

# Identification and Characterization of a Novel HIV-1 Circulating Recombinant (CRF59\_01B) Identified among Men-Who-Have-Sex-with-Men

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Phylogenetic Analysis of HIV-1 CRF65_CPX Reveals Yunnan Province Is Still a Source Contributing to the Spread of HIV-1 in China. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2015, 70, e120-e122.	2.1	11
2	Novel HIV-1 Recombinant Identified in a Foreign Heterosexual Resident in Japan: Relatedness to Recently Reported CRF69_01B, Detected Primarily among Japanese Men Who Have Sex with Men. <i>Genome Announcements</i> , 2015, 3, .	0.8	4
3	HIV-1 Genetic Diversity and Its Impact on Baseline CD4+T Cells and Viral Loads among Recently Infected Men Who Have Sex with Men in Shanghai, China. <i>PLoS ONE</i> , 2015, 10, e0129559.	2.5	37
4	HIV-1 diversity in infected individuals in Suzhou and Suqian, China. <i>SpringerPlus</i> , 2016, 5, 886.	1.2	6
5	Characterization of the Near Full-Length Genome of a Novel HIV-1 CRF01_AE/CRF07_BC Recombinant in an Injection Drug User from Southern Taiwan. <i>AIDS Research and Human Retroviruses</i> , 2016, 32, 588-594.	1.1	3
6	Nationwide Trends in Molecular Epidemiology of HIV-1 in China. <i>AIDS Research and Human Retroviruses</i> , 2016, 32, 851-859.	1.1	81
7	Identification of a Novel HIV-1 Circulating Recombinant Form (CRF85_BC) in Sichuan, China. <i>AIDS Research and Human Retroviruses</i> , 2016, 32, 895-899.	1.1	12
8	The Epidemic Dynamics of Four Major Lineages of HIV-1 CRF01_AE Strains After Their Introduction into China. <i>AIDS Research and Human Retroviruses</i> , 2016, 32, 420-426.	1.1	16
9	Phylogenetic and temporal dynamics of human immunodeficiency virus type 1 CRF01_AE and CRF07_BC among recently infected antiretroviral therapy-naïve men who have sex with men in Jiangsu province, China, 2012 to 2015. <i>Medicine (United States)</i> , 2018, 97, e9826.	1.0	14
10	Surging HIV-1 CRF07_BC epidemic among recently infected men who have sex with men in Fujian, China. <i>Journal of Medical Virology</i> , 2018, 90, 1210-1221.	5.0	15
11	Identification of a Novel HIV-1 Unique Recombinant Form Comprising CRF01_AE, Subtype B <sup>1</sup> , and CRF65_cpx Among Men Who Have Sex with Men in Jilin, China. <i>AIDS Research and Human Retroviruses</i> , 2018, 34, 714-718.	1.1	4
12	Phylogenetic transmission clusters among newly diagnosed antiretroviral drug-naïve patients with human immunodeficiency virus-1 in Korea: A study from 1999 to 2012. <i>PLoS ONE</i> , 2019, 14, e0217817.	2.5	5
13	Local Epidemics Gone Viral: Evolution and Diffusion of the Italian HIV-1 Recombinant Form CRF60_BC. <i>Frontiers in Microbiology</i> , 2019, 10, 769.	3.5	6
14	Rapid and complicated HIV genotype expansion among high-risk groups in Guangdong Province, China. <i>BMC Infectious Diseases</i> , 2019, 19, 185.	2.9	18
15	Brief Report: Onward Transmission of Multiple HIV-1 Unique Recombinant Forms Among Men Who Have Sex With Men in Beijing, China. <i>Journal of Acquired Immune Deficiency Syndromes</i> (1999), 2019, 81, 1-4.	2.1	13
16	Characterization of a Novel HIV-1 Circulating Recombinant Form (CRF80_0107) Among Men Who Have Sex with Men in China. <i>AIDS Research and Human Retroviruses</i> , 2019, 35, 419-423.	1.1	25
17	HIV-1 genetic diversity and transmitted drug resistance among newly diagnosed HIV-1 individuals in Jiangmen, China. <i>Journal of Medical Virology</i> , 2020, 92, 3209-3218.	5.0	1
18	First complete-genome documentation of HIV-1 intersubtype superinfection with transmissions of diverse recombinants over time to five recipients. <i>PLoS Pathogens</i> , 2021, 17, e1009258.	4.7	9

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19	HIV-1 molecular transmission network among sexually transmitted populations in Liaoning Province, China. <i>Medicine (United States)</i> , 2021, 100, e26640.	1.0	4
20	Evaluation of a Novel In-house HIV-1 Genotype Drug Resistance Assay using Clinical Samples in China. <i>Current HIV Research</i> , 2022, 20, 32-41.	0.5	1
21	The Influence of Social Support on Quality of Life of Men Who Have Sex with Men in China: A Preliminary Study. <i>PLoS ONE</i> , 2015, 10, e0127644.	2.5	10
22	HIV-1 molecular epidemiology among newly diagnosed HIV-1 individuals in Hebei, a low HIV prevalence province in China. <i>PLoS ONE</i> , 2017, 12, e0171481.	2.5	37
23	Geographically-stratified HIV-1 group M pol subtype and circulating recombinant form sequences. <i>Scientific Data</i> , 2018, 5, 180148.	5.3	7
24	Identification of Two Novel HIV-1 Unique Recombinant Forms in Shenzhen, China. <i>AIDS Research and Human Retroviruses</i> , 2022, , .	1.1	0
25	OUP accepted manuscript. <i>Journal of Antimicrobial Chemotherapy</i> , 2022, , .	3.0	0
27	Identification of a New HIV-1 Circulating Recombinant Form CRF112_01B Strain in Baoding city, Hebei Province, China. <i>Current HIV Research</i> , 2022, 21, .	0.5	2
28	Molecular networks reveal complex interactions with MSM in heterosexual women living with HIV-1 who play peripheral roles in Guangzhou, China. <i>Acta Tropica</i> , 2023, 245, 106953.	2.0	0