# Comparing ABL and numerical eGFR

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## 1 Introduction

TODO: together with Alex

TODO: outwrite GFR: (Wikipedia) Glomerular filtration rate (GFR) de-

scribes the flow rate of filtered fluid through the kidney.

# 2 Method

### 2.1 Data

Data from 780 patients (16-97 years, 340 female and 440 male) was used for the analysis. All patients were consecutively referred for determination of GFR by two different methods. TODO: describe the methods and their measurement uncertainty, call them (A) and (B) or so.

According to TODO: *put ref* we group the GFR values into six groups G1, G2, G3a, G3b, G4, and G5 as shown in table 1.

Table 1: GFR-Stadium used for categorize the chronically kidney diseases.

Category	Description	Range $(ml/min/1.73m^2)$
G1	Normal	90+
G2	Mildly decreased	60-89
G3a	Mildly to moderately decreased	45-59
G3b	Moderately to severely decreased	30-44
G4	Severely decreased	15-29
G5	Kidney failure	<15

Table 2: TODO: Caption

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	GFR (Opus ABL)	eGFR (CKD-EPI)		
Min	5	5		
Max	171	165		
Mean	89.93	89.13		
$\operatorname{Std}$	28.29	28.11		
CV	0.3146	0.3154		
$p_{\mathrm{G1}}$	0.5603	0.5474		
$p_{\mathrm{G2}}$	0.2949	0.3051		
$p_{\mathrm{G3a}}$	0.0564	0.059		
$p_{\mathrm{G3b}}$	0.059	0.059		
$p_{\mathrm{G4}}$	0.0167	0.0179		
$p_{ m G5}$	0.0128	0.0115		

### 2.2 Evaluation

The equation for GFR is

GFR = 
$$\beta \min(S_{cr}/k, 1)^{\alpha} \max(S_{cr}/k, 1)^{-1.209} 0.993^{age} * 1.018$$
(if female)

where  $\alpha$  is -0.329 for females and -0.411 for males and  $\beta$  is defined based on the origin and the sex of the patient as in table 3.

Table 3: Definition of  $\beta$  based on the origin and the sex of the patient.

sex	black	white or other
female	$\beta = 166$	$\beta = 144$
male	$\beta = 163$	$\beta = 141$

# 3 Results and Discussion

#### 3.1 Correlation

The correlation between two variables X and Y (also called Pearson's Correlation Coefficient (PCC) TODO:  $put\ ref\ here$ ) is defined as

$$\rho_{X,Y} = \frac{\text{cov}(X,Y)}{\sigma_X \sigma_Y}$$

where cov(X, Y) is the covariance between X and Y and  $\sigma_X$  (resp.  $\sigma_Y$ ) denotes the standard deviation of X (resp. Y).

Assuming that the two samples X and Y are drawn from independent normal distributions, the probability density function of the sample correlation coefficient  $\rho$  reads [Student, 1908]

$$f(\rho) = \frac{(1 - \rho^2)(n/2 - 2)}{B(1/2, n/2 - 1)} \tag{1}$$

where  $B(\cdot, \cdot)$  is the beta function and n the number of samples. This function can be used to approximately compute a confidence interval as well as probability value (p-value) that uncorrelated and normally distributed samples give a correlation values at least as extreme as  $\rho_{X,Y}$ .

The data described above produced correlation coefficient of  $\rho_{X,Y} = 0.9908$  with p-value  $< 10^{-10}$  and confidence interval between 0.9894 and 0.992.

## 3.2 Regression

Deming Regression (DR) is commonly used for fitting a linear spline to two-dimensional samples where both variables, X and Y, are measured with error TODO: put ref here. DR assumes that the error ratio (denoted by  $\lambda$ ) is constant. For the following analysis we assumed the error ratio to be equal to 1 in which case DR gives the same result as orthogonal regression. The left figure of 1 shows the Deming regression (dashed red) and its uncertainty region (magenta) between GFR (Opus ABL) and eGFR (CKD-EPI), while the right part shows the corresponding Bland-Altman-Plot. TODO: put analysis and explanations of the plots here

## 3.3 Prediction

The Deming regression line from figure 1 can be used for predicting the G-category (according to table 1) for eGFR (CKD-EPI) from the values of GFR (Opus ABL). Figure 2 shows the confusion matrix that was obtained by a 10-fold cross-validation technique. The x-axis shows the predicted and the y-axis the true categories. In total, there are 725 (p = 0.9295) subjects that are classified correctly what leaves 55 (p = 0.0705) misassignmenets. We observe that both, the true as well as the predicted distribution highly emphasize the categories G1 and G2. This becomes even more clear in Table 4, where assignment frequencies for both distributions are presented.

TODO: explain the measures in table 5, explain what the vaues of Cohen's kappa and Fleiss' kappa mean

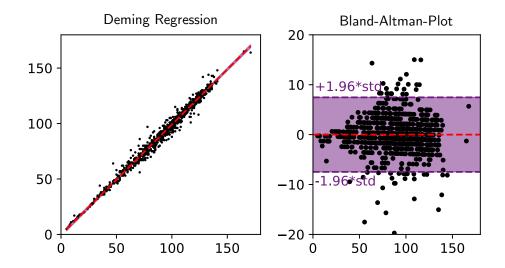


Figure 1: Deming regression line and the corresponding Bland-Altman-Plot of the two data sets.

Table 4: Assignment frequencies for the true and the predicted categories.

Category	True		Predicted	
G1	427	(p = 0.5474)	428	(p = 0.5487)
G2	238	(p = 0.3051)	231	(p = 0.2961)
G3a	46	(p = 0.059)	50	(p = 0.0641)
G3b	46	(p = 0.059)	43	(p = 0.0551)
G4	14	(p = 0.0179)	16	(p = 0.0205)
G5	9	(p = 0.0115)	12	(p = 0.0154)

# 4 Conclusion

# References

Student. Probable error of a correlation coefficient. *Biometrika*, pages 302–310, 1908.

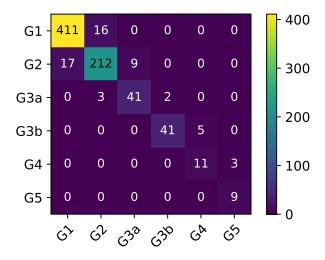


Figure 2: 10-fold cross-validation confusion matrix for the G-category prediction by a Deming regression line.

Table 5: Correlation measures when predicting the G-category of eGFR (CKD-EPI) from the value of GFR (Opus ABL) using a 10-fold cross-validation scheme.

Measure	Correlation		
Baseline	0.3983		
Cohen's $\kappa$	0.8828		
Fleiss' $\kappa$	0.881		