

concordance=TRUE

```
## Warning in des.print(dat = MyData, create = "knitr", var.names =  
c("sex", : group is missing! index and group.miss were set to FALSE  
and which.col were set to "total"  
  
## % latex table generated in R 3.5.3 by xtable 1.8-4 package  
## % Tue May 28 15:06:07 2019  
## \begin{longtable}{lc}  
## \caption{Beispiel Auswertung} \\  
## \hline & Total & \& (n = 25) \\  
## \hline\\  
## \endhead  
## \hline\\endfoot sex (sex) & \\  
## \hspace{2ex} N & 25 \\  
## \hspace{2ex} Missing & 0 \\  
## \hspace{2ex} Mean & 0.200 \\  
## \hspace{2ex} SD & 0.408248 \\  
## \hspace{2ex} Median & 0 \\  
## \hspace{2ex} Q1 -- Q3 & 0 -- 0 \\  
## \hspace{2ex} Min. -- Max. & 0.0 -- 1.0 \\  
## & \\  
## jaw (jaw) & \\  
## \hspace{2ex} N & 25 \\  
## \hspace{2ex} Missing & 0 \\  
## \hspace{2ex} Mean & 0.520 \\  
## \hspace{2ex} SD & 0.509902 \\  
## \hspace{2ex} Median & 1 \\  
## \hspace{2ex} Q1 -- Q3 & 0 -- 1 \\  
## \hspace{2ex} Min. -- Max. & 0.0 -- 1.0 \\  
## & \\  
## front (front) & \\  
## \hspace{2ex} N & 25 \\  
## \hspace{2ex} Missing & 0 \\  
## \hspace{2ex} Mean & 0.960 \\  
## \hspace{2ex} SD & 0.734847 \\  
## \hspace{2ex} Median & 1 \\  
## \hspace{2ex} Q1 -- Q3 & 0 -- 1 \\  
## \hspace{2ex} Min. -- Max. & 0.0 -- 2.0 \\  
## & \\  
## premolar (premolar) & \\  
## \hspace{2ex} N & 25 \\  
## \hspace{2ex} Missing & 0 \\  
## \hspace{2ex} Mean & 0.240 \\  
## \hspace{2ex} SD & 0.435890 \\  
## \hspace{2ex} Median & 0 \\  

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## \hspace{2ex} Q1 -- Q3 & 0 -- 0 \\
## \hspace{2ex} Min. -- Max. & 0.0 -- 1.0 \\
## & \\
## anchor\_tooth (molar) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 0.280 \\
## \hspace{2ex} SD & 0.458258 \\
## \hspace{2ex} Median & 0 \\
## \hspace{2ex} Q1 -- Q3 & 0 -- 1 \\
## \hspace{2ex} Min. -- Max. & 0.0 -- 1.0 \\
## & \\
## \pagebreak furcation (anchor\_tooth) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 0.280 \\
## \hspace{2ex} SD & 0.458258 \\
## \hspace{2ex} Median & 0 \\
## \hspace{2ex} Q1 -- Q3 & 0 -- 1 \\
## \hspace{2ex} Min. -- Max. & 0.0 -- 1.0 \\
## & \\
## GTR (furcation) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 0.480 \\
## \hspace{2ex} SD & 0.714143 \\
## \hspace{2ex} Median & 0 \\
## \hspace{2ex} Q1 -- Q3 & 0 -- 1 \\
## \hspace{2ex} Min. -- Max. & 0.0 -- 2.0 \\
## & \\
## bone\_level (GTR) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 0.120 \\
## \hspace{2ex} SD & 0.331662 \\
## \hspace{2ex} Median & 0 \\
## \hspace{2ex} Q1 -- Q3 & 0 -- 0 \\
## \hspace{2ex} Min. -- Max. & 0.0 -- 1.0 \\
## & \\
## prognosis (bone\_level) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 1.720 \\
## \hspace{2ex} SD & 0.890693 \\
## \hspace{2ex} Median & 1

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## \hspace{2ex} Q1 -- Q3 & 1 -- 2 \\
## \hspace{2ex} Min. -- Max. & 1.0 -- 4.0 \\
## & \\
## age (prognosis) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 1.120 \\
## \hspace{2ex} SD & 0.331662 \\
## \hspace{2ex} Median & 1 \\
## \hspace{2ex} Q1 -- Q3 & 1 -- 1 \\
## \hspace{2ex} Min. -- Max. & 1.0 -- 2.0 \\
## & \\
## \pagebreak NA (age) & \\
## \hspace{2ex} N & 25 \\
## \hspace{2ex} Missing & 0 \\
## \hspace{2ex} Mean & 52.800 \\
## \hspace{2ex} SD & 12.529964 \\
## \hspace{2ex} Median & 54 \\
## \hspace{2ex} Q1 -- Q3 & 43 -- 61 \\
## \hspace{2ex} Min. -- Max. & 31.0 -- 76.0 \\
## & \\
## \hline
## \end{longtable}

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