

## 探討大量飲酒、ALDH2 rs671基因多型性和B型肝炎感染在肝硬化患者的 臨床組-醫師 肝癌發生率及整體死亡率的相關性

## Association of Heavy Alcohol Intake and ALDH2 rs671 Polymorphism With Hepatocellular **Carcinoma and Mortality in Patients With Hepatitis B Virus-Related Cirrhosis**

Pojen Hsiao MD<sup>1,5</sup>, Pei-Min Hsieh, MD<sup>3,4</sup>, Yu-Chan Li<sup>1,5</sup>, Yaw-Sen Chen MD<sup>3,5</sup>, Chih-Wen Lin, MD, PhD<sup>1,2,5</sup> <sup>1</sup>Division of Gastroenterology and Hepatology, E-Da Dachang Hospital, I-Shou University, Kaohsiung; <sup>2</sup>Division of Gastroenterology and Hepatology, Department of Medicine, E-Da Hospital, I-Shou University, Kaohsiung; <sup>3</sup>Department of Surgery, E-Da Hospital, I-Shou University, Kaohsiung; <sup>4</sup>Department of Surgery, E-Da Cancer Hospital, I-Shou University, Kaohsiung; <sup>5</sup>School of Medicine, College of Medicine, I-Shou University, Kaohsiung; Taiwan.

## **INTRODUCTION METHOD** Study flowchart for the inclusion of participants Hepatitis B virus (HBV) infection and alcoholism are risk factors for hepatocellular • We followed these patients carcinoma (HCC) and mortality. Cirrhotic patients at E-Da Hospital, Kaohsiung Chang Gung Memorial Hospital, until June 2021. The synergistic effect of alcohol intake and HBV infection on clinical prognosis has and Cathay General Hospital from January 2005 to December 2020 (n=5168) • All patients were followed up been reported. Exclude for more than 6 months. Our previous study demonstrated that heavy alcohol consumption promoted the -HCV-positive (n=1302) -Alcohol intake of < 80 g/day and • Heavy alcohol intake was incidence of HCC in HBV-related cirrhosis and that antiviral therapy reduced the risk for <5 years (n=2096) -Patients with HCC at inclusion (n=219) defined as consuming more of HCC in cirrhotic patients with HBV infection and alcoholism. -Data incomplete (n=36) than 80g of ethanol each day About 45% of Han Chinese (East Asians) carry a single point mutation in aldehyde Cirrhotic patients with HBV infection for at least 5 years. or/and heavy alcoholism (n=1515) dehydrogenase 2\*2 (ALDH2). • The primary endpoint was The ALDH2 polymorphism is an effect of the development of HCC in alcoholic liver • newly developed HCC and Cirrhotic patients with Cirrhotic patients with HBV Cirrhotic patients with disease. HBV infection alone heavy alcoholism alone infection and heavy alcoholism the secondary endpoint was (n=342) (n=796) (n=377) AIMS overall mortality after more This study aims to investigate the impact of heavy alcohol intake, ALDH2 polymorphism than six months of follow-up. 746 of 1515 patients were prospectively enrolled and and HBV infection on the clinical prognosis of cirrhotic patients blood samples were collected for ALDH2 polymorphism RESULTS Table 1 Demographic data of all cirrhotic patients

ruble 1. Demographie data of an entitotic patients											
Characteristics	Total cohort (n=1515)	HBV + alcoholism (n =342)	HBV (n =796)	Alcoholism $(n = 377)$	P-value*						
Gender (male)	1277 (84.3)	314 (91.8)a	647 (81.3)	316 (83.8)c	<0.001						
Age (years)	49 (21-77)	44 (21-72)a	49 (30-77)	49 (30-75)c	< 0.001						





BMI (kg/m <sup>2</sup> )	23.6 (16.8-41.9)	25.1 (18.5-41.9)a	23.3 (16.8-37.3)	23.3 (16.8-40.8)c	< 0.001				
Alcohol intake (g/day)	12 (0-600)	160 (80-350)a	0 (0 <b>-</b> 20)b	150 (80-600)c	< 0.001				
Alcohol intake duration (years)	0 (0-30)	20 (5-30)a	0 (0-5)b	16 (5-30)	<0.001 <0.001				
Abstinence	408 (56.7)	171 (50)	N/A	237 (62.9)c					
ALDH2 polymorphisms									
GG	329 (44.1)	157 (55.4)a	81 (33.1)b	81 (44.0)c	< 0.001				
GA/AA	417 (55.9)	137 (44.6)	164 (66.9)	116 (56.0)					
AST (IU/L)	110 (18-1133)	86 (18-1133)a	114 (18-491)	113 (24-491)c	< 0.001				
ALT (IU/L)	43 (10-488)	45 (13-278)	43 (10-488)	41 (10-252)	0.706				
Total bilirubin (mg/dL)	2.1 (0.3-37.9)	1.7 (0.3-19.9)a	2.1 (0.4-37.9)	2.1 (0.5-37.9)c	< 0.001				
Alkaline phosphatase (IU/L)	298 (33-1484)	311 (88-1484)a	275 (60-889)b	324 (33-1092)c	< 0.001				
γ-GT (IU/L)	214 (11-1992)	179 (14-1992)	213 (11-1240)	287 (11-1240)c	0.71				
Albumin (g/dL)	3.2 (1.8-4.8)	3.2 (2.0-4.8)	3.2 (1.8-4.8)	3.3 (1.8-4.6)	0.58				
Platelet count (x10 <sup>3</sup> /mL)	92 (33-633)	76 (17-286)a	99 (33-633)	94 (33-633)c	< 0.001				
Characteristics	Total cohort (n=1515)	HBV + alcoholism (n =342)	HBV (n =796)	Alcoholism $(n = 377)$	<i>P</i> -value*				
INR	1.2 (0.88-2.9)	1.1 (0.9-2.3)a	1.2 (0.9-2.9)	1.2 (0.9-2.9)	0.043				
α-fetoprotein (ng/mL)	5 (1-866)	6.5 (1-886)a	5 (1-660)	5 (1-660)	0.088				
HBsAg-positive	1138 (75.1)	342 (100)	796 (100)b	0 (0)c	< 0.001				
HBeAg-positive	360 (23.8)	90 (26.3)a	270 (33.9)b	0 (0)c	< 0.001				
Baseline HBV DNA (log <sup>10</sup>									
IU/mL)	3.8 (0-9.0)	4.1 (0-9.0)	5.3 (0-9.0)b	0 (0)c	< 0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL)	3.8 (0-9.0) 449 (29.6)	4.1 (0-9.0) 136 (39.8)	5.3 (0-9.0)b 313 (39.3)b	0 (0)c 0 (0)c	<0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive	3.8 (0-9.0)   449 (29.6)   975 (64.5)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b	0 (0)c 0 (0)c 0 (0)c	<0.001 <0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class	3.8 (0-9.0)   449 (29.6)   975 (64.5)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b	0 (0)c 0 (0)c 0 (0)c	<0.001 <0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A	3.8 (0-9.0) 449 (29.6) 975 (64.5) 644 (42.5)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8)	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c	<0.001 <0.001 <0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A B	3.8 (0-9.0) 449 (29.6)   975 (64.5) 553 (36.5)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a 85 (24.9)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8) 316 (39.7)	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c 152 (40.3)c	<0.001 <0.001 <0.001 <0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A B C	3.8 (0-9.0) 449 (29.6) 975 (64.5) 644 (42.5) 553 (36.5) 318 (21)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a 85 (24.9)a 85 (24.7)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8) 316 (39.7) 163 (20.5)	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c 152 (40.3)c 70 (18.6)c	<0.001 <0.001 <0.001 <0.001 <0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A B C Follow up time (year)	3.8 (0-9.0) 449 (29.6) 975 (64.5) 644 (42.5) 553 (36.5) 318 (21) 5.0 (0.5-15)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a 85 (24.9)a 85 (24.7)a 2.8 (0.5-15)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8) 316 (39.7) 163 (20.5) 5.0 (0.5-15)	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c 152 (40.3)c 70 (18.6)c 5.1 (0.5-15)	<0.001 <0.001 <0.001 <0.001 <0.001 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A B C Follow up time (year) HCC development	3.8 (0-9.0) 449 (29.6)   975 (64.5) 553 (36.5)   644 (42.5) 553 (36.5)   318 (21) 50 (0.5-15)   270 (17.8) 51	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a 85 (24.9)a 85 (24.7)a 2.8 (0.5-15)a 81 (23.7)a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8) 316 (39.7) 163 (20.5) 5.0 (0.5-15) 134 (16.8)	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c 152 (40.3)c 70 (18.6)c 5.1 (0.5-15) 55 (14.6)c	<0.001 <0.001 <0.001 <0.001 <0.001 <0.001 0.004				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A B C Follow up time (year) HCC development Annual HCC incidence (%/year)	3.8 (0-9.0) 449 (29.6)   975 (64.5) 553 (36.5)   644 (42.5) 553 (36.5)   318 (21) 5.0 (0.5-15)   270 (17.8) 3.5	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a 85 (24.9)a 85 (24.7)a 2.8 (0.5-15)a 81 (23.7)a 5.9a	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8) 316 (39.7) 163 (20.5) 5.0 (0.5-15) 134 (16.8) 3.6	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c 152 (40.3)c 70 (18.6)c 5.1 (0.5-15) 55 (14.6)c 2.9c	<0.001 <0.001 <0.001 <0.001 <0.001 <0.001 0.004 <0.001				
IU/mL) Baseline HBV DNA (≥5 log <sup>10</sup> IU/mL) Antiviral NUCs therapy - positive Child-Pugh class A B C Follow up time (year) HCC development Annual HCC incidence (%/year) Mortality	3.8 (0-9.0) 449 (29.6)   975 (64.5) 5   644 (42.5) 5   553 (36.5) 3   318 (21) 5   5.0 (0.5-15) 2   270 (17.8) 3   3.5 627 (41.4)	4.1 (0-9.0) 136 (39.8) 308 (90.1)a 172 (50.3)a 85 (24.9)a 85 (24.7)a 2.8 (0.5-15)a 81 (23.7)a 5.9a 155 (45.3)	5.3 (0-9.0)b 313 (39.3)b 667 (83.8)b 317 (39.8) 316 (39.7) 163 (20.5) 5.0 (0.5-15) 134 (16.8) 3.6 322 (40.5)	0 (0)c 0 (0)c 0 (0)c 155 (41.1)c 152 (40.3)c 70 (18.6)c 5.1 (0.5-15) 55 (14.6)c 2.9c 150 (39.8)	<0.001 <0.001 <0.001 <0.001 <0.001 <0.001 0.004 <0.001 0.238				



Characteristics	ALDH2 polymorphisms	Total cohort (n=746)	HBV + alcoholism (n =294)	HBV (n =245)	Alcoholism (n = 207)	P- value*
Genotypes	GG	329 (44.1)	157 (55.4)a	81 (33.1)b	81 (44.0)c	< 0.001
	GA/AA	417 (55.9)	137 (44.6)	164 (66.9)	116 (56.0)	
HCC development	GG	36 (10.9)	7/157 (4.5)a	23/81 (28.4)b	6/91 (6.6)c	<0.001
	GA/AA	147 (35.3)	66/137 (48.2)	50/164 (30.5)	31/116 (26.7)	
	HR (95% CI)	12.6 (5.80- 27.6)	10.5 (4.80- 22.8)	1.13 (0.68- 1.91)	4.68 (1.94- 11.2)	
	P-value#	<0.001	<0.001	0.62	0.001	
Mortality	GG	109 (33.1)	51/157 (42.5)a	36/81 (44.4)b	22/91 (24.2)c	< 0.001
	GA/AA	250/417 (60.0)	91/137 (66.4)	82/164 (50.0)	77/116 (66.4)	
	HR (95% CI)	1.46 (1.04- 2.05)	1.55 (1.09- 2.19)	0.99 (0.67- 1.47)	3.01 (1.87- 4.86)	
	P-value#	0.027	0.012	0.969	<0.001	



alcoholism than in those with HBV infection alone or alcoholism alone after after propensity score matching.

The cumulative incidences of HCC and mortality according to alcohol intake with the ALDH2 polymorphism and serum HBV DNA levels with antiviral NUC therapy.

Patients with the GA/AA allele with alcohol intake >160 g/day were remarkably correlated with increased incidences of HCC and mortality compared with patients with the GG allele with alcohol intake 80-160 g/day (A and C).

Patients with high serum HBV DNA without antiviral NUC therapy were remarkably correlated with increased incidences of HCC and mortality compared with patients with low serum HBV DNA with antiviral NUC therapy (B and D).

a: P-value < 0.05, HBV and alcoholism vs. HBV; b: P-value < 0.05, HBV and alcoholism vs. alcoholism;</p> c: P-value < 0.05, HBV vs. alcoholism; P-value is used by Chi-squared tests. \*: P-value is used by one-way ANOVA test among three groups. #: P-value is used by Cox regression analyses.

Defense 1	Year of follow-up											Defects of the	Year of follow-up																				
Patients at risk	Patients at risk   0   1   2   3   4   5   6   7   8   9   10   11   12   13   14   15	0	1	2	3	4	5	6	7	8	9	10	11	12	13	14	1																
HBV DNA (≥5 log <sup>10</sup> IU/mL) without antiviral therapy	136	111	73	66	57	45	26	18	13	7	2	0	0	0	0	0	HBV DNA (≥5 log <sup>10</sup> IU/mL) without antiviral therapy	136	123	83	72	63	57	33	20	13	10	3	1	1	0	0	(
HBV DNA (≥5 log <sup>10</sup> IU/mL) with antiviral therapy	34	21	13	10	10	9	3	2	2	2	0	0	0	0	0	0	HBV DNA (≥5 log <sup>10</sup> IU/mL) with antiviral therapy	34	21	16	13	13	13	11	5	2	2	0	0	0	0	0	(
HBV DNA (<5 log <sup>10</sup> IU/mL) with antiviral therapy	172	124	99	81	72	69	53	37	25	23	13	7	5	3	1	1	HBV DNA (<5 log <sup>10</sup> IU/mL) with antiviral therapy	172	129	101	82	73	67	51	37	25	22	14	7	5	4	3	ļ

The cumulative incidences of HCC (A) and mortality (B) were higher in cirrhotic patients with concomitant HBV infection and alcoholism than those with HBV infection alone or alcoholism alone.

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## **CONCLUSION**

- **Cirrhotic patients with concomitant HBV infection and heavy** alcoholism had significantly higher incidences of HCC development and mortality than those with HBV infection alone or alcoholism before and after PSM.
- Heavy alcohol intake and ALDH2 rs671 polymorphism significantly ٠ increased the incidence and risk of HCC and mortality in HBVrelated cirrhotic patients.

 $\diamond$ **Financial disclosure:** This study was partially supported by EDAHP109044  $\cdot$ EDAHP110036 · EDAHP111025.