

Home > Origin of hepatitis C virus genotype 3 in Africa as estimated through an evolutionary analysis of the full-length genomes of nine subtypes, including the newly sequenced 3d and 3e

Origin of hepatitis C virus genotype 3 in Africa as estimated through an evolutionary analysis of the full-length genomes of nine subtypes, including the newly sequenced 3d and 3e ^[1]

Li, Chunhua; Lu, Ling; Murphy, Donald G.; Negro, Francesco; Okamoto, Hiroaki
The Journal of General Virology
August, 2014
10.1099/vir.0.065128-0

Extra (Zotero):

PMID: 24795446 PMCID: PMC4103067

Québec 

© 2001-2019 Gouvernement du Québec

Source URL (modified on 10/04/2018 - 16:02): <https://www.inspq.qc.ca/en/publications/autres/c6p3y7wi>

Links

[1] <https://www.inspq.qc.ca/en/publications/autres/c6p3y7wi>

