

Last Updated July 2023

Refereed original research:

indicates co-first authorship

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1. Liu C, Mokashi NV, Darville T, Sun X, O'Connell CM, Hufnagel K, Waterboer T, and **Zheng X***. A machine learning based analytic pipeline applied to clinical and serum IgG immunoproteome data to predict *Chlamydia trachomatis* genital tract ascension and incident infection in women. *Microbiol Spectr*. 2023 Jun 15:e0468922. doi: 10.1128/spectrum.04689-22. Epub ahead of print. PMID: 37318345.
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