



Prevalence of Gene Mutations in Dental Agenesis and Colorectal Cancer

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Objective

To describe the available evidence about the prevalence of KRAS, PIK3CA, BRAF and AXIN2 mutations in colorectal cancer individuals and its possible association with dental agenesis.

Methods

Study design. a systematic review was conducted according to Preferred Reporting Items for Systematic Reviews and Meta-Analyses statement (PRISMA).

Selection criteria. Observational studies (english or spanish). prevalence of KRAS, PIK3CA, BRAF and AXIN2 mutation/polymorphism in primary or metastatic colorectal cancer. Interval of seven years.

Literature searches.



Terms: "Genes", "RAS", "Kras", "PIK3CA", "BRAF", "AXIN2"; in combination with "Prevalence", "Mutation" or "Polymorphism" and "Colorectal Neoplasms" or "Colorectal Cancer". In addition, "dental agenesis".

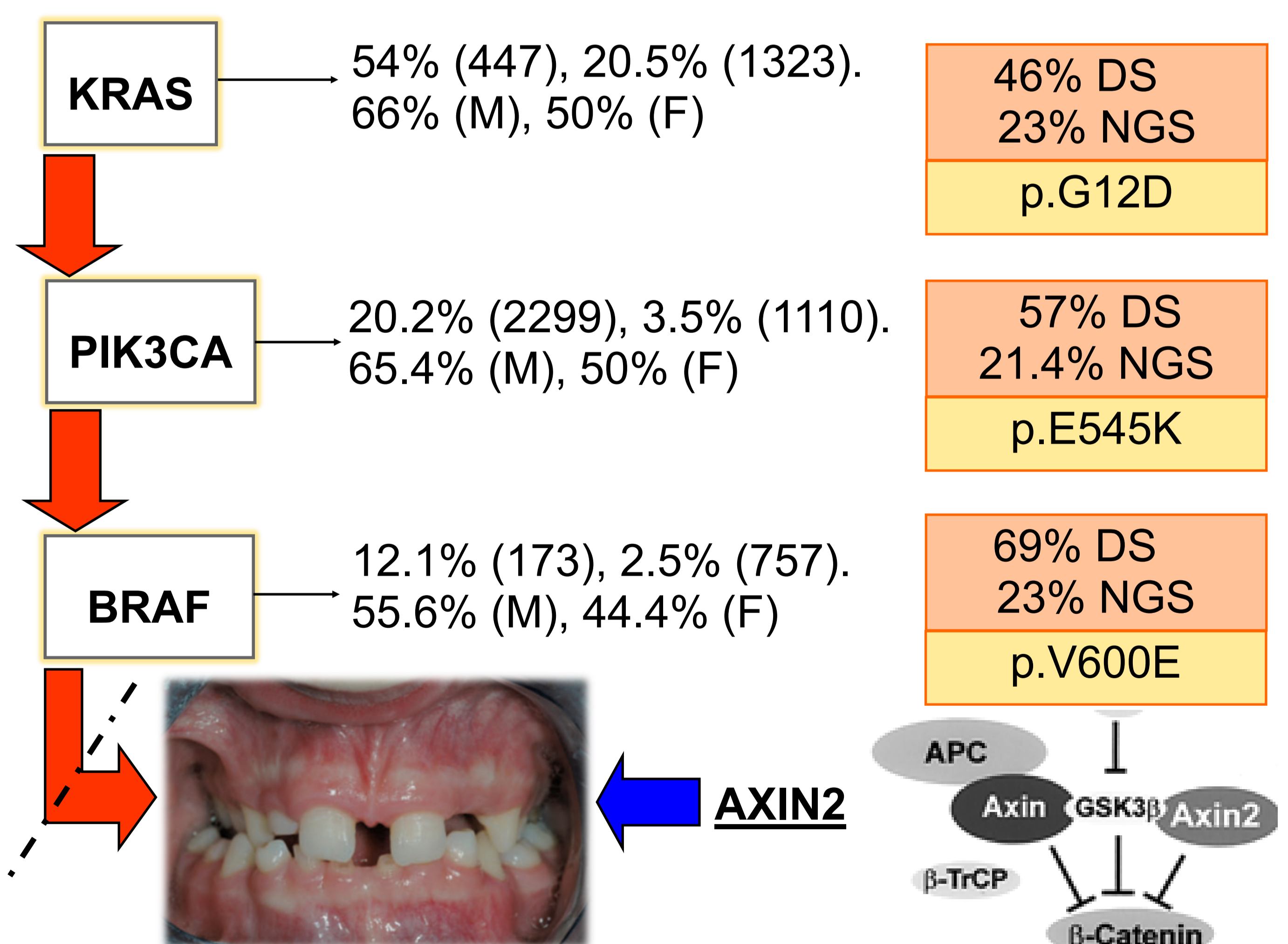
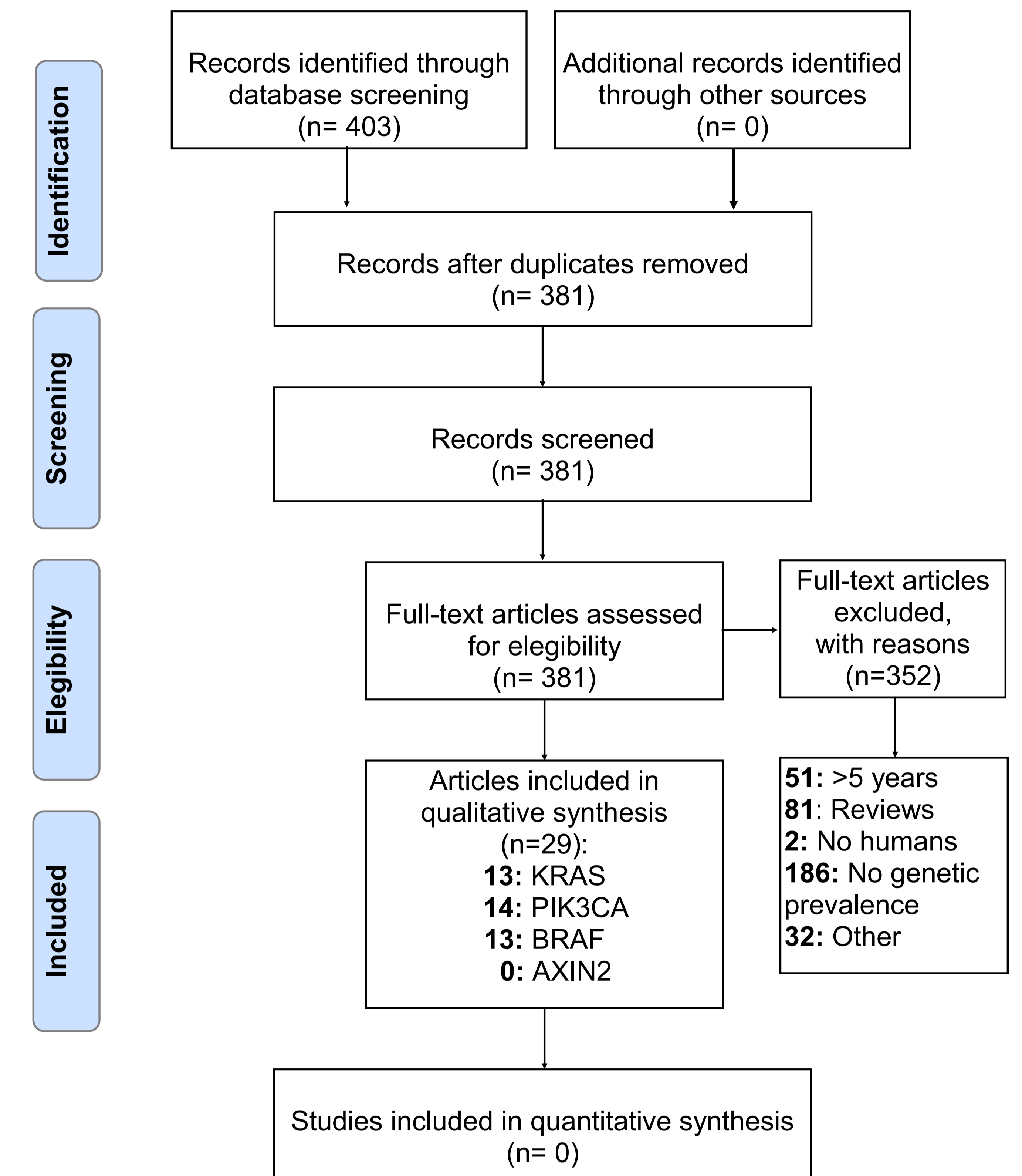
Study selection. EndNote® (Version X8, Thomson Reuters).

Methodological quality assessment. "Aroot for the critical appraisal of epidemiological cross-sectional studies" Berra S. *et al.*, 2008.

≥ 108 + Good internal validity	→	High MQ
81—107 + Regular internal validity	→	Median MQ

Data extraction. author, publication year, geographic región where the study was performed, sample size, general prevalence of the mutation, mutation prevalence by sex, aminoacids changes and sequencing techniques.

Results



Conclusion

KRAS mutations were the most prevalent, although, there is a lacking evidence on relation of dental agenesis and the prevalence of colorectal cancer mutations.

References

Henson BS, et al., Collection, storage, and processing of saliva samples for down-stream molecular applications. In Oral Biology. 2010:21-30.
 Jessie K, et al., Protein precipitation method for salivary proteins and rehydration