

LETTERS

***Candida auris* can be identified accurately**

I read with interest the recent *CMAJ* article by Schwartz and Dingle, which gives a good synopsis of *Candida auris*.¹ However, the authors' claim that certain matrix-assisted laser desorption/ionization time of flight (MALDI-TOF) mass spectrometry identification systems can misidentify *C. auris* is somewhat misleading.

The article by Tsay and colleagues² cited by Schwartz and Dingle states that accurate identification for *C. auris* can be performed by VITEK MS and Bruker Biotyper MALDI-TOF mass spectrometry. Public Health Ontario also states that these 2 methods can identify *C. auris*.³ Furthermore, a recent study of 50 molecularly confirmed *C. auris* isolates from different geographical areas showed 100% accurate identification by MALDI-TOF mass spectrometry.⁴ Despite the small sample, that is more than the 19 cases of *C. auris* identified in Canada since 2012.¹

I would like to gently caution that Schwartz and Dingle's misleading claim could lead to clinicians and laboratorians unnecessarily questioning the identification

of *Candida* species by MALDI-TOF mass spectrometry, and consequently overtesting and misusing resources. Patients may also be misled, which is not helpful in establishing good physician–patient rapport.

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