

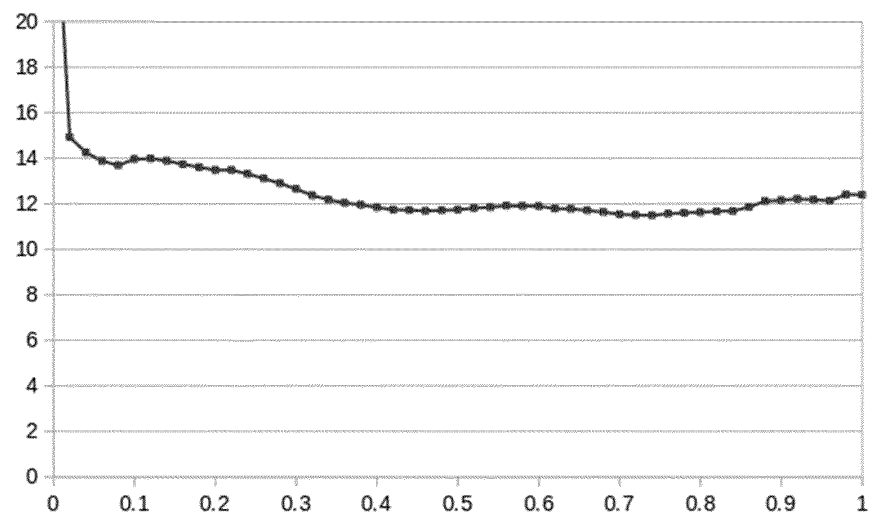
Figures

Fig. 1.

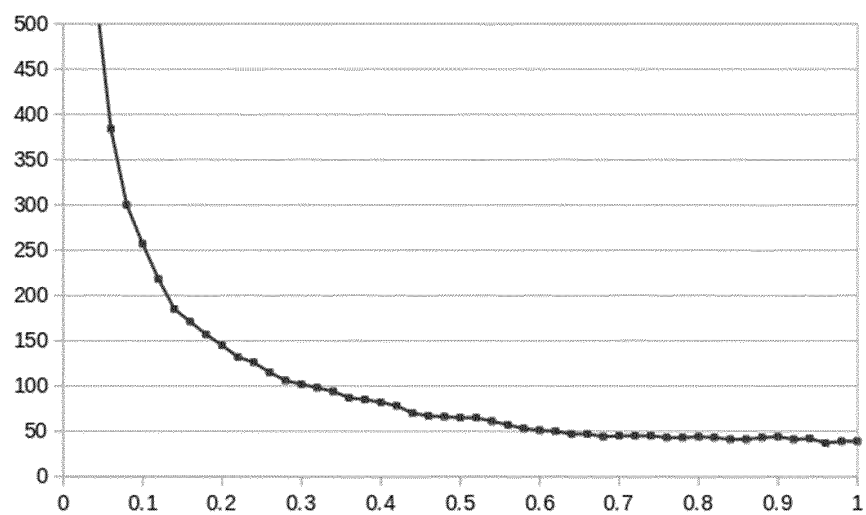


Fig. 2.

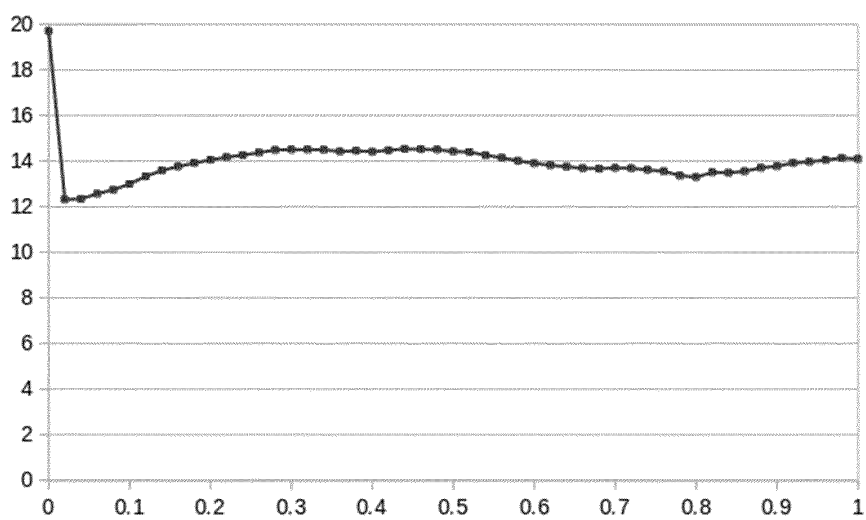


Fig. 3.

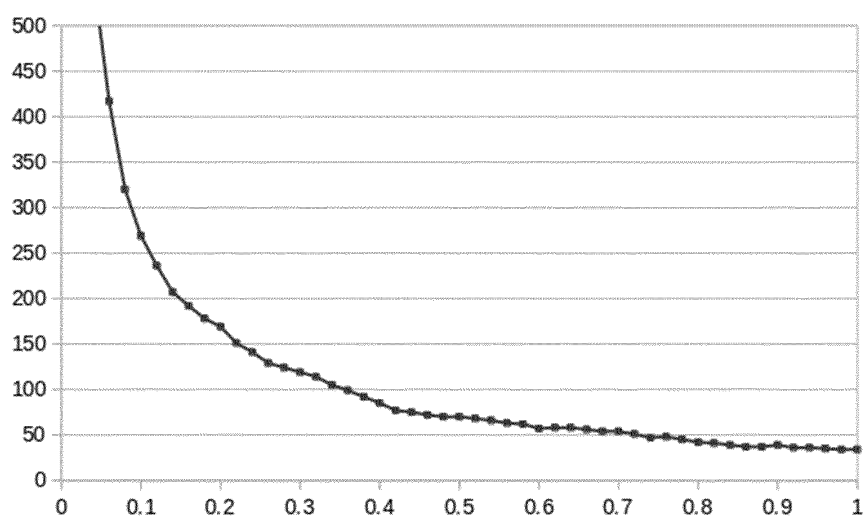


Fig. 4.

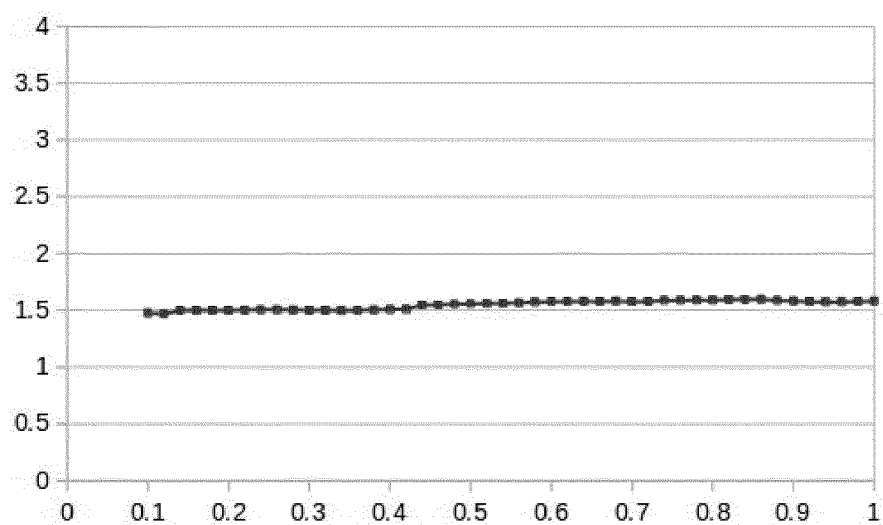


Fig. 5.

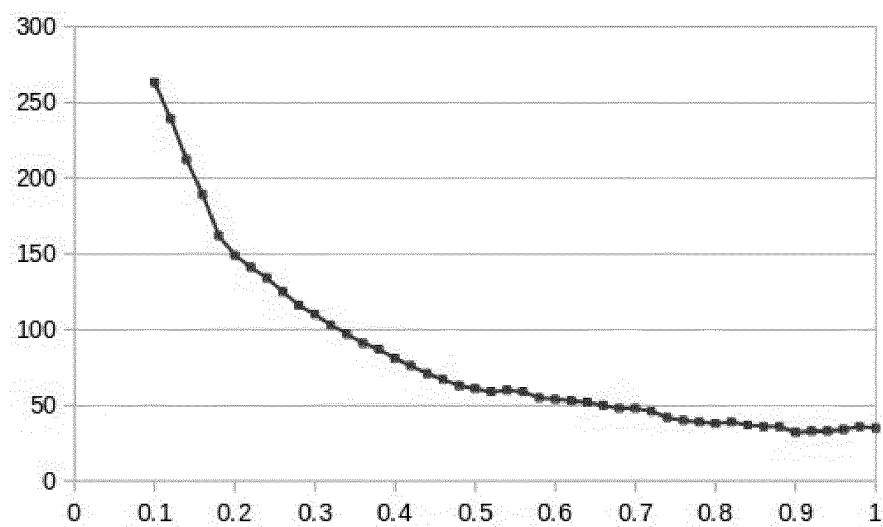


Fig. 6.

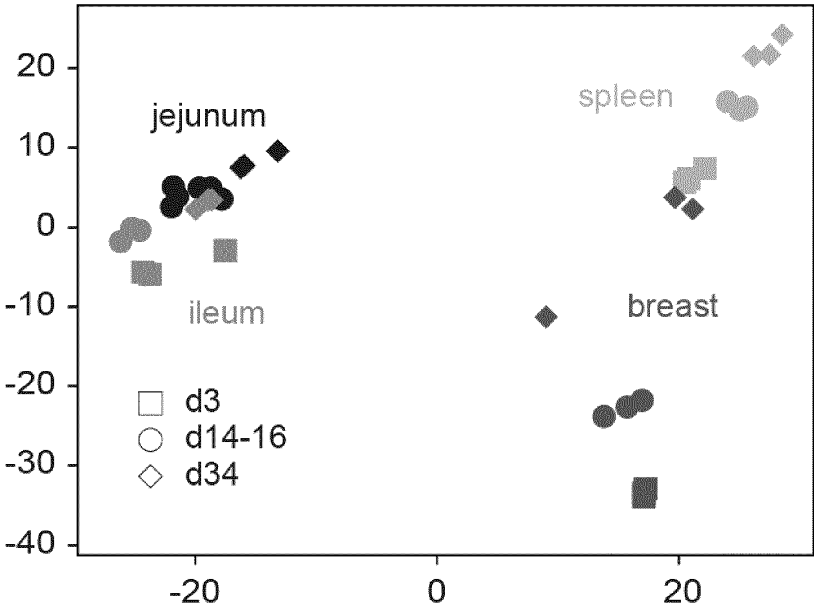


Fig. 7.: PCA analysis based on the DNA methylation patterns of 67,651 LMRs in the training set (N=36 samples).

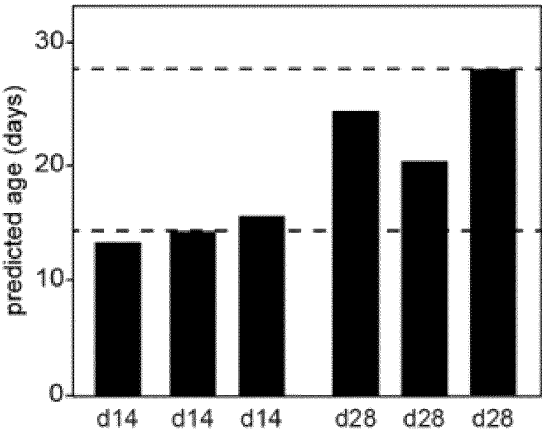


Fig. 8.

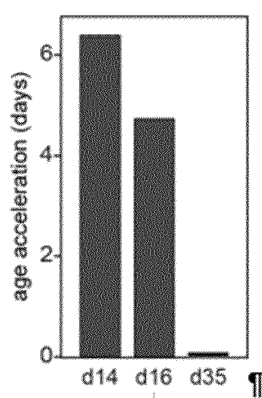


Fig. 9.