

WNT/β-catenin pathway activation via Wnt1 overexpression and Axin1 downregulation correlates with cadherin-catenin complex disruption and increased lymph node involvement in micropapillary-predominant lung adenocarcinoma

Liang Zhu^{1,2,3}, Shifeng Yang^{1,2,3}, Linfeng Zheng^{1,2,3}, Gu Zhang^{1,2,3}, Guoping Cheng^{1,2,3}

¹Institute of Cancer and Basic Medicine (ICBM), Chinese Academy of Sciences, Hangzhou, China; ²Department of Pathology, Cancer Hospital of the University of Chinese Academy of Sciences, Hangzhou, China; ³Department of Pathology, Zhejiang Cancer Hospital, Hangzhou, China *Contributions:* (I) Conception and design: L Zhu, G Cheng; (II) Administrative support: G Zhang; (III) Provision of study materials or patients: S Yang; (IV) Collection and assembly of data: S Yang, L Zheng; (V) Data analysis and interpretation: L Zhu, G Cheng; (VI) Manuscript writing: All authors; (VII) Final approval of manuscript: All authors.

Correspondence to: Guoping Cheng. Department of Pathology, Cancer Hospital of the University of Chinese Academy of Sciences, Hangzhou 310022, China. Email: chenggp@zjcc.org.cn.

Background: Micropapillary-predominant adenocarcinoma (MPA) of the lung is associated with extensive lymph node involvement and rapid terminal metastasis. However, this subtype has been recognized for only a few years, and there have been few studies of the molecular mechanisms associated with its highly invasive behaviors.

Methods: The present study utilized immunohistochemical staining of surgically resected tissue blocks of MPA and lepidic-predominant lung adenocarcinoma to quantify the expression of specific biological markers in the WNT/ β -catenin pathway and evaluate their influence on the lymph nodes invasion of these two types of lung adenocarcinomas.

Results: Our findings revealed that disruption of the cell membrane cadherin-catenin complex, which weakens the tumor cell adherence of MPA, was caused by the dissociation of β -catenin from the cadherin-catenin complex and the subsequent accumulation of β -catenin in the cytoplasm. This caused abnormal activation of the WNT/ β -catenin pathway. We also found that Wnt-1-specific overexpression and Axin1 inhibition in MPA could explain the redistribution and cytoplasmic retention of β -catenin. Collectively, these findings suggest that an abnormality in the WNT/ β -catenin pathway could enhance the invasiveness of MPA through the overexpression of Wnt-1 and downregulation of Axin1 molecules.

Conclusions: Our data support the need for further research regarding the WNT/ β -catenin pathway and the need to develop novel targeted therapies for restoration of tumor cell adherence and improvement of the prognosis of MPA.

Keywords: Micropapillary-predominant adenocarcinoma (MPA); WNT/β-catenin; Axin1; lymph node invasion

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Introduction

Morbidity and mortality from non-small cell lung cancer (NSCLC), which constitutes approximately 80% of all lung cancers, have been increasing steadily since the 1930s

(1,2). In recent years, morbidity from lung adenocarcinoma has continued to increase in the Chinese population. Lung adenocarcinoma now become the most prevalence one within the NSCLC (3). Most fatal event of lung adenocarcinoma remains to be the terminal metastasis in

the late stage of the disease (4).

Micropapillary-predominant adenocarcinoma (MPA) was first described by Silver and Askin in 1997 (5). Notable aggressive biological behaviors, including lymph node involvement and terminal metastasis, are common in this particular subtype. MPA is a poorly differentiated subtype of lung adenocarcinomas. Patients who have MPA, or only have micropapillary components in their lesions, will experience poorer clinical outcomes (6). Histological identification of MPA depending on its unique morphology under the microscopy, which has been defined by the International Association for the Study of Lung Cancer, the American Thoracic Society, and the European Respiratory Society (7).

Activation of the WNT signaling cascade is a critical molecular event in the embryonic development of mammals, it plays a role in hemostatic processes in tissues and stem cell regulation (8,9). Thus, this molecular mechanism can accelerate tissue development and regeneration. However, it can also be utilized by cancer cells to promote uncontrollable cancer growth within the host bodies and lead to development of malignant lesions, such as leukemia, liver cancer, endometrial cancer, osteosarcoma, and lung cancer (10-13).

The WNT/ β -catenin pathway is regarded as the canonical WNT signaling pathway. It is triggered when Wnt-family proteins binding to specific cell membrane receptors, like Frizzled family of seven-transmembrane domain receptors and the low-density lipoprotein-receptorrelated proteins (14). Combination of these molecules can inhibit the axis inhibition protein 1 (Axin1) complex, which is competent in β -catenin degradation, resulting in cytosol accumulation and nuclear translocation of β-catenin. Nuclear β-catenin can heterodimerization with transcription factors of the lymphoid enhancer-binding factor/T cell factor family and upregulate specific oncogenes (e.g., Cyclin-D1 and c-Myc) (15). Recent studies demonstrated that enhanced WNT/β-catenin signaling activity in breast, liver, and colorectal cancers were positively correlated with epithelial-to-mesenchymal transition (EMT) status and terminal metastasis (16-18).

The β -catenin destruction complex, as described above, plays a key role in determining the fate of β -catenin. This complex is composed of the scaffold protein Axin1 and adenomatous polyposis coli (APC) together with some kinases, including glycogen synthase kinase 3 β (GSK-3 β) and casein kinase 1 α (CK-1 α) (19). In the absence of Wnt ligand protein, cytoplasmic β -catenin can be phosphorylated by this complex, then undergo degradation (20). When Wnt/ β -catenin pathway is activated, the receptors are phosphorylated and inactivate the β -catenin destruction complex, thereby aiding in the accumulation and nuclear translocation of β -catenin (19).

In our study, we found that in MPA, upregulated Wnt-1 and downregulation of Axin1 can over-activate WNT/ β -catenin signaling pathway and enhance the expression of downstream oncogene cyclinD1. The cadherin-catenin complex was also disrupted. These findings may provide us with a novel perspective on the molecular mechanisms associated with the aggressive biological behavior and poor clinical outcomes of MPA. The WNT/ β -catenin pathway may serve as a molecular target in improved treatments for MPA of the lung. We present the following article in accordance with the MDAR reporting checklist (available at http://dx.doi.org/10.21037/jtd-20-1495).

Methods

Patients and specimens

Twenty cases of lung adenocarcinoma tissue blocks for each MPA group and lepidic group were collected from the Department of Pathology at the Cancer Hospital of the University of Chinese Academy of Sciences between 2015 and 2018, the clinicopathological features of the patients were presented in Table 1. The study was conducted in accordance with the Declaration of Helsinki (as revised in 2013). The study was approved by the ethics committee of Zhejiang Cancer Hospital (No. IRB-2016-90). Written informed consent was provided by the patients from whom the tissue blocks were taken. For each case, hematoxylin and eosin-stained slides were reviewed under a microscope to confirm histological type. A case was classified as micropapillary or lepidic pattern predominant if the indicated component comprised more than 50% of the total neoplastic area.

Reagents and antibodies

Monoclonal primary antibodies against human α -catenin (Cat#2028-1), β -catenin (Cat#1247-1), and E-cadherin (Cat#1702-1) were purchased from Epitomics (Burlingame, CA, USA). Anti-Axin1 (Cat#ab115205), anti-Wnt-1 (Cat#ab15251), anti-Wnt-6 (Cat#ab150588), anti-Wnt10a (Cat#ab106522) and anti-Cyclin-D1 (Cat#ab16663) antibodies were purchased from Abcam (Cambridge, MA,

Table 1 Clinicopathological features of the patients

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Group	Num.
MPA group	
Gender	
Male	11
Female	9
Age (years)	
Male	58 [33–76]
Female	52 [25–79]
Differentiation	Poor
T staging*	
T1	10
T2	7
ТЗ	3
N staging	
NO	9
N1	5
N2	6
M staging	
MO	19
M1	1
Lepidic group	
Gender	
Male	7
Female	13
Age (years)	
Male	61 [36–72]
Female	55 [31-69]
Differentiation	Well
T staging	
T1	18
T2	2
ТЗ	0
N staging	
NO	20
N1	0
N2	0
M staging	-
MO	20
M1	0
	0

 $^{\star},$ the TNM stage were classified according to the WHO Classification of Tumors of the Lung, Pleura, Thymus and Heart (4 $^{\rm th}$ edition, 2015).

USA). The fluorescent conjugated secondary antibody Goat Anti-Rabbit IgG H&L (Alexa Fluor[®] 488) was also obtained from Abcam (ab150081).

Immunobistochemistry staining

Tissue blocks were collected for each case. Paraffin sections (4 um thick) of either tissue or cell blocks were deparaffinized and rehydrated in xylene and a graded alcohol series. Endogenous peroxidase activity was blocked by incubation in 0.3% hydrogen peroxide, and the sections were blocked for 20 minutes with 10% goat serum. The sections were then incubated with primary antibodies including Wnt1 (1:100), Wnt6 (1:200), Wnt10a (1:400), Axin1 (1:100), E-cadherin (1:100), α -catenin (1:100), β -catenin (1:200), and cyclinD1 (1:400) for 90 minutes at room temperature. After slides had been washed in phosphate-buffered saline, the slides were developed using 3,3'-diaminobenzidine (Dako Cytomation, Carpinteria, CA, USA), in accordance with the manufacturer's instructions, and counterstained with hematoxylin. Negative control slides were incubated with phosphate-buffered saline, instead of the primary antibody.

Immunobistochemical staining evaluation

All immunohistochemically stained slides were observed under the microscope. For each slide, hot spots of positive staining were identified at 100× power. Tumor positive scores (TPS) were calculated using 10 high-power phase of view fields (400×) within the hot spots, using the equation TPS = (number of positive tumor cells/number of total tumor cells) ×100%. Positive expression of a protein was defined as reliable and complete membrane or nuclear staining at any intensity when viewed under a magnification of 400×.

Statistical analyses

Comparisons of TPS for the MPA and lepidic groups were performed using the two-tailed unpaired Student's *t*-test. All statistical analyses were performed using GraphPad Prism ver.8.0 (GraphPad Software, La Jolla, CA, USA).

Results

The cadherin-catenin complex was disrupted in MPA

The rate of metastasis and lymph node invasion in MPA

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Figure 1 Disruption of cadherin-catenin complex in MPA of lung. (A) Cancer tissues of the MPA and lepidic groups were applied with H&E staining and IHC staining to show expression or the intracellular distribution of the three key molecules which compose the cadherin-catenin complex (E-cadherin, α -catenin and β -catenin). The pictures were taken under 400× power phase microscopically. Red arrows indicated typical negative tumor cells in the MPA group and the green ones indicated typical positively stained tumor cells in the lepidic group. (B) For each slide from both groups, the TPS of E-cadherin and α -catenin were calculated by counting the positive rate of tumor cells in each 10 views at 400× power phases under microscope. Any convincing membrane staining of the cancer cells were considered to be positive. The data were then analyzed by application of two-tailed unpaired student's *t*-tests. ***, indicated significant difference (P<0.0001). H&E, hematoxylin and eosin; IHC, immunohistochemistry; MPA, Micropapillary-predominant adenocarcinoma; TPS, tumor positive score.

suggested reduced intercellular adhesion, which has been described by other researchers (11,21). This characteristic of MPA prompting us to evaluate the intact of the cadherincatenin complex of the cancer cells. Immunohistochemical staining with anti-E-cadherin (TPS: 31.6% *vs.* 81.9%, P<0.0001) and anti- α -catenin (TPS: 25.3% *vs.* 74.7%, P<0.0001) antibodies indicated diminished protein expression, in terms of both intensity and percentage, in the cell membrane of MPA cases, compared to lepidic cases (*Figures 1A,B,S1,S2*). Concurrently, β -catenin was dissociated from the cell membrane in the MPA group (*Figure 1A*). These data suggested that destruction of the cadherin-catenin complex may have contributed to the poor intercellular adherence of MPA.



Figure 2 Cytoplasmic accumulation of β -catenin enhanced the WNT/ β -catenin signaling activity in the MPA of lung. (A) Cancer tissues of the MPA and lepidic groups were applied with H&E staining and IHC staining with fluorescent conjugated secondary antibody to show the intracellular distribution of β -catenin. The pictures were taken under microscope at 400x and 1,000x (oil lens) power phase. The red arrows indicated interstitial cells. (B) Cyclin-D1 expression in both groups were evaluated by IHC staining to show the activity of WNT/ β -catenin signaling activity in cancer cells. The pictures were taken under microscope at 400x power phase. (C,D) Within both of the groups, cytoplasm positive rates of β -catenin were calculated by counting tumor cells with dramatic cytoplasmic but no clear membrane fluorescent signal (C) and Cyclin-D1 TPS were calculated by counting tumor cells with convincing nuclear staining (D). For each slide, 10 views at 400x power phases were analyzed. The data were then analyzed by application of two-tailed unpaired student's *t*-tests. Three asterisks indicated significant difference (P<0.0001). H&E, hematoxylin and eosin; IHC, immunohistochemistry; MPA, Micropapillary-predominant adenocarcinoma; TPS, tumor positive score.

Cytoplasmic accumulation of β -catenin induced canonical Wnt signaling and may have contributed to cadherincatenin complex disruption in MPA

We then sought to identify the mechanism by which cadherin-catenin complex disruption was initiated in MPA. It has been reported that activation of the Wnt/ β -catenin pathway may favor the degradation of E-cadherin (22). Based on this finding, together with the dissociation of β -catenin from the cell membrane shown in *Figure 1A*, we proceeded to perform immunofluorescent staining of β -catenin to allow observation of the intracellular distribution of β -catenin at high resolution. The results were presented in *Figure 2A,B*. Accumulated cytoplasmic β -catenin was more common in MPA than in the lepidic subtype (77.2% vs. 27.3%, P<0.001). To verify downstream Wnt/ β -catenin signaling activity, the expression of Cyclin-D1, a key product of this pathway, was evaluated in typical components from both groups. As shown in *Figure 2C*, a higher expression rate of Cyclin-D1 was observed in the MPA group (87.9% vs. 35.5%, P<0.001, *Figure 2D*), indicating more active Wnt/ β -catenin signaling activity. In conclusion, cytoplasmic accumulation of β -catenin disrupted the cell membrane cadherin-catenin complex, leading to low intercellular adherence. In contrast, it enhanced Wnt/ β -catenin signaling and increased the production of relevant oncogenes in MPA.

Wnt-1 expression was specifically increased in MPA

Abnormal Wnt/ β -catenin signaling resulting from the overexpression of various Wnt ligands has been reported in NSCLC. However, details of the mechanism by which excessive Wnt/ β -catenin signaling is triggered in MPA have been unclear. We analyzed the expression levels of three canonical Wnt pathway ligands—Wnt-1, Wnt-6, and Wnt-10a—in cancer cells. The results indicated a moderate increase in the percentage of cells with Wnt-1 expression (90.6% *vs.* 71.4%, P<0.05) and a dramatically enhanced staining intensity (*Figure 3*) in the MPA group. In contrast, Wnt-6 (82.3% *vs.* 85.6, P=0.134) and Wnt-10a (18.7% *vs.* 22.4%, P=0.11) were present at similar levels in the two groups (*Figure 3*), indicating that Wnt/ β -catenin signaling in MPA was triggered by a specific type of Wnt molecule.

Axin1 expression was downregulated in MPA, especially in cases with lymph node invasion

Axin1 protein is a scaffold protein in the cytoplasmic β-catenin-destruction complex. Axin1 deficiency has been shown to cause overactive Wnt/β-catenin signaling in various malignant neoplasms. We explored Axin1 expression status in the MPA and lepidic groups. After immunohistochemical staining had been performed, slides from the two groups were evaluated for staining intensity and TPS of Axin1 in the histologically typical components. The results, shown in Figure 4A, indicated that almost all tumor cells in the lepidic group were strongly positive; in the MPA group, the majority of tumor components were only weakly to moderately positive and the percentage with positive expression was smaller than the percentage in the lepidic group (63.5% vs. 95.0%, P<0.001). These findings demonstrate that MPA expressed less Axin1 than the lepidic subtype of lung adenocarcinoma. We also evaluated the TPS of Axin1 in both lymph node invasion-negative and lymph node invasion-positive MPA cases to explore whether Axin1 expression was essential for lymph node invasion. The results are presented in Figure 4B. Axin1 was differentially expressed in the lymph node invasionpositive and lymph node invasion-negative groups (61.46% *vs.* 68.04%, P=0.018). Taken together, these results indicate that diminished Axin1 expression may affect the invasiveness of lung adenocarcinoma, in addition to its histological differentiation.

Discussion

According to the histologic classification of the International Association for the Study of Lung Cancer, American Thoracic Society, and European Respiratory Society, lung adenocarcinoma has five predominant subtypes: lepidic, acinar, papillary, micropapillary, and solid (7,23). The micropapillary subtype behaves more aggressively and has poorer clinical outcomes than the other subtypes. Pathologically, MPA is defined with a tissue structure that is organized in delicate papillary tufts and lacks fibrovascular cores (24). Clinically, it induces higher metastatic rate and more rapid tumor proliferation (25,26). However, because MPA was only recently identified, its biological behavior has not been explained. Our study attempts to explore the mechanism by which MPA derived more invasiveness and we find abnormal Wnt-1 and Axin1 expression in MPA, which may interpret as the reason for elevated activity of the WNT/ β -catenin pathway. These molecular events may promote lymph nodes invasion of MPA in our cases.

In some type of cancers, activation of WNT signaling can coupled with dissociation of β -catenin from the cell membrane and break down the cadherin-catenin complex, which is essential for maintenance of intercellular adherence (9). Our data presented downregulation of E-cadherin and α -catenin on the cell membrane (*Figure 1*). This may explain the low intercellular adhesion of MPA, which frequently leads to intra-alveolar tumor spread (27). However, the role of WNT pathway ligands in NSCLC is presumably complicated. Some ligands, such as Wnt-1, Wnt-2, and Wnt-3, can activate the canonical WNT pathway and directly promote cancer growth (23-26). Others, including Wnt-5a and Wnt-11, serve as non-canonical WNT pathway activators. They can block canonical WNT signaling as well as inducing other malignant behavior (such as EMT) by β -catenin-independent mechanisms (28,29). These findings led us to hypothesize that an expression panel of Wnt family members would interact with each other in cancer progression. Our data were partially consisted with this interpretation and we showed that Wnt-1 expression was enhanced in MPA (Figure 3). However, the expression levels of other Wnt members, such as Wnt-6 and Wnt-



Figure 3 Dramatic and specific Wnt-1 over-expression in the MPA of lung. (A) Cancer tissues of the MPA and lepidic groups were applied with H&E staining and IHC staining to show the expression of Wnt-1, Wnt-6 and Wnt-10a in cancer cells. The pictures were taken under microscope at 400× power phase. (B) For each slide from both groups, the TPS of Wnt-1, Wnt-6 and Wnt-10a were calculated by counting the positive rate of tumor cells in each 10 views at 400× power phases under microscope. Any convincing cytoplasmic staining of the cancer cells were considered to be positive. The data were then analyzed by application of two-tailed unpaired student's *t*-tests. *, indicated statistical difference (P<0.05). H&E, hematoxylin and eosin; IHC, immunohistochemistry; MPA, Micropapillary-predominant adenocarcinoma; TPS, tumor positive score.

10a, did not show significant difference between the MPA and lepidic groups. These results suggest the existence of complicated and dynamic Wnt-regulation networks in different pathologic patterns of lung adenocarcinoma.

 β -catenin expression has been identified in up to 51% of lung adenocarcinomas (30). Jin *et al.* reported that the expression of β -catenin in the cell membrane was associated with a higher survival rate [hazard ratio (HR): 0.53; 95% confidence interval (CI): 0.32–0.87].

In contrast, β -catenin expression patterns in the cytoplasm and nucleus were associated with a poor survival rate, with HRs of 1.63 (95% CI: 1.34–1.99) and 3.15 (95% CI: 1.97–5.05), respectively (31). These data demonstrated that excessive intracellular β -catenin may help cancer cells become more invasive. Nuclear β -catenin accumulation was also correlated with epidermal growth factor receptor (EGFR) mutations and resistance to gefitinib (32,33), which cause these cancers

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Figure 4 Expression of WNT/ β -catenin signaling negative regulator Axin1 was decreased in MPA of lung, especially in cases with lymph node invasion. (A) Cancer tissues of the MPA and lepidic groups were applied with H&E staining and IHC staining to show the intension of cytoplasmic Axin1. The pictures were taken under microscope at 400× power phase. (B) For each slide from both groups, the TPS of Axin1 was calculated by counting the positive rate of tumor cells in each 10 views at 400× power phases under microscope. Any convincing cytoplasmic staining of the cancer cells were considered to be positive. The data were then analyzed by application of two-tailed unpaired student's *t*-tests. *, indicated statistical difference (P<0.05); ***, indicated significant difference (P<0.001). (C) Within the MPA group, MPA expression comparison between cases with or without lymph node invasion by the same method indicated lower TPS in the ones with lymph node invasion.

to be insensitive to treatment with EGFR-tyrosine kinase inhibitor. However, the difference in β -catenin distribution between well-differentiated and poorly differentiated lung adenocarcinomas remains unknown. Our study firstly discovers that poorly differentiated lung adenocarcinomas, such as MPA, involve more extensive intracellular β -catenin translocation, compared to well-differentiated lung adenocarcinomas (*Figure 2A,C*). Our data provide a possible explanation for highly aggressive phenotype of this particular histological subtype of lung adenocarcinoma.

The Axin1 protein acts as a scaffold in a destruction complex, whereby β -catenin is phosphorylated by glycogen synthase kinase 3 β and undergoes ubiquitination-dependent degradation. Disruption of this complex will lead to excessive accumulation of β -catenin and downstream oncogene transcription. Axin1 protein dysfunction caused by gene mutations has been reported in various tumors, including hepatocellular, colorectal, prostate, and gastric carcinomas (34-37). In our study, we found that Axin1 expression was lower in the MPA subtype than in the welldifferentiated lepidic subtype in terms of both percentage and staining intensity (*Figure 4A,B*). Evidence for a similar relationship between a silent Axin1 mutation and prognosis has been demonstrated in esophageal cancer (38), whereas the role of Axin1 in the biological behavior of lung cancers remains a mystery. Our results suggested that reduced Axin1 expression and a dysfunctional β -catenin destruction complex serve as a potential mechanism for cytoplasmic accumulation of β -catenin and activation of downstream signal transduction in MPA, which can further facilitate lymph nodes invasion of the MPA.

Conclusions

In summary, our results indicate that over activation of the WNT/ β -catenin signaling pathway leads to disruption of the cadherin-catenin complex and causes decreased intercellular adherence in MPA of the lung. Elevated expression of Wnt-1 and downregulation of Axin1 may play key roles in activation of this pathway. Our investigation may provide the potential target for development of medical strategies focused on the WNT/ β -catenin pathway to improving the prognosis of MPA of the lung in the future.

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Footnote

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Figure S1 Expression of E-cadherin, b-catenin and a-catenin in the MPA. M1-6 stands for MPA case 1-6. All the pictures were taken under 40x power phase.

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Figure S2 Expression of E-cadherin, b-catenin and a-catenin in the lepidic adenocarcinoma. L1-6 stands for lepidic case 1-6. All the pictures were taken under 40× power phase.