



## Christian Hammer

### Contact

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## Publications (3)

Rüeger S, Fellay J, Zdobnov E, Breuer J, Morfopoulou S, Depledge D, Rauch A, Kahlert C, Günthard H, Cavassini M, Bernasconi E, Khanna N, Naret O, Lawless D, McLaren P, Loetscher A, Hammer C, Swiss HIV Cohort Study. Author Correction: The influence of human genetic variation on Epstein-Barr virus sequence diversity. *Sci Rep* 2021; 11:8946.

Rüeger S, Fellay J, Zdobnov E, Breuer J, Morfopoulou S, Depledge D, Rauch A, Kahlert C, Günthard H, Cavassini M, Bernasconi E, Khanna N, Naret O, Lawless D, McLaren P, Loetscher A, Hammer C, Swiss HIV Cohort Study. The influence of human genetic variation on Epstein-Barr virus sequence diversity. *Sci Rep* 2021; 11:4586.

Thorball C, Rauch A, Battegay M, Hoffmann M, Schmid P, Bernasconi E, Günthard H, Mohammadi P, McLaren P, Rabkin C, Besson C, Cavassini M, Theodorou I, Oudot-Mellakh T, Ehsan N, Hammer C, Santoni F, Niay J, Costagliola D, Goujard C, Meyer L, Wang S, Hussain S, Fellay J. Genetic variation near CXCL12 is associated with susceptibility to HIV-related non-Hodgkin lymphoma. *Haematologica* 2020

## Projects (0)

No results found.

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