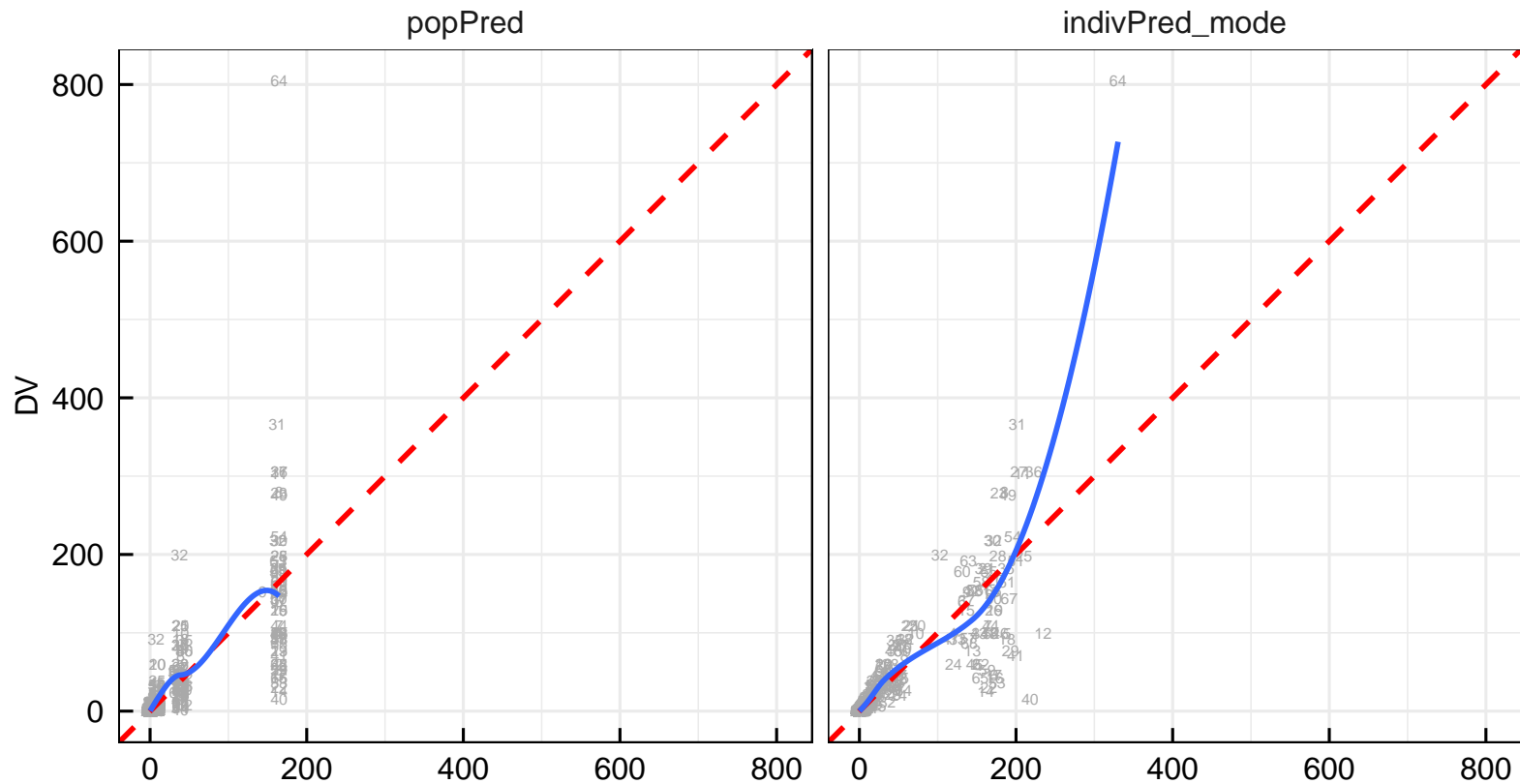
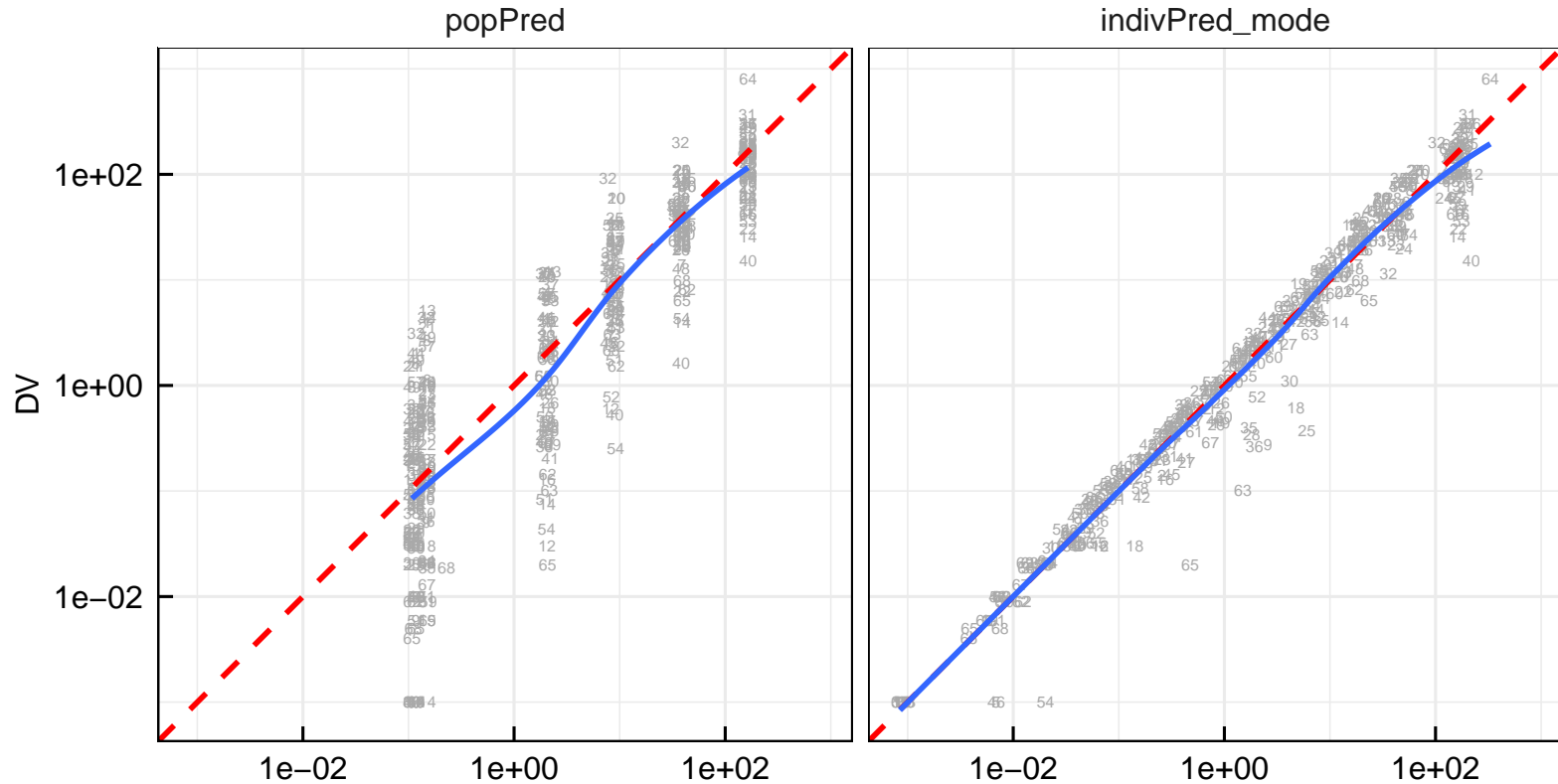


Output Nr. 1: BCRABL [prct]
Observation vs. Population and Individual Predictions
Linear Axes



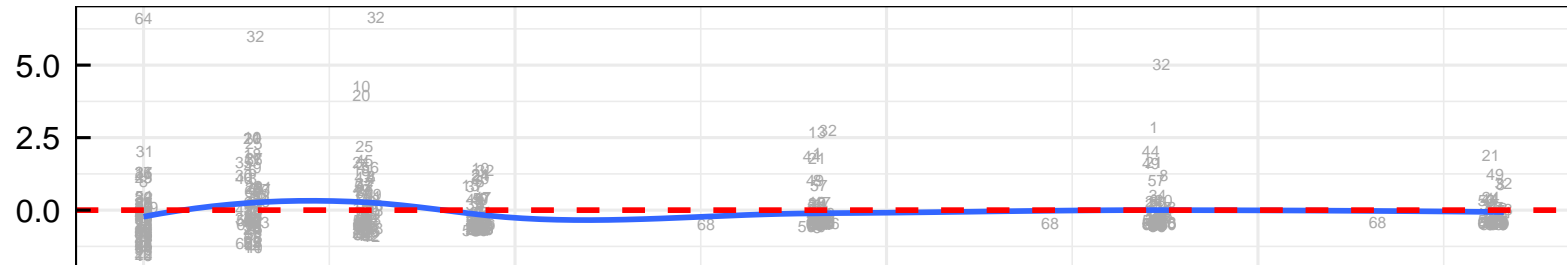
Output Nr. 1: BCRABL [prct]
Observation vs. Population and Individual Predictions
Log Axes (zero values plotted on axis)



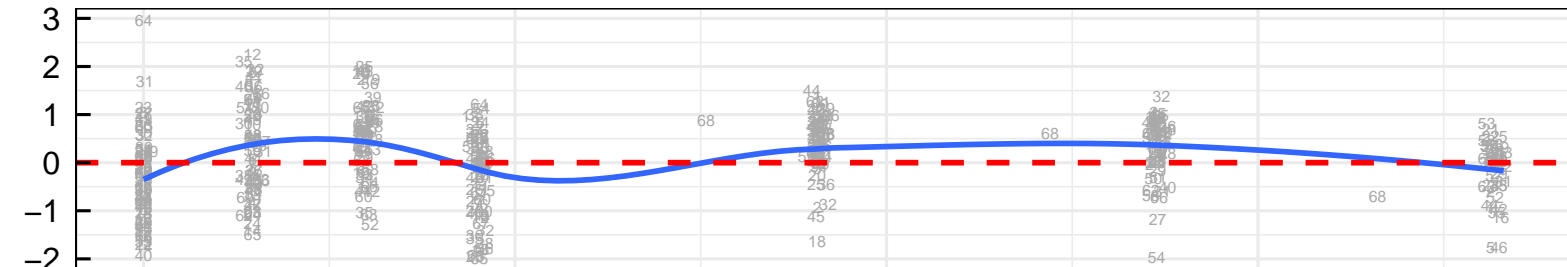
Output Nr. 1: BCRABL [prct]

Weighted Individual and Population Residuals and NPDE vs. Time

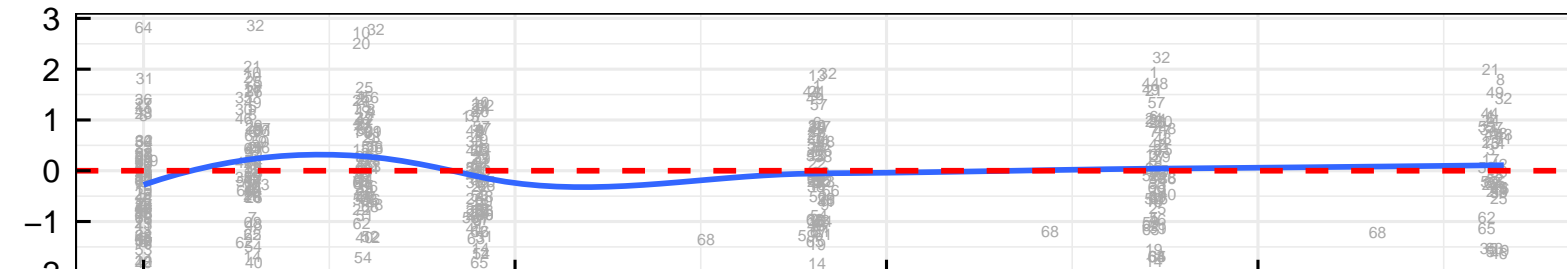
pwRes



indWRes_mode



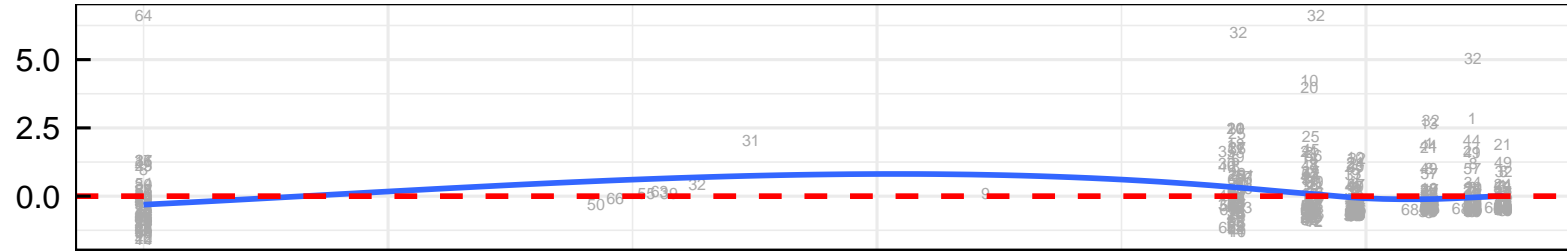
NPDE



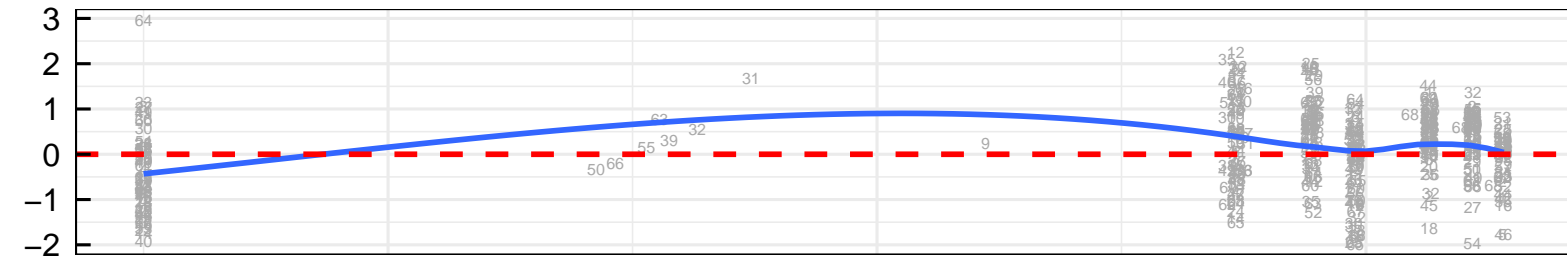
Time [MONTHS]

Output Nr. 1: BCRABL [prct]
 Weighted Individual and Population Residuals and NPDE vs. Time
 Log TIME axis (values for time = 0 plotted on axis)

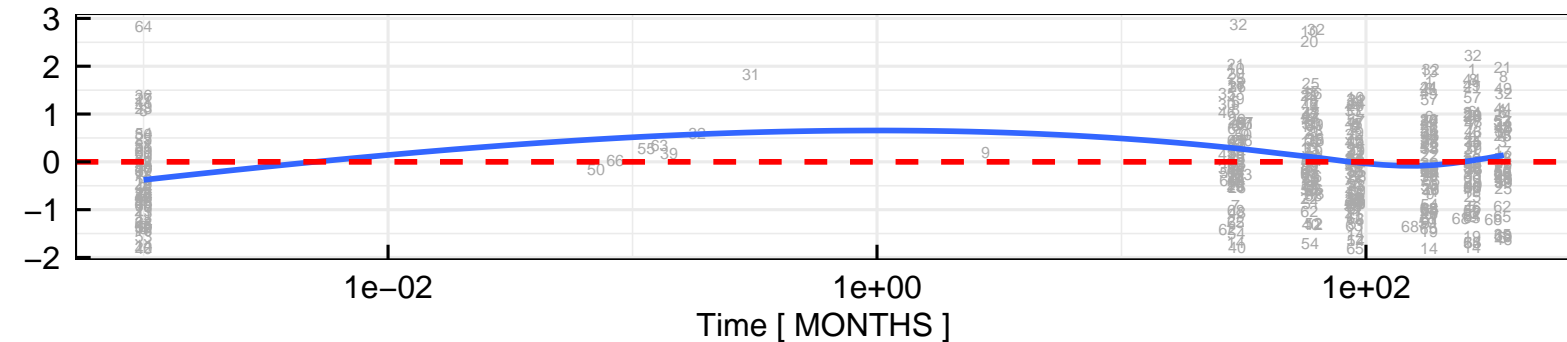
pwRes



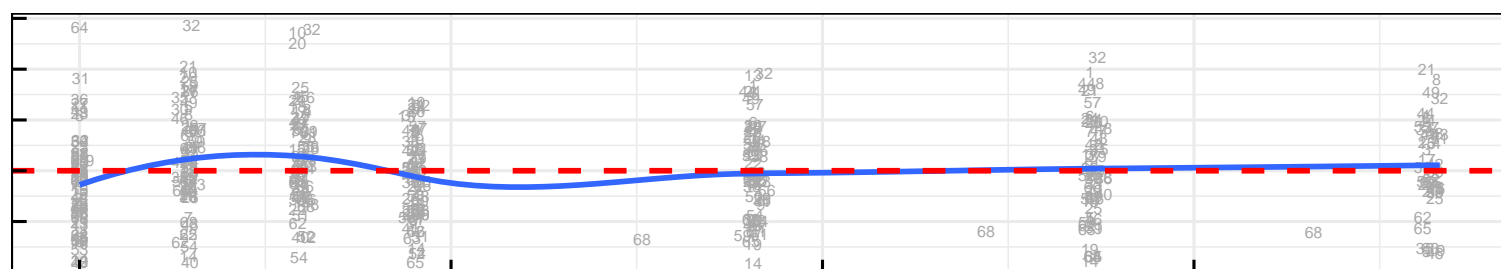
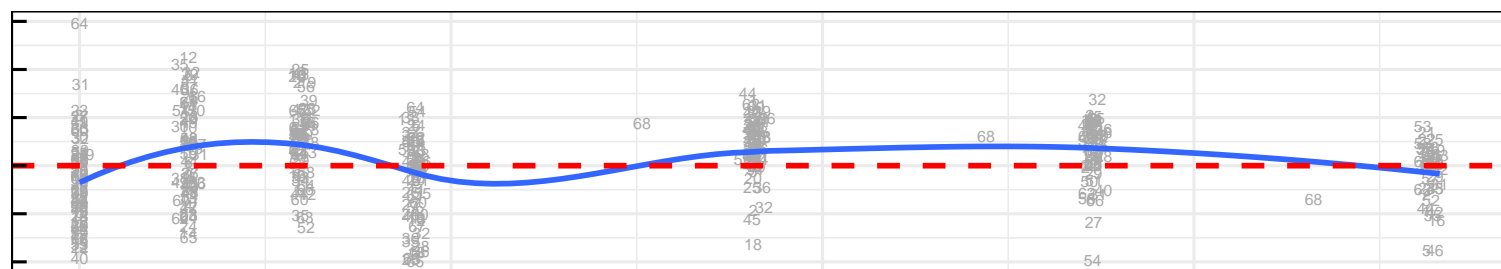
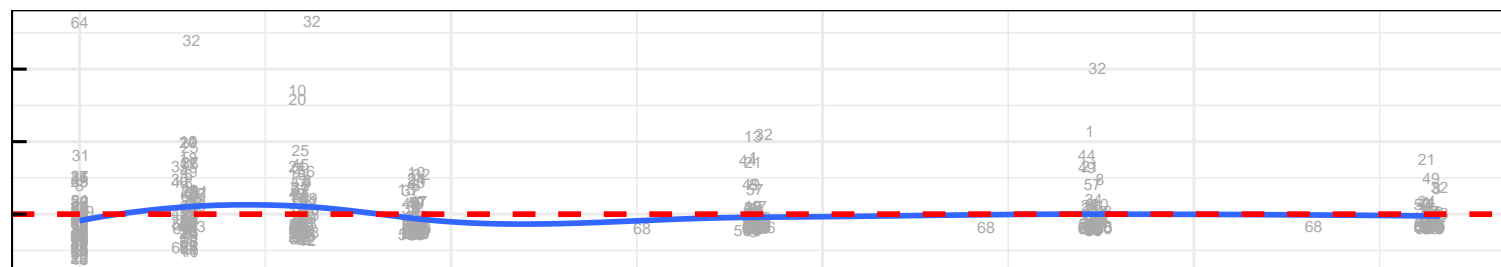
indWRes_mode



NPDE

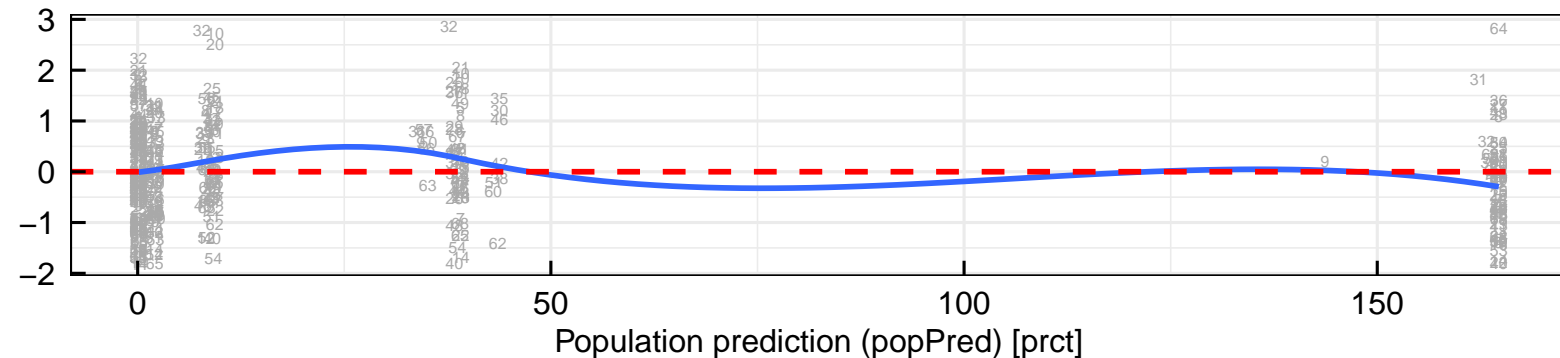
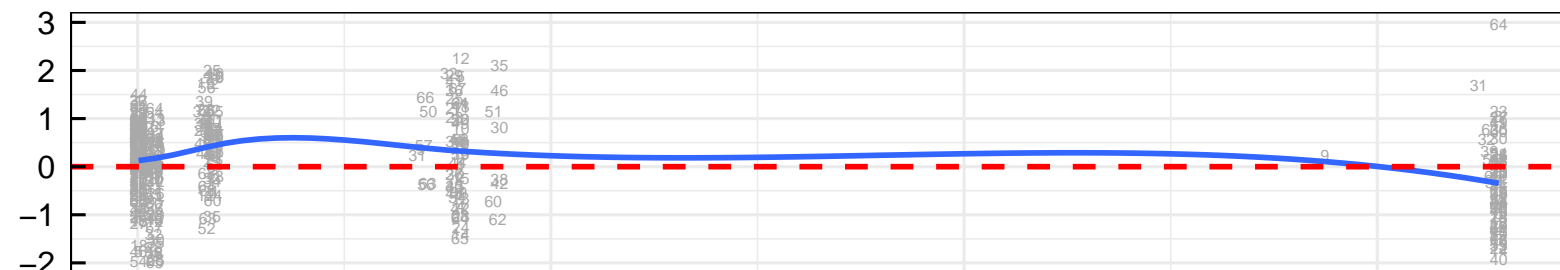
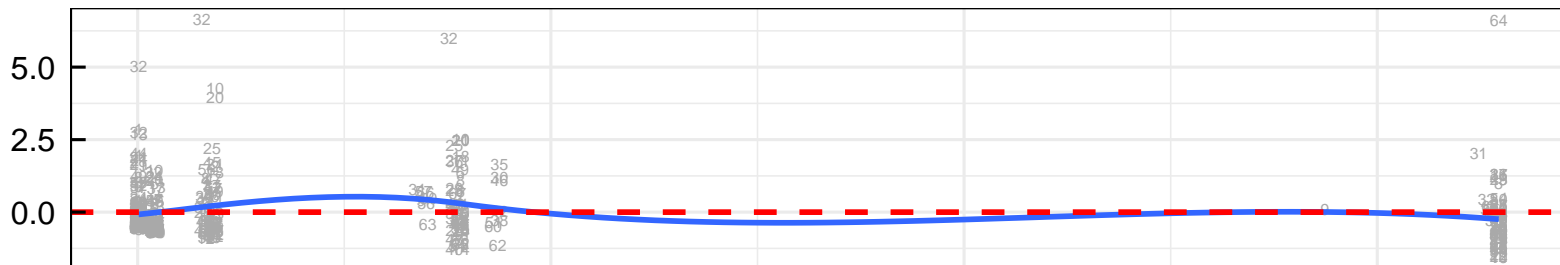


Weighted Individual and Population Residuals and NPDE vs. TAD



Time after previous dose (TAD) [MONTHS]

Weighted Individual and Population Residuals and NPDE vs. Population Prediction

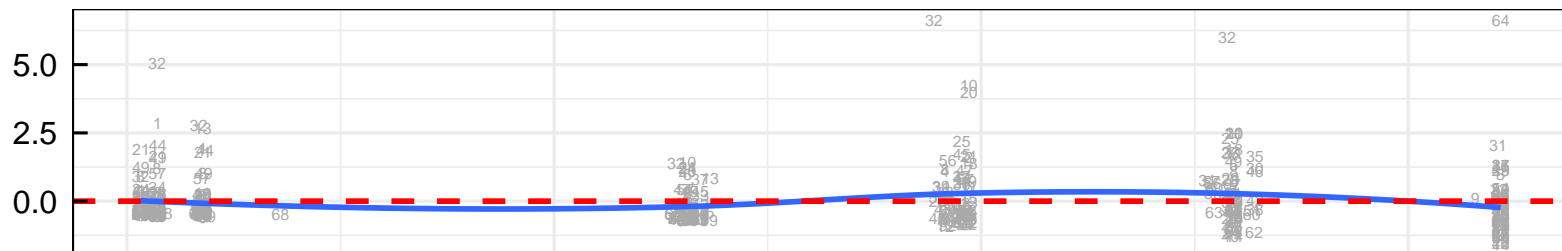


Output Nr. 1: BCRABL [prct]

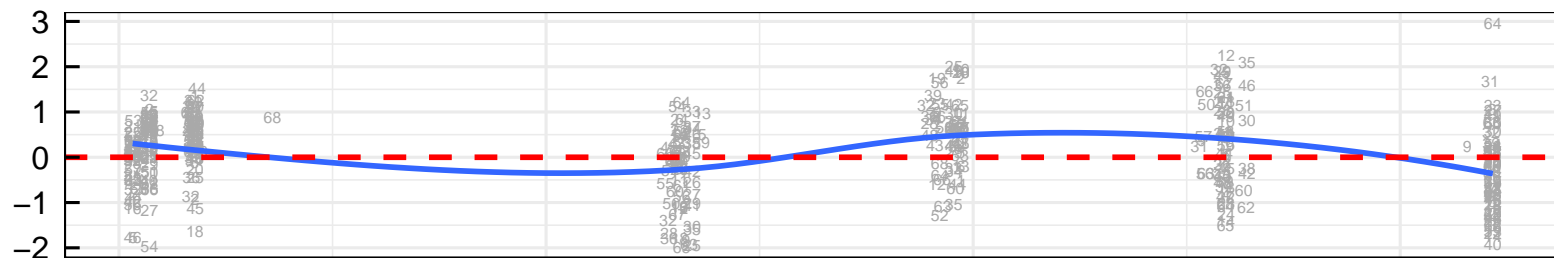
Weighted Individual and Population Residuals and NPDE vs. Population Prediction

Log Prediction axis (values for prediction = 0 plotted on axis)

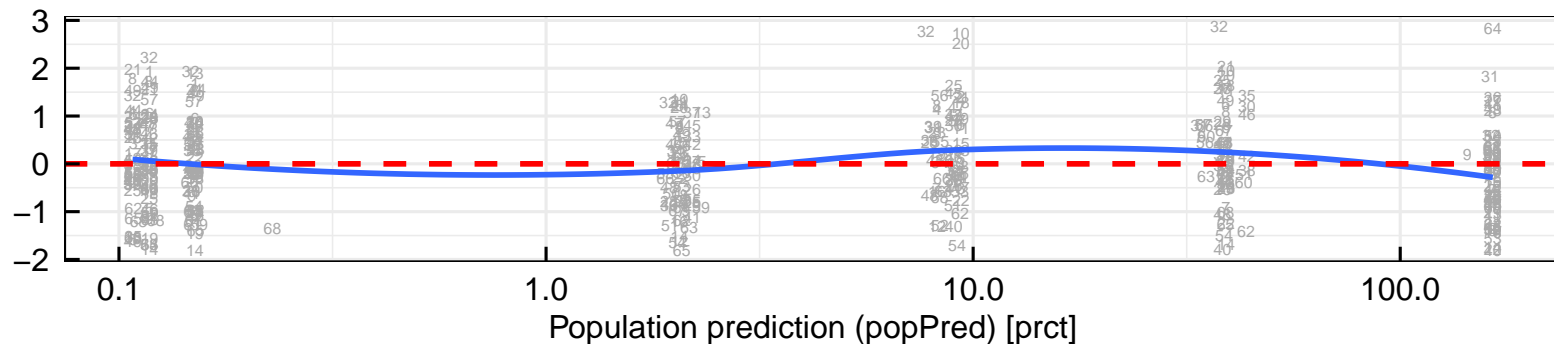
pwRes



indWRes_mode



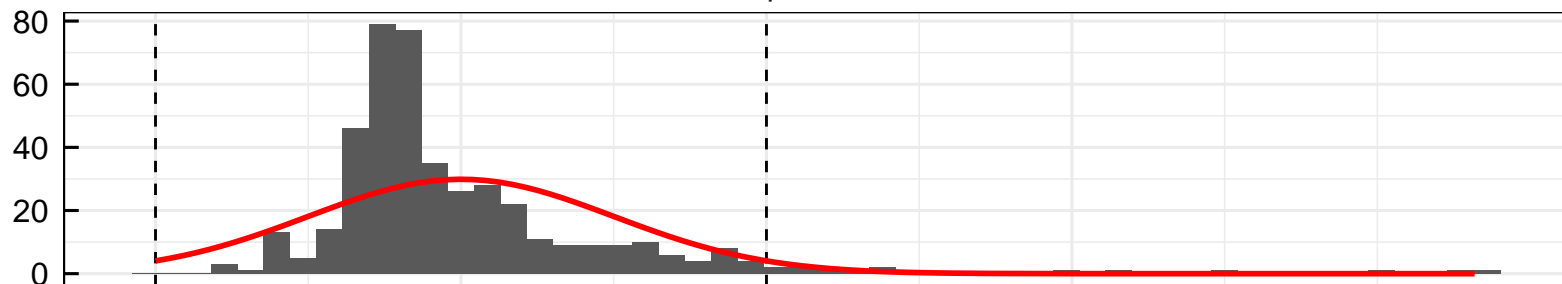
NPDE



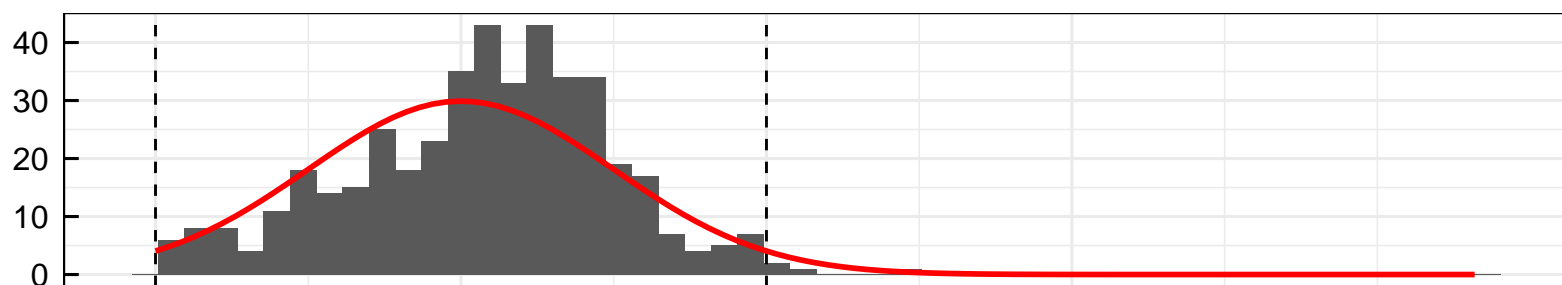
Output Nr. 1: BCRABL [prct]

Histograms of Residuals compared to Standard Normal

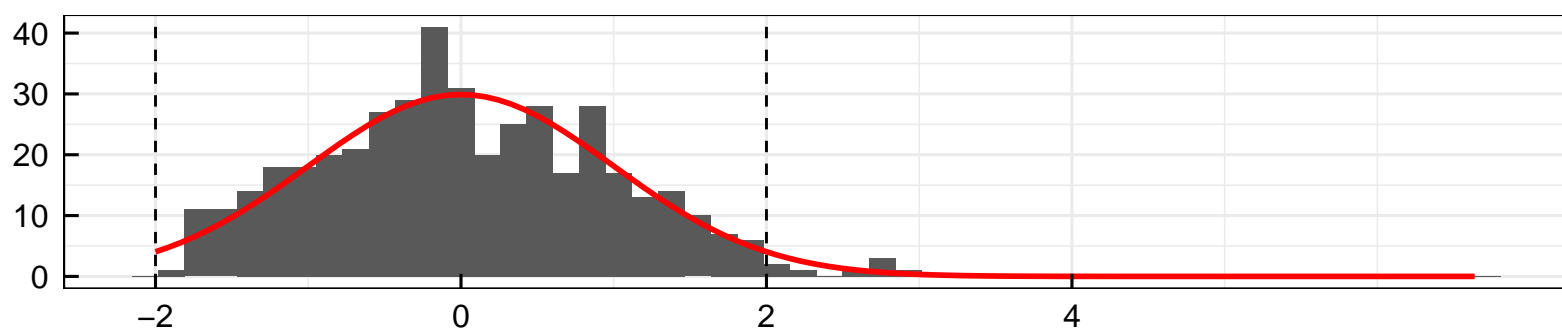
pwRes



indWRes_mode

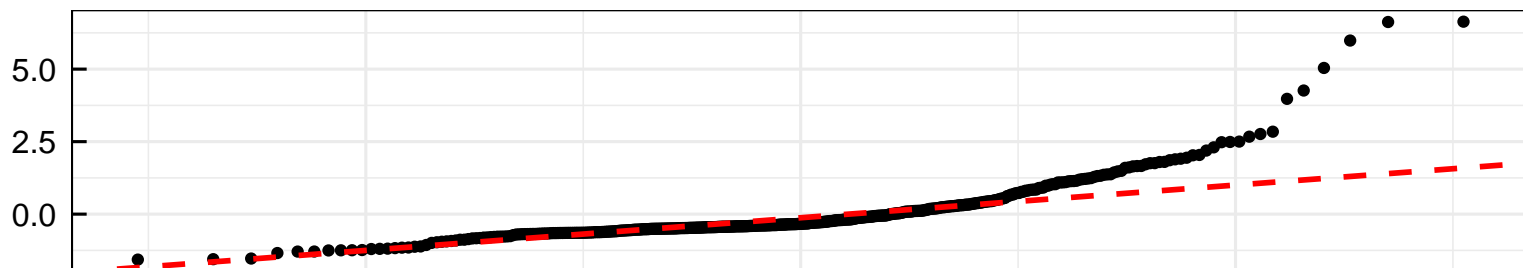


NPDE

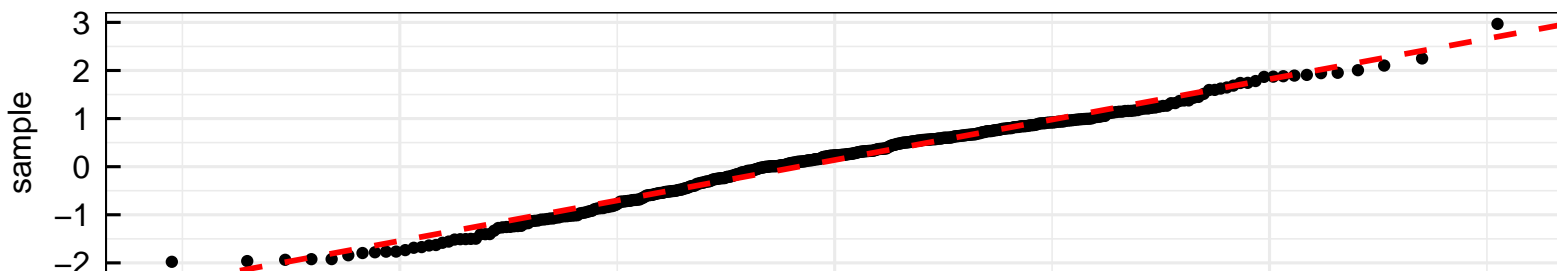


Output Nr. 1: BCRABL [prct]
QQ-Plots of Residuals

pwRes



indWRes_mode



NPDE

