New wiki on salivary proteins may transform diagnostic testing and personalized medicine
25 May 2021, by Marcene Robinson

A diagram that shows the interconnectedness of the thousands of salivary proteins originating from blood plasma, parotid glands, and submandibular and sublingual glands. The diagram is one of many tools available to researchers and clinicians through the Human Salivary Proteome Wiki. Credit: University at Buffalo

To improve the development of new saliva-based diagnostic tests and personalized medicine, the National Institute of Dental and Craniofacial Research (NIDCR) has supported the development of the Human Salivary Proteome Wiki, the first public platform that catalogs and curates data on each of the thousands of proteins within our saliva.

A challenge in saliva's study, however, is the high level of variability in the thousands of salivary proteins. Therefore, normal ranges for the composition of saliva need to be established to define the status of health, says Ruhl, who in 2020, co-authored research on how saliva is made, pinpointing the origins of proteins in human saliva back to its sources.

"Saliva has become an attractive body fluid for on-site, remote and real-time monitoring of oral and systemic health. The scientific community needs a saliva-centered information platform that keeps pace with the rapid accumulation of new data and knowledge," says William Lau, Ph.D., of the
National Institutes of Health (NIH) and first author of the study.

"The Human Salivary Proteome Wiki will improve salivary sciences, saliva-based diagnostics, precision medicine and dentistry, and ultimately facilitate personalized treatment for both oral and systemic diseases," says Preethi Chander, Ph.D., program director of the NIDCR Salivary Biology and Immunology Program.

Modernized to 'harness the full potential of the salivary proteome'

Funded by NIDCR and developed in collaboration with the NIH Center for Information Technology, the Human Salivary Proteome Wiki was released in 2019 as a user-friendly public database to provide rich analytical, curation, and annotation tools for the biomedical research community.

The database compiles information and research on the salivary proteome, genome, transcriptome (types and levels of mRNA expressed in salivary gland tissue), and glycome (array of sugars attached to salivary glycoproteins).

Users may submit proposals to add, and remove or modify data to an interdisciplinary team of curators who ensure information is accurate and supported by sound scientific evidence. Annotations are also credited to the researchers and clinicians who submit them to the wiki, helping drive references to critical research.

The researchers will also host a series of activities that bring contributors together to inspire novel ideas and foster cross-discipline collaborations.


Provided by University at Buffalo