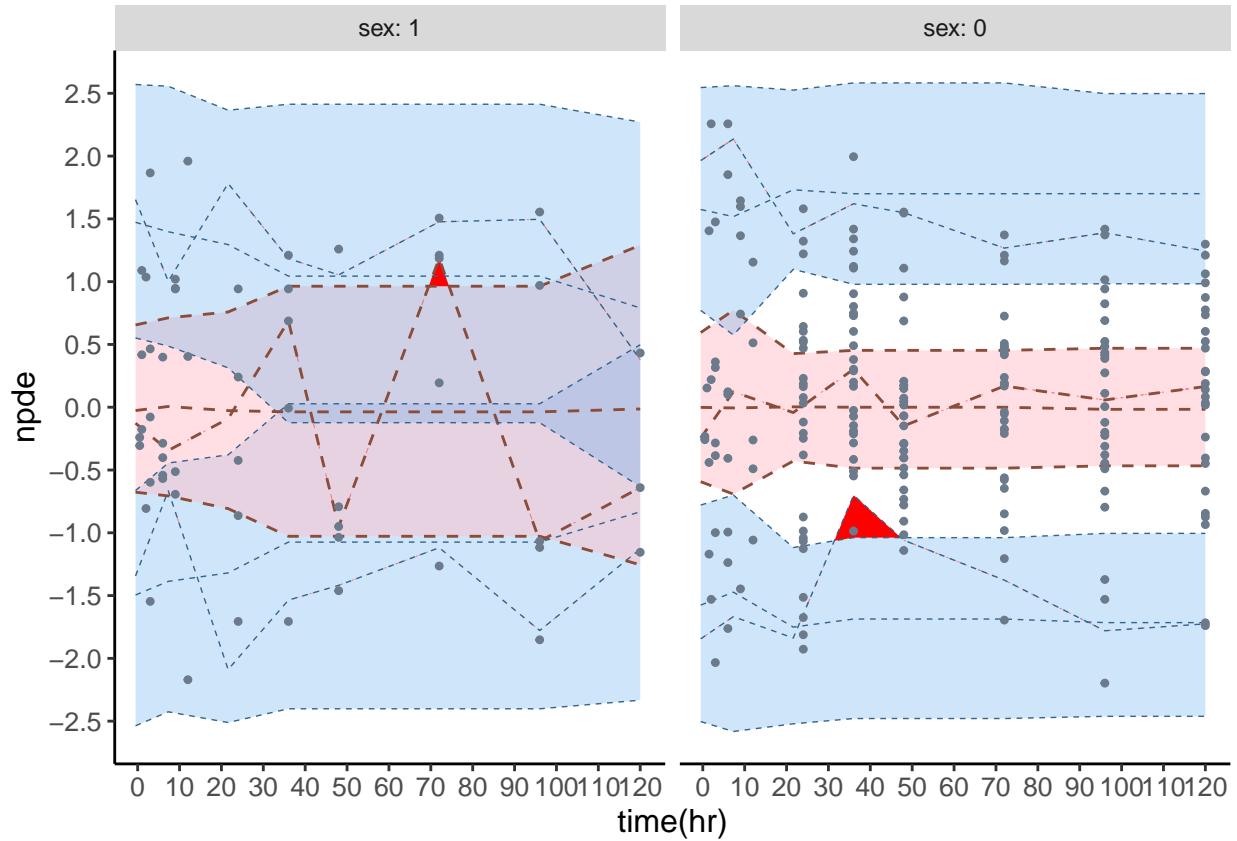


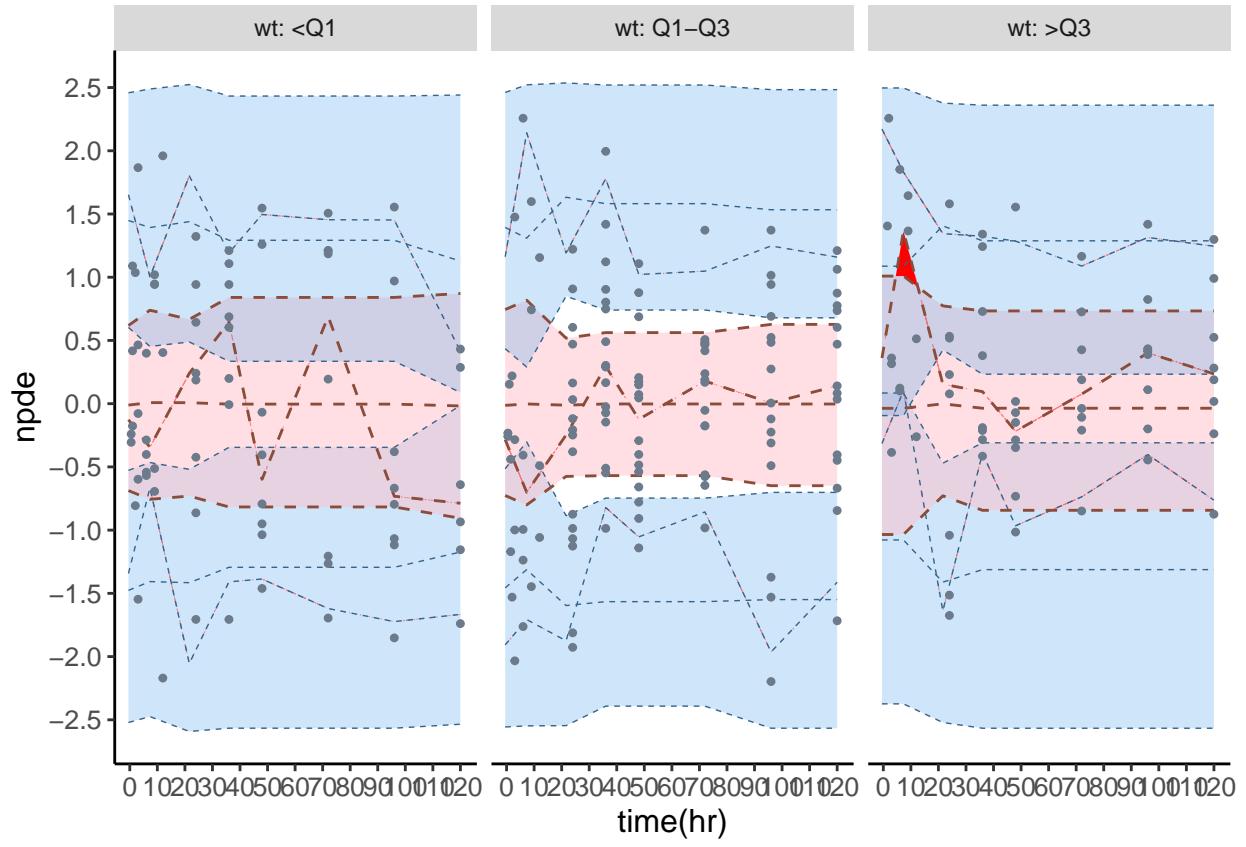


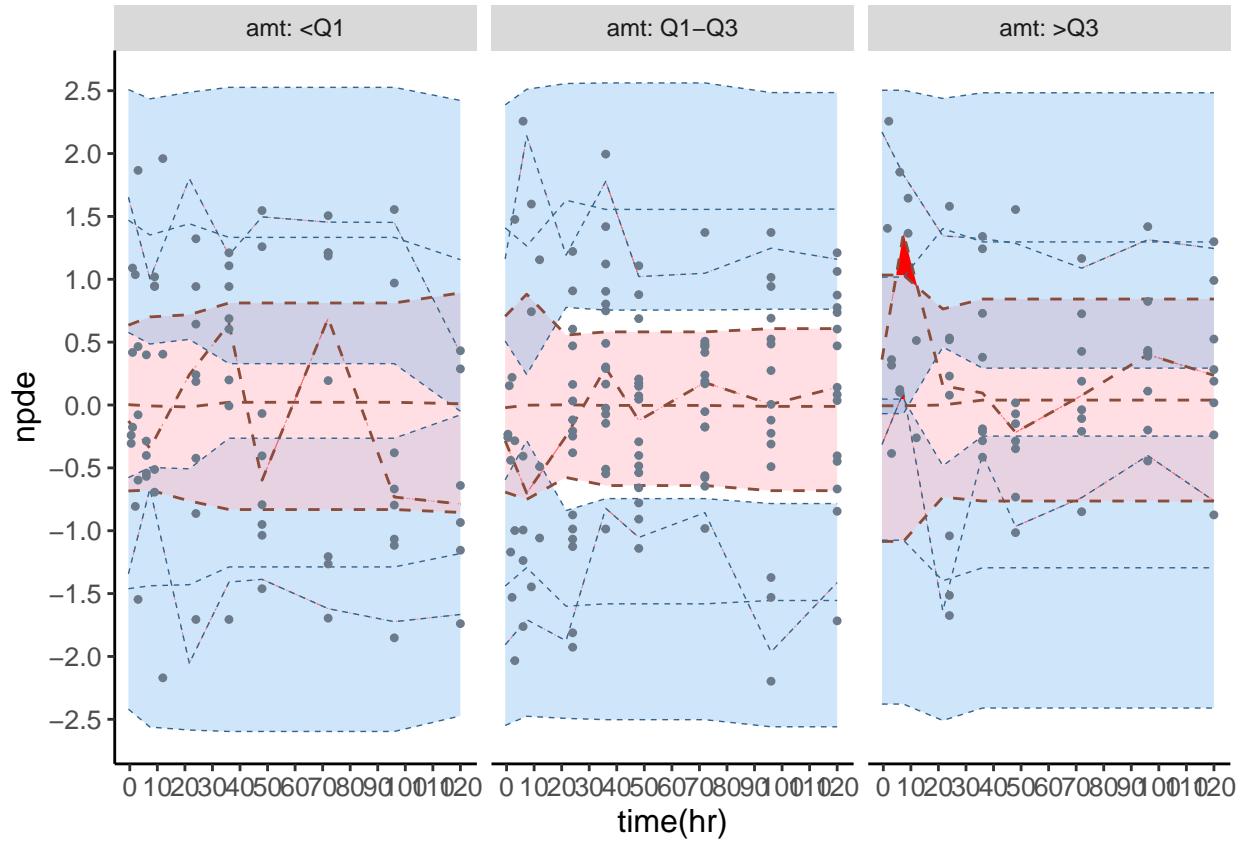


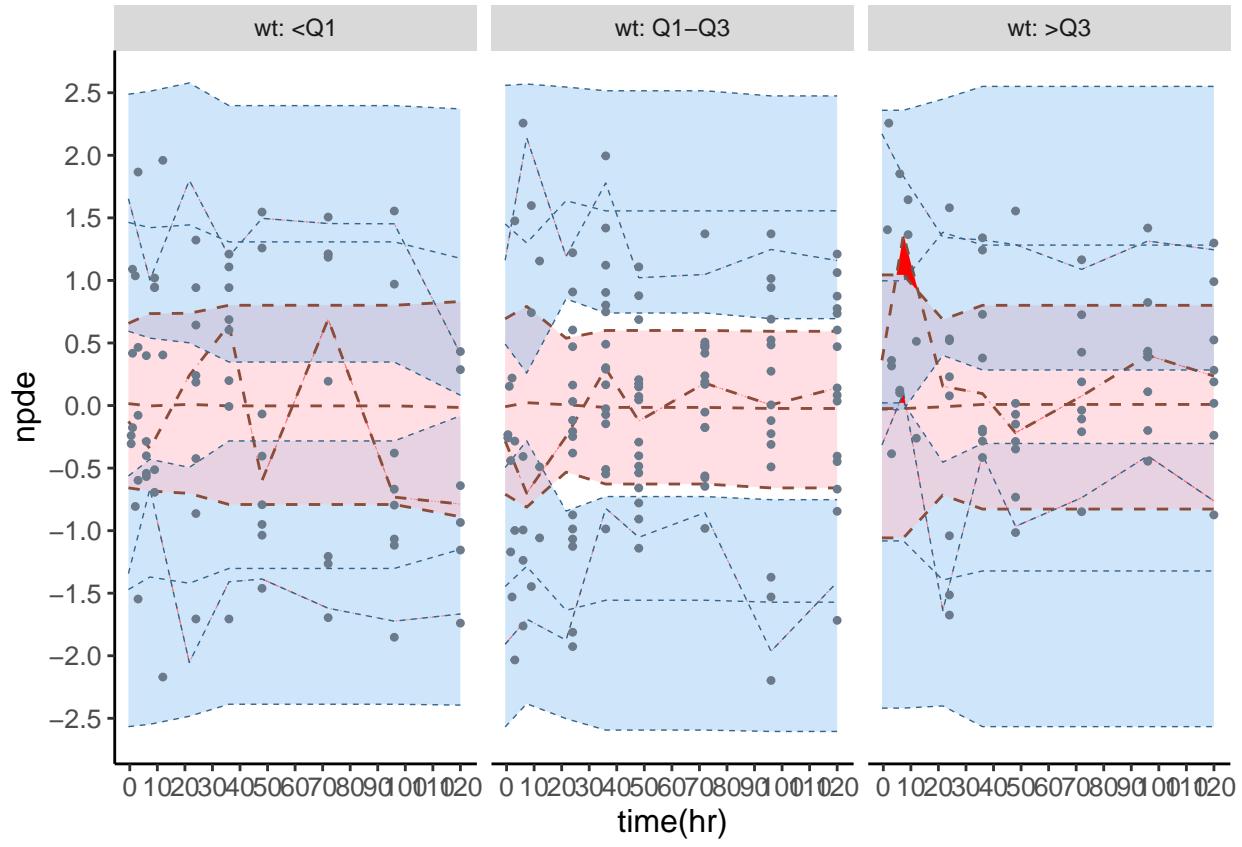


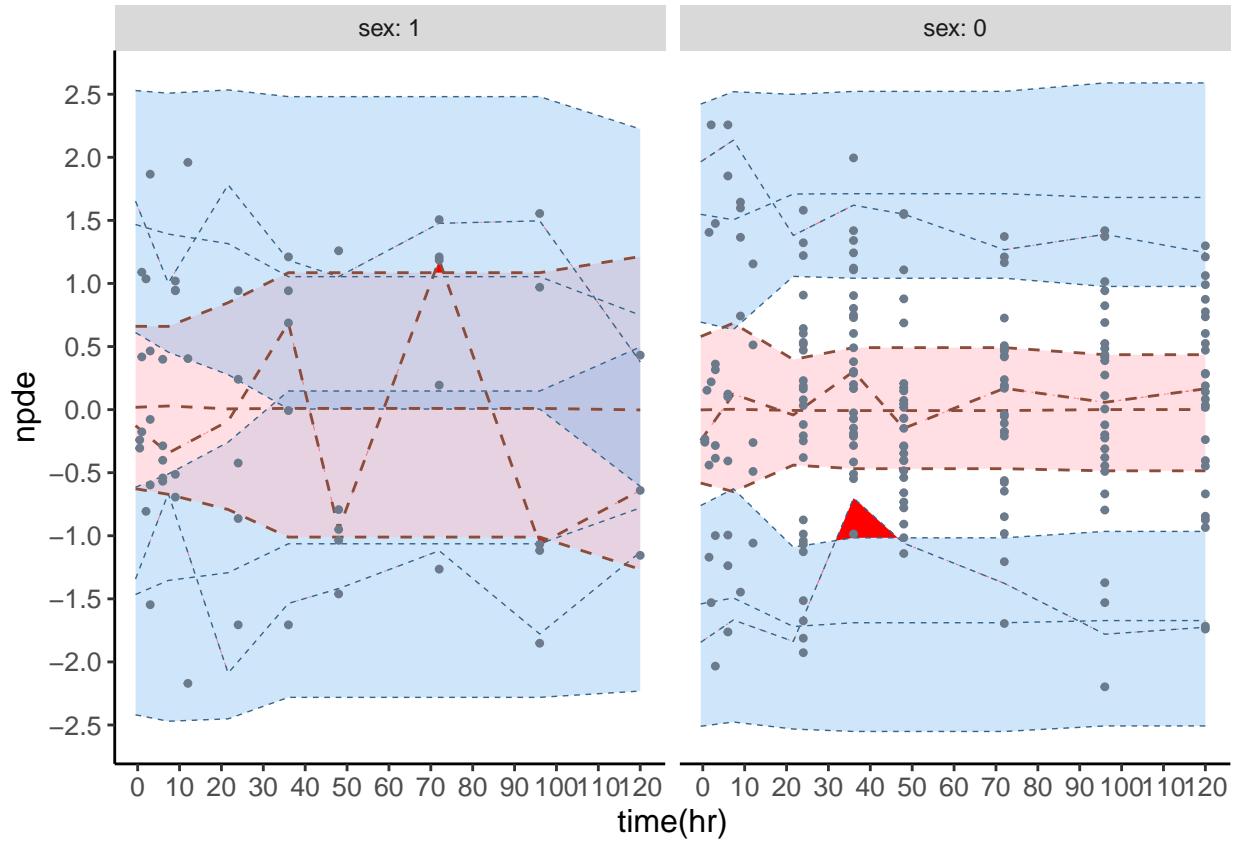


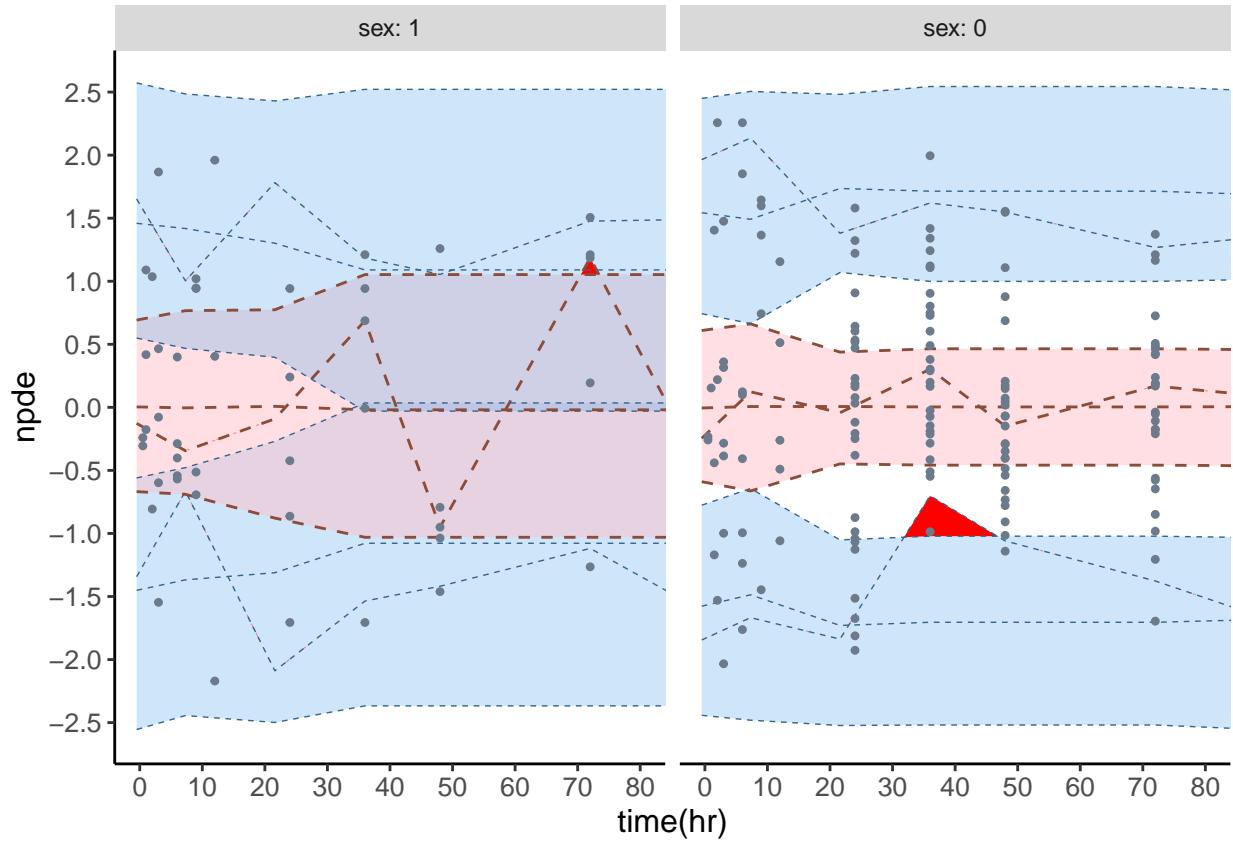


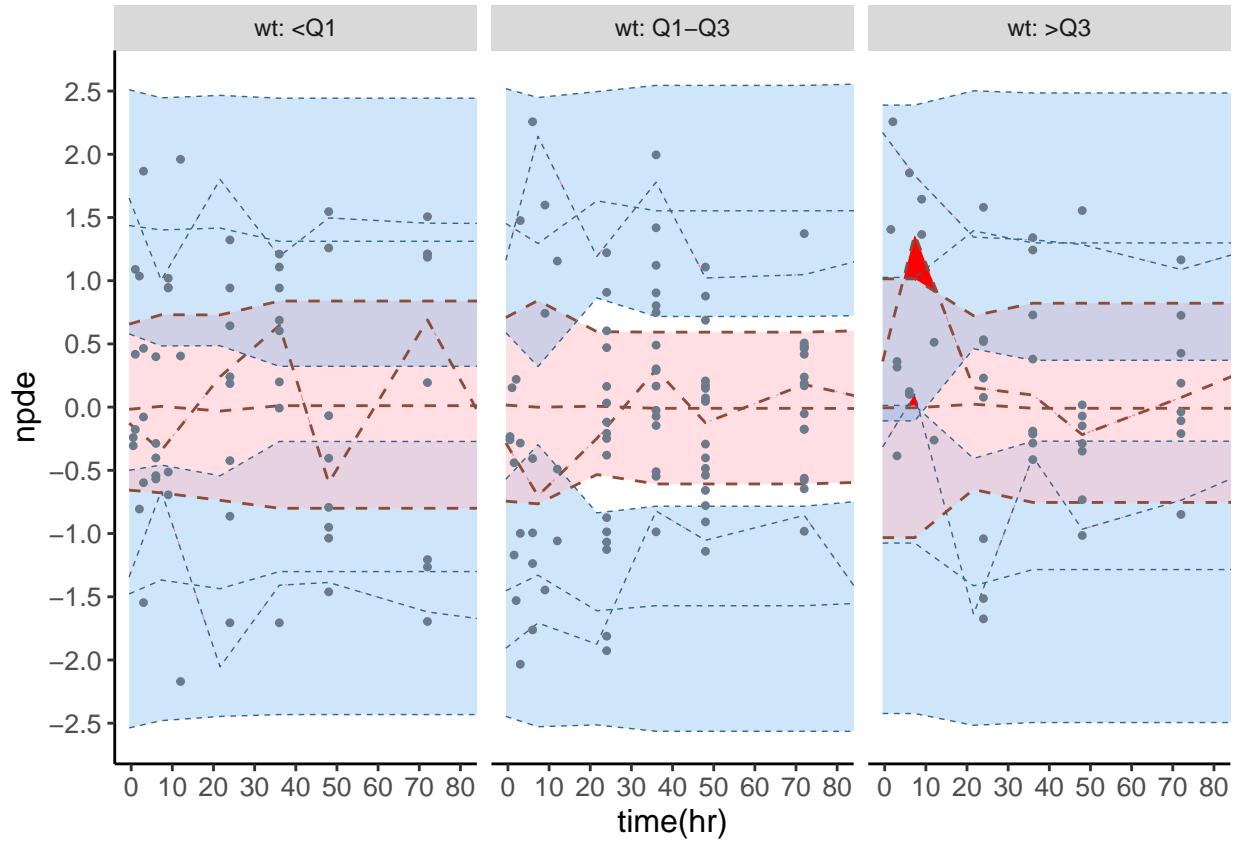


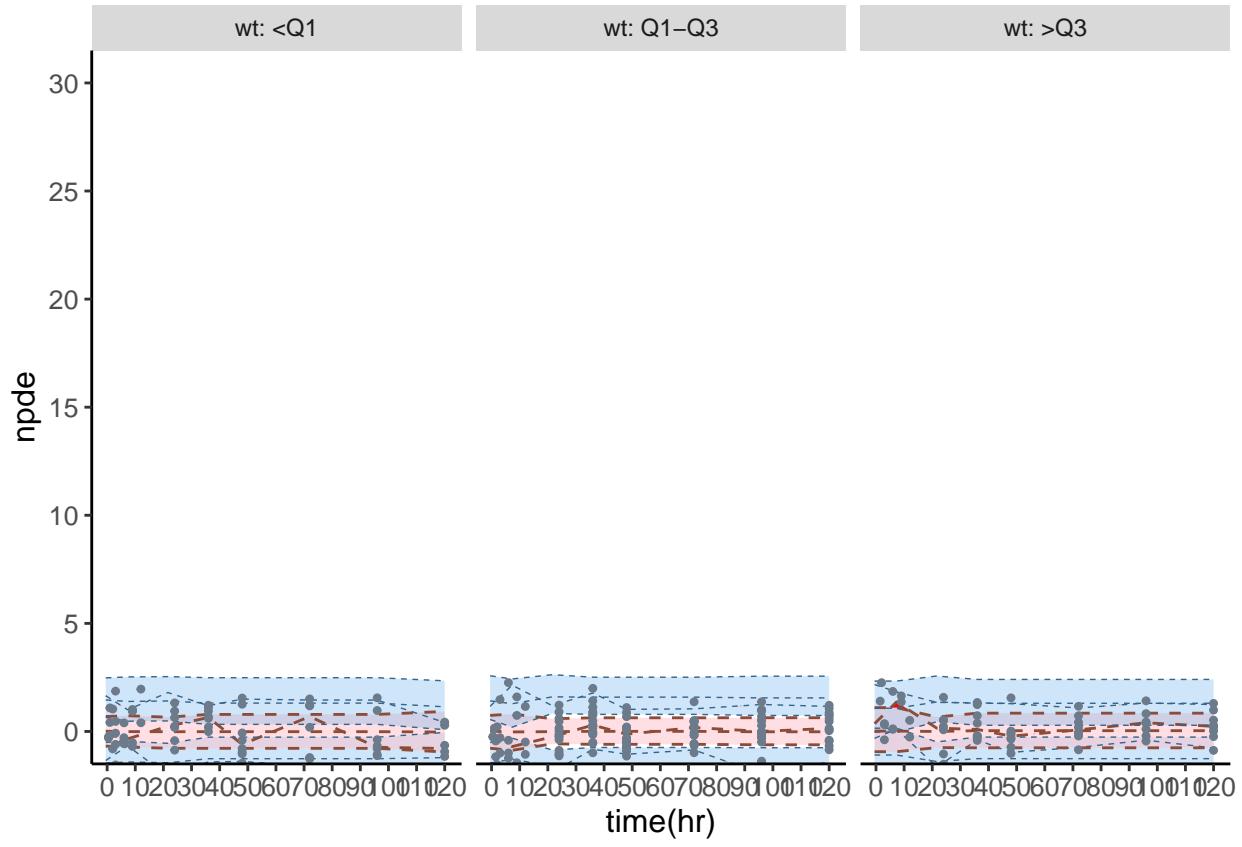


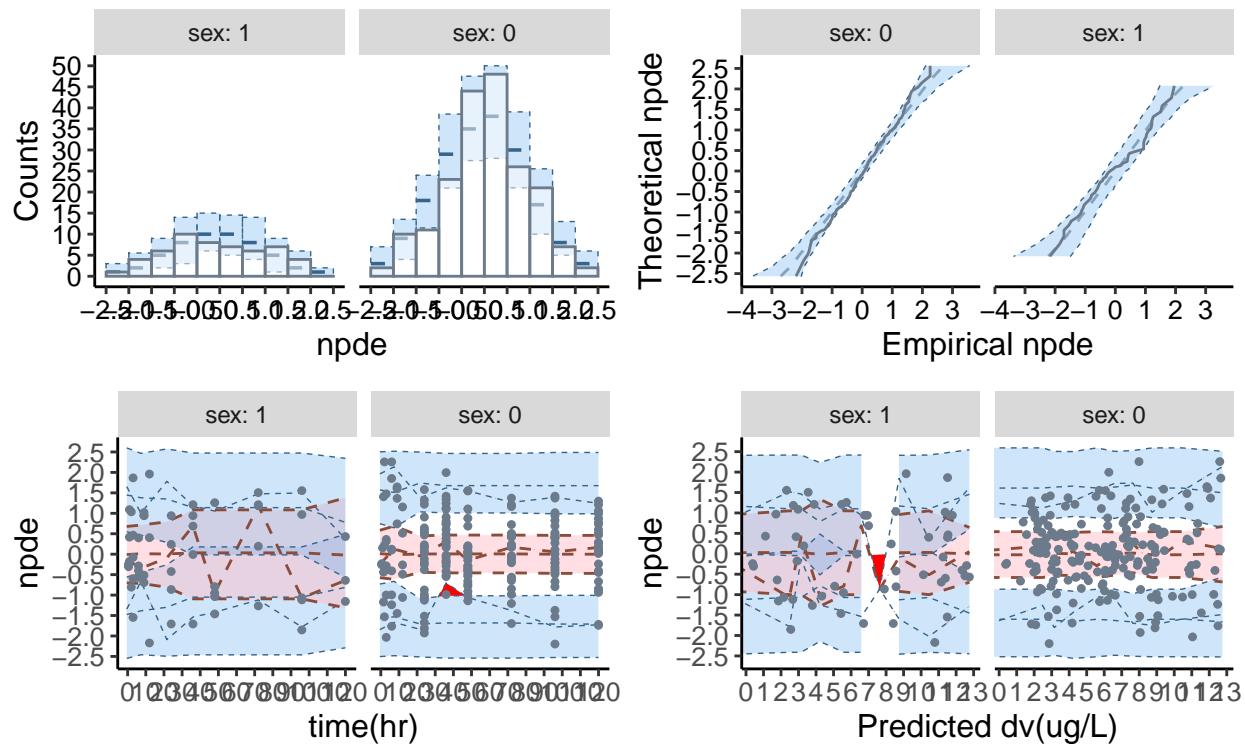


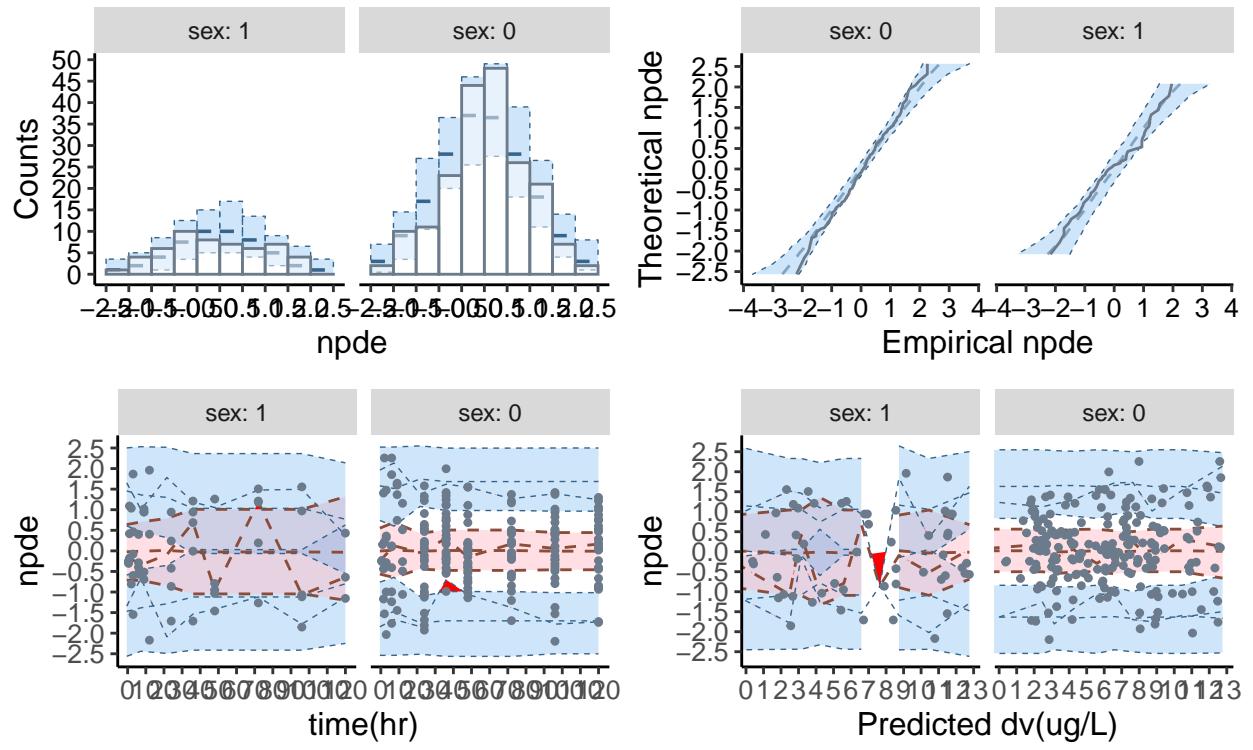


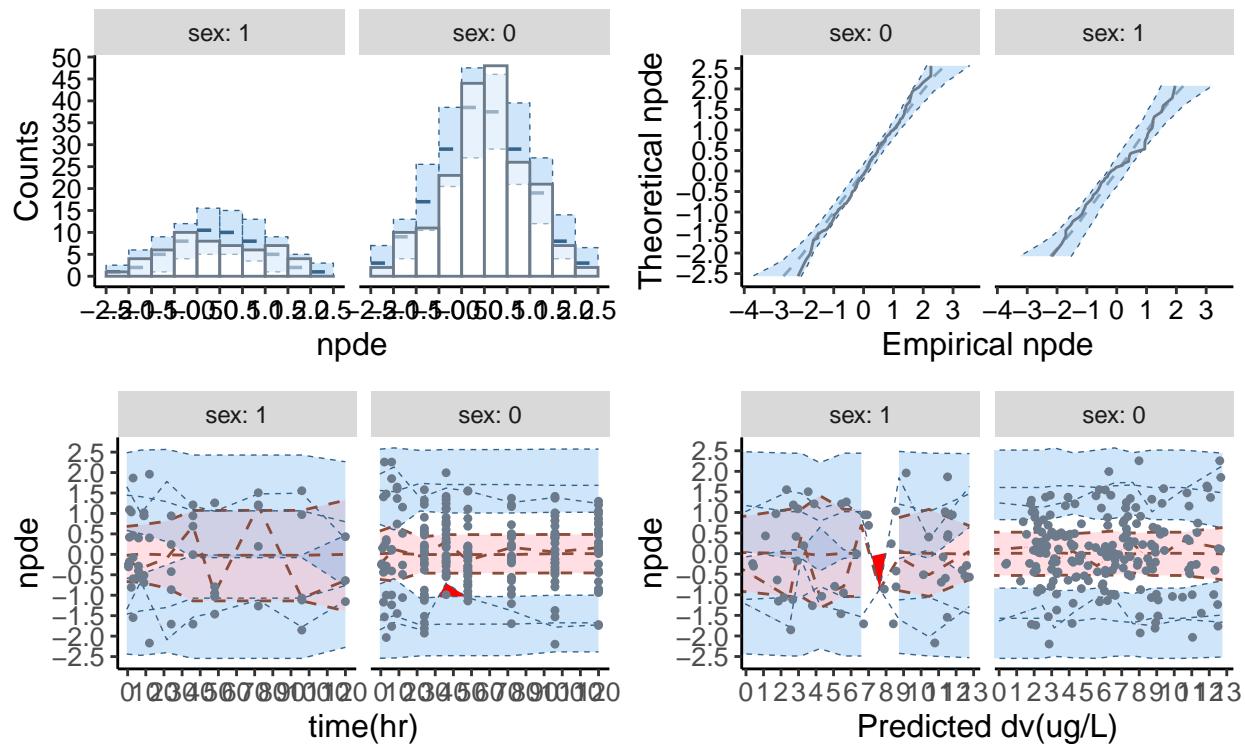




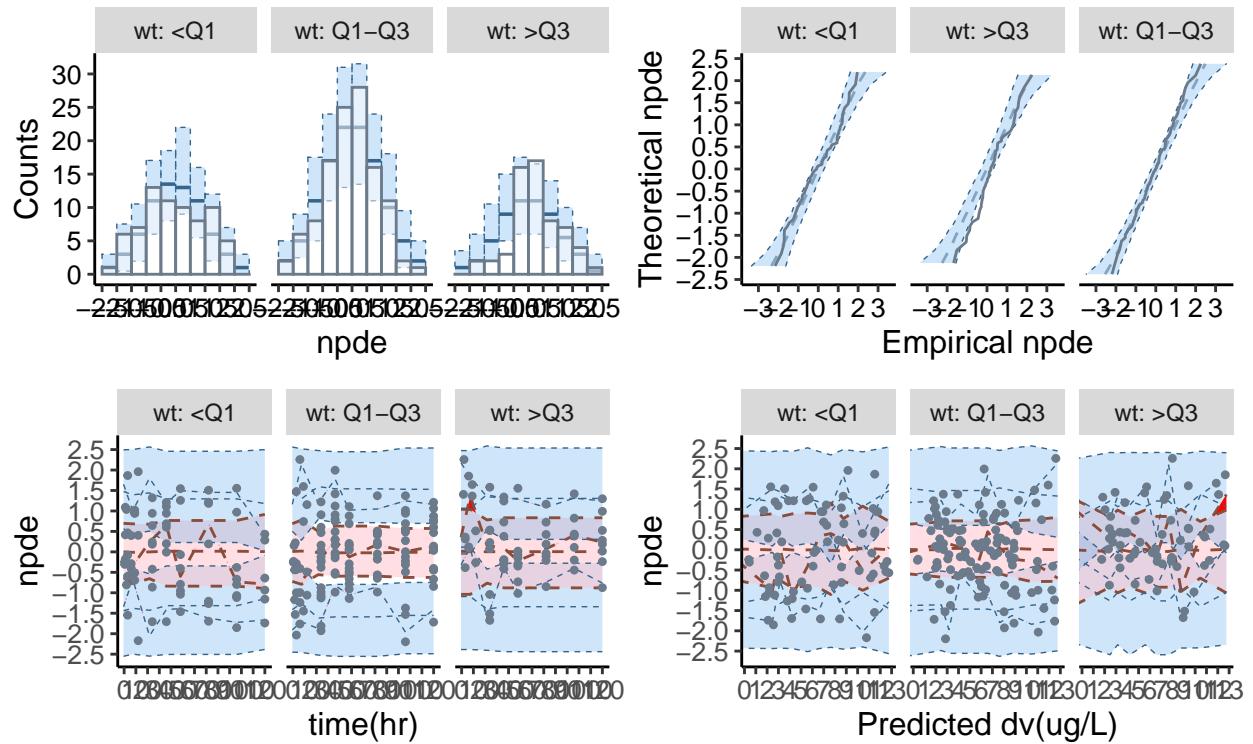


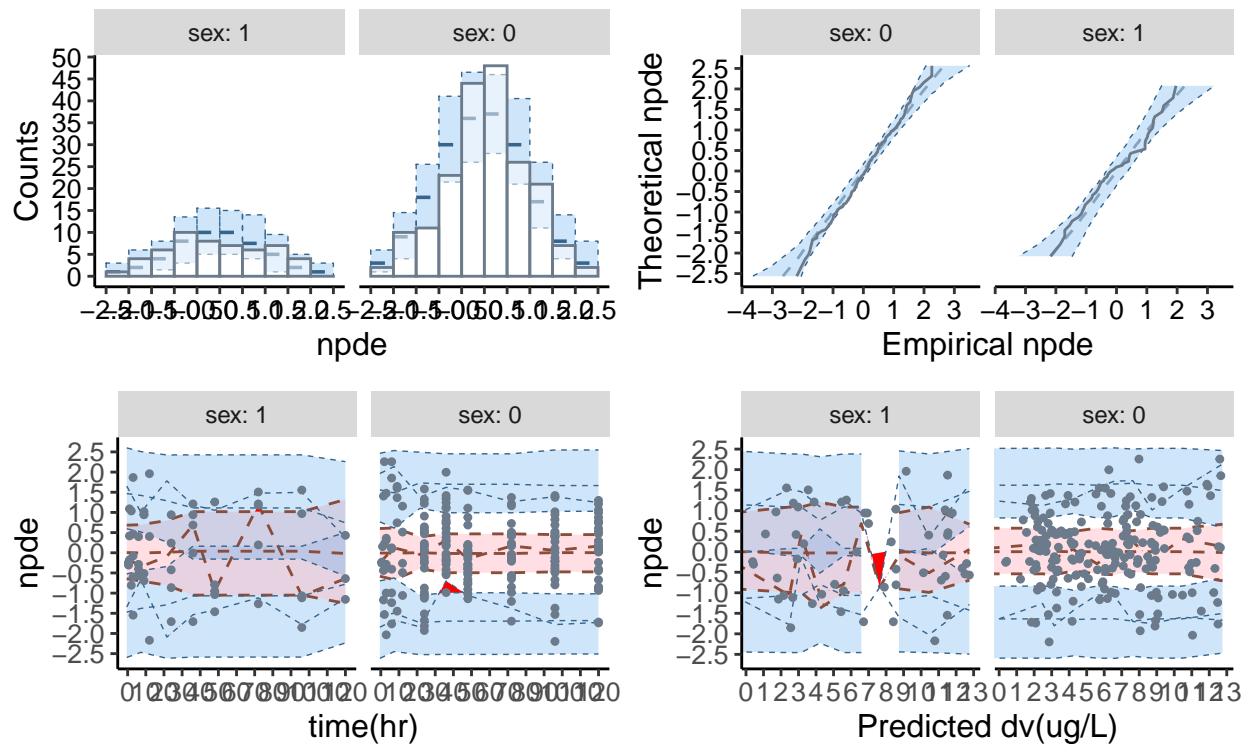




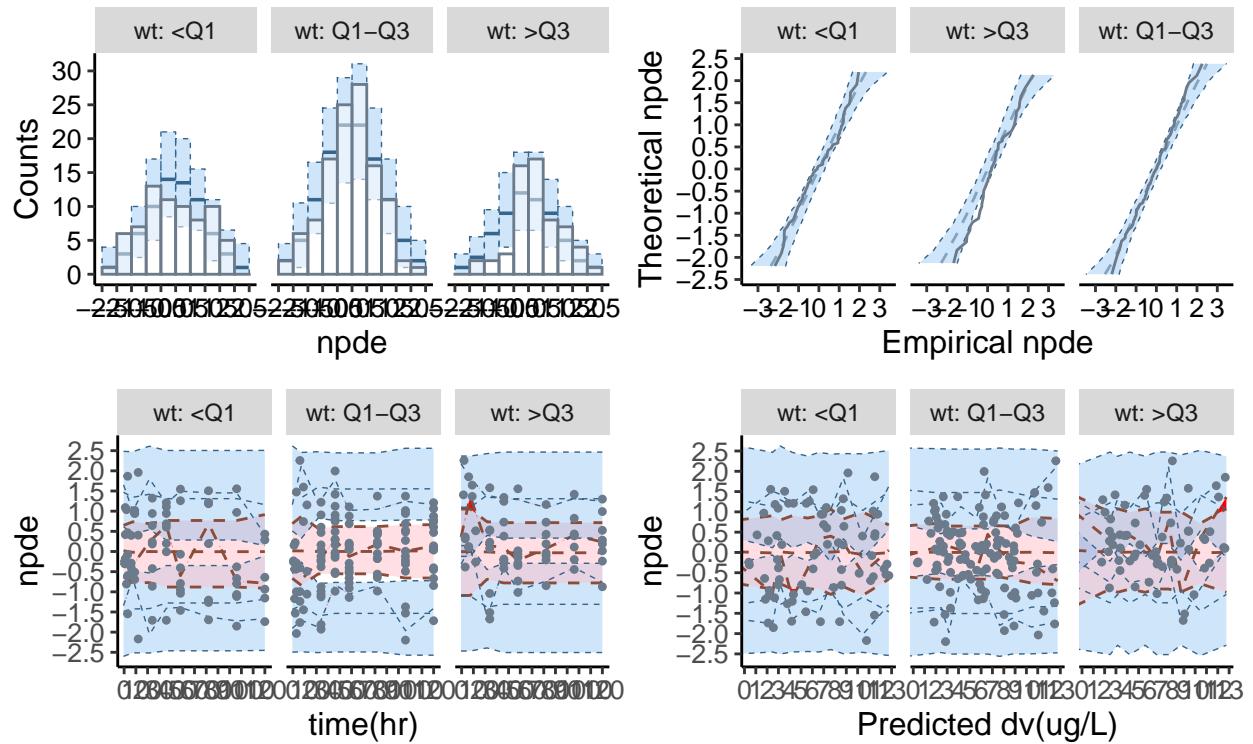


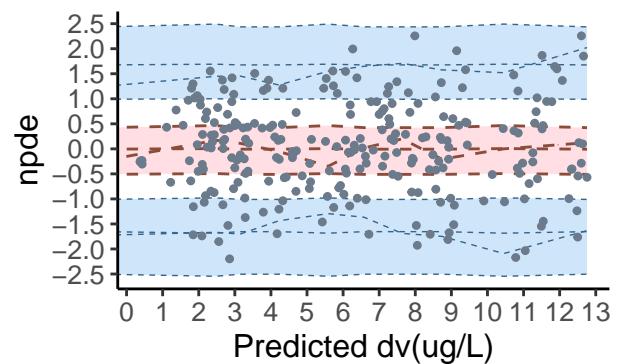
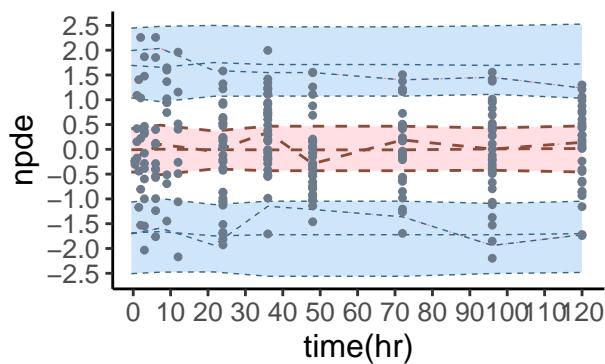
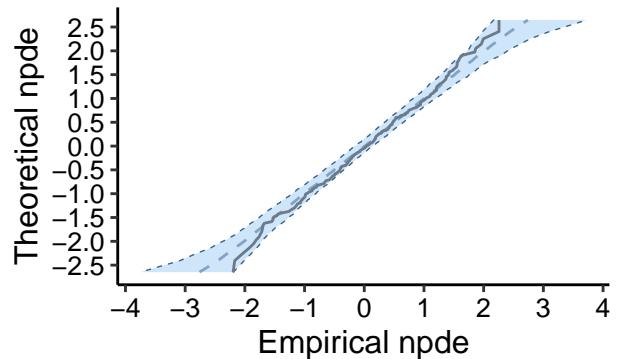
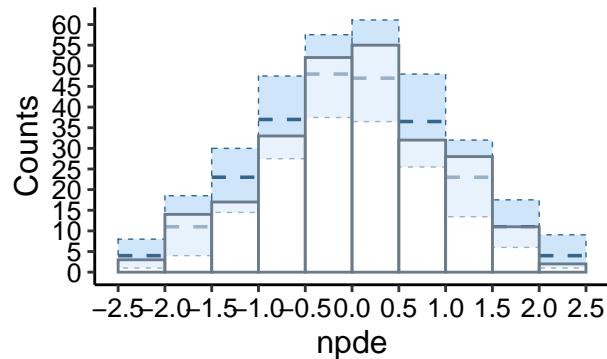
NA



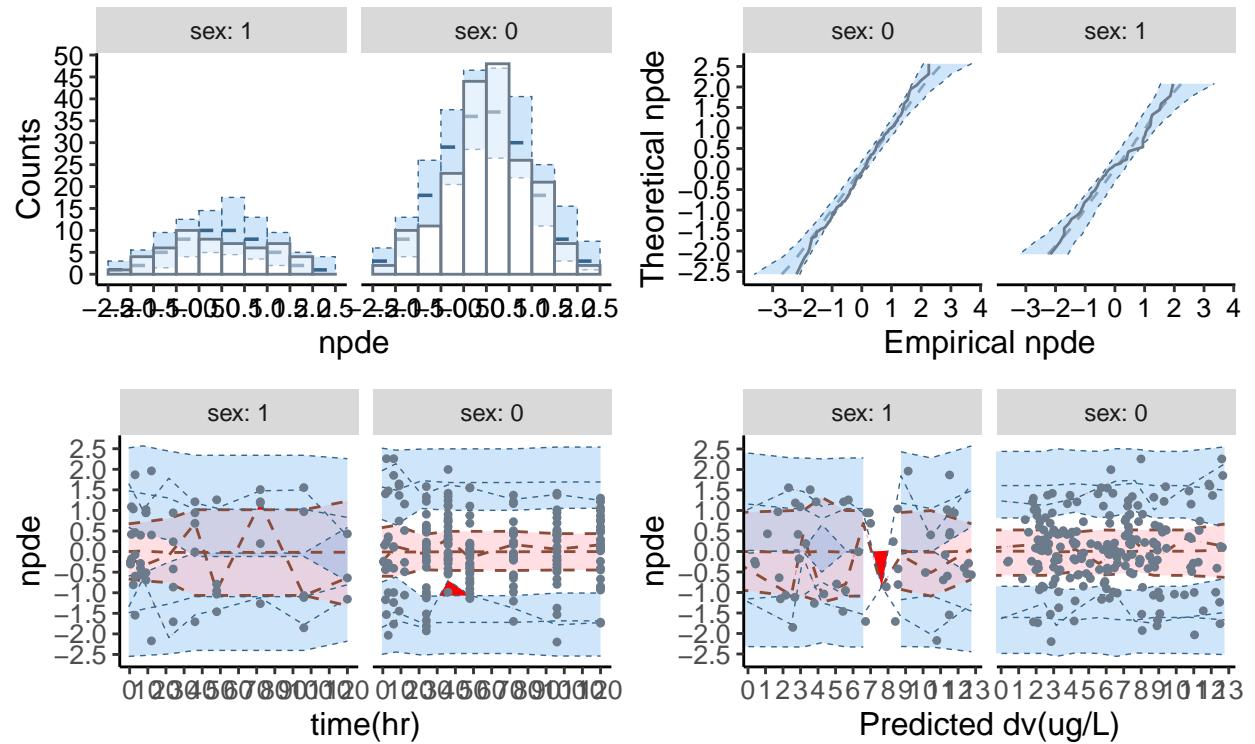


NA

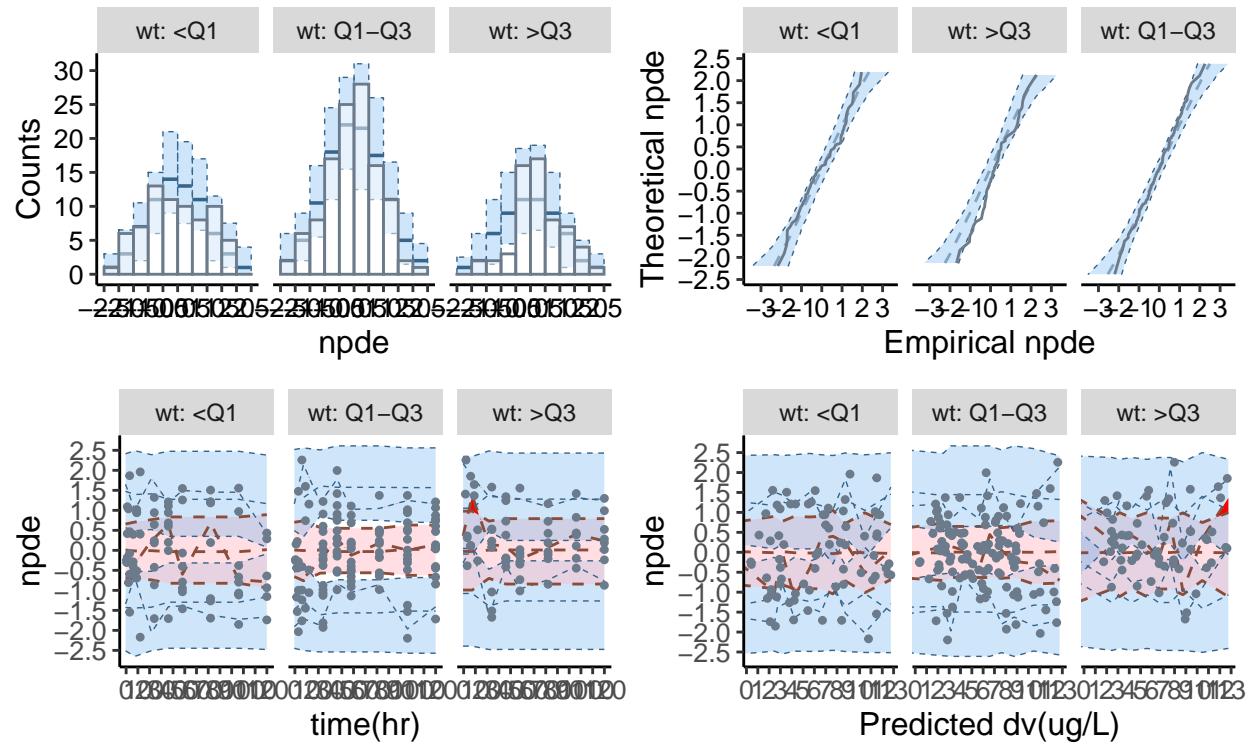




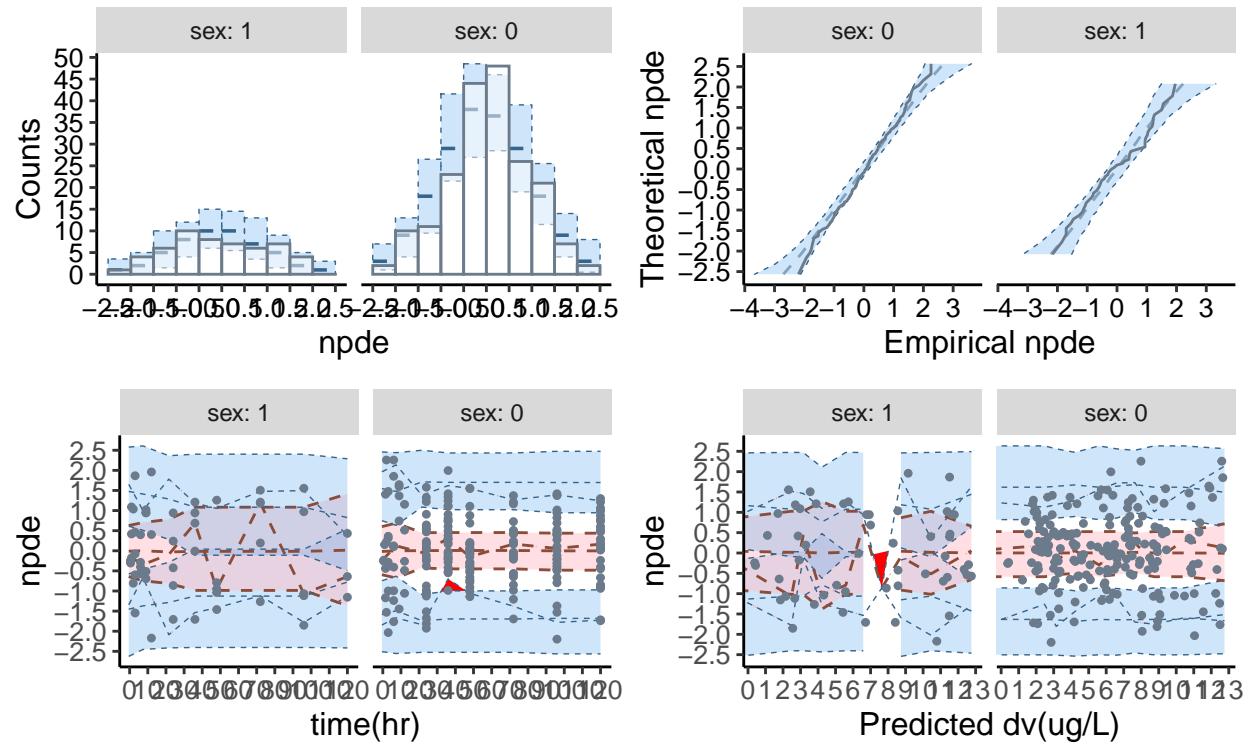
waffle plot : Sex



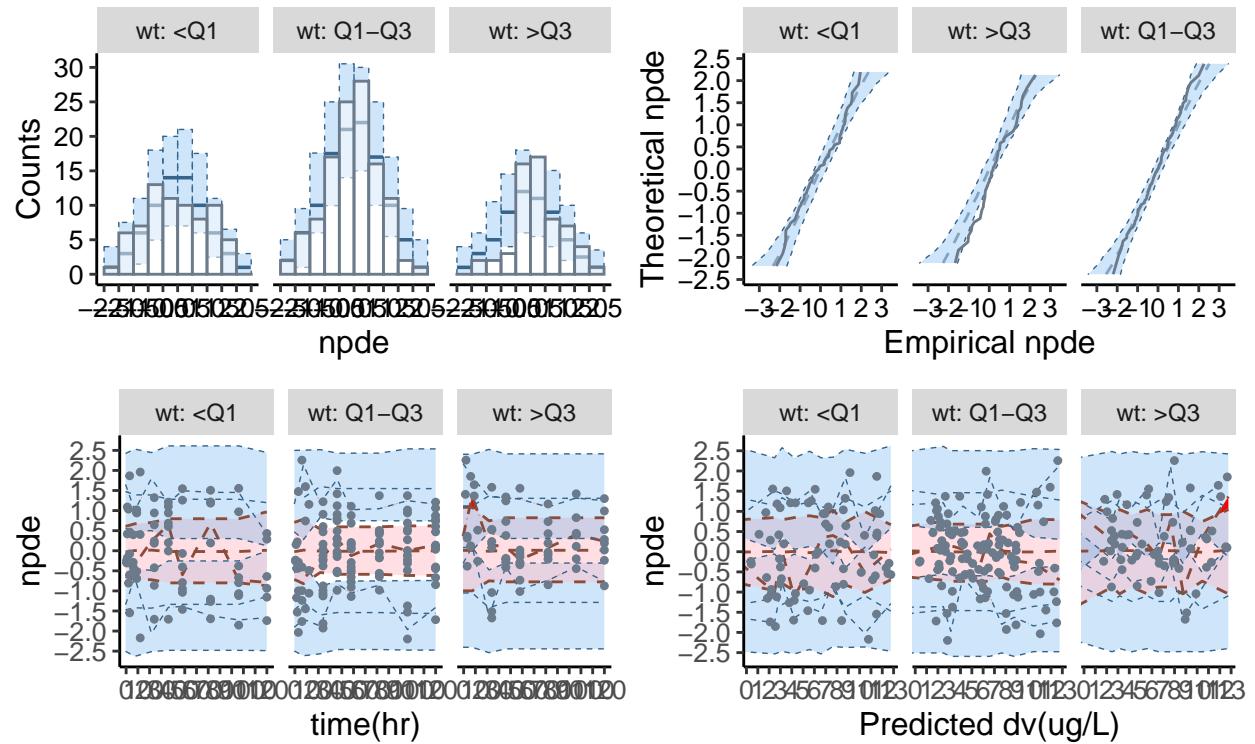
waffle plot : Wt



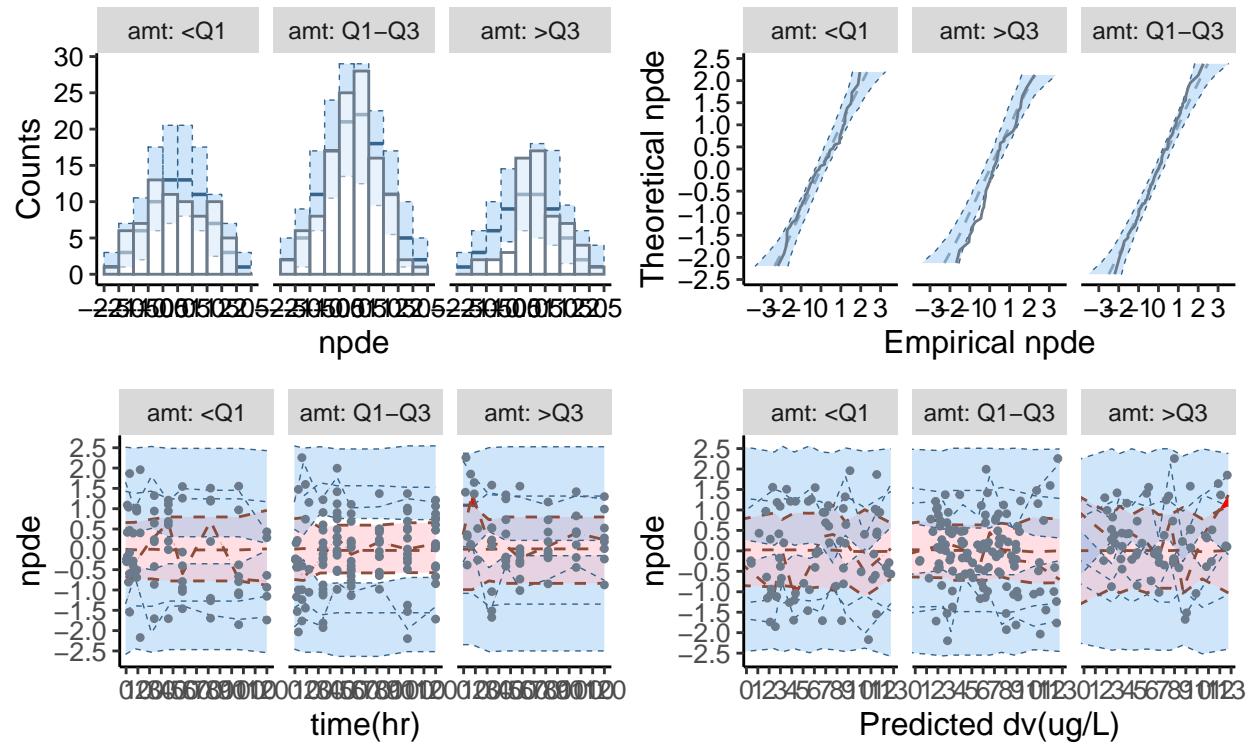
waffle plot : Sex



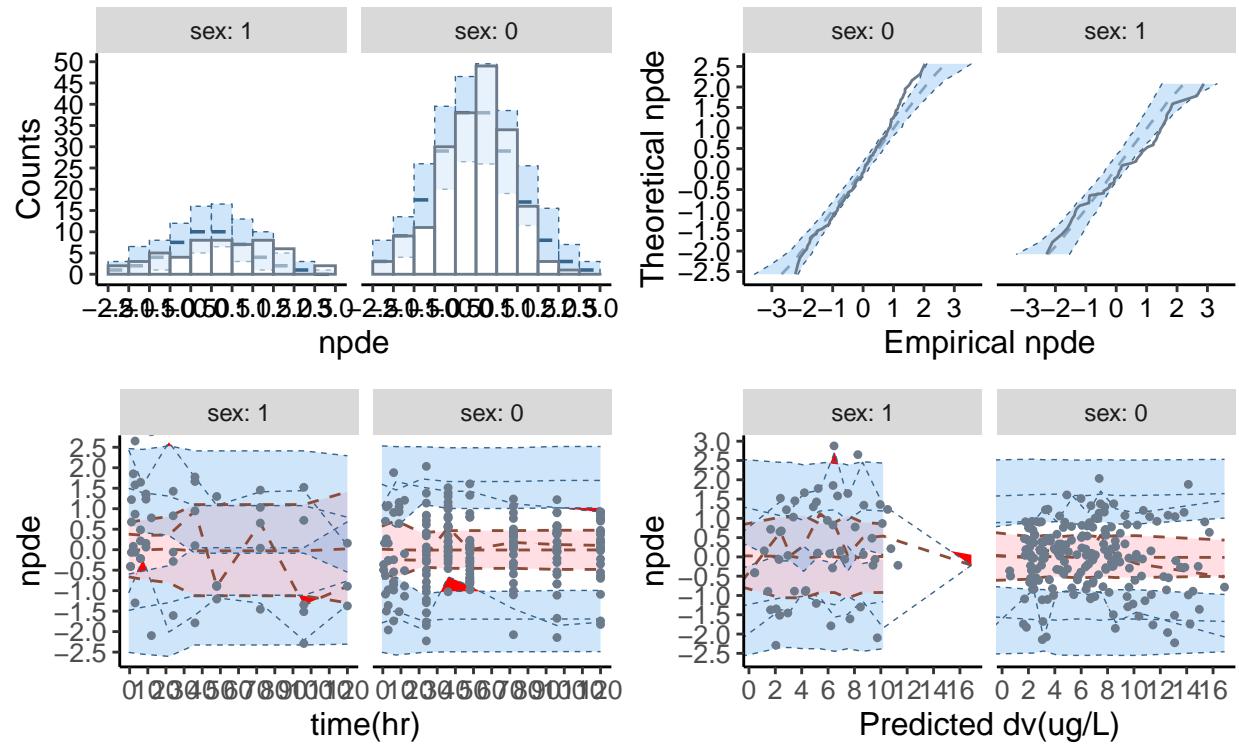
waffle plot : Wt



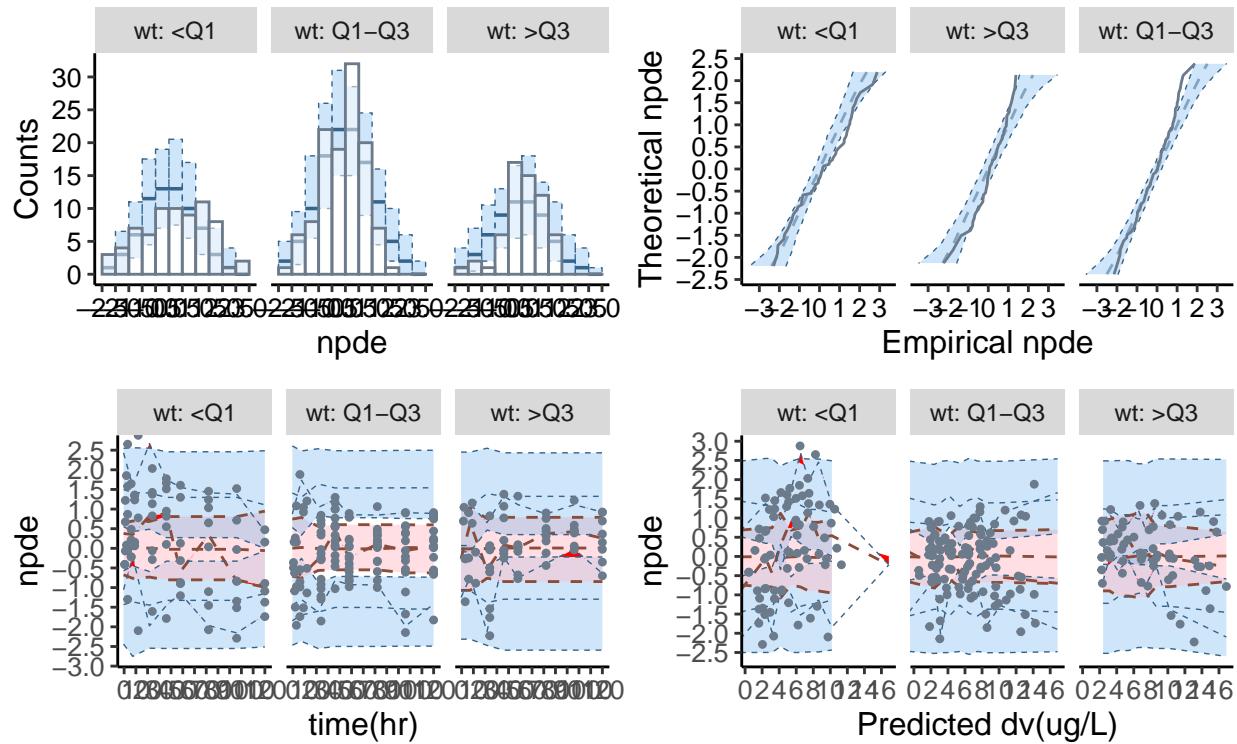
waffle plot : amt



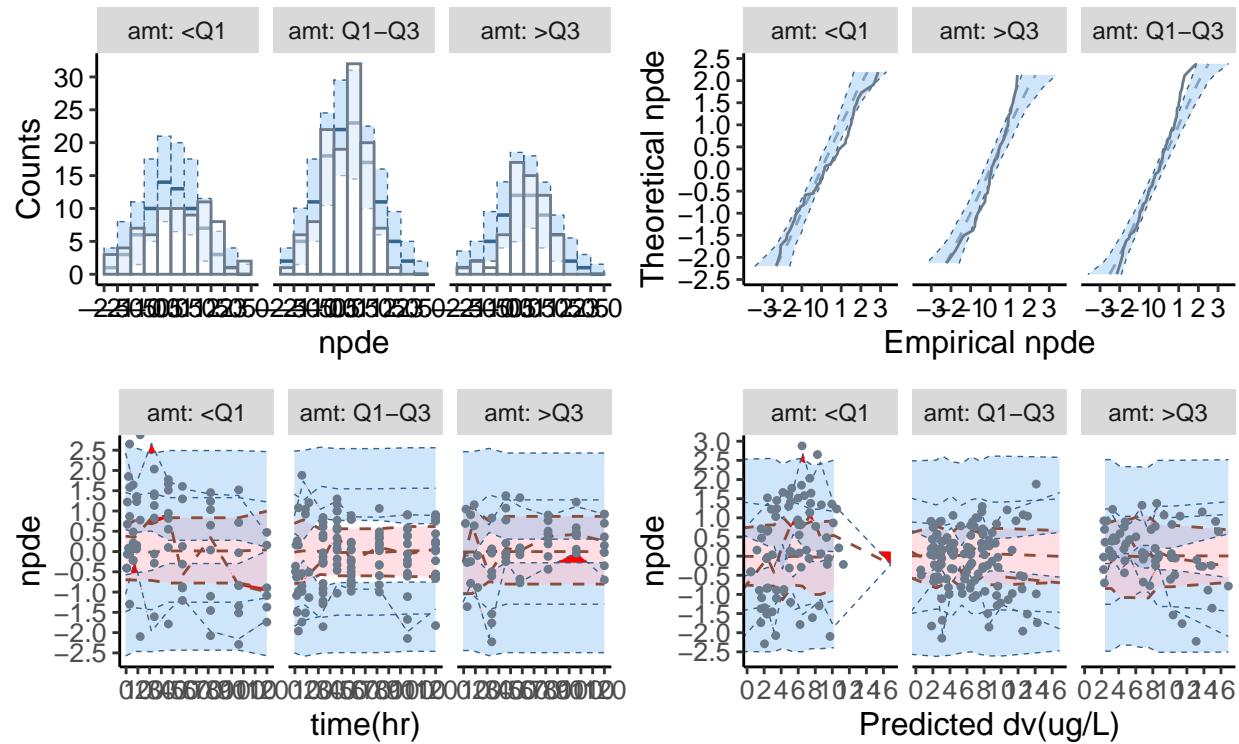
waffle plot : Sex



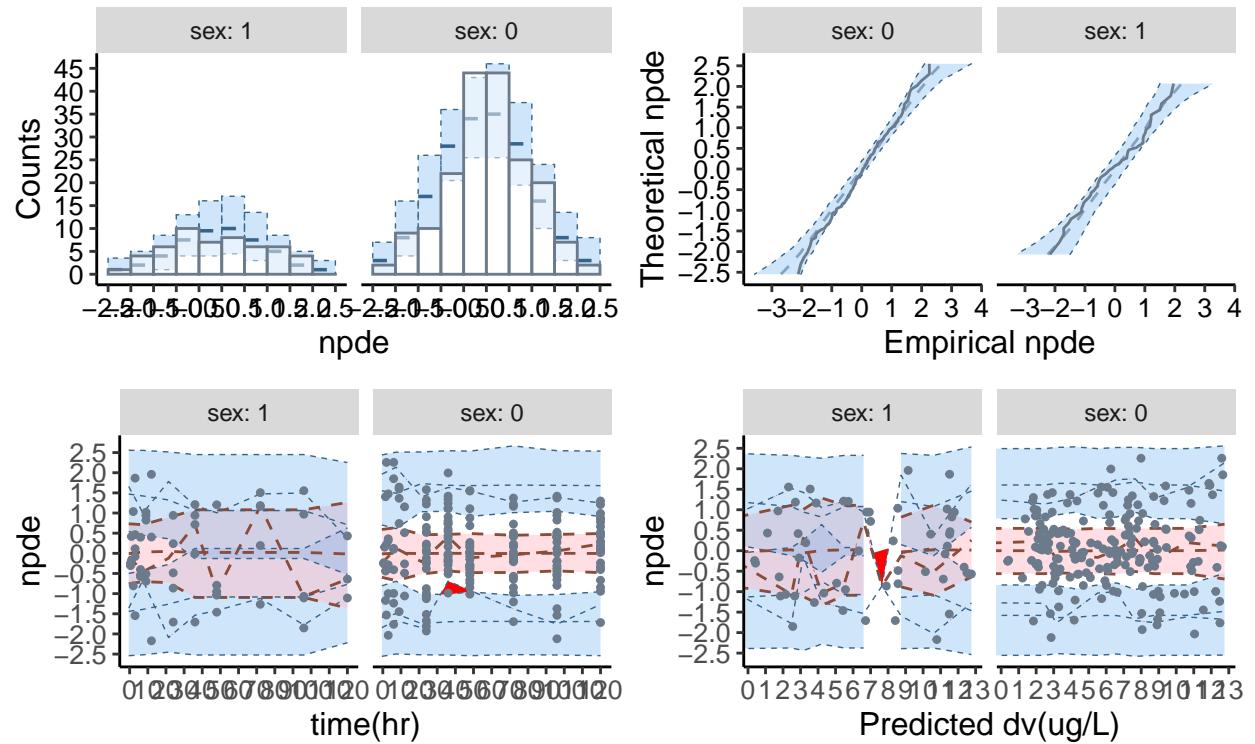
waffle plot : Wt



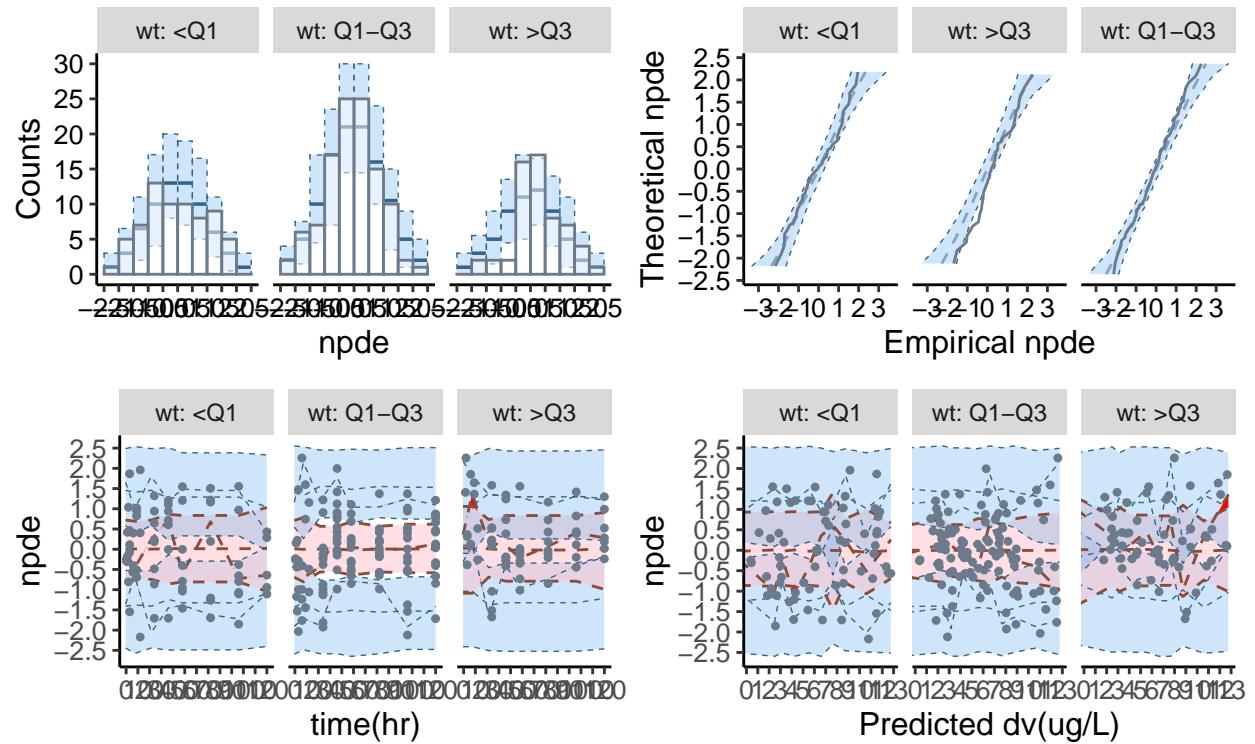
waffle plot : amt



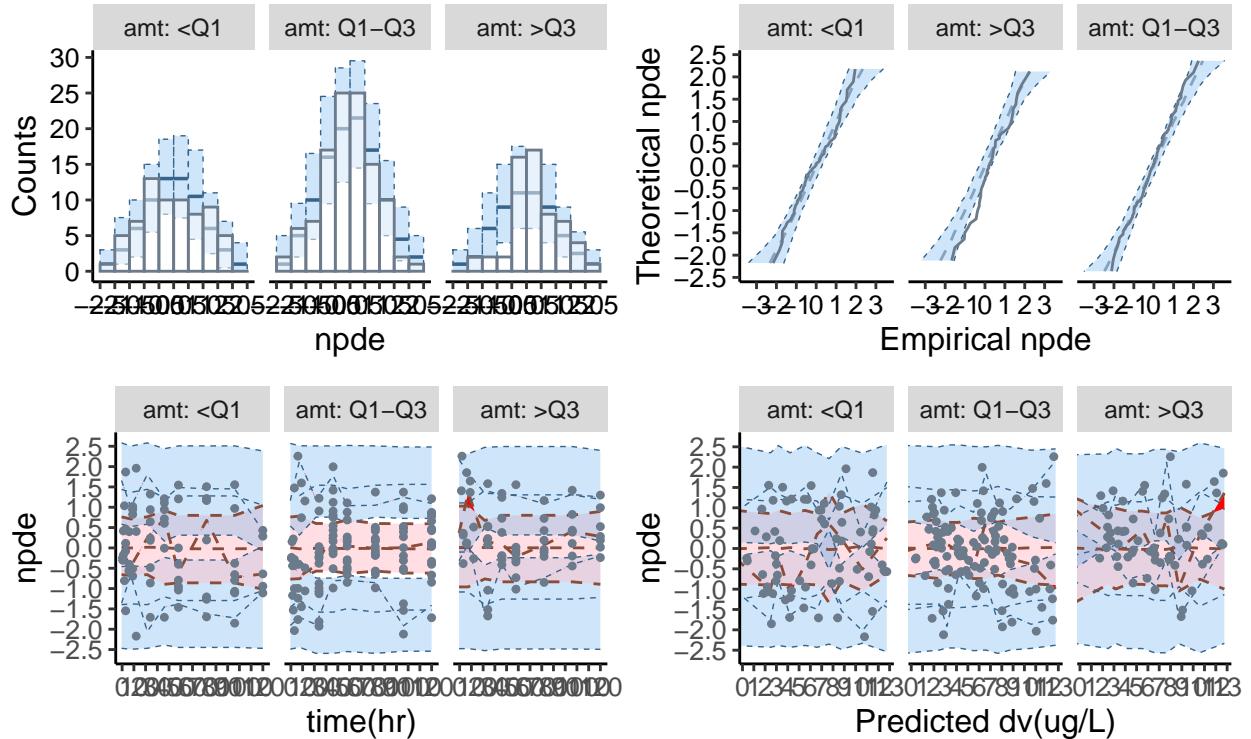
waffle plot : Sex



waffle plot : Wt

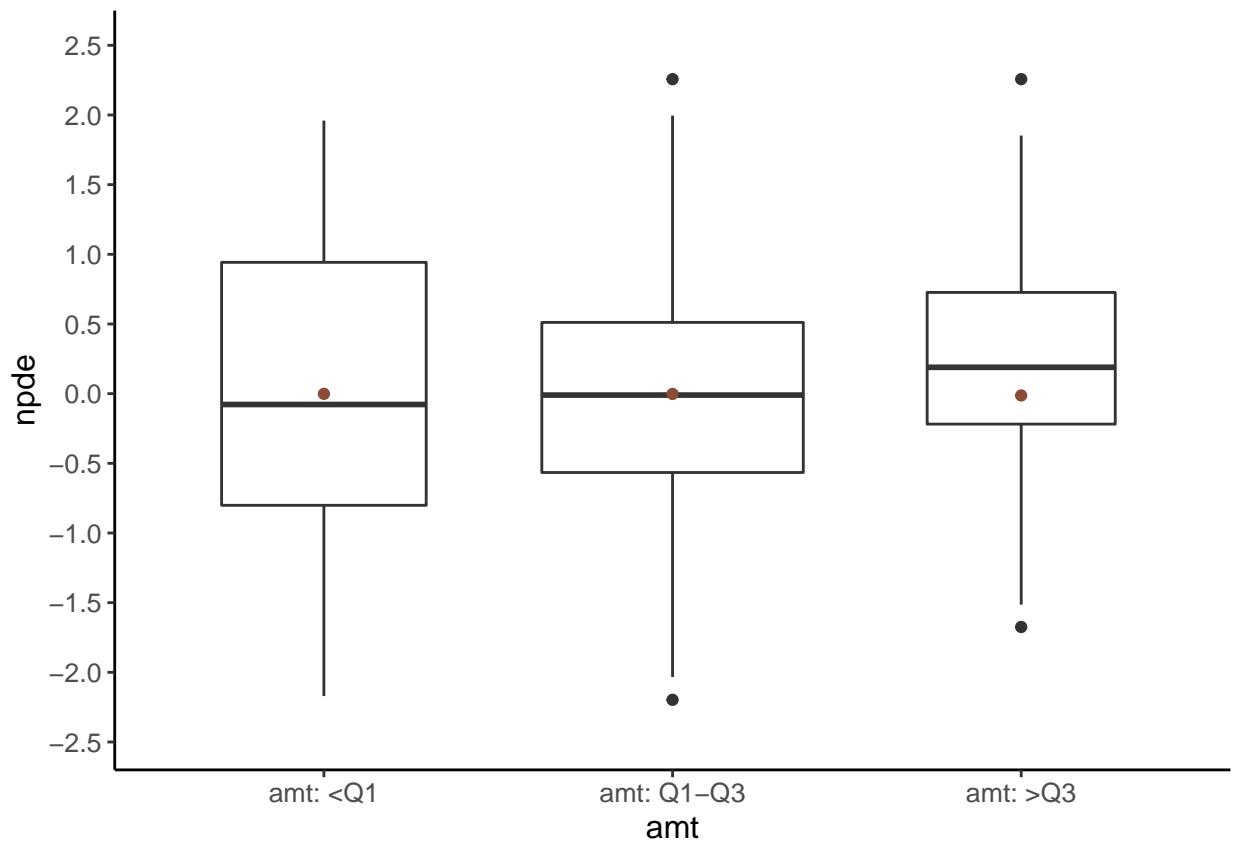


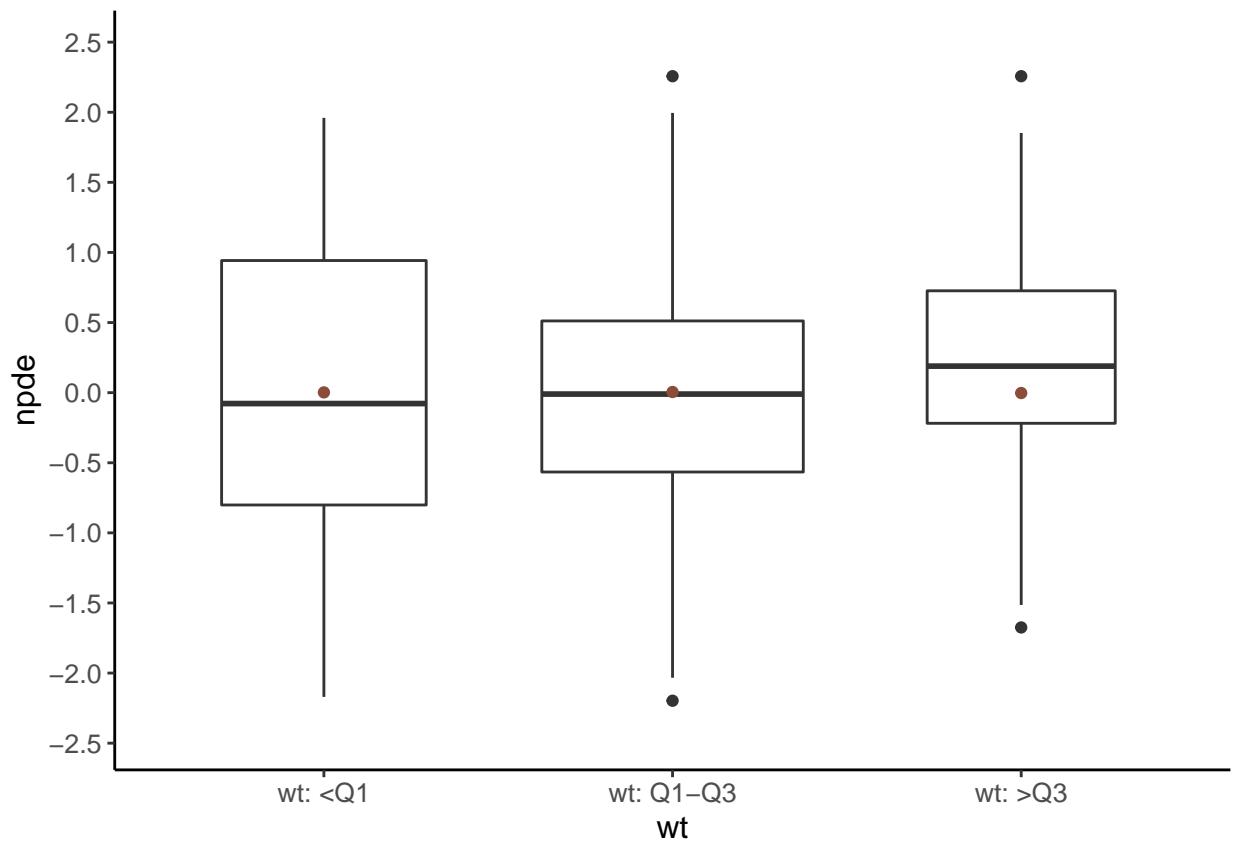
waffle plot : amt

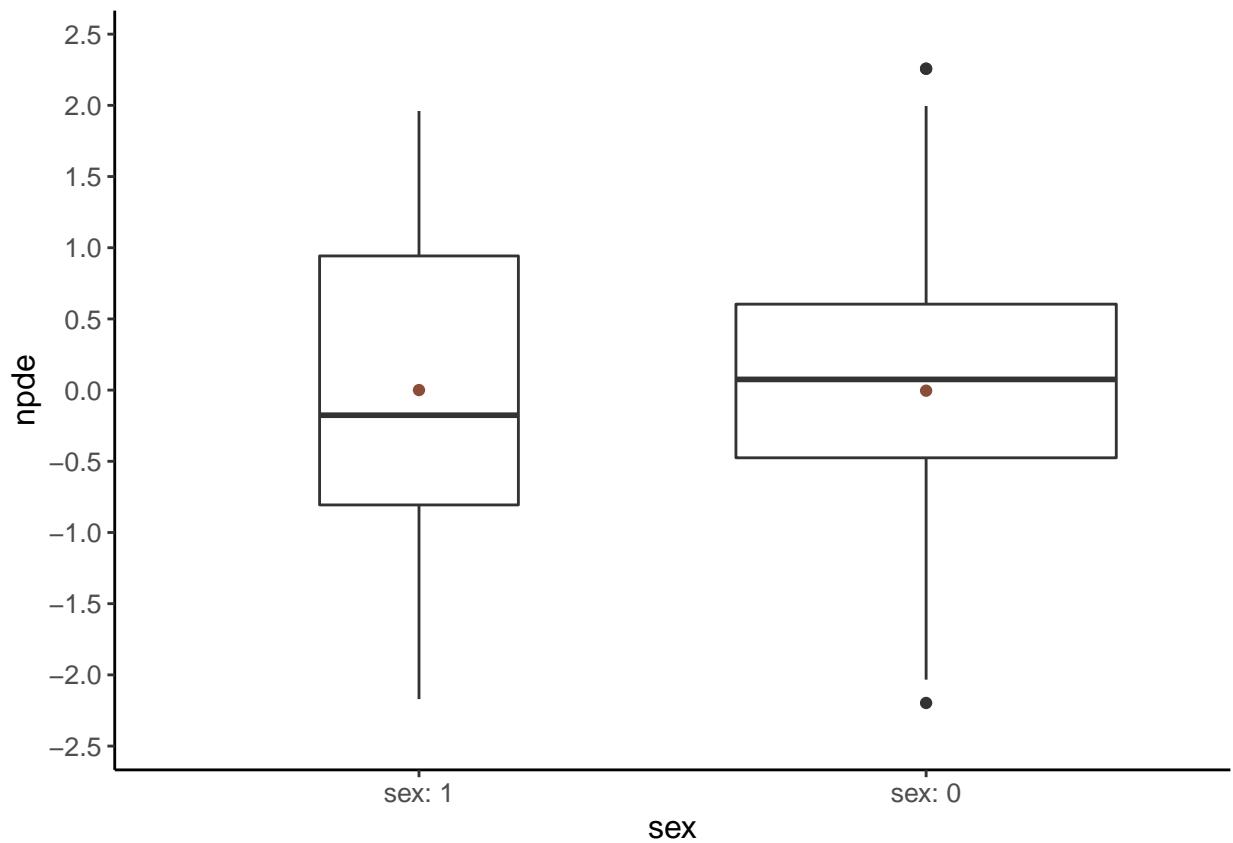


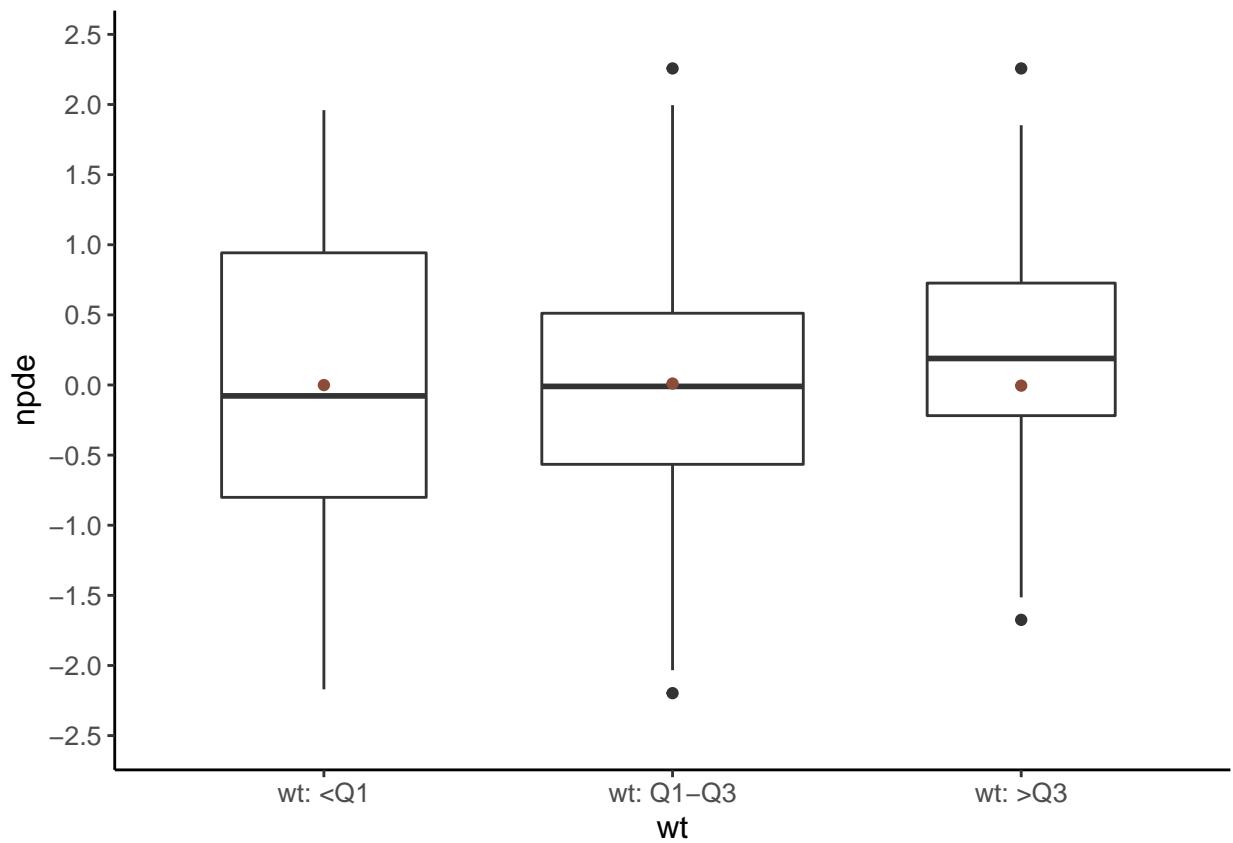
Parameters versus covariates

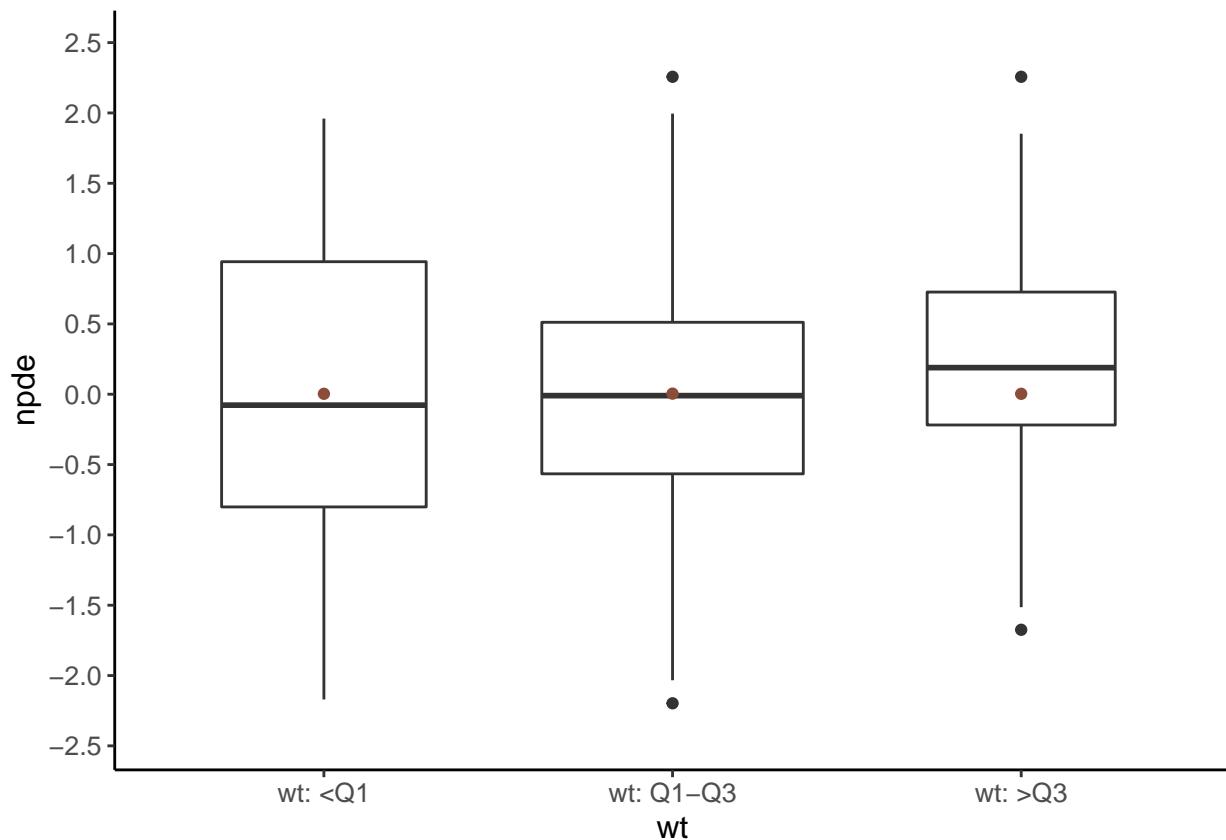
- **Problems (à résoudre Romain)**
 - solved yes here we really need to extend PI to cover the whole X-range for x.scatter and pred.scatter (not VPC)
 - * solved just the PI (not the observed percentiles or outlier bands)
 - solved cov.scatter (eg npde versus covariates)
 - * boxplot for categorical covariates (overlaid on PI)
 - * here two different behaviours ?



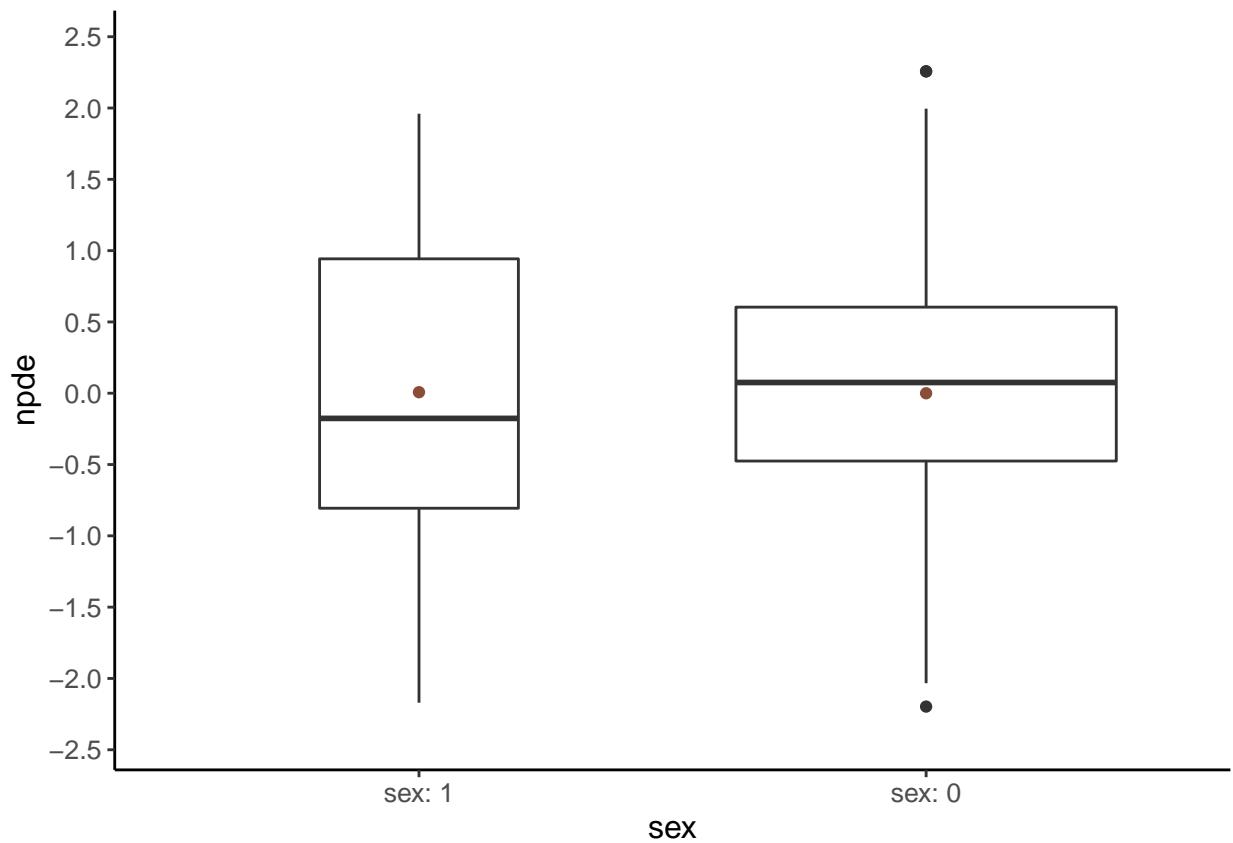


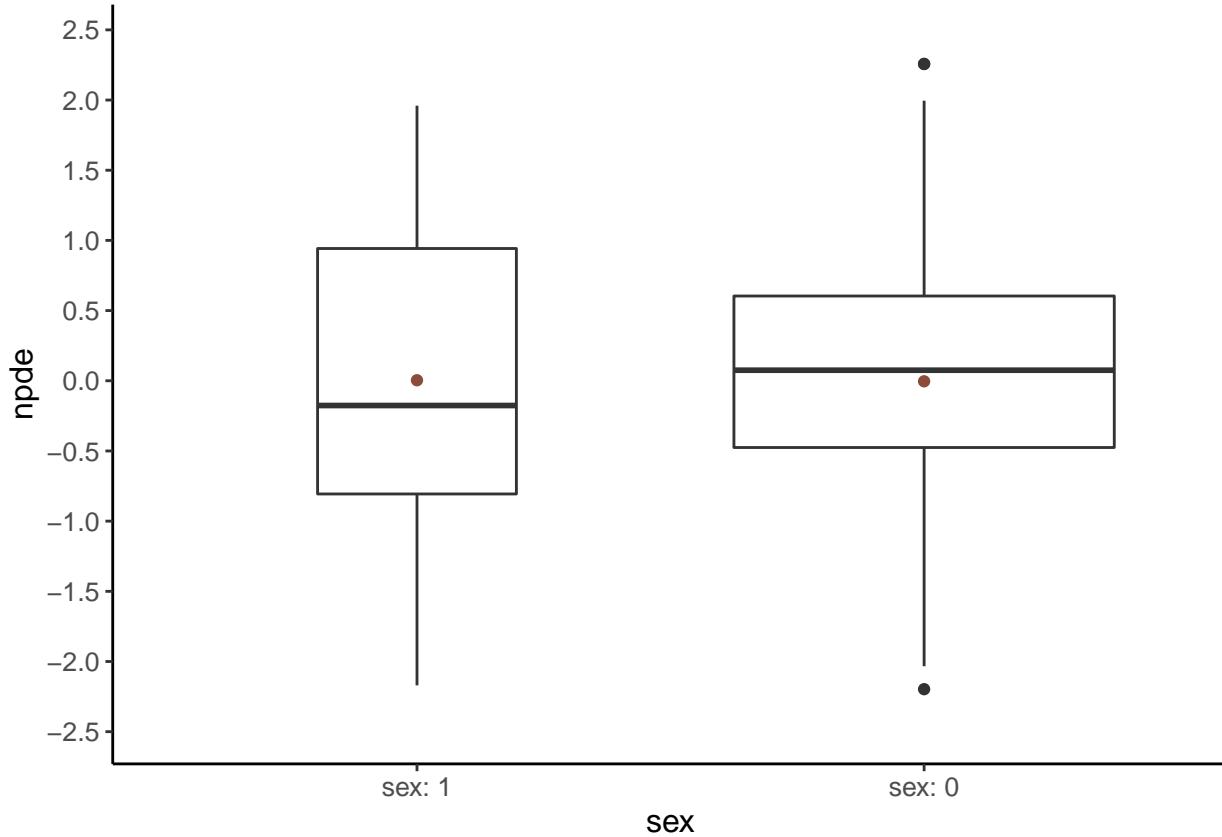






```
## Test failed, not the expected behaviour
```





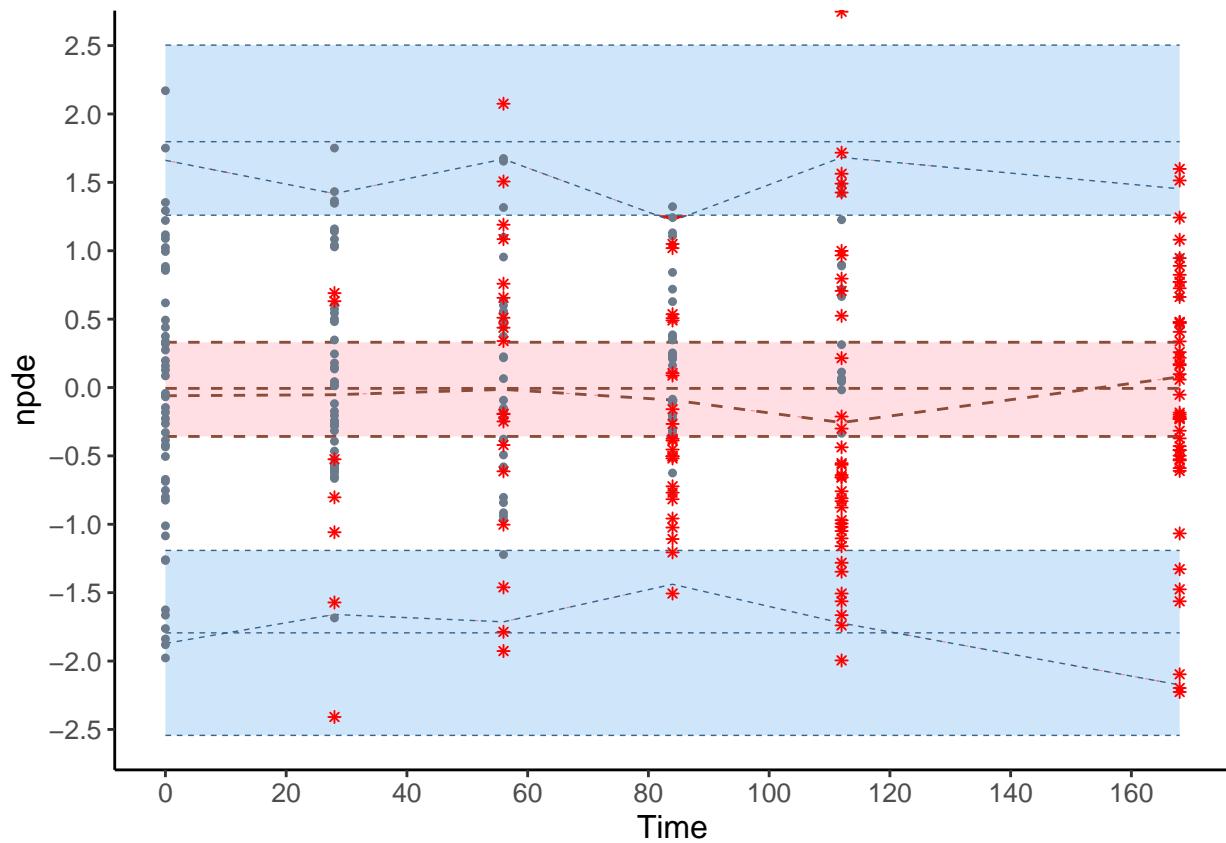
Creating the plots for the user guide

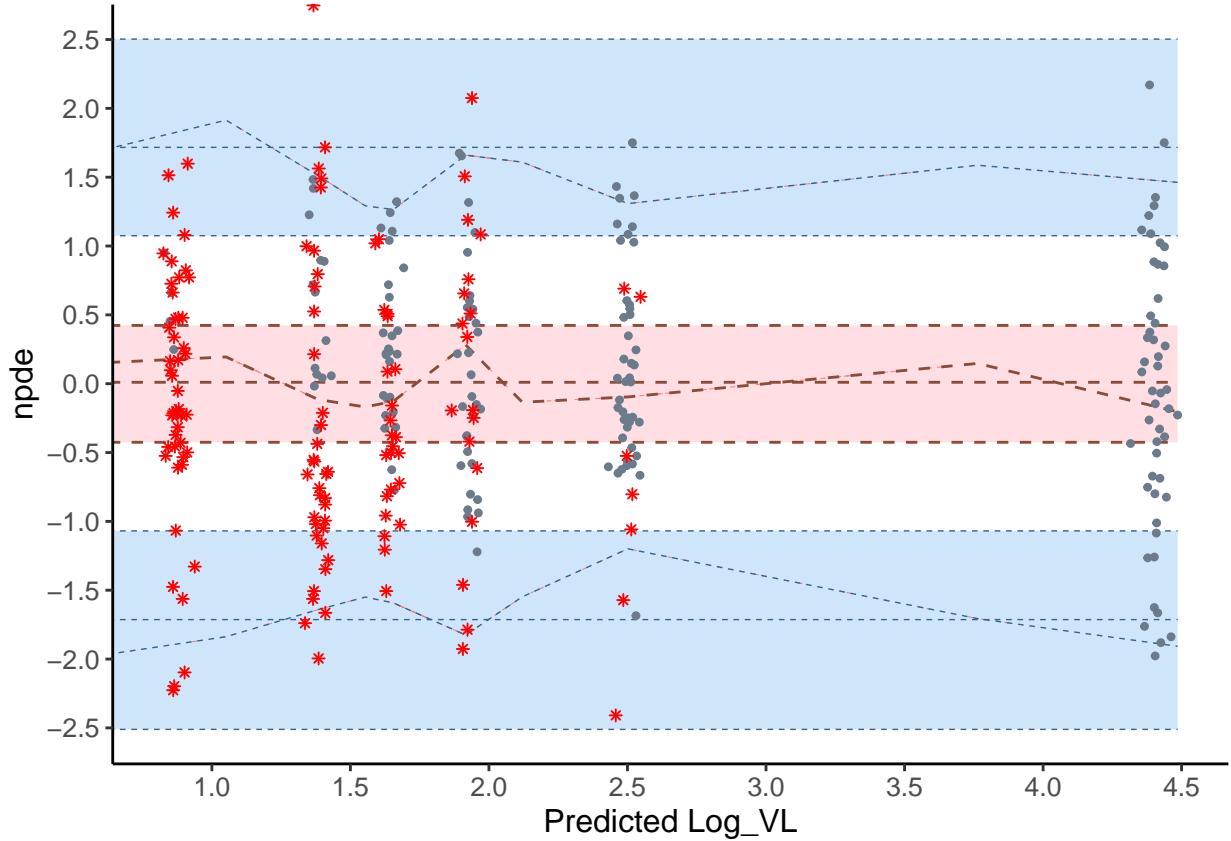
Theophylline

- Default plots and VPC: **OK pour doc**
- Problem with plot of data (see above)

Viral load data

- **TODO**
 - plot data
 - est-ce que les fonctions graphiques ‘de base’ pourraient renvoyer les graphes comme objet pour pouvoir ensuite en faire un *grid.arrange()* (on avait parlé de cette option pour pouvoir faire ensuite ses propres graphes comme avant) ?
 - **note**
 - Maintenant les fonctions graphiques renvoient `objet=NULL` pour les plots, avant non. Peut-être faire remonter le return des objets ggplots jusque dans la fonction “plot” générale?
 - * 2 graphes de données (voir 3)
 - * 2 graphes de VPC côté à côté (\Rightarrow donner code correspondant dans la doc)
 - * **note** si c'est l'option `plot.default` il faudrait en changer le nom (pas du tout explicite), en un truc comme `return` ou `return.plot...`
 - corriger le graphe de VPC





```
## Error in mapply(child_vp, vp_name = vpname(x$layout), t = .subset2(x$layout, :
##   des entrées de longueur nulle ne peuvent être mélangés avec celles de longueur non nulle
```

End of file, deactivating development mode

```
## Dev mode: OFF
```