# Investigating the Causal Links between COVID-19 and Pancreatitis by Bidirectional Mendelian Randomization

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#### Abstract

Objective The immune response is a double-edged sword, and COVID-19 shares similarities with pancreatitis in terms of natural immune response, immune storm, and multi-organ involvement. However, whether a causal association between them remained unclear. This study aimed to investigate the potential causal association between COVID-19 and pancreatitis using a bidirectional Mendelian Randomization (MR) approach. Methods The study analyzed three variables related to COVID-19 (severity, hospitalization, and susceptibility) with a sample size ranging from approximately 1,059,456 to 1,557,411. Additionally, four types of pancreatitis (acute, chronic, alcohol-induced acute, and chronic) were examined, with a sample size ranging from 337,126 to 377,277. Causal associations were estimated using inverse-variance weighted (IVW), median weighted, and MR-Egger methods. Results The IVW model indicated potential causal associations between genetic susceptibility to severe and hospitalized COVID-19 and a decreased risk of acute pancreatitis (OR = 0.914, p = 0.01; OR = 0.884, p = 0.008) and alcoholinduced chronic pancreatitis (OR = 0.852, p = 0.013; OR = 0.768, p = 0.002), including chronic pancreatitis. Inconsistent associations were observed between IVW and sensitivity analyses in acute and chronic pancreatitis of severe and hospitalized COVID-19. Conversely, no significant associations were found between pancreatitis traits and COVID-19-related variables in reverse MR analysis. No heterogeneity or pleiotropy was found. Conclusions Host genetic liability to severe and hospitalized COVID-19 was causally associated with declining risk of alcohol-induced chronic pancreatitis, while no significant association was observed for pancreatitis on COVID-19 outcomes. This study has significant implications for unraveling their pathogenesis and guiding clinical management.

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Outcome	Method	nSNP	р			OR(95%cl)	Heterogeneity.testp	MR_Egger.intercept.testp	MR_PRESSO.global.testp
Acute pancreatitis	IVW	5	0.685			1.05(0.82 to 1.34)	0.19	0.572	0.287(raw, 0 outliers)
	Weighted median	5	0.960			0.99(0.72 to 1.37)			
	MR Egger	5	0.665			0.81(0.34 to 1.93)			
Chronic pancreatitis	IVW	5	0.283	· · · · · · · · · · · · · · · · · · ·		1.31(0.80 to 2.17)	0.037	0.662	0.101(raw, 0 outliers)
	Weighted median	5	0.933 -			1.02(0.62 to 1.68)			
	MR Egger	5	0.487			+ 1.90(0.39 to 9.34)			
Alcohol-induced acute pancreatitis	IVW	5	0.775	· · · · ·		0.91(0.49 to 1.70)	0.074	0.636	0.149(raw, 0 outliers)
	Weighted median	5	0.806			0.89(0.37 to 2.15)			
	MR Egger	5	0.616			* 0.46(0.03 to 6.98)			
Alcohol-induced chronic pancreatitis	IVW	5	0.405			0.83(0.53 to 1.30)	0.107	0.798	0.227(raw, 0 outliers)
	Weighted median	5	0.900			0.96(0.49 to 1.88)			
	MR Egger	5	0.957	-		+ 1.06(0.17 to 6.72)			
p<0.05 was considered statistics	ally significant			1 2	3	4			
			protective facto	r risk factor					



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