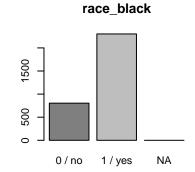
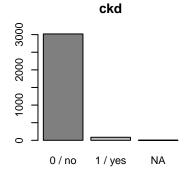


3106 records; 1936 '0', 1170 '1', 0 'NA'

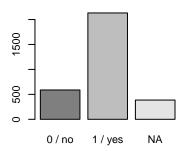


3106 records; 804 '0', 2302 '1', 0 'NA'



3106 records; 3019 '0', 86 '1', 1 'NA'

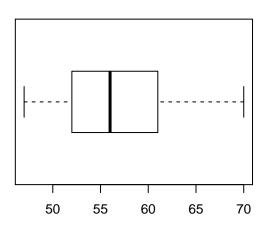
microalbuminuria



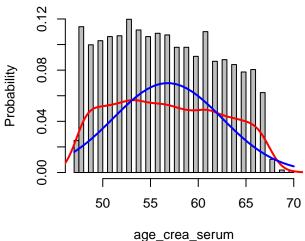
3106 records; 588 '0', 2132 '1', 386 'NA'

age_crea_serum

Box plot



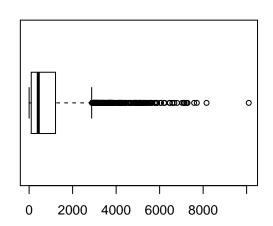
min: 47, q1: 52, med: 56, mean: 56.79, q3: 61, max: 70, na: 0



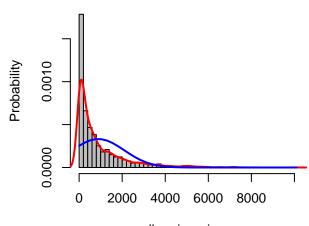
3106 non-missing records (0 NA)

albumin_urinary

Box plot



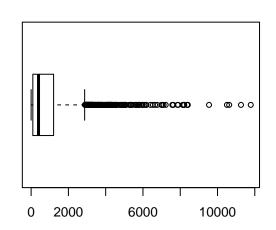
min: 0.000147, q1: 97.38, med: 420.3, mean: 888.9, q3: 1214, max: 10100, na: 0



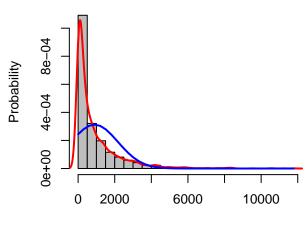
albumin_urinary 3106 non-missing records (0 NA)

creatinine_urinary

Box plot



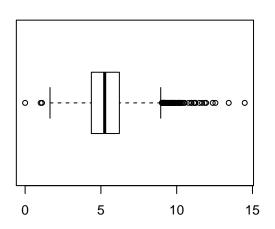
min: 7.028e-05, q1: 92.61, med: 397, mean: 899.2, q3: 1206, max: 11780, na: 0



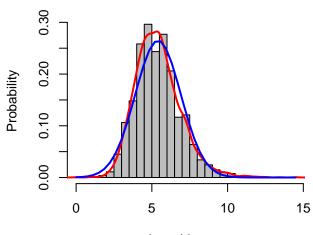
creatinine_urinary 3106 non-missing records (0 NA)

uric_acid_serum

Box plot

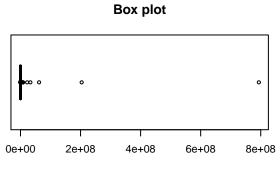


min: 0, q1: 4.37, med: 5.25, mean: 5.409, q3: 6.218, max: 14.49, na: 1

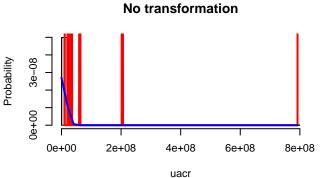


uric_acid_serum
3105 non-missing records (1 NA)

uacr

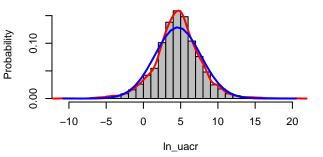


min: 2.126e–05, q1: 18.14, med: 102.3, mean: 381900, q3: 594, max: 793700000, na: 0

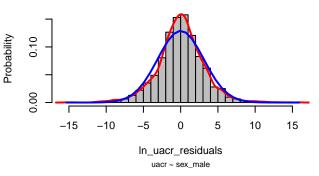


3106 non–missing records (0 NA)

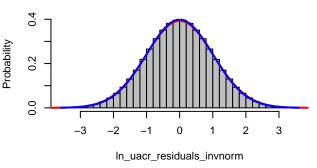
Logarithmic transformation



Residuals

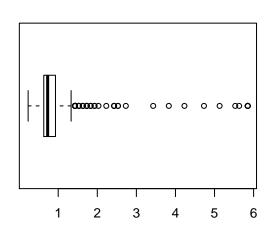


Normalized residuals



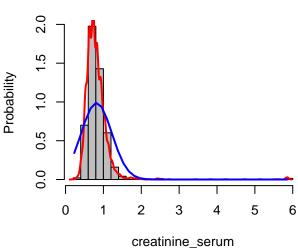
creatinine_serum





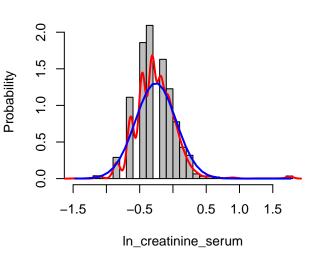
min: 0.23, q1: 0.63, med: 0.73, mean: 0.8179, q3: 0.93, max: 5.85, na: 1

No transformation

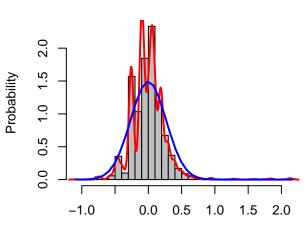


creatinine_serum
3105 non-missing records (1 NA)

Logarithmic transformation



Residuals

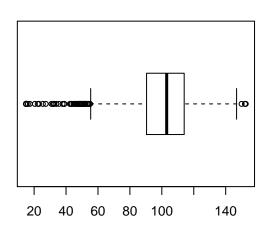


In_creatinine_serum_residuals

creatinine_serum ~ age_crea_serum + sex_male

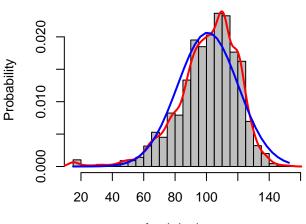
egfr_ckdepi_creat





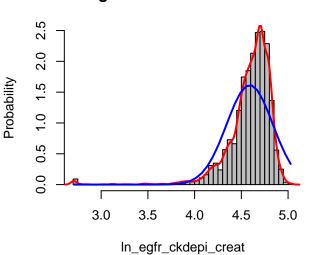
min: 15, q1: 90.44, med: 103, mean: 100.9, q3: 114, max: 152.6, na: 1

No transformation

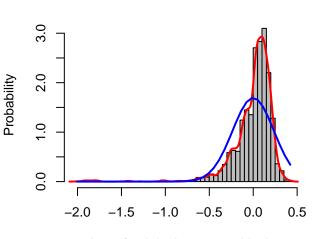


egfr_ckdepi_creat 3105 non-missing records (1 NA)

Logarithmic transformation



Residuals



In_egfr_ckdepi_creat_residuals
egfr_ckdepi_creat ~ age_crea_serum + sex_male