

SalivaPrint – A tool for patient stratification in oral health

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Introduction

It is known that saliva reflects many systemic and oral pathologies (Rosa *et al* 2012). At SalivaTec we are interested in using this fluid in diagnostics and patient stratification and have been analyzing saliva from several individuals characterizes as to their oral and systemic health.

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Goals: Demonstrate that total salivary protein profile determined by capillary electrophoresis (SalivaPrint) may be used to direct the search for biomarkers.



Methods

SalivaPrints were obtained for **26 individuals** with **periodontal disease**. These total salivary protein profiles were obtained through automated microfluidic Experion electrophoresis system (Bio-Rad, PT).

Individuals included participate in a larger study relating oral health with different systemic factors and are volunteers from a group of seniors participating in and exercise program of the Município de Viseu. Their profiles were compared with **SalivaPrints** from **healthy individuals** and the most representative features for separation of the two groups were selected.



Figure 1. Data and saliva collection locations spanning 12 locations in the Viseu district

Results



Table I. Proteins and respective molecular weights in SalivaPrint (Rosa et al 2016) Molecular Molecular SalivaTecDB Weight Entry Protein names Weight condition Range (kDa) identification (kDa) Periodontitis/ P31947 14-3-3 protein sigma 28 Healthy 28 - 29 P06870 Kallikrein-1 29 Healthy P30740 Leukocyte elastase inhibitor 43 Healthy 42 - 43 Protein-glutamine gamma-Q08188 77 Healthy glutamyltransferase E 77 - 78 Periodontitis/ P02788 Lactotransferrin 78 Healthy

Five molecular weight ranges of SalivaPrint seem to be important to separate the individuals with periodontal disease from the healthy individuals. These molecular weight ranges (28-29, 42-43, and 77-78 kDa) have been shown to include several proteins presented in Table I (Rosa et al 2016).

Figure 2. Molecular weight ranges (kDa) contributing to discrimination of health *vs.* periodontal disease.



Some of these proteins are functionally related to processes deregulated in oral and systemic disease.

Proteins such as P31947 (14-3-3 protein sigma) are related to the mTor pathway involved in **glucose resistance**, a condition related to **obesity** and **diabetes**. This is consistent with a **high prevalence of these pathologies in the individuals with periodontal disease included**.

Figure 3. Metabolic pathway of mTOR, related with the glucose regulation (insulin resistant, diabetes and obesity). Adapted from Kegg

Conclusion

SalivaPrint can provide information on which molecules should be used to distinguish between individuals with oral health and periodontal disease.

Although the diagnostics of periodontal disease through salivary markers is an interesting approach, explored by our laboratory, the results presented here indicate that metabolic deregulations such as diabetes and obesity have to be considered if the quantification of salivary markers is to be used for diagnostics.

