



Louisa A Carlisle

Contact

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Publications (2)

Carlisle L, Günthard H, Bernasconi E, Vernazza P, Stoeckle M, Cavassini M, Nguyen A, Yerly S, Rauch A, Salazar-Vizcaya L, Fehr J, Braun D, Huber M, Böni J, Shah C, Mbunkah H, Metzner K, Turk T, Kouyos R. HCV Genetic Diversity Can Be Used to Infer Infection Recency and Time since Infection. *Viruses* 2020; 12

Carlisle L, Braun D, Rauch A, Calmy A, Cavassini M, Battegay M, Vernazza P, Bernasconi E, Günthard H, Kouyos R, Perreau M, Klimkait T, Yerly S, Turk T, Kusejko K, Metzner K, Leemann C, Schenkel C, Bachmann N, Posada S, Beerwinkler N, Böni J, Swiss HIV Cohort Study. Viral diversity from next-generation sequencing of HIV-1 samples provides precise estimates of infection recency and time since infection. *J Infect Dis* 2019

Projects (0)

No results found.

Kantonsspital St.Gallen

Rorschacher Strasse 95

CH-9007 St.Gallen

T: +41 71 494 11 11

support.forschung@kssg.ch