Anti-apoptotic N-cadherin signaling and its prognostic implication in human hepatocellular carcinomas

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Abstract. N-cadherin signaling has recently been implicated in the progression of certain epithelial tumors by promoting invasion and dissemination of cancer cells. N-cadherin has also been reported to exert an anti-apoptotic effect. In this study, we attempted to evaluate the participation of this adhesion molecule in the progression of human hepatocellular carcinomas (HCCs) by analyzing its anti-apoptotic signaling as well as its prognostic implication in HCC patients. N-cadherin was found to be expressed in human HCCs. We established a stable human HCC cell line expressing a truncated N-cadherin, NCad\(Delta\)C, with a dominant-negative action. NCad\(Delta\)C-expressing cells were more susceptible to bile acid-induced apoptosis than control cells. N-cadherin was found to complex with procaspase-8, and this association was diminished in NCadΔC-expressing cells, leading to enhanced procaspase-8 recruitment to death-inducing signaling complex following bile acid treatment. A clinicopathological analysis in patients who had undergone surgical resection for HCC revealed that tumoral N-cadherin up-regulation was significantly related to poor recurrencefree and overall survival. Our findings implicate N-cadherin signaling as contributing to HCC progression by exerting anti-apoptotic effects. Thus, we suggest that the selective interruption of this signaling may have therapeutic potential.

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Abbreviations: HCC, hepatocellular carcinoma; DC, deoxycholate; DAPI, 4',6-diamidino-2-phenylindole dihydrochloride; FADD, Fas associated death domain; RFS, recurrence-free survival; OS, overall survival; DISC, death-inducing signaling complex

Key words: N-cadherin, hepatocellular carcinoma, apoptosis, caspase-8, recurrence, survival

Introduction

Hepatocellular carcinoma (HCC) development is a major cause of mortality in chronic liver disease patients. Massforming or even locally advanced HCCs can be efficiently eradicated by surgical resection or locoregional therapies. However, the long-term prognosis for these patients remains unsatisfactory due to high rates of tumor recurrence, and there is therefore a need to understand the clinicopathological features of HCCs in the broadest sense. In recent years, the development of cellular and molecular biological techniques has allowed extensive studies to be conducted on many factors potentially related to cancer cell invasion, metastasis, recurrence and patient survival. These include alterations in DNA ploidy, cellular proliferation markers, tumor suppressor and promoter genes, cell cycle regulators, proteinases that degrade the extracellular matrix, angiogenic factors, and adhesion molecules (1).

Alterations in the cadherin/catenin intercellular adhesion system are known to be involved in the invasiveness and metastatic behavior of certain tumors. E-cadherin is representative of the classic cadherins, and its downregulation has frequently been related to invasiveness, metastasis, and a poor prognosis in a variety of human cancers, including HCC (2,3). On the other hand, β-catenin, which was originally identified as a submembrane component in the cadherin-mediated cell-cell adhesion system, functions as a downstream transcriptional activator of the Wnt signaling pathway and forms a complex with the DNA-binding proteins, T-cell factor and lymphoidenhancer factor 1 (4). Moreover, unlike E-cadherin, β-catenin expression correlates positively with HCC grade and is highest in poorly differentiated HCCs (2).

In addition to E-cadherin/β-catenin deregulation, N-cadherin signaling has also been shown to participate in the progression of certain epithelial tumors (5,6). N-cadherin expression is usually observed in neural tissue and fibroblasts (7-9), where it is believed to mediate a less stable and more dynamic form of cell-cell adhesion (10). Therefore, aberrant N-cadherin expression in tumors may lead to the acquisition of a migrating and invasive phenotype by cancer cells. In addition to promoting cancer cell invasion, it has also been suggested that N-cadherin signaling may contribute to the progression

of tumors by exerting an anti-apoptotic effect (11). However, the participation of N-cadherin signaling in the progression of human HCC has not yet been clarified.

In this study, we hypothesized that anti-apoptotic N-cadherin signaling is involved in HCC progression. To test this hypothesis, we formulated the following questions: i) Is N-cadherin expressed in human HCCs? ii) Does N-cadherin signaling affect cellular apoptosis? If so, iii) what is the anti-apoptotic mechanism mediated by N-cadherin? And finally, iv) does N-cadherin signaling have prognostic implications in HCC patients? Collectively, the results of the current study demonstrate that N-cadherin signaling is involved in HCC cell apoptosis, and tumoral N-cadherin up-regulation is associated with tumor recurrence and a poor outcome in HCC patients following surgical resection. Therefore, our findings implicate N-cadherin signaling during HCC progression, and suggest that interrupting this signaling may have therapeutic potential.

Materials and methods

Cell culture and reagents. Huh-BAT cells (Huh-7 cells, a well-differentiated human HCC cell line, stably transfected with a bile acid transporter) were grown in DMEM supplemented with 10% fetal bovine serum, penicillin 100,000 U/l, and streptomycin 100 mg/l (12). Two other human HCC cell lines with poorly differentiated phenotypes, SNU-475 (13) and SNU-761 (14), were grown in RPMI-1640 supplemented with 10% fetal bovine serum, 100,000 U/l penicillin and 100 mg/l streptomycin. In all experiments performed in this study, cells were serum-starved overnight to avoid the effects of serum-induced signaling. Deoxycholate (DC) was obtained from Sigma Chemicals Co. (St. Louis, MO).

Plasmids. The NCadΔC construct, cloned in the pSP72 (pSP72NCadΔC) vector, was obtained from Dr Jeffery Gordon (Washington University, St. Louis, MO) with the permission of Dr Chris Kintner (Salk Institute, San Diego, CA). This construct, derived from an in-frame deletion of most of the extracellular domain of Xenopus N-cadherin, has been shown to function as a dominant negative cadherin (15,16). A hemagglutinin epitope was incorporated at the C-terminal of NCadΔC by PCR, then ligated into the Bg/III/EcoRI site of pMSCV-IRES-GFP retroviral vector (kind gift from Neil A. Clipstone, Northwestern University, Chicago, IL) upstream of IRES, giving pMSCV-NCadΔC-IRES-GFP.

Retrovirus generation and transduction. For the transient generation of VSV-G pseudo-typed retrovirus, 293T cells were transfected with pMD-gag-pol, pMD-VSVG [both obtained from Dr Richard C. Mulligan (Harvard Medical School, Boston, MA)] and the retroviral vector, pMSCV-NCad Δ C-IRES-GFP or pMSCV-IRES-GFP, using LipofectAmine Plus reagents (Invitrogen, Carlsbad, CA). Huh-BAT cells were transduced with virus-containing supernatants in the presence of 5 μ g/ml polybrene for 5 h. Cells were collected 48-72 h later, and GFP-positive fractions were FACS-sorted using a BD FACS Vantage Cell Sorter (Franklin Lakes, NJ).

Apoptosis. Apoptosis was assessed by examining the nuclear changes associated with apoptosis (i.e. chromatin condensation

Table I. The baseline characteristics of HCC patients.

	*
Variable	n (%)
Age (years)	
<50	31 (37)
≥ 50	53 (63)
Sex	
Male	74 (88)
Female	10 (12)
Child class	
A	82 (98)
В	2 (2)
Viral hepatitis	
В	60 (71)
C	11 (13)
B+C	4 (5)
NBNC	9 (11)
UICC T	
T1	59 (70)
T2/3	25 (30)
Multiplicity	
Yes	6 (7)
No	78 (93)
Vascular invasion	
Yes	23 (27)
No	61 (73)
Grade	
Low (I-II)	47 (56)
High (III-IV)	37 (44)

and nuclear fragmentation) using the nuclear binding dye, 4',6-diamidino-2-phenylindole dihydrochloride (DAPI), and fluorescence microscopy (Zeiss, Germany).

Immunoblotting. Cells were lysed for 20 min on ice with lysis buffer (50 mM Tris-HCl, pH 7.4; 1% Nonidet P-40; 0.25% sodium deoxycholate; 150 mM NaCl; 1 mM EDTA; 1 mM PMSF; 1 µg/ml aprotinin, leupetin, pepstatin; 1 mM Na₃VO₄; and 1 mM NaF) and centrifuged at 14,000 x g for 10 min at 4°C. Samples were resolved with 10% or 12% SDS-PAGE, transferred to PVDF membranes, and blotted using appropriate primary antibodies and peroxidase-conjugated secondary antibodies (Biosource International, Camarillo, CA). Bound antibodies were visualized using chemiluminescent substrate (ECL; Amersham, Arlington Heights, IL) and exposed to Kodak X-OMAT film. The primary antibodies used were mouse anti-N-cadherin from BD Transduction Laboratories (San Jose, CA); rabbit anti-caspase-10 from Cell Signaling Technology, Inc. (Beverly, MA); rabbit anti-caspase-8, rabbit anti-caspase-3 and mouse anti-Fas-associated death domain (FADD) from Pharmingen (San Diego, CA); goat anti-TRAIL-

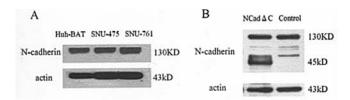


Figure 1. Establishment of an HCC cell line expressing dominant-negative N-cadherin. (A) Cell lysates of human HCC cell lines, i.e. Huh-BAT, SNU-475 and SNU-761 cells, were immunoblotted with anti-N-cadherin and anti-actin antibodies. (B) Huh-BAT cells were infected with either a retroviral vector pMSCV-NCad Δ C-IRES-GFP encoding NCad Δ C (a truncated N-cadherin with a dominant-negative action) or a control vector. NCad Δ C and control cell lysates were immunoblotted with anti-N-cadherin and anti-actin antibodies.

R2/DR5 and rat anti-cFLIP from Alexis (San Diego, CA); goat anti-bid from R&D Systems (Minneapolis, MN); and rabbit anti-Bax, mouse anti-Bcl-2 and goat anti-actin from Santa Cruz Biotechnology Inc. (Santa Cruz, CA).

Immunoprecipitation. Cells were lysed in lysis buffer as described above, and the lysates were incubated with either mouse anti-caspase-8 (Upstate USA, Inc., Chicago, IL) or mouse anti-FADD antibody (Pharmingen) overnight at 4°C. Immune complexes were precipitated with 50 μl of protein A/G Plus agarose beads (Santa Cruz Biotechnology Inc.) for 2 h at 4°C and then washed five times with lysis buffer. Polypeptides were eluted from the beads by boiling for 5 min in 2X Laemmli sample buffer, and the immunoprecipitates were then immunoblotted with anti-N-cadherin, anti-caspase-8, anti-TRAIL-R2/DR5 or anti-FADD antibody.

Patients. Eighty-four patients (74 men and 10 women; mean age, 53±10 years) who had undergone surgical resection at Seoul National University Hospital between June 1994 and December 1998 were retrospectively included in this study. Baseline characteristics of the study population are detailed in Table I. During follow-up periods of 1-108 months (median 48 months), HCC recurred in 58 (69%) patients [median recurrence-free survival (RFS) time following resection, 22 months; 95% confidence interval (CI), 13-31 months], and 52 (62%) patients died [median overall survival (OS) time following resection, 48 months; 95% CI, 39-57 months]. Causes of death were HCC progression in 25 cases, hepatic failure in 13 cases, gastrointestinal bleeding in 2 cases, spontaneous bacterial peritonitis in 1 case, hepatorenal syndrome in 1 case, other causes in 7 cases, and unknown causes in 3 cases. The study protocol was approved by the Seoul National University Hospital Institutional Review Board.

Tissue array and immunohistochemistry. HCC specimens from each patient were processed into 10% neutral formalin-fixed, paraffin-embedded blocks. All tumors were histologically diagnosed, graded according to Edmonson's scale, and grouped as low (I-II) or high grade (III-IV). Tumors were also stratified according to multiplicity and the presence or absence of microscopic vascular invasion. These pathological findings are summarized in Table I.

For immunohistochemical staining, all specimens were evaluated using a tissue-array method. Core tissue biopsies

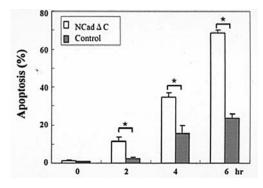


Figure 2. N-cadherin-dependent modulation of HCC cell apoptosis. NCad Δ C and control cells were incubated with deoxycholate (200 μ M) for the indicated times. Apoptosis was quantitated by DAPI staining and fluorescence microscopy. Data are expressed as the means \pm SD of three individual experiments. *p<0.050, vs. control.

(2 mm in diameter) were taken from individual paraffinembedded tissues (donor blocks) and arranged in a new recipient paraffin block (tissue array block) using a trephine apparatus (Superbiochips Laboratories, Seoul, Korea). Immunohistochemical staining was performed using E-cadherin, \(\beta\)-catenin (BD Transduction Laboratories) and N-cadherin (Zymed Laboratories Inc., San Francisco, CA) antibodies, and using a streptavidin peroxidase-based procedure after microwave antigen retrieval. The intensity of immunoreactivity in each case was scored as none to weak, moderate, or strong, and the extent of immunoreactivity was scored as less than one-third, one-third to two-thirds, or more than two-thirds. The histological examination was carried out by a pathologist (J.J. Jang) who was blinded to any clinical information.

Data analysis. All in vitro experimental data represent at least three independent experiments using cells from a minimum of three separate isolations and are expressed as means ± SD. Differences between groups were compared using the two-tailed Student's t-test. RFS and OS were estimated using the Kaplan-Meier method for the clinicopathological variables of age, sex, viral hepatitis, child classification, multiplicity, vascular invasion, stage and grade, or the biological markers of E-cadherin, β-catenin, and N-cadherin, and were compared using the log-rank test. To identify independent prognostic factors affecting RFS and OS, multivariate analysis was performed using the Cox proportional hazard model for variables with p-values <0.100 by univariate analysis. More generous p-value criteria were used to include characteristics with a marginal association. All statistical analyses were performed using SPSS version 12.0 (SPSS, Chicago, IL). p-values <0.050 were considered statistically significant.

Results

Establishment of a human HCC cell line expressing dominantnegative N-cadherin. Initially, we examined N-cadherin expression in human HCC cell lines. As shown in Fig. 1A, N-cadherin was expressed in all three human HCC cell lines examined, including one well-differentiated and two poorly

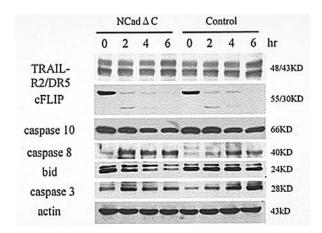


Figure 3. Potentiation of apoptosis in HCC cells expressing dominant-negative N-cadherin. NCad Δ C and control cells were incubated with deoxycholate (200 μ M) for the indicated times. Cell lysates were immunoblotted with each indicated antibody.

differentiated. Among these, the Huh-BAT cell line was selected for the establishment of dominant-negative N-cadherin-expressing cells, since these cells are susceptible to bile acid-induced apoptosis (12). NCad Δ C, a truncated N-cadherin mutant lacking extracellular domains, was found to prevent the lateral clustering and dimerization of endogenous N-cadherins, and thus inhibit the formation of the adhesion complex (17,18). Thus, we established an NCad Δ C-expressing cell line, i.e. Huh-BAT cells expressing a truncated (45 kDa) N-cadherin, following transduction of retroviruses containing the Ncad Δ C construct (Fig. 1B).

Does the inhibition of N-cadherin signaling affect HCC cell survival? We next evaluated the effect of N-cadherin signaling on HCC cell survival by comparing apoptosis in Ncad Δ C and control cells. For this purpose, cellular apoptosis was induced by treating these cells with bile acid. As shown in Fig. 2, NCad Δ C cells were more susceptible to bile acid-induced apoptosis than control cells, suggesting the antiapoptotic function of N-cadherin signaling.

We further explored the nature of N-cadherin antiapoptotic signaling by comparing bile acid-induced apoptotic signaling cascades in NCad∆C and control cells. Bile acid-induced TRAIL-R2/DR5, c-FLIP and caspase-10 modulations

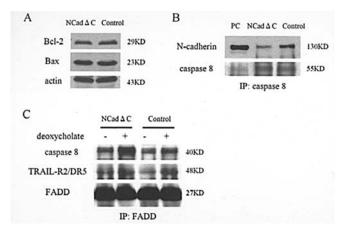


Figure 4. N-cadherin-dependent anti-apoptotic signaling. (A) Cell lysates of NCad Δ C and control cells were immunoblotted with anti-Bcl-2, anti-Bax, and anti-actin antibodies. (B) Caspase-8 was immunoprecipitated from both cell lysates. The immunoprecipitates were immunoblotted using anti-N-cadherin and anti-caspase-8 antibodies. PC, positive control. (C) NCad Δ C and control cells were treated with deoxycholate (200 μ M) or media for 2 h. FADD was immunoprecipitated from cell lysates, and immunoblotted using each indicated antibody.

were similar in both cell lines, whereas caspase-8 and its downstream signal activation were significantly enhanced in NCad Δ C cells (Fig. 3). The expression levels of Bcl-2 and Bax were similar in both cell lines (Fig. 4A). However, N-cadherin formed a complex with procaspase-8, and this association was diminished in NCad Δ C cells versus the control (Fig. 4B). Accordingly, bile acid-induced procaspase-8 recruitment to FADD was significantly enhanced in NCad Δ C cells versus the control (Fig. 4C). Therefore, these results collectively demonstrate that N-cadherin has an anti-apoptotic function in HCC cells, which involves complexing with procaspase-8, and reduced procaspase-8 recruitment to death-inducing signaling complex (DISC).

Does N-cadherin expression in HCC predict prognosis? We finally investigated whether tumoral N-cadherin expression is associated with HCC recurrence or postoperative survival in HCC patients. Of 84 HCC specimens examined, 31, 33 and 20 showed none to weak, moderate, or strong N-cadherin immunoreactivity at cell membranes, respectively. Fig. 5 shows representative cases.

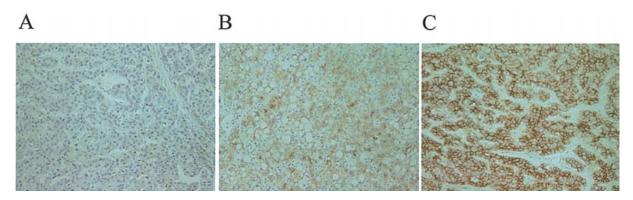


Figure 5. N-cadherin immunoreactivity in human HCC tissue. Representative immunohistochemical staining of (A) none to weak, (B) moderate, and (C) strong N-cadherin immunoreactivity in human HCC tissue. Original magnification, x 200.

Table II. Univariate analysis of clinicopathological and biological variables with respect to RFS and OS.

	RFS		OS			
	n	(mo ^a)	p-value	(mo ^a)	p-value	
Age (years)		-0				
<50	31	29	0.475	43	0.818	
≥ 50	53	22		50		
Sex						
Male	74	21	0.570	48	0.762	
Female	10	33		53		
Child class						
A	82	23	0.163	50	0.864	
В	2	9		19		
Viral hepatitis						
В	60	26	0.680	48	0.753	
C	11	22		104		
В+С	4	14		39		
NBNC	9	20		53		
UICC T						
T1	59	33	0.031	53	0.039	
T2/3	25	7		24		
Multiplicity						
Yes	6	9	0.633	19	0.192	
No	78	26	0.022	51	0.172	
E-cadherin						
intensity						
None to weak	32	38	0.139	50	0.710	
Moderate Moderate	31	20	0.137	44	0.710	
Strong	21	18		56		
_						
E-cadherin extent						
<1/3	32	18	0.237	46	0.420	
$\geq 1/3, <2/3$	21	17	0.237	46	0.420	
$\geq 1/3, \langle 2/3 \rangle$ $\geq 2/3$	31	33		104		
	31	33		101		
N-cadherin						
intensity	21	41	0.014	104	0.000	
None to weak Moderate	31 33	41 20	0.014	104 50	0.090	
Strong	20	12		39		
_	20	12		37		
N-cadherin						
extent	25	27	0.214	40	0.064	
<1/3	35	27 7	0.214	48 44	0.864	
$\geq 1/3, <2/3$ $\geq 2/3$	26 23	34		65		
	23	JĦ		03		
ß-catenin						
Name to week	20	<i>E</i> 1	0.041	02	0.424	
None to weak Moderate	20 40	51 18	0.041	83 46	0.434	
Strong	24	20		48		
Strong	∠4	20		40		

Table II. Continued.

	RFS			OS		
	n	(mo ^a)	p-value	(mo ^a)	p-value	
ß-catenin						
extent						
<1/3	24	29	0.403	53	0.821	
$\geq 1/3, <2/3$	31	14		44		
≥ 2/3	29	33		48		
Vascular						
invasion						
Yes	23	7	0.164	24	0.110	
No	61	27		51		
Grade						
Low (I-II)	47	21	0.353	53	0.181	
High (III-IV)	37	26		43		
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^aMedian survival time in months.

Univariate analysis indicated that RFS significantly correlated with UICC T stage and N-cadherin and \(\beta\)-catenin intensities (p=0.031, 0.014 and 0.041, respectively), whereas OS was significantly associated with UICC T stage (p=0.039) (Table II). Multivariate analysis employing parameters that showed p<0.100 by univariate analysis revealed that both UICC T stage and N-cadherin intensity independently predicted RFS (p=0.025 and 0.018, respectively) as well as OS (p=0.013 and 0.034, respectively) (Table III). In the UICC T1 and T2/3 groups, median RFS times were 33 and 7 months, respectively (Fig. 6A), and median OS times were 53 and 24 months, respectively (Fig. 6B). In the none to weak, moderate, and strong N-cadherin intensity groups, median RFS times were 41, 20, and 12 months, respectively (Fig. 7A), and median OS times were 104, 50, and 39 months, respectively (Fig. 7B). No significant association was found between N-cadherin expression and UICC T staging in these patients (p=0.645). Therefore, these results collectively imply that advanced UICC T stage and tumoral N-cadherin up-regulation are independent predictors of poor RFS and OS in HCC patients.

Discussion

The principal finding of this study relates to N-cadherin signaling in HCC. Collectively, our results demonstrate that N-cadherin-dependent signaling attenuates apoptosis in human HCC cells. In addition, tumoral N-cadherin up-regulation was associated with poor RFS and OS in HCC patients following resection. These findings implicate N-cadherin signaling in the progression of HCCs. Each of these findings will be discussed below.

The present study demonstrated N-cadherin expression in cultured HCC cells as well as in resected human HCC tissue. N-cadherin expression has been reported to cause tumor cells to become motile, invasive, and metastatic (5,6). This invasive effect of N-cadherin results, at least in part, from a functional

Table III. Multivariate analysis of factors affecting RFS and OS.

Varia	ble	Odds ratio	95% CI	p-value
RFS	UICC T			0.025
	T1	1.000		
	T2/3	1.930	1.087-3.426	
	N-cadherin intensity			0.018
	None to weak	1.000		
	Moderate	1.727	0.905-3.293	
	Strong	2.792	1.373-5.679	
	β-catenin intensity			NS
OS	UICC T			0.013
	T1	1.000		
	T2/3	2.153	1.179-3.931	
	N-cadherin intensity			0.034
	None to weak	1.000		
	Moderate	1.507	0.773-2.940	
	Strong	2.606	1.264-5.375	

NS, not significant.

interaction with the FGF receptor at the cell membrane (19,20). A complex between N-cadherin and FGFR-1 results in enhanced FGFR tyrosine phosphorylation, and sustained MAPK-ERK activation, leading to MMP-9 gene transcription and hence cellular invasion. N-cadherin is also likely to facilitate the systemic dissemination of tumor cells by enabling these cells to easily associate with stroma or endothelium at distant loci (6,21).

In addition to promoting invasion and dissemination, N-cadherin has been reported to exert an anti-apoptotic effect by recruiting phosphatidylinositol 3-kinase to the N-cadherin/catenin/actin complex and thus increase the Bcl-2/ Bax protein ratio (11). However, in the present study, the expression levels of Bcl-2 and Bax were no different between NCadΔC and control cells. Therefore, the present study describes an additional form of anti-apoptotic N-cadherin related signaling in HCCs. In the present study, apoptosis was induced in HCC cells by treating with bile acid, which mediates its effect by activating death receptor-dependent pro-apoptotic signals. The Huh-BAT cells used in this study do not express Fas, and thus bile acid-induced apoptosis proceeds via the TRAIL-R2/DR5-dependent pathway (12). We observed that bile acid-induced caspase-8 and its downstream signal activation were significantly up-regulated in NCad∆C cells versus control cells. This up-regulated caspase-8 activation is either due to enhanced procaspase-8 cleavage or increased procaspase-8 recruitment to DISC. However, bile acid-induced TRAIL-R2/DR5, c-FLIP and caspase-10 modulation occurred in a similar way in NCadΔC and control cells. Therefore, it is likely that procaspase-8 is more efficiently recruited to DISC in NCad∆C cells versus

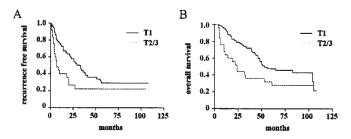


Figure 6. UICC T stage as an independent predictor of RFS and OS in HCC patients. (A) RFS curves according to UICC T stage are shown. Median RFS times were 33 and 7 months for T1 and T2/3 patients, respectively. (B) OS curves according to UICC T stage are shown. Median OS times were 53 and 24 months for T1 and T2/3 patients, respectively.

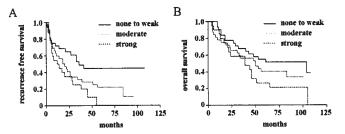


Figure 7. N-cadherin intensity as an independent predictor of RFS and OS in HCC patients. (A) RFS curves according to N-cadherin intensity are shown. Median RFS times were 41, 20, and 12 months in the none to weak, moderate, and strong N-cadherin intensity groups, respectively. (B) OS curves with respect to N-cadherin intensity are shown. Median OS times were 104, 50, and 39 months in the none to weak, moderate, and strong intensity groups, respectively.

control cells. Indeed, the present study demonstrates that N-cadherin complexes with procaspase-8, and this association is diminished more so in NCadΔC cells than in controls. Moreover, bile acid-induced procaspase-8 recruitment to FADD was significantly enhanced in NCadΔC cells. Therefore, these findings collectively imply that N-cadherin signaling exerts an anti-apoptotic function in HCC cells by complexing with procaspase-8, and this is followed by reduced procaspase-8 recruitment to DISC.

Based on this anti-apoptotic activity of N-cadherin, we evaluated whether tumoral N-cadherin expression is associated with tumor recurrence or postoperative survival in HCC patients. In our evaluation of prognostic factors in these patients, we included the clinicopathological variables of age, sex, viral hepatitis, child classification, staging system, multiplicity, vascular invasion and grade, and the cadherin signaling parameters of E-cadherin, \(\beta\)-catenin and N-cadherin expression. Of these, tumoral N-cadherin up-regulation and advanced UICC T stage were found to be independently related to poor RFS and OS by multivariate analysis. Therefore, our findings suggest that N-cadherin up-regulation is an independent prognostic factor in HCC patients.

In conclusion, the current study demonstrates that N-cadherin signaling exerts an anti-apoptotic effect in human HCCs, and thus probably participates in HCC progression. Therefore, we propose that N-cadherin up-regulation in tumor tissue should be considered a prognostic marker of

HCC, and the interruption of this signaling may have therapeutic potential.

Acknowledgements

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