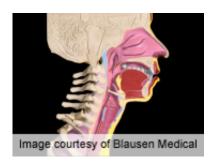


Gene signature validated for oral cancer metastases

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A multigene signature effectively predicts the presence of lymph node metastases in squamous cell carcinoma of the oral cavity and oropharynx, according to a study published online Oct. 8 in the *Journal of Clinical Oncology*.

(HealthDay)—A multigene signature effectively predicts the presence of lymph node metastases in squamous cell carcinoma of the oral cavity (OSCC) and oropharynx (OPSCC), according to a study published online Oct. 8 in the *Journal of Clinical Oncology*.

Sander R. van Hooff, from the University Medical Center Utrecht in the Netherlands, and colleagues used a diagnostic DNA microarray to evaluate a previously reported multigene signature for distinguishing metastasizing from non-metastasizing (N0) OSCC and OPSCC in 222 patients from all head and neck oncology centers in the Netherlands.

The researchers found that the negative predictive value was 72 percent



for the diagnostic signature in the entire validation cohort. In a subset of early-stage (cT1 to T2N0) OSCC (101 patients), the signature performed well, with a negative predictive value of 89 percent.

"Combining current <u>clinical assessment</u> with the expression signature would decrease the rate of undetected nodal metastases from 28 to 11 percent in early-stage OSCC. This should be sufficient to enable clinicians to refrain from elective neck treatment," the authors write. "A new clinical decision model that incorporates the expression signature is therefore proposed for testing in a prospective study, which could substantially improve treatment for this group of patients."

One author disclosed **financial ties** to the diagnostic company Agendia.

More information: Abstract

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Editorial

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