



**Nadine Bachmann**

**Kontakt**

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## Publikationen (6)

Thorball C, Klimkait T, Yerly S, Battegay M, Rauch A, Schmid P, Bernasconi E, Cavassini M, Kouyos R, Günthard H, Metzner K, Fellay J, Perreau M, Böni J, Wieser M, Borghesi A, Bachmann N, Von Siebenthal C, Vongrad V, Turk T, Neumann K, Beerenwinkel N, Bogojeska J, Roth V, Kok Y, Parbhoo S, Swiss HIV Cohort Study. Host Genomics of the HIV-1 Reservoir Size and Its Decay Rate During Suppressive Antiretroviral Treatment. *J Acquir Immune Defic Syndr* 2020; 85:517-524.

Wan C, Yerly S, Battegay M, Waliti L, Calmy A, Vernazza P, Bernasconi E, Cavassini M, Metzner K, Günthard H, Kouyos R, Klimkait T, Perreau M, Böni J, Bachmann N, Mitov V, Blanquart F, Céspedes S, Turk T, Neumann K, Beerenwinkel N, Bogojeska J, Fellay J, Roth V, Swiss HIV Cohort Study. Heritability of the HIV-1 reservoir size and decay under long-term suppressive ART. *Nat Commun* 2020; 11:5542.

Thorball C, Klimkait T, Yerly S, Battegay M, Rauch A, Schmid P, Bernasconi E, Cavassini M, Kouyos R, Günthard H, Metzner K, Fellay J, Perreau M, Böni J, Wieser M, Borghesi A, Bachmann N, Von Siebenthal C, Vongrad V, Turk T, Neumann K, Beerenwinkel N, Bogojeska J, Roth V, Kok Y, Parbhoo S, Swiss HIV Cohort Study. Host genomics of the HIV-1 reservoir size and its decay rate during suppressive antiretroviral treatment. *J Acquir Immune Defic Syndr* 2020

Bachmann N, Kouyos R, Günthard H, Cavassini M, Bernasconi E, Vernazza P, Ramette A, Rauch A, Battegay M, Yerly S, Klimkait T, Perreau M, Böni J, Turk T, Kadelka C, Chaudron S, Nguyen H, Kusejko K, Swiss HIV Cohort Study. Phylogenetic Cluster Analysis Identifies Virological and Behavioral Drivers of HIV Transmission in MSM. *Clin Infect Dis* 2020

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, Beerenwinkel 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

Yerly S, Marzolini C, Martinez de Tejada B, Martinetti G, Ledergerber B, Kovari H, Kouyos R, Klimkait T, Keiser O, Kaiser L, Kahlert C, Hösl I, Hoffmann M, Hirsch H, Hasse B, Metzner K, Müller N, Weber R, Wandeler G, Vernazza P, Trkola A, Tarr P, Stöckle M, Speck R, Schmid P, Scherrer A, Rudin C, Rauch A, Paioni P, Pantaleo G, Nicca D, Haerry D, Günthard H, Fux C, Kouyos R, Günthard H, Hoffmann M, Furrer H, Cavassini M, Calmy A, Bernasconi E, Battegay M, Klimkait T, Aubert V, Yerly S, Böni J, Kadelka C, Bachmann N, Swiss HIV Cohort Study, Aubert V, Furrer H, Fellay J, Fehr J, Elzi L, Egger M, Dollenmaier G, Ciuffi A, Cavassini M, Calmy A, Bucher H, Braun D, Böni J, Bernasconi E, Battegay M, Turk T. Assessing the danger of self-sustained HIV epidemics in heterosexuals by population based phylogenetic cluster analysis. *Elife* 2017; 6

## Projekte (0)

Keine Resultate gefunden.

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