# Code merge - Distribution plots

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# **Summary**

#### Objective

Recode aux.plot.hist and aux.plot.dist to simplify their structure.

#### Current issue

Sorting the covariates in the facet plot

### Requirements

- main function npde.plot.dist()
  - input: npdeObject, which (variable to plot), dist.type (plot type, one of hist, qqplot, ecdf), "..."
  - output: one plot
- auxiliary functions aux.npdeplot.hist() and aux.npdeplot.dist()
  - input: both functions should take as input
    - \* a dataframe obsmat: matrix with the data to plot, with columns
      - · x; variable to plot
      - · category: covariate category (if "all", overall plot)
    - \* graphical options: plot.opt
    - \* distrib: reference distribution plot (one of norm, unif)
    - \* sim.ypl: if given, a vector of simulated data for the variable to plot
    - \* for aux.npdeplot.hist(), nclass (nb of classes)
  - output
    - \* distribution plots: histogram (aux.plot.hist), ecdf or qqplot (aux.plot.dist)
    - \* note: QQ-plot for uniform variable is similar to ecdf (theoretical quantiles correspond to the cumulative density function)

#### Defining folders, loading libraries

Loading functions

Run npde

Prepare objects

# Debug auxiliary plots

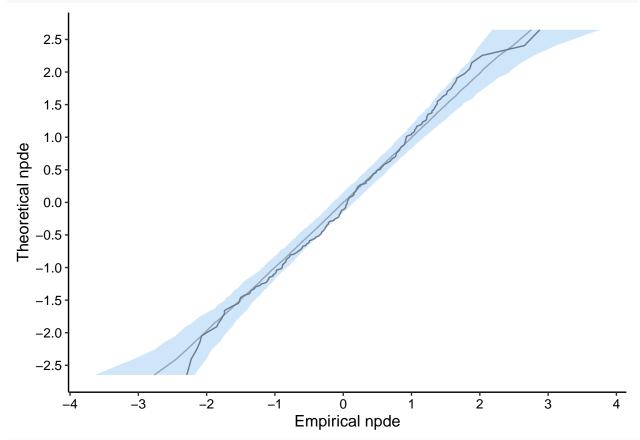
#### Histogram

- 3 plots: no covariate, sex in 2 categories, weight in 3 categories
- problem with sorting the categories

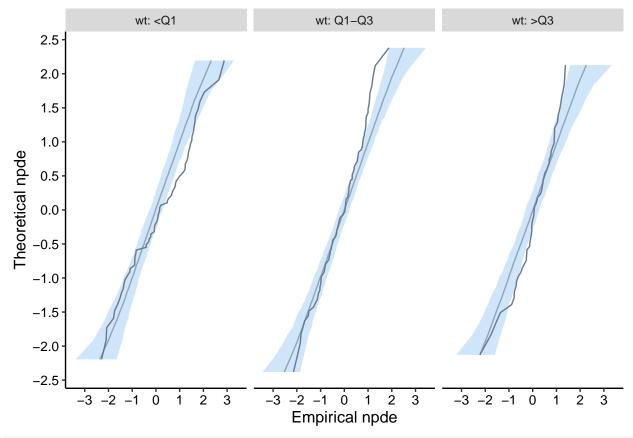
Distribution plot: ecdf Distribution plot: qqplot

# Debug main distribution plot function

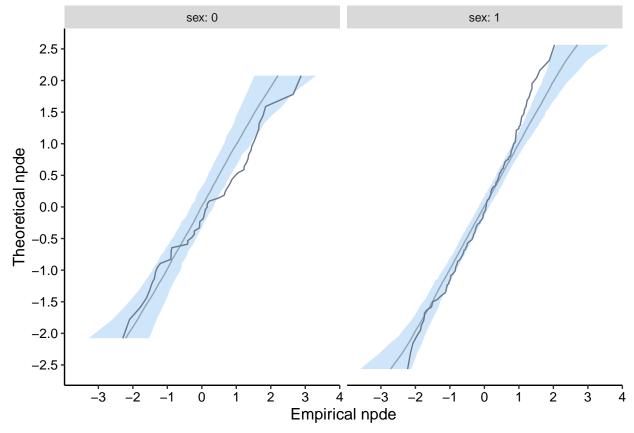
source(file.path(workDir, "npde", "R", "plotNpde-distributionPlot.R")) # renamed from plotNpde-unitFunctio
npde.plot.dist(wbase, which="npde", dist.type="qqplot")



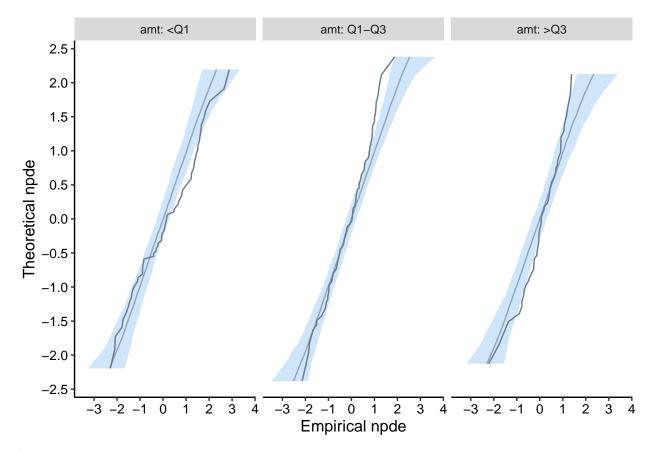
npde.plot.dist(wbase, which="npde", dist.type="qqplot", covsplit=TRUE, which.cov=c("wt"))



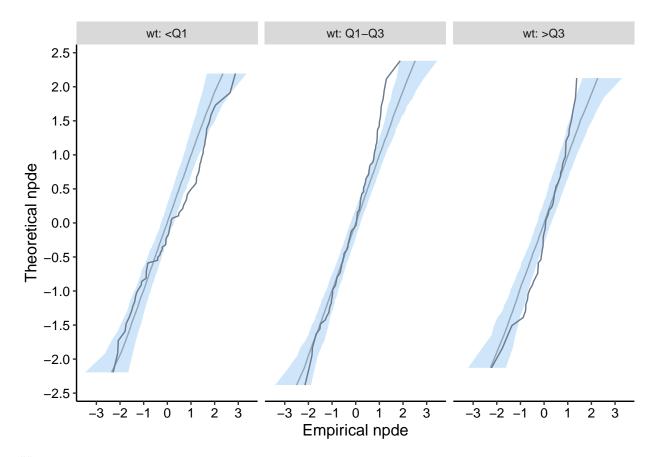
npde.plot.dist(wbase, which="npde", dist.type="qqplot", covsplit=TRUE, which.cov=c("sex"))



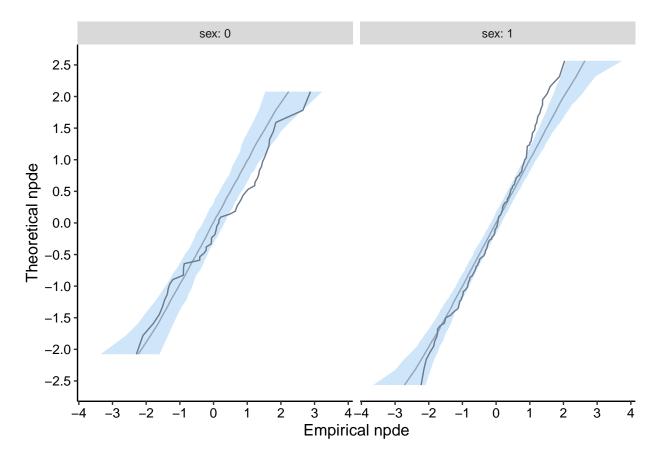
npde.plot.dist(wbase, which="npde", dist.type="qqplot", covsplit=TRUE, which.cov=c("all"))
## [[1]]



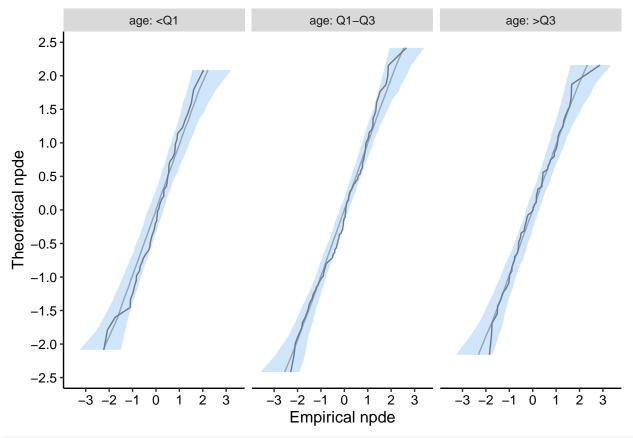
## ## [[2]]



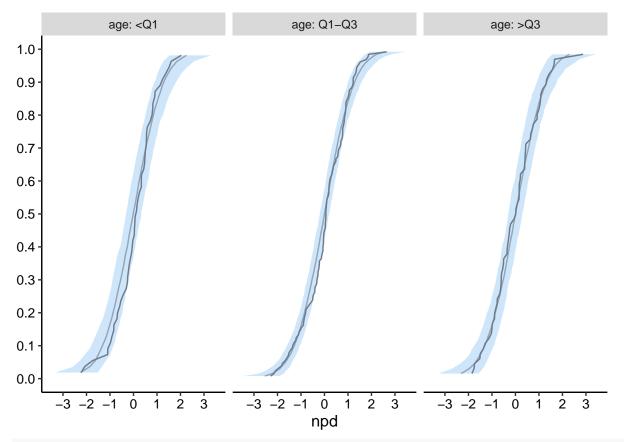
## ## [[3]]



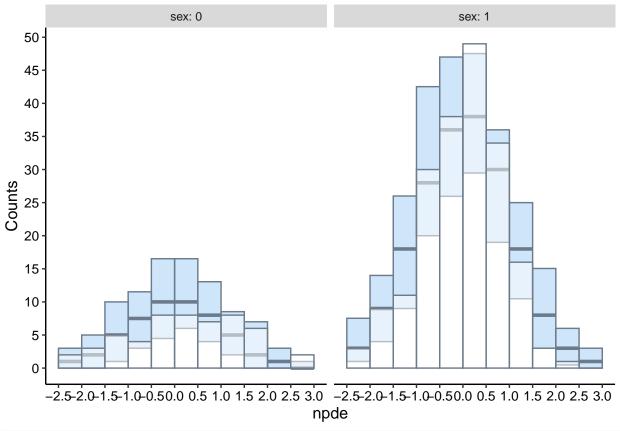
## ## [[4]]



npde.plot.dist(wbase, which="npde", dist.type="ecdf", covsplit=TRUE, which.cov=c("age"))



npde.plot.dist(wbase, which="npde", dist.type="hist", covsplit=TRUE, which.cov=c("sex"))



```
# debut
if(FALSE) {
    npdeObj<-wbase
    plot.opt<-npdeObjectOprefs
    which<-"npde"
    plot.opt$covsplit<-TRUE
    plot.opt$which.cov<-c("wt")
    dist.type="ecdf"
}</pre>
```