

The Small Heal Shock Protein α A-Crystallin Is Expressed In Pancreas and Acts as Negative Regulator of Carcinogenesis

Mi Deng, Pei-Chao Chen, Sisi Xie, Junqiong Zhao, Lili Gong, Jinping Liu, Lan Zhang, Shuming Sun, Jiao Liu, Haili Ma, et al.

▶ To cite this version:

Mi Deng, Pei-Chao Chen, Sisi Xie, Junqiong Zhao, Lili Gong, et al.. The Small Heal Shock Protein α A-Crystallin Is Expressed In Pancreas and Acts as Negative Regulator of Carcinogenesis. Biochimica et Biophysica Acta - Molecular Basis of Disease, 2010, 1802 (7-8), pp.621. 10.1016/j.bbadis.2010.04.004 . hal-00600124

HAL Id: hal-00600124

https://hal.science/hal-00600124

Submitted on 14 Jun 2011

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Accepted Manuscript

The Small Heal Shock Protein α A-Crystallin Is Expressed In Pancreas and Acts as Negative Regulator of Carcinogenesis

Mi Deng, Pei-Chao Chen, Sisi Xie, Junqiong Zhao, Lili Gong, Jinping Liu, Lan Zhang, Shuming Sun, Jiao Liu, Haili Ma, Surinder Batra, David Wan-Cheng Li

PII: S0925-4439(10)00083-9

DOI: doi: 10.1016/j.bbadis.2010.04.004

Reference: BBADIS 63091

To appear in: BBA - Molecular Basis of Disease

Received date: 16 January 2010 Revised date: 14 April 2010 Accepted date: 16 April 2010



Please cite this article as: Mi Deng, Pei-Chao Chen, Sisi Xie, Junqiong Zhao, Lili Gong, Jinping Liu, Lan Zhang, Shuming Sun, Jiao Liu, Haili Ma, Surinder Batra, David Wan-Cheng Li, The Small Heal Shock Protein α A-Crystallin Is Expressed In Pancreas and Acts as Negative Regulator of Carcinogenesis, *BBA - Molecular Basis of Disease* (2010), doi: 10.1016/j.bbadis.2010.04.004

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.

The Small Heal Shock Protein AlphaA-Crystallin Is Expressed In Pancreas and Acts as A Negative Regulator of Carcinogenesis

(Short Title: αA-Crystallin Is Down-Regulated in Pancreatic carcinoma)

Mi Deng^{1*}, Pei-Chao Chen^{2*}, Sisi Xie², Junqiong Zhao², Lili Gong¹, Jinping Liu¹, Lan Zhang^{1, 2}, Shuming Sun^{1, 2}, Jiao Liu^{1, 2}, Haili Ma¹, Surinder Batra¹, and David Wan-Cheng Li^{1, 2, 3**}

*Equal Contribution Authors

¹Department of Biochemistry and Molecular Biology, ³Department of Ophthalmology
& Visual Sciences, University of Nebraska Medical Center,
Omaha, NE 68198-5870, USA;

²key Laboratory of Protein Chemistry and Developmental Biology of National Education Ministry of China, College of Life Sciences, Hunan Normal University, Changsha, Hunan 410081, China.

**Correspondence: Dr. David Wan-Cheng Li, Department of Biochemistry and Molecular Biology, University of Nebraska Medical Center, 985870 Nebraska Medical Center, Omaha, NE 68198-5870, USA. Tel: 402-559-5073, Fax: 402-559-6650, E-mail: dwli1689@yahoo.com

CCEPTED MANUS

ABSTRACT

The small heat shock protein αA-crystallin is a structural protein in the

ocular lens. In addition, recent studies have also revealed that it is a molecular

chaperone, an autokinase and a strong anti-apoptotic regulator. Besides its

lenticular distribution, a previous study demonstrates that a detectable level of

αA-crystallin is found in other tissues including thymus and spleen.

present study, we have re-examined the distribution of α A-crystallin in various

normal human and mouse tissues and found that the normal pancreas expresses

a moderate level of aA-crystallin. Moreover, aA-crystallin is found

significantly downregulated in 60 cases of pancreatic carcinoma of different

types than it is in 11 normal human pancreas samples. In addition, we

demonstrate that αA -crystallin can enhance activity of the activating protein-1

(AP-1) through modulating the function of the MAP kinase, and also up-

regulates components of TGF β pathway. Finally, expression of α A-crystallin in

a pancreatic cancer cell line, miapaca, results in retarded cell migration.

Together, these results suggest that αA-crystallin seems to negatively regulate

pancreatic carcinogenesis.

Key words: Small heat shock protein, αA, Pancreas, AP-1, Smad2/3/5, TGFβ

Pancreatic Cancer, cell migration

2

Introduction

Heat-shock proteins (HSPs) are implicated in multiple cellular functions including signaling transduction, protein degradation, survival promotion, exocytosis and endocytosis [1-5]. Recent studies have revealed that HSPs also play an important role in tumorigenesis [6-8]. The lens structural proteins, α A- and α B-crystallins are characterized as HSPs [9]. Besides their robust expression in mammalian lens [10-11], the two α -crystallins are also expressed in non-lenticular tissues. α B-crystallin is significantly expressed in retina, muscle, heart and brain, and acts to prevent neural degradation, heart fail, but promote carcinogenesis [4-5, 12-15]. α A-crystallin is also expressed in some non-lenticular tissues, such as retina, spleen and thymus [16]. However, the function of non-lenticular α A-crystallin remains unknown.

Pancreatic cancer in human poses clinical challenges in diagnosis and treatment [17-18]. Among the various types of pancreatic malignancies, pancreatic duct adenocarcinoma is dominant and in addition, the mucinous adenocarcinoma, islet adenocarcinoma and acinic cell carcinoma are also observed with less frequency of occurrence [19-20]. At the molecular level, development of pancreatic malignancies is associated with canonical oncogenes and tumor suppressor genes, such as Ki-Ras, p16, p53, Smad4 and BRCA2, which are involved in several cross-talking cellular signaling pathways including TGFβ/SMAD, PI3K/AKT, and MAPK pathways [20-22].

In the present study, we present evidence to show that αA -crystallin is expressed in the normal human and mouse pancreas at a moderate level. Moreover, analysis of αA -crystallin in the tissue samples from normal human pancreas and 60 cases of pancreatic carcinoma reveals significant difference. αA -crystallin is downregulated more than 10-fold in the pancreatic carcinoma of various types than

that in normal pancreas. To explore the possible role of αA -crystallin in pancreatic carcinoma, we demonstrate here for the first time that αA -crystallin positively regulates the DNA binding and transactivity of the activating protein-1 (AP-1). This upregulation of AP-1 activity is derived from changed MAPK activity, which can positively regulate c-Jun and c-Fos. Moreover, expression of αA -crystallin in pancreatic cells enhances TGF β signaling pathway. Finally, expression of αA -crystallin in pancreatic cancer cells leads to retarded cell migration. Together, these results suggest that αA -crystallin appears to suppress pancreatic carcinogenesis.

2. Materials and Methods

2.1. Reagents and antibodies

Anti-αA/B-crystallin antibodies were described before [23]. ERK2 kinase assay kit, and anti-phosphorylated Elk antibody were purchased from Cell Signaling, Inc. Anti-c-Jun antibody was a kind gift from Dr. Tom Curran's Laboratory. Anti-c-Fos, anti-Lamin B, and anti-β-actin antibodies were purchased from Santa Cruz Biotechnology. Human normal pancreas and pancreatic carcinoma tissue samples were obtained from US Biomax, Inc.

2.2. Immunohistochemistry analysis of α A-Crystallin, measure of fluorescence intensity and HE staining

Immunohistochemistry analysis was conducted as recently described [24]. Briefly, the sections were rehydrated in graded alcohols, heated up to boiling for 5 min in citrate buffer at pH 6. The sections were washed three times with PBS. For non-specific blocking, each section was incubated in 500 µl of 5% normal goat serum (Sigma-Aldrich, CA) in PBS for 1 h at room temperature and then incubated

overnight in 400 μl of diluted anti-αA-crystallin antibody (1:100) in a humidified chamber at 4 °C. The sections were then washed with PBS three times (5 min each) followed by incubation with 400 μl of FITC-linked secondary antibody and DAPI (Vector Laboratories, CA) in blocking solution for 1 h in the absence of visible light. After incubation, sections were washed with PBS at room temperature 6 times (5 min each) and then observed under a Carl Zeiss Laser Scanning Confocal microscope. For negative controls, the sections were treated in the same way except that the primary antibody was replaced with normal serum IgG.

A quantitative measure of fluorescence intensity was conducted using confocal microscopy for assessment of the level of αA -crystallin expression. During analysis of the obtained results, the designated values of 0 to 10, 10 to 100, 100 to 1000 and more than 1000 represent negative background, weak, moderate and strong expression of αA -crystallin, respectively. The same immunohistochemical quantitative system was used for tissue microarray analysis by others [25]. Each core was measured individually. If more than one core was evaluated from each tissue the mean intensity and standard deviation were calculated and shown as Mean \pm SD.

For histology analysis, hematoxylin and eosin (HE) staining was conducted as recently described [26].

2.3. Cell culture and cell fraction extraction

The preparation of the pEGFP-C3 and pEGFP-C3- α A/ α B-crystallin expression constructs, and establishment of stable transfected cell lines were conducted as previously described [23-24]. The pEGFP- α TN4-1, pEGFP- α A- α TN4-1, pEGFP- α B-Miapaca stable

clones were grown in Dulbecco's Modified Eagle's Minimal Essential Medium (DMEM) containing 10% fetal bovine serum, 50 units/ml penicillin and streptomycin and 400 μ g/ml neomycin as described before [23-24]. All cells were kept at 37 °C and 5% CO2 gas phase. Cell fractions were extracted by differential sucrose gradient as previously described [23]. The cytoplasmic and nuclear extracts of vector (pEGFP)-, α A-crystallin (pEGFP- α A)- or α B-crystallin (pEGFP- α B)-transfected cells were used for western blot analysis. The nuclear extracts from parent, vector (pEGFP)-, α A-crystallin (pEGFP- α A)- or α B-crystallin (pEGFP- α B)-transfected cells were used for gel mobility shifting assays.

2.4 Reverse transcription-linked polymerase chain reaction (RT-PCR)

Total RNA samples were extracted from αA-crystallin-transfected cells (pEGFP-αA-Miapaca), αB-crystallin-transfected cells (pEGFP-αB-Miapaca), vector transfected cells (pEGFP-Miapaca) and control parent pancreatic cancer cells (Miapaca) using TRIZOL reagent. Five μg of total RNA from each sample was used for cDNA synthesis. Several components of TGFβ signaling pathway were examined by RT-PCR using specific primer pairs listed in table 1. The PCR reaction is: 94°C for 5 min; 94°C for 30 sec, 55°C for 30 sec, and 72°C for 40 sec, in 30 cycles; 72°C for 5 min; keep at 4°C. The PCR products were examined by agarose gel electrophoresis and visualized under UV illumination.

2.5. Gel mobility shifting assay

Gel mobility shifting assays were conducted as previously described [24, 27-29]. The following oligos were used: 5'-GTCCTTCATTACGTCACGCATAG-3' for conserved AP-1 binding site, 5'-GTCCTTCATTCATTCACGCATAG-3' for mutated

AP-1 binding site; Twenty μg of nuclear extracts or cytoplasmic extracts prepared from the pEGFP-αTN4-1, pEGFP-αA-αTN4-1, pEGFP-Miapaca, and pEGFP-αA-Miapaca cells were incubated with 1x10⁵ cpm of ³²P-labeled double-stranded synthetic oligonucleotides for 30 minutes at 37 °C in binding shifting buffer [27-29]. For competition experiments, 50-fold of the non-labeled wild type or mutant double-stranded synthetic oligonucleotides were pre-incubated with the nuclear extracts for 30 minutes before the labeled probe was added into the reaction. For the pre-cleared experiments, 20 μg of nuclear extract prepared from pEGFP-αTN4-1, pEGFP-αA-αTN4-1, pEGFP-Miapaca, and pEGFP-αA-Miapaca cells was pre-incubated with 10 μg antibody against c-Jun or/and c-Fos for 1 hour on ice, then mixed with protein A/G agarose and rotated for additional 1 hour at 4 °C, and then the supernatant was recovered for incubation with 1x10⁵ cpm of ³²P-labeled double-stranded synthetic AP-1 oligonucleotides for 30 minutes at 37 °C in a binding shifting buffer. After the binding reactions, the mixtures were loaded onto 6% native PAGE and detected by autoradiography.

2.6. Kinase activity assay

Substrate Elk protein was incubated with different doses of ERK2 kinase, 0.1, 1, 5 or 10 ng or mock kinase, GSK3 β in the kinase buffer for 30 min at 37 °C in the presence of γ -³²P-ATP. For the effect of α A-crystallin on ERK kinase, differential concentrations (0, 1 and 10 μ g) of purified bovine α A-crystallin were added to the kinase assay reactions. After reaction, the mixtures were loaded to SDS-PAGE and detected by western blot analysis using anti-phosphorylated-Elk antibody.

2.7 Western blot analysis

Preparation of total proteins from mouse tissues and various transfected cells and Western blot analysis of different protein samples were conducted as previously described [30-32].

2.8. Wound healing assay.

The three types of stable clones of pEGFP-Miapaca, pEGFP-αA-Miapaca and pEGFP-αB-Miapaca cells were seed in 6-well plates and cultured until 100% confluent. A straight scratch was made by using a 1 ml blue pipette tip to simulate the wound in each well. After PBS washing for 2 times, new DMEM medium was added for a continuous growth of another 48 hours. The wound healing process was recorded daily using the Leica Inverted Phase Contrast Fluorescence Microscopy with a 10x objective.

2.9. Quantitation and Statistical analysis

The mRNA gels were analyzed using the UN-SCAN-IT software from Silk Scientific Corporation (Orem, UT, USA). Total pixel data were averaged from three different groups of samples of each species after normalization against the background. The total pixels from the target band were divided by the total pixels from the corresponding β -actin. The student t-test was used to compare the medians of two unpaired groups. P < 0.05 was considered significant [31].

3. Results

3.1. Expression of αA -crystallin in human non-lenticular tissues

Recent studies have shown that αB -crystallin is implicated in promotion of carcinogenesis (6, 15, 33). Whether αA -crystallin has similar functions remains to be

explored. For this reason, we have re-examined the tissue-specific expression of α A-crystallin in human non-lenticular tissues using immunohistochemistry analysis of human multi-organ tissue array samples and confocal microscope measurement of the relative fluorescence density derived from immunohistochemistry staining. As shown in Fig. 1A, a strong lenticular expression of α A-crystallin in the developing mouse eye was observed, which is consistent with previously reported result (10). In addition, we observed a moderate level of α A-crystallin expression in human pancreas and to a less degree, in human liver (Fig. 1A). Similar pancreatic level of α A-crystallin was observed in 7 out of 11 human samples examined (Fig. 1B). In pancreas, α A-crystallin was specifically expressed in islets, acinus and interlobular tubes (Fig. 1A). In other non-lenticular ocular tissues, α A-crystallin was also expressed at moderate level in retina and ciliary processes (data not shown). The relative mean values of fluorescence intensities in various non-lenticular tissues were measured with confocal microscope coupled with related software and were presented in Fig. 1B.

3.2. Expression of α A-crystallin in mouse tissues

To validate the observed results of αA -crystallin expression in human samples, we next dissected 10 different tissues from adult mouse: lens, heart, thymus, kidney, brain, spleen, pancreas, lung, liver and testis, and analyzed the expression level of αA -crystallin in these tissues. As shown in the top panel of Fig. 2, Western blot analysis revealed that besides its robust expression in the ocular lens (lane 1 contains only 0.5 μg protein, rest of the lanes 100 μg proteins), αA -crystallin was detected in thymus and pancreas at a relatively strong level, then to a less degree in brain. The αA -crystallin level in heart, kidney, spleen and liver was much lower than that in

thymus and pancreas. Finally, a trace of αA -crystallin was also observed in testis. No αA -crystallin was detected in lung. As comparison, we also examined the tissue distribution of αB -crystallin in mouse. Besides the strongest expression in lens, αB -crystallin was highly expressed in heart, thymus and brain, moderately expressed in kidney and lung. However, different from the tissue specific expression patterns of αA -crystallin, αB -crystallin was barely detectable in pancreas and spleen, and absent in liver and testis examined. Thus, our results confirm that the pancreas expresses a moderate level of αA -crystallin. In addition, we also show that several other non-lenticular tissues also express low level of αA -crystallin. These results are slightly different from the pattern of αA -crystallin detected in different tissues from rat [16, 34].

3.3. Down-regulation of α A-crystallin in pancreatic tumors

Since both Hsp27 and α B-crystallin are implicated in carcinogenesis [6-8, 15, 33, 36-37], we reason that α A-crystallin may be also involved in regulation of carcinogenesis. To test this possibility, we compared the expression pattern of α A-crystallin in human normal pancreatic tissues and various types of pancreatic carcinoma using the tissue array samples. As shown in Fig. 3A, expression of α A-crystallin in human pancreatic cancer tissues was lower than that in human normal pancreas. Among the 11 normal pancreas samples examined, 7 samples displayed moderate expression level of α A-crystallin (normal pancreas A in Fig.3A) and 4 samples relatively weak expression (normal pancreas B in Fig.3A). In contrast, among the 60 cases of pancreatic cancer samples, 4 had weak expression of α A-crystallin signal (shown as pancreatic cancer #A of Fig.3A), 17 displayed barely

detectable αA-crystallin signal and the remaining 39 samples displayed no detectable signals (shown as pancreatic cancer #B of Fig.3A).

To further characterize the expression of αA -crystallin in different human pancreatic cancer, we divided the pancreatic cancer samples into 4 different types: duct adenocarcinoma (n=48), mucinous adenocarcinoma (n=3), islet adenocarcinoma (n=6) and acinic cell carcinoma (n=3). The fluorescence signal in each individual sample and the statistic data with the mean value of fluorescence intensity of αA -crystallin expression in each type of pancreatic carcinoma were presented in Fig. 3B. The expression level of each normal pancreas and the statistical data derived from the normal pancreas were also included for comparison (Fig. 3B). Together, our results reveal that αA -crystallin is significantly down-regulated in human pancreatic carcinoma comparing with normal pancreas.

3.4. Expression of αA -crystallin in miapaca cells upregulates expression of c-Jun and c-Fos.

It is well established that the transcriptional factor complex, activating protein-1 (AP-1) plays a significant role in tumorigenesis [38-42]. AP-1 is either a homodimer containing two Jun family members (c-Jun, Jun-B or Jun D) or a heterodimer containing one member from the Jun family and another from the Fos family (C-Fos, Fra-1 and Fra-2) [38]. To explore the possible function of α A-crystallin in human pancreatic cancer, we overexpressed α A-crystallin in both pancreatic cancer cells, miapaca and non-cancer cells, α TN4-1, a SV40 large-T-transformed mouse lens epithelial cell line expressing very little endogenous α A-crystallin. Then, we examined the AP-1 expression and activity in these transfected cells. As shown in Figure 4, expression and phosphorylation of c-Jun were significantly upregulated in

the nucleus of the miapaca cells expressing αA -crystallin than in the nucleus of the vector-transfected cells, expression of c-Fos was also upregulated, though to a much less degree, in both nucleus and cytoplasm of the miapaca cells expressing αA -crystallin than in the same fractions of the vector-transfected cells. Similar results were also obtained in the nucleus of the $\alpha TN4$ -1 cells expressing αA -crystallin (Supplementary Fig. 1).

3.5. Expression of αA -crystallin in miapaca cells upregulates AP-1 activity.

To determine whether the upregulated c-Jun and c-Fos actually led to increased AP-1 activity, we first conducted DNA binding assays using the oligos contained either the conserved AP-1 binding site (TGAGTCA) or the mutated site (ATCGGTA). As shown in Fig. 5A, the nuclear extract prepared from either parent miapaca cells (lane 13) or vector pEGFP transfected Miapaca cells can bind to the wild type AP-1 oligo (lane 2) which was competed off by cold wild type oligo (lane 3) but much less by the mutated oligo (Lane 4). The nuclear extracts pre-cleared by antibodies against c-Jun (lane 5), c-Fos (Lane 6) or both antibodies (lane 7) lost most of the AP-1 activity so that the formed AP-1 complex was either lost or appeared very weak. In contrast, the nuclear extracts pre-cleared by normal IgG only slightly attenuated the AP-1 complex (Lane 8). These results suggest that the specific band we detected between the nuclear extract from pEGFP-Miapaca cells and the wild type oligo is the AP-1/DNA complex. Next, we compared the AP-1 DNA binding activity in vector- and αAcrystallin-transfected cells. As shown in lane 9 and 11 of Fig. 5A, miapaca cells expressing αA-crystallin display significant upregulation of AP-1 activity. Similar results were obtained with the lens epithelial cells, αTN4-1 expressing either only EGFP or EGFP-αA-crystallin (Supplementary Fig. 2). As a control for the specific

binding to the AP-1 site, we conducted the reaction using purified p53 protein and AP-1 oligo, lane 12 of Fig. 5A clearly showed that p53 did not bind to AP-1 oligo. To further confirm that the transactivity of AP-1 was indeed upregulated in miapaca cells expressing α A-crystallin, we introduced two reporter genes into the two types of stable transfectants: pEGFP-Miapaca and pEGFP- α A-Miapaca. As shown in Fig. 5B, assayed with either pJun-Luc (the luciferase gene was driven by the Jun promoter), or pProlactin-1xAP-1-Luc (the luciferase gene was driven by a mini-prolactin gene promoter containing a copy of AP-1 binding site), the miapaca cells expressing α A-crystallin showed statistically significant upregulation of luciferase activity, indicating the upregulation of AP-1 activity in α A-crystallin expression miapaca cells (Fig. 5B). Thus, expression of α A-crystallin in pancreatic cancer cells positively regulates AP-1 activity.

3.6. \(\alpha A\)-crystallin positively reglates ERK2 MAPK.

It has been shown that AP-1 expression and activity can be enhanced by MAP kinases [44-48]. To explore how might αA -crystallin regulate AP-1, we examined the possible modulation of ERK kinase activity by αA -crystallin since ERK is a major MAP Kinase regulating AP-1 components in both expression and activation [44-48]. As shown in Fig. 6A, we found that Elk was dose-dependently phosphorylated by ERK2 in vitro. Moreover, in the presence of αA -crystallin, phosphorylation of Elk by 0.1 or 1 ng ERK2 was significantly enhanced. Addition of 1 µg of αA -crystallin into the ERK2 kinase reaction with low level of ERK2 (1 ng) enhanced more than 10-fold Elk phosphorylation, reflecting the corresponding enhancement of ERK2 activity. Together, these results reveal that αA -crystallin can positively regulate AP-1 activity through modulation of ERK activity. To further

confirm that αA -crystallin enhances ERK2 kinase activity, we analyzed the ERK1/2 expression and activity in vector- and αA -crystallin transfected stable lines of cells. As shown in Fig. 6C, both expression and activity of ERK2 were clearly upregulated in miapaca cells expressing αA -crystallin than vector-transfected cells.

3.7. αA-crystallin reglates TGF-β Signaling Pathway

To determine whether αA -crystallin actively regulates TGF β signaling pathway which plays key role in pancreatic carcinogenesis [20], we first examined expression of several genes in the TGF β signaling pathway at the mRNA level. As shown in Fig. 7, TGF β was increased about 2-fold when αA -crystallin was overexpressed in Miapaca (Fig. 7A and 7B, similar result was observed in αA -crystallin-transfected Capan-1 cells, data not shown). For the receptor, TGF β receptor 1 and 2 are not detected and TGF β receptor 3 displayed little change in the three types of cells (Fig. 7A and 7C). To further confirm the modulation of TGF β signaling pathway by αA -crystallin, we examined the protein expression levels of TGF β and the downstream smads, as well as the phosphorylation status of the smads. As shown in Fig. 8, Western blot analysis revealed that TGF β was also upregulated at the protein level. In addition, expression and phosphorylation of Smad 3 and Smad 5 were significantly upregulated. Thus, αA -crystallin actively enhances TGF β signaling pathway.

3.8. Retardation of cell migration in α A-crystallin-overexpressed cells

To explore how αA -crystallin may regulate carcinogenesis, we conducted wound healing assay using the established stable cell lines: pEGFP-miapaca, pEGFP- αA -Miapaca and pEGFP- αB -Miapaca. When these cells with similar expression levels

were selected to conduct wound-healing assays (Fig. 9), it was found that miapaca cells expressing αA -Crystallin displayed statistically significant retardation in cell migration. In contrast, the same cells expressing either the vector or αB -Crystallin did not show such effect. Thus, αA -Crystallin retards migration of pancreatic cancer cells, which is consistent with its downregulation in various types of pancreatic carcinoma (Fig. 3).

4. Discussion

In the present study, we have demonstrated the following: 1) αA -crystallin is expressed at moderate level in pancreas from human and mouse, a non-ocular tissue previously not documented; 2) αA -crystallin is significantly downregulated in the pancreatic tumors than in the normal human pancreas; 3) Expression of αA -crystallin in pancreatic carcinoma cells induces significant upregulation of the AP-1 components and activity, 4) αA -crystallin can enhance activity of ERK2 MAP kinase, through which it can regulate AP-1 expression and activity; 5) Expression of αA -crystallin in pancreatic carcinoma cells significantly enhanced the TGF β signaling pathways; and 6) expression of αA -crystallin in pancreatic cells led to obvious retardation of the cell migration. Together, our results suggest that αA -crystallin appear to negatively regulate pancreatic carcinogenesis, which is contrast to the function of αB -crystallin in breast cancer.

αA-crystallin Is Expressed in Normal Pancreas Besides Other non-Lenticular Tissues

15

αA-crystallin is first known as a lens structural protein [10-11]. It shares about 60% identity in amino acid sequence with αB-crystallin, another lens structure protein [10-11]. Several lines of evidence have shown that although the two genes encoding αA - and αB -crystallins may arise from gene duplication, they have diverged significantly. First, during murine development, the initial turning on of the two genes occurs at different time. While αB mRNA is first becoming detectable at E9.5, expression of αA mRNA appears at E10.5 [10]. Such differential turning on time reflects the different control mechanisms in the promoters of the two α -crystallin Second, the two genes display distinct tissue-specific expression patterns. While αA is highly restricted to lens during mouse embryonic developmental process, αB-crystallin is expressed in the developing heart, nasal epithelium, and retinal pigment epithelium [10]. In the adult vertebrates, although both αA and αB are abundantly expressed in the lens, they display significantly difference in nonlenticular tissue expressions. aB is strongly expressed in heart, skeletal muscle, kidney and brain [11-13]. In contrast, αA is reported to be expressed at very low level in some non-lenticular tissues including spleen, thymus, heart, brain and liver [16-17]. In the present study, our results not only confirm the presence of a low level of α A-crystallin in kidney and liver, but also reveal a moderate level of α A-crystallin expression in a novel non-lenticular tissue, the normal human pancreas. Among the 11 normal human pancreas samples examined, 7 samples display consistently moderate level of αA-crystallin expression and the remaining 4 samples also showed a low level of α A-crystallin expression (Fig. 1). The pancreatic distribution of α Acrystallin is further confirmed from analysis of mouse tissue samples (Fig.2).

In a previous study, Kato et al. [16] compared the relative abundance αA -crystallin in both lens and non-lenticular tissues (ng αA -crystallin/per mg total protein in organ). These authors found that in the adult rat (8–16 weeks), the lens contains 244 μg αA -crystallin/per mg total lens protein, and the spleen contains 16.6 ng αA -crystallin/per mg total spleen protein, so the relative abundance of lens αA -crystallin is about 14,698-fold. Using confocal fluorescence microscopy, we quantitated the relative abundance of αA -crystallin in the developing mouse lens and the pancreatic tissue, and found the corresponding values are 807+/-48 and 283+/-67, respectively. Therefore, the human adult pancreatic signal of αA -crystallin is about 1/3 of the developing mouse eye lens (Fig. 1). In mouse tissue, we estimate that αA -crystallin in lens is about 500-fold higher than that in pancreas (see Fig.2, 200-fold difference in sample load x 2.5-fold difference in αA -crystallin band in lens verse in pancreas). Together, our results demonstrate that the pancreas expresses a moderate level of αA -crystallin.

αA-Crystallin Is Implicated in Carcinogenesis

It is well established that small heat shock proteins are implicated in carcinogenesis [8]. One of the most studied members is Hsp27, which is involved in carcinogenesis of many different types of tissues [49-57]. Another member of the HSP family implicated in carcinogenesis is αB -crystallin. Iwaki and Tateishi [58] first demonstrated the existence of αB -crystallin in hamartomas of tuberous sclerosis. Then, it was found that concentrations of αB -crystallin in prostatic carcinoma tissues were significantly higher than in benign prostatic hyperplasia [59]. In breast cancer cells, αB -crystallin was found expressing constitutively in certain breast carcinoma

cell lines, and increased intensity of αB -crystallin expression was correlated with shorter survival [33]. Moreover, expression of αB -crystallin results in transformation of immortalized human mammary epithelial cells, induction of EGF- and anchorage-independent growth, and enhancement of cell migration and invasion [15]. Thus, αB -crystallin is considered as a novel oncoprotein [15].

Both αA - and αB -Crystallins belong to the heat shock protein (Hsp) family [2]. Whether α A-crystallin is also in implicated in carcinogenesis remains to be explored. Compared with α B-Crystallin, the limited distribution of α A-crystallin in nonlenticular tissues may restrict its function in carcinogenesis [10-14]. Nevertheless, several recent studies suggest that α A-crystallin may be also implicated in tumor development. First, in 4 cases of human sebaceous carcinoma of the eyelid examined, both α A-Crystallin and α B-Crystallin were found highly expressed [60]. Second, in the retinocytoma, αA-crystallin was expressed in the cytoplasm of all tumor cells and the apoptotic index was significantly higher in those cases with weak signal of α Acrystallin than in those that were strongly positive [61]. These results suggest that αA-crystallin, acting like αB-crystallin, seems to promote carcinogenesis. On the other hand, a recent study revealed that in 6 cases of retinoblastoma, preoperative chemotherapy induced strong expression of only Hsp27 and αB-Crystallin but not αA-Crystallin [62]. Moreover, the viable tumor cells survived contained high levels of Hsp27 and α B-Crystallin but not α A-Crystallin. Therefore, these results indicate that α A-crystallin does not seem to act the same way as Hsp27 and α B-Crystallin do in promoting carcinogenesis. Our present finding that in 60 different cases of pancreatic carcinoma, the expression level of αA-crystallin was consistently downregulated than that in 11 normal human pancreas samples also support the negative

regulation of carcinogenesis by αA -crystallin. In addition, our results also suggest that αA -crystallin may be used as one of the biomarkers for pancreatic cancer diagnosis.

αA-crystallin May negatively Regulates Development of Pancreatic Tumors

Pancreatic malignancy develops from intraepithelial neoplasia (PanIN), progressing through PanIN-1a, PanIN-1b, PanIN-2, and PanIN-3 to ductal adenocarcinomas (PDAC) [63-64]. During this process, key oncogenes and tumor suppressors such as Ki-RAS, p16^{INK4a}, P53, DPC/SMAD4, and BRCA2 play very important roles [20-22, 65-68]. The carcinogenesis signals are mediated by major signaling pathways including MAPK pathways, sonic hedgehog and notch signaling pathways, and also TGFβ signaling pathways [20-22, 65-68]. In the present study, we present several lines of evidence to show that αA-crystallin seems to negatively regulate development of pancreatic tumors. First, we demonstrate that compared with the physiological level in normal human pancreas, α A-crystallin is significantly downregulated in 60 cases of pancreatic carcinomas of various types. Second, we show that pancreatic cancer cells expressing α A-Crystallin display retarded cell migration. Finally, we show that pancreatic cancer cells expressing α A-crystallin have enhanced expression levels of c-Jun and c-Fos, increased phosphorylation of c-Jun oncoprotein, and elevated level of AP-1 activities in both DNA binding and transactivation, which is derived from enhanced activity of the ERK2 kinase promoted by α A-Crystallin (Fig. 6). As a result, α A-crystallin positively regulates TGFβ signaling pathway (Figs. 7 and 8) through MAP kinase, and thus negatively regulate development of pancreatic malignancy.

Acknowledgements

We thank Drs. J. Horvitz for the $\alpha A/\alpha B$ -Crystallin antibodies and Dr. Tom Curran for the C-Jun antibody. This work is supported in part by the NIH grants 1R01 EY018380, and 1R01EY015765, and the Lotus Scholar Program Funds from Hunan Province Government and Hunan Normal University, and the Changjiang Scholar Team Award from the National Education Ministry of China to DWL.

References

- [1] R. Benndorf, M.J.Welsh. Shocking degeneration, Nat Genet 36 (2004) 547-548.
- [2] Y.Sun, T.H.MacRae. The small heat shock proteins and their role in human disease, FEBS J. 272 (2005) 2613-2627.
- [3] F.Bennardini, A.Wrzosek, M.Chiesi. Alpha B-crystallin in cardiac tissue. Association with actin and desmin filaments, Circ Res. 71 (1992) 288-294.
- [4] D.W.Li, L. Gong, M. Deng, J.P. Liu, M. Liu, Y.W.Mao. The two lens structural proteins, αA- and αB-crystallins, prevent stress-induced apoptosis through regulation of multiple signaling transduction pathways. In: Arrigo, Simon (eds). *Small Stress Proteins and Human Diseases*. Nova Science Publisher, Inc NY, USA. 2009. *In Press*.
- [5] I.J.Benjamin, Y.Guo, S.Srinivasan, S.Boudina, R.P.Taylor, N.S. Rajasekaran, R. Gottlieb, E.F. Wawrousek, E.D. Abel, R.Bolli. CRYAB and HSPB2 deficiency alters cardiac metabolism and paradoxically confers protection against myocardial ischemia in aging mice, Am J Physiol Heart Circ Physiol 293 (2007) H3201-3209.
- [6] M. Sherman, G. Multhoff. Heat shock proteins in cancer. Ann N Y Acad Sci 1113 (2007) 192-201.
- [7] C.Didelot, D.Lanneau, M.Brunet, A.L.Joly. A.De Thonel, G.Chiosis, C.Garrido. Anti-cancer therapeutic approaches based on intracellular and extracellular heat shock proteins. Current Medicinal Chemistry 14 (2007) 2839-2847.
- [8] D.R.Ciocca, S.K.Calderwood. (2005) Heat shock proteins in cancer: diagnostic, prognostic, predictive, and treatment implications. Cell Stress Chaperon 10, 86-103.
- [9] G.B.Reddy, P.A.Kumar, M.S. Kumar. Chaperone-like activity and hydrophobicity of alpha-crystallin. Iubmb Life 58 (2006) 632-641.
- [10] M. L.Robinson, P.A.Overbeek. Differential expression of alpha A- and alpha B-crystallin during murine ocular development. Invest Ophth Vis Sci 37 (1996) 2276-2284.
- [11] J.Piatigorsky. Lens crystallins and their genes: diversity and tissue-specific expression. FASEB J 3 (1989) 1933-1940.
- [12] S.P.Bhat, C.N.Nagineni. alpha B subunit of lens-specific protein alpha-crystallin is present in other ocular and non-ocular tissues. Biochem Biophys Res Commun 158 (1989) 319-325.

- [13] R.A.Dubin, E.F.Wawrousek, J.Piatigorsky. Expression of the murine alpha B-crystallin gene is not restricted to the lens. Mol Cell Biol 9 (1989) 1083-1091.
- [14] S. Kase, J.G.Parikh, N.A.Rao. Expression of alpha-crystallin in retinoblastoma. Arch Ophthalmol 127 (2009) 187-192.
- [15] J.V.Moyano, J.R Evans, F.Chen, M.Lu, M.E.Werner, F.Yehiely, L.K.Diaz, D. Turbin, G.Karaca, E.Wiley, T.O.Nielsen, C.M.Perou, V.L.Cryns. AlphaB-crystallin is a novel oncoprotein that predicts poor clinical outcome in breast cancer. J Clin Invest 116 (2006) 261-270.
- [16] K.Kato, H.Shinohara, N.Kurobe, S.Goto, Y.Inaguma, K.Ohshima, Immunoreactive alpha A crystallin in rat non-lenticular tissues detected with a sensitive immunoassay method. Biochim Biophys Acta 1080 (1991) 173-180.
- [17] M.B. Ujiki, M.S. Talamonti. Surgical management of pancreatic cancer. Semin Radiat Oncol 15 (2005) 218-225.
- [18] J. E. Niederhuber, M. F. Brennan, H.R.Menck. The National Cancer Data Base report on pancreatic cancer. Cancer 76 (1995) 1671-1677.
- [19] R.Pezzilli. Pancreatic ductal carcinoma: from the bench to the bedside. JOP 7 (2006) 508-510.
- [20] A.F.Hezel, A.C.Kimmelman, B.Z.Stanger, N.Bardeesy, R.A.Depinho. Genetics and biology of pancreatic ductal adenocarcinoma. Genes Dev 20 (2006) 1218-1249.
- [21] N.Bardeesy, R.A.DePinho. Pancreatic cancer biology and genetics. Nat Rev Cancer 2 (2002) 897-909.
- [22] A.K.Rustgi. The molecular pathogenesis of pancreatic cancer: clarifying a complex circuitry. Genes Dev 20 (2006) 3049-3053.
- [23] Y-W.Mao, J.P.Liu, H.Xiang, D.W-C.Li, Human alphaA- and alphaB-crystallins bind to Bax and Bcl-X(S) to sequester their translocation during staurosporine-induced apoptosis. Cell Death Differ 11 (2004) 512-526.
- [24] L Xiao, L-L Gong, D Yuan, M Deng, J-P Liu, L-L Chen, L Zhang, S-M Sun, J Liu, X-H Hu, H-L Ma, C-B Zheng, H Fu, P-Q Chen, J-Q Zhao, S-S Xie, X-M Zeng, L-J Zou, Y-M Xiao, W-B Liu, J Zhang, Y Liu, D W-C Li (2009). Protein Phosphatase-1 regulates Akt1 signaling pathway to control gene expression, cell survival and differentiation. Cell Death & Differentiation. In Press.
- [25] M.Marinov, A.Ziogas, O.E.Pardo, L.T.Tan, T. Dhillon, F.A.Mauri, H.A.Lane, N. R.Lemoine, U.Zangemeister-Wittke, M.J.Seckl, A.Arcaro. AKT/mTOR pathway activation and BCL-2 family proteins modulate the sensitivity of

- human small cell lung cancer cells to RAD001. Clin Cancer Res 15 (2009) 1277-1287.
- [26] Y-M. Xiao, L.Chen, J.Liu, L-J. Zou, W-B. Liu, H.G. Chen, Y.Liu D.W-C. Li. JNK1 downregulation is associated with sex reversal of the ricefield eel, J Exp Zool. Part B. Vol. 314B. (2009) In Press.
- [27] H.Feng, H.Xiang, Y-W.Mao, J.Wang, J-P.Liu, X.Q. Huang, Y.Liu, S-J.Liu, C. Luo, X-J.Zhang, Y.Liu, D.W-C.Li. Human Bcl-2 activates RAF/MEK/ERK pathway and regulates AP-1 activity and downstream genes in rabbit Lens epithelial cells. Oncogene. 23 (2004) 7310-7321.
- [28] J-C. Qin, H-G. Chen, Q.Yan, M.Deng, J-P.Liu S. Doerge, W-Y.Ma, Z. Dong, D. W-C. Li. Protein phosphatase-2A is a target of epigallocatechin-3-gallate and modulates the p53-Bak apoptotic pathway. Cancer Res. 68(11) (2008) 4150-4162.
- [29] H-G.Chen, W-J.Han, M. Deng, J.Qin, D.Yuan, J-P.Liu, L.Xiao, L.Gong, S.Liang, J.Zhang, Y.Liu, D.W-C.Li. Transcriptional regulation of PP2A-A alpha is mediated by multiple factors including AP-2alpha, CREB, ETS-1, and SP-1. PLoS One 4 (2009) e7019.
- [30] D.W-C.Li, J.R.Kuszak, K.Dunn, R-R.Wang, W-C.Ma, G-M.Wang, A.Spector, M.Leib, A.M.Cotliar, M.Weiss, J.Espy, G.Howard, R.L.Farris, J.Auran, A.Donn, A.Hofeldt, C.Mackay, J.Merriam, R.Mittl, T.R.Smith. Lens epithelial cell apoptosis appears to be a common cellular basis for non-congenital cataract formation in humans and animals. J. Cell Biol. 130 (1995) 169-181.
- [31] D.W-C.Li, Jin-Ping Liu, J. Wang, Y-W.Mao, Li-Hui Hou. Protein expression patterns of the signaling molecules for MAPK pathways in human, bovine and rat lenses. Invest. Ophthalmol. Vis. Sci. 44 (2003) 5277-5286.
- [32] D. W-C. Li, Liu, J-P., Mao Y. W., H. Xiang, J. Wang, W-Y Ma, Z. Dong, H. M. Pike, Rhoderick E. Brown and J. C., Reed. Calcium-Activated RAF/MEK/ERK pathway mediates p53-dependent apoptosis and is abrogated by αB-crystallin through Inhibition of Ras Activation. Mol. Biol. Cell. 16 (2005) 4437-4453.
- [33] Chelouche-Lev, D., Kluger HM, Berger AJ, Rimm DL, Price JE. alphaB-crystallin as a marker of lymph node in breast carcinoma. Cancer. 100 (2004) 2543-2548.
- [34] Srinivasan AN, Nagineni CN, Bhat SP. αA-Crystallin is expressed in non-ocular tissues. J. Biol. Chem. 267 (1992) 23337-23341.
- [35] G.L.Vegh, V.Fulop, Y.Liu, S.W.Ng, Z.S.Tuncer, D.R.Genest, P.Paldi-Haris, J.Foldi, S.C.Mok, R.S.Berkowitz. Differential gene expression pattern between normal human trophoblast and choriocarcinoma cell lines: downregulation of heat shock protein-27 in choriocarcinoma in vitro and in vivo. Gynecol Oncol 75 (1999) 391-396.

- [36] K.Ou, K.Yu, D.Kesuma, M.Hooi, N.Huang, W.Chen, S.Y.Lee, X.P.Goh, L. K.Tan, J.Liu, S.Y.Soon, S.Bin Abdul Rashid, T.C. Putti, H.Jikuya, T.Ichikawa, O.Nishimura, M.Salto-Tellez, P.Tan. Novel breast cancer biomarkers identified by integrative proteomic and gene expression mapping. J Proteome Res 7 (2008) 1518-1528.
- [37] S.K.Gruvberger-Saal, R.Parsons. Is the small heat shock protein alphaB-crystallin an oncogene? J Clin Invest 116 (2006) 30-32.
- [38] T.Curran, B.R.Franza, Jr. Fos and Jun: the AP-1 connection. Cell 55 (1988) 395-397.
- [39] J.L.Imler, B.Wasylyk. AP1, a composite transcription factor implicated in abnormal growth control. Prog Growth Factor Res 1 (1989) 69-77.
- [40] H.E. Varmus. Oncogenes and transcriptional control. Science 238 (1987) 1337-1339.
- [41] Durchdewald M, Angel P, Hess J. The transcription factor Fos: a Janus-type regulator in health and disease. Histol Histopathol. 24(2009):1451-61.
- [42] Karin M, Gallagher E. From JNK to pay dirt: jun kinases, their biochemistry, physiology and clinical importance. IUBMB Life. 57(2005) 283-295.
- [43] D.A.Leonard, N.Rajaram, T.K.Kerppola. Structural basis of DNA bending and oriented heterodimer binding by the basic leucine zipper domains of Fos and Jun. Proc Natl Acad Sci U S A 94 (1997) 4913-4918.
- [44] G. Gutierrez-Venegas, R. Castillo-Aleman. Characterization of the transduction pathway involved in c-fos and c-jun expression induced by Aggregatibacter actinomycetemcomitans lipopolysaccharides in human gingival fibroblasts. Int Immunopharmacol 8 (2008) 1513-1523.
- [45] D.B.Chen, J.S.Davis. Epidermal growth factor induces c-fos and c-jun mRNA via Raf-1/MEK1/ERK-dependent and -independent pathways in bovine luteal cells. Mol Cell Endocrinol 200 (2003) 141-154.
- [46] M.Cavigelli, F.Dolfi, F.X.Claret, M.Karin. Induction of c-fos expression through JNK-mediated TCF/Elk-1 phosphorylation. EMBO J 14 (1995) 5957-5964.
- [47] P.Monje, M.J.Marinissen, J.S.Gutkind. Phosphorylation of the carboxylterminal transactivation domain of c-Fos by extracellular signal-regulated kinase mediates the transcriptional activation of AP-1 and cellular transformation induced by platelet-derived growth factor. Mol Cell Biol 23 (2003) 7030-7043.
- [48] G.J.Babu, M.J.Lalli, M.A.Sussman, J.Sadoshima, M.Periasamy. Phosphorylation of elk-1 by MEK/ERK pathway is necessary for c-fos gene

- activation during cardiac myocyte hypertrophy. J Mol Cell Cardiol 32 (2000) 1447-1457.
- [49] Arrigo AP, Simon S, Gibert B, Kretz-Remy C, Nivon M, Czekalla A, Guillet D, Moulin M, Diaz-Latoud C, Vicart P. Hsp27 (HspB1) and alphaB-crystallin (HspB5) as therapeutic targets. FEBS Lett. 581(2007)3665-3674.
- [50] Kostenko S, Moens U. Heat shock protein 27 phosphorylation: kinases, phosphatases, functions and pathology. Cell Mol Life Sci. 66(2009)3289-3307.
- [51] Storm FK, Gilchrist KW, Warner TF, Mahvi DM. Distribution of Hsp-27 and HER-2/neu in in situ and invasive ductal breast carcinomas. Ann Surg Oncol. 2(1995) 43-48.
- [52] Porter W, Wang F, Duan R, Qin C, Castro-Rivera E, Kim K, Safe S. Transcriptional activation of heat shock protein 27 gene expression by 17beta-estradiol and modulation by antiestrogens and aryl hydrocarbon receptor agonists. J Mol Endocrinol. 26 (2001) 31-42.
- [53] Ciocca DR, Stati AO, Amprino de Castro MM. Colocalization of estrogen and progesterone receptors with an estrogen-regulated heat shock protein in paraffin sections of human breast and endometrial cancer tissue. Breast Cancer Res Treat. 16(1990)243-251.
- [54] Ciocca DR, Puy LA, Edwards DP, Adams DJ, McGuire WL. The presence of an estrogen-regulated protein detected by monoclonal antibody in abnormal human endometrium. J Clin Endocrinol Metab. 60(1985)137-143.
- [55] Navarro D, Cabrera JJ, León L, Chirino R, Fernández L, López A, Rivero JF, Fernández P, Falcón O, Jiménez P, et al. Endometrial stromal sarcoma expression of estrogen receptors, progesterone receptors and estrogen-induced srp27 (24K) suggests hormone responsiveness. J Steroid Biochem Mol Biol. 41(1992)589-596.
- [56] Langdon SP, Rabiasz GJ, Hirst GL, King RJ, Hawkins RA, Smyth JF, Miller WR. Expression of the heat shock protein HSP27 in human ovarian cancer. Clin Cancer Res. 1995 Dec;1(12):1603-9.
- [57] Foster CS, Dodson AR, Ambroisine L, Fisher G, Møller H, Clark J, Attard G, De-Bono J, Scardino P, Reuter VE, Cooper CS, Berney DM, Cuzick J. Hsp-27 expression at diagnosis predicts poor clinical outcome in prostate cancer independent of ETS-gene rearrangement. Br J Cancer. 101(2009)1137-1144.
- [58] Iwaki T, Tateishi J. Immunohistochemical demonstration od alphaB-crystallin in hamartomas of tuberous sclerosis. Am J Pathol. 139 (1991) 1303-1308.
- [59] Takashi M, Katsuno S, Sakata T, Ohshima S, Kato K. Different concentrations of two small stress proteins, alphaB crystallin and HSP27 in human urological tumor tissues. Urol Res. 26(1998) 395-399.

- [60] Rigas PK, Kase S, Rao NA. Expression of alpha-crystallins in human sebaceous carcinoma of the eyelid. Eur J Ophthalmol. 19(2009)702-707.
- [61] Kase S, Parikh JG, Rao NA. Expression of alpha-crystallin in retinoblastoma. Arch Ophthalmol. 127(2009)187-192.
- [62] Kase S, Parikh JG, Rao NA. Expression of heat shock protein 27 and alphacrystallins in human retinoblastoma after chemoreduction. Br J Ophthalmol. 93(2009):541-544.
- [63] Hruban RH, Adsay NV, Albores-Saavedra J, Compton C, Garrett ES, Goodman SN, Kern SE, Klimstra DS, Klöppel G, Longnecker DS, Lüttges J, Offerhaus GJ. Pancreatic intraepithelial neoplasia: a new nomenclature and classification system for pancreatic duct lesions. Am J Surg Pathol. 25(2001)579-586.
- [64] Aguirre, A.J., Bardeesy, N., Sinha, M., Lopez, L., Tuveson, D.A., Horner, J., Redston, M.S., and DePinho, R.A. 2003. Activated Kras and Ink4a/Arf deficiency cooperate to produce metastatic pancreatic ductal adenocarcinoma. *Genes & Dev.* 17(2003)3112–3126.
- [65] Abraham, S.C., Wu, T.T., Klimstra, D.S., Finn, L.S., Lee, J.H., Yeo, C.J., Cameron, J.L., and Hruban, R.H. Distinctive molecular genetic alterations in sporadic and familial adenomatous polyposis-associated pancreatoblastomas: Frequent alterations in the APC/-catenin pathway and chromosome 11p. Am. J. Pathol. 159 (2001)1619–1627.
- [66] Bardeesy, N., Cheng, K.-h., Berger, J.H., Chu, G.C., Pahler, J., Olson, P., Hezel, A.F., Horner, J., Lauwers, G.Y., Hanahan, D., et al. 2006. *Smad4* is dispensable for normal pancreas development yet critical in progression and tumor biology of pancreas cancer. Genes & Dev. 20(2006) 3130-3146.
- [67] Ijichi, H., Chytil, A., Gorska, A.E., Aakre, M.E., Fujitani, Y., Fujitani, S., Wright, C.V.E., and Moses, H.L. 2006. Aggressive pancreatic ductal adenocarcinoma in mice caused by pancreas-specific blockade of transforming growth factor-signaling in cooperation with active Kras expression. Genes & Dev. 20(2006) 3147-3160.
- [68] Pasca di Magliano, M., Sekine, S., Ermilov, A., Ferris, J., Dlugosz, A.A., and Hebrok, M. 2006. Hedgehog/Ras interactions regulate early stages of pancreatic cancer. Genes & Dev. 20(2006) 3161-3173.

Figure Legends:

Figure 1. Expression of αA-crystallin in human normal tissues. (A) Immunohistochemistry of αA-crystallin in human lens, pancreas, liver, kidney, brain, heart, lung, esophagus, testis, ovary, and lymph node. Hematoxylin-Eosin staining (H.E. staining) results of various tissues were included for comparison (column 1). Nuclei were visualized by Hoechst staining (blue color in column 2). αA-crystallin stained in green was derived from FITC (Column 3). Merge showed combination of Hoechst in blue and αA-crystallin in green (Column 4). Mouse lens section from embryonic day 11.5 was stained as positive control for αA-crystallin expression. Primary antibody was replaced by normal IgG in negative control (last row). Note that normal human pancreas displays moderate level of αA-crystallin signal. Liver also displays αA-crystallin signal but the intensity is less (see Fig. 1B for other tissues). Scale bar: 200 μm. (B) Quantification of fluorescence intensity. The mean intensity of fluorescence was measured by Carl Zeiss Laser Scanning System LSM 510.

Figure 2. Western blot analysis of αA - and αB -crystallins in 11 tissues from normal adult mouse (4 weeks). Various tissues were dissected and total proteins were extracted from the 11 tissues indicated. For the lens sample, 0.5 μg of total protein was used for Western blot analysis. For other samples, 100 μg of total protein were used. The blot was probed with specific antibodies against αA -crystallin, αB -crystallin antibody, β -tubulin or β -Actin at a dilution of 1:1000 for the primary antibody and at a dilution of 1:1500 for the secondary antibody. Note that the mouse pancreas expresses a distinct level of αA -crystallin but only a trace of αB -Crystallin.

Both β -actin and β -tubulin are absent in lane 1 (lens) and lane 2 (heart) due to limited sample loaded (only 0.5 μ g lens protein in lane 1) or due to tissue characteristics (lacking β -actin and β -tubulin).

Figure 3. Expression of α A-crystallin in human pancreatic cancer tissues. (A) Immunohistochemistry results of α A-crystallin in human pancreatic cancer. As comparison, immunohistochemistry results of α A-crystallin in two human normal pancreas samples were included. Hematoxylin-Eosin staining (H.E. staining) results of two normal and two cancerous pancreatic sections were included for comparison (column 1). Nuclei were visualized by Hoechst staining (blue color in column 2). αA-crystallin stained in green was derived from FITC (Column 3). Merge showed combination of Hoechst in blue and αA-crystallin in green (Column 4). Mouse lens section from embryonic day 11.5 was stained as positive control for αA-crystallin expression. Primary antibody was replaced by normal IgG in negative control (last row). Scale bar: 200 µm. Note that normal human pancreas sample A displays moderate level of α A-crystallin signal, and normal human pancreas sample B shows weaker signals than the sample A. In the cancerous sample A, the level of α Acrystallin signal was much weaker than that in normal human sample A and also slightly weaker than that in normal human sample B. (B) Quantification of fluorescence intensity. The fluorescence intensity was measured by Carl Zeiss Laser Scanning System LSM 510. Each spot indicates the value of fluorescence intensity from a single tissue sample. Each short solid line indicates mean value of fluorescence intensity of one type of pancreatic cancer. Specifically, mean intensity of normal pancreas is 283±267 (n=11 among which 3 samples have a value less than 100 and 8 samples have a value more than 200); mean intensity of pancreatic duct

adenocarcinoma is 9 ± 16 (n=48); mean intensity of pancreatic mucinous adenocarcinoma is 3 ± 5 (n=3); mean intensity of pancreatic islet adenocarcinoma is 56 ± 69 (n=6) and mean intensity of pancreatic acinic cell carcinoma is 128 ± 55 (n=3).

Figure 4. Expression of αA-crystallin in pancreatic cancer cell line, Miapaca, upregulates expression of c-Jun and c-Fos. The stable clones of miapaca cells expressing pEGFP, pEGFP-HαA or pEGFP-HαB were grown to 95% confluence in MEM with 10% FBS and 400 μ g/ml G418, and then harvested for extraction of total proteins which were subjected to Western blot analysis for AP-1 components, c-Jun and c-Fos, in two different cellular fractions: cytoplasmic and nuclear portions. Note that miapaca cells expressing αA-crystallin display upregulated levels of c-Jun and c-Fos protein. In addition, the phospho-c-Jun level was also enhanced in pEGFP-HαA-Miapaca cells.

Figure 5. Expression of αA-crystallin in pancreatic cancer cell line, Miapaca, upregulates AP-1 DNA binding activity (A) and transactivity (B). A. Gel mobility shifting assay demonstrates the existence of AP-1 DNA binding activity in parent miapaca cell nuclear extract (Lane 13), pEGFP-Miapaca cell nuclear extracts (Lane 2 to 9), which was much lower than that in Miapaca cell nuclear extracts expressing αA-crystallin (Lane 11) or αB-crystallin (Lane 10). Nuclear extracts prepared from three types of cells were incubated with γ -32P-ATP-labeled oligos containing wild-type AP-1 binding site (up panel of Fig. 5A left part) under various conditions shown in the figure. Lane 1, gel mobility shifting assay with labeled wild type AP-1 oligo. Lane 2, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-Miapaca nuclear extract. Lane 3, the same assay as in lane 2 except that 50-fold of

non-labeled wild-type AP-1 oligos was added into the reaction. Lane 4, the same assay as in lane 3 except that the non-labeled competing oligo contains a mutated AP-1 binding site (bottom panel of Fig. 5A left part). Lane 5, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-Miapaca nuclear extract pre-cleared by anti-c-Jun antibody. Lane 6, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-Miapaca nuclear extract pre-cleared by anti-c-Fos antibody. Lane 7, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-Miapaca nuclear extract pre-cleared by anti-c-Jun antibody and anti-c-Fos antibody. Lane 8, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-Miapaca nuclear extract pre-cleared by normal IgG. Lane 9, the same assay as in lane 2. Lane 10, the same assay as in lane 2 except that the nuclear extracts were from the pEGFP-HαB-Miapaca cells. Lane 11, the same assay as in lane 2 except that the nuclear extracts were from the pEGFP-HαA-Miapaca cells. Lane 12 (mock for AP-1 site specificity), the same assay as in lane 2 except that the nuclear extract was replaced with in vitro synthesized p53 protein. Note that p53 does not bind to AP-1 site. B. Luciferase assays to demonstrate that the AP-1 transactivity in pEGFP-HαA-Miapaca cells was significantly higher than that in pEGFP-Miapaca cells. The luciferase reporter gene driven by a mini-prolactin gene promoter without enhancer (pProlactin-Luc), or with 1 copy of AP-1 binding site (pProlactin-1xAP-1-Luc), or driven by the c-Jun promoter (pJun-Luc), together with a reference plasmid were separately transfected into pEGFP-Miapaca cells, pEGFP-HαA-Miapaca cells, or pEGFP-HαB-Miapaca cells. After 48 hours, the transfected cells were harvested for the assaying of luciferase activity. After calibration with the luciferase activity from the reference plasmid, the transactivity of AP-1 in each type of cells was presented. Note that as assayed by both constructs, pProlactin-1xAP-1-Luc and pJun-Luc, the AP-1 activity

in pEGFP-Miapaca cells were significantly lower than that in pEGFP-H α A-Miapaca cells (p < 0.005 for pJun-Luc construct, and p < 0.05 for pProlactin-1xAP-1-Luc construct). In contrast, the AP-1 activity in pEGFP-Miapaca cells is only slightly lower than that in pEGFP-H α B-Miapaca cells (p > 0.05 in both cases).

Figure 6. αA-crystallin enhances MAPK (ERK2) activity in the in vitro kinase assay. A. The substrate of Elk-1 protein was phosphorylated by ERK2 kinase in the in vitro assay in the presence of ATP. Phosphorylation of Elk was detected by Western blot analysis as described above. Results of two different exposures were shown. Kinase reactions with GSK3β as kinase were also included as control for ERK2. Quantitative results of the Elk phosphorylation under different conditions. The relative pixel of each band was quantitated using the software from Silk Scientific Inc. Note that at the concentration of 0.1 and 1ng ERK2 kinase conditions, addition of 1 or 10 μg of αA-crystallin significantly enhances the ERK2 activity as reflected by the differential intensity of Elk phosphorylation. C. Western blot analysis of expression and activities of ERK1/2 in vector- or α A-transfected miapaca cells. Western blot analysis was conducted as described in Fig.2. Note that both expression and activity of ERK2 was upregulated in miapaca cells expressing αA-crystallin than vectortransfected cells.

Figure 7. αA-crystallin modulates expression of TGF β as demonstrated by RT-PCR analysis. **A.** RT-PCR. Total cellular RNAs were extracted from vector-, αA- or αB-crystallin-transfected cells, then were used for RT-PCR assays as described in the experimental procedures. B. Quantitation of TGF β mRNA level in vector-, αA- or αB-crystallin-transfected miapaca cells. C. Quantitation of TGF β receptor 3 mRNA

level in vector-, αA - or αB -crystallin-transfected miapaca cells. Note that αA crystallin enhances expression of TGF β mRNA but has little effect on TGF receptors.

Figure 8. αA-crystallin modulates protein expression of TGF β , and Smad3/5, as well as the phosphorylation status of Smad3/5. Western blot analysis was conducted as described in Fig. 2. Note that αA-crystallin enhances expression of TGF β , Smad3 and Smad5. Moreover, αA-crystallin up-regulates the phosphorylation status of Smad3/5.

Figure 9. Demonstration that the pancreatic cancer cell line, miapaca, expressing the exogenous HαA-Crystallin display retarded cell migration. The stable transfected cell lines, pEGFP-Miapaca, pEGFP-HαA-Miapaca or pEGFP-HαB-Miapaca cells were seeded in 6-well plates and grown until 95% confluent. Then, two straight scratches were made by using a 1 ml blue pipette tip to simulate the wound healing in each well. After PBS washing for 2 times, new DMEM medium was added for a continuous growth of another 48 hours. The wound healing process was recorded daily using the Leica Inverted Phase Contrast Fluorescence Microscopy with a 10x objective. Note that the pEGFP-HαA-Miapaca cells display slower healing process than pEGFP-Miapaca and pEGFP-HαB-Miapaca cells, suggesting both slower cell migration and proliferation of pEGFP-HαA-Miapaca cells.

Supplementary Figure 1. Expression of α A-crystallin in mouse lens epithelial cell line, α TN4-1, also up-regulates expression of c-Jun and c-Fos. The stable clones of α TN4-1 cells expressing pEGFP, or pEGFP-H α A were grown to 95% confluence in

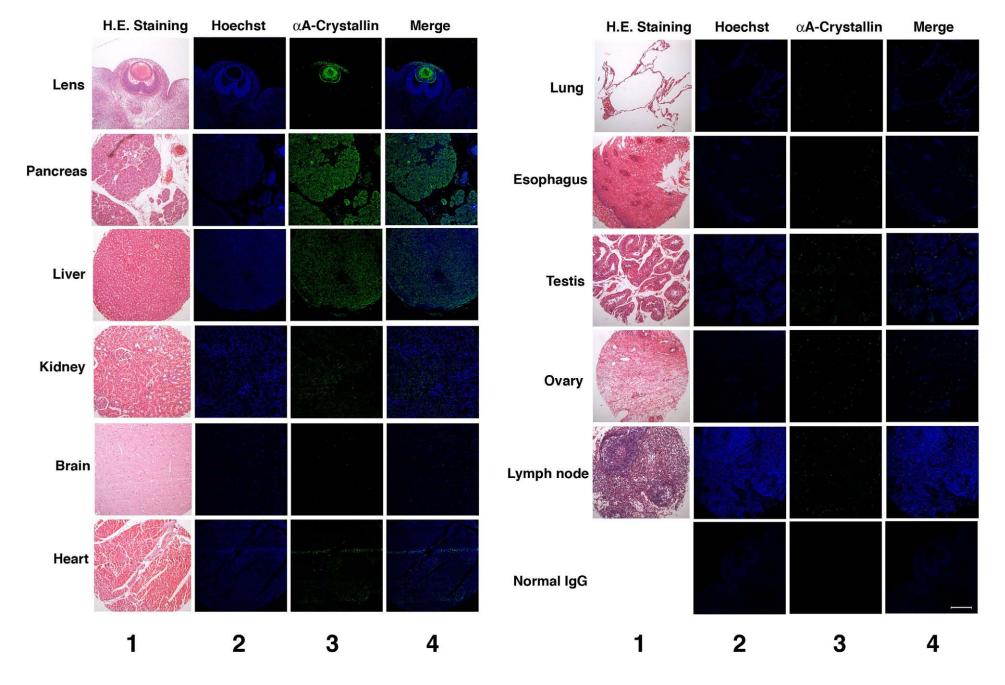
MEM with 10% FBS and 400 μ g/ml G418, and then harvested for extraction of total proteins which were subjected to Western blot analysis for AP-1 components, c-Jun and c-Fos, in two different cellular fractions: cytoplasmic and nuclear portions. Note that α TN4-1 cells expressing α A-crystallin display upregulated levels of c-Jun and c-Fos protein.

Supplementary Figure 2. Expression of α A-crystallin in mouse lens epithelial cell line, αTN4-1 upregulates AP-1 DNA binding activity. Gel mobility shifting assay demonstrates the existence of AP-1 DNA binding activity in pEGFP–αTN4-1 cell nuclear extracts (Lane 2 to 8), which was much lower than that in pEGFP- $H\alpha A - \alpha TN4-1$ nuclear extracts (Lane 9). Nuclear extracts prepared from either pEGFP- α TN4-1 cells or pEGFP-H α A- α TN4-1 cells were incubated with γ -³²P-ATP-labeled oligos containing wild-type AP-1 binding site (up panel of Fig. 5A left part) under various conditions shown in the figure. Lane 1, gel mobility shifting assay with labeled wild type AP-1 oligo. Lane 2, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-αTN4-1 nuclear extract. Lane 3, the same assay as in lane 2 except that 50-fold of non-labeled wild-type AP-1 oligos was added into the reaction. Lane 4, the same assay as in lane 3 except that the non-labeled competing oligos contain a mutated AP-1 binding site (bottom panel of Fig. 5A left part). Lane 5, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-αTN4-1 nuclear extract pre-cleared by anti-c-Jun antibody. Lane 6, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-αTN4-1 nuclear extract pre-cleared by anti-c-Fos antibody. Lane 7, gel mobility shifting assay with labeled wild type AP-1 oligo and pEGFP-αTN4-1 nuclear extract pre-cleared by anti-c-Jun antibody and anti-c-Fos antibody. Lane 8, the same assay as in lane 2. Lane

9, the same assay as in lane 2 except that the nuclear extracts were from the pEGFP-H α A- α TN4-1 cells.



Fig. 1A Deng et al.



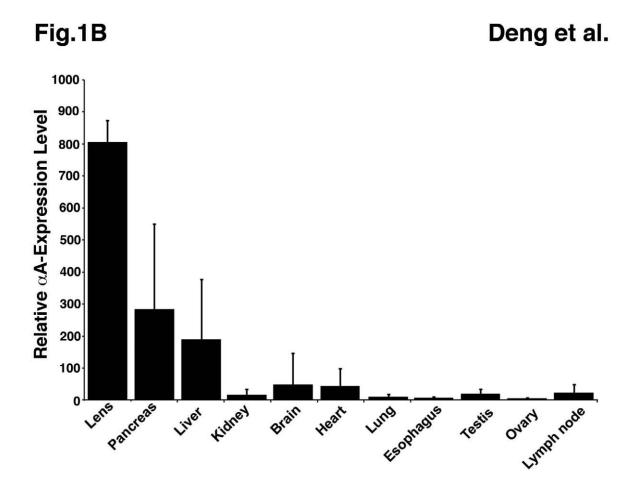


Fig.2 Deng et al.

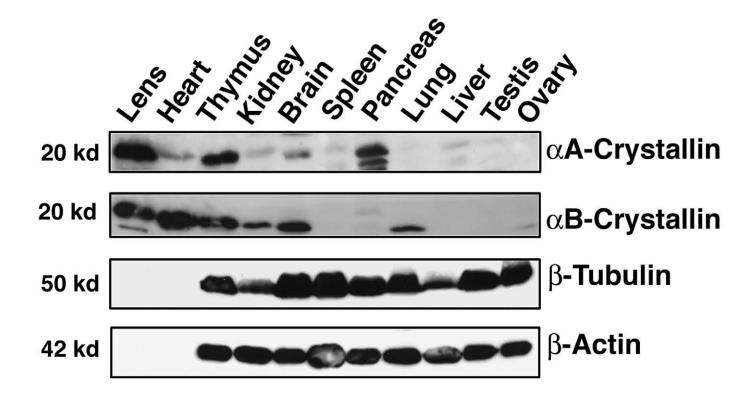
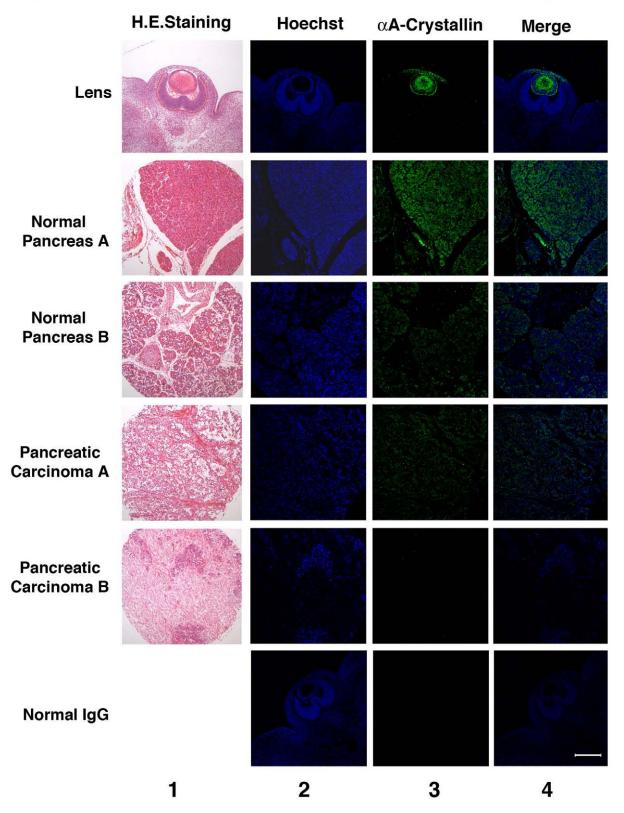


Fig. 3A Deng et al.



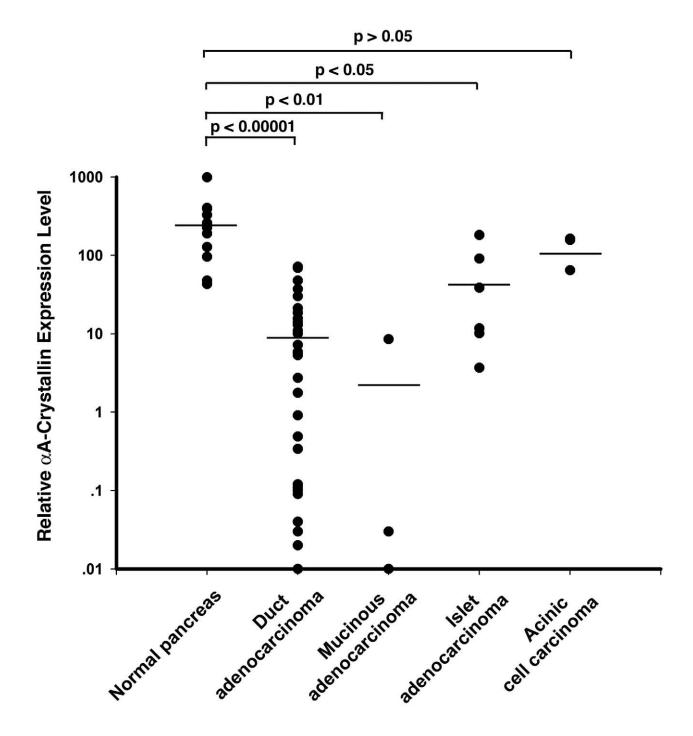
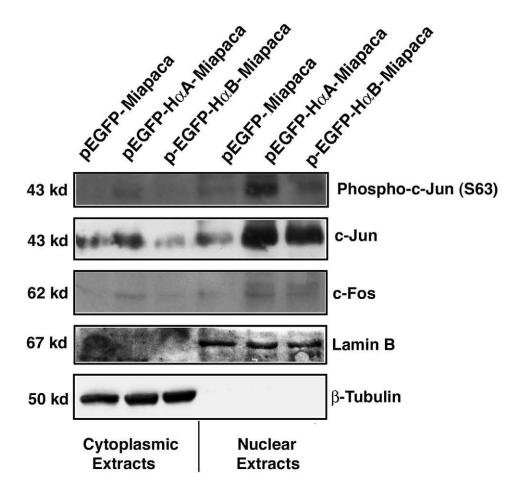
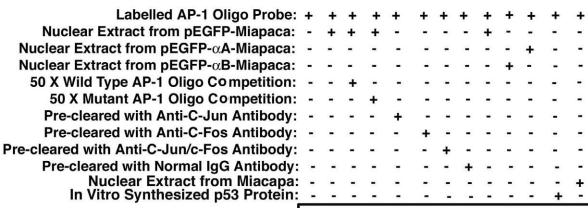
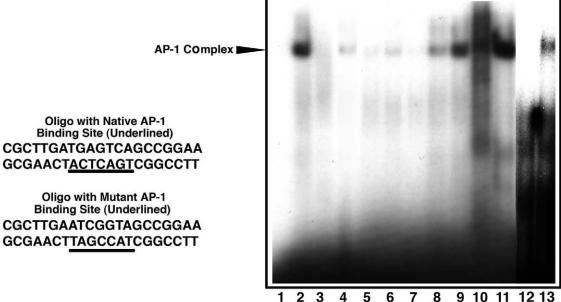


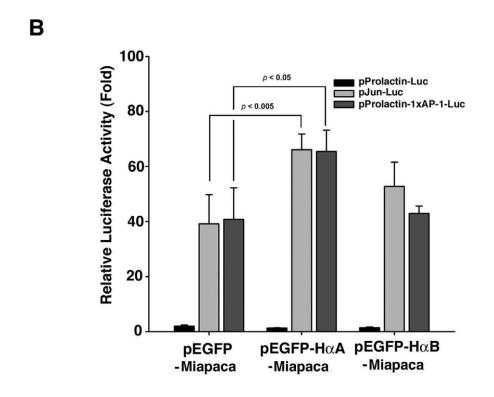
Fig. 4 Deng et al.



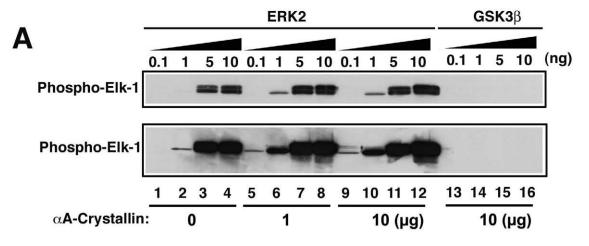
Deng et al.

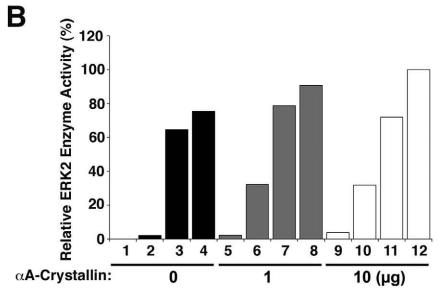


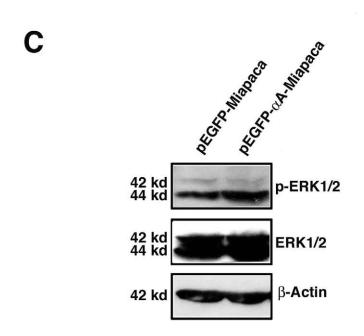




Deng et al.







Deng et al.

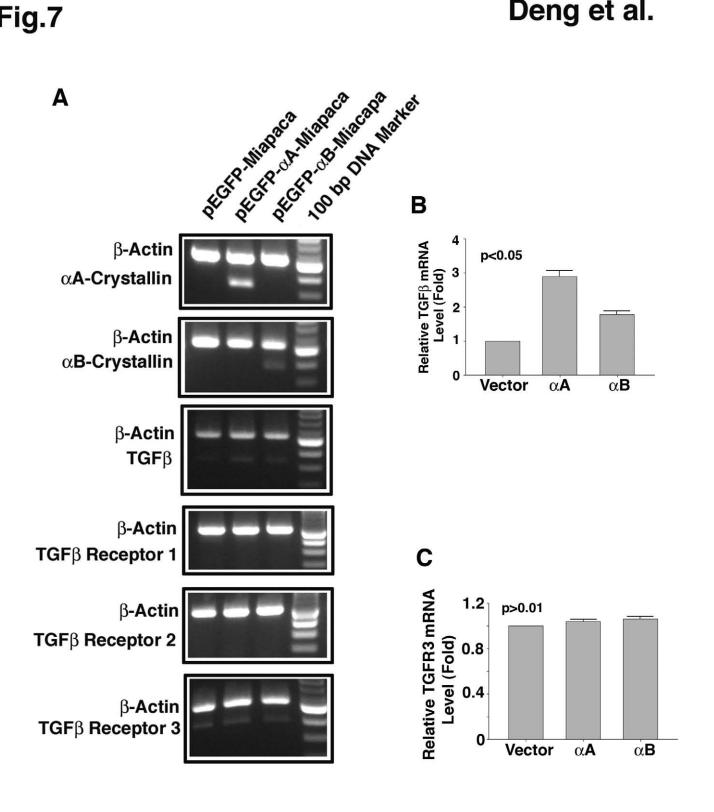


Fig.8 Deng et al.

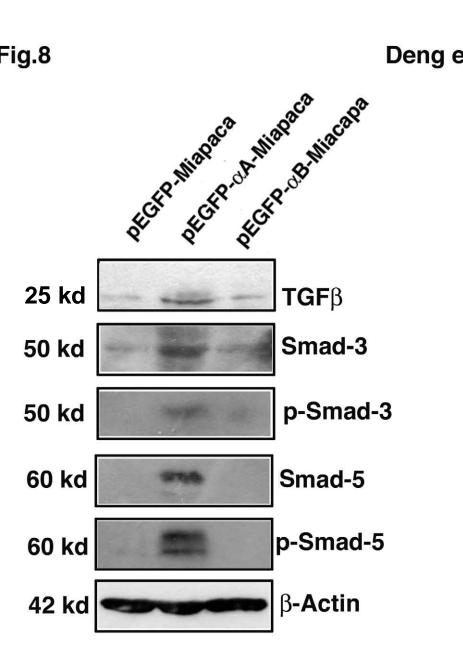
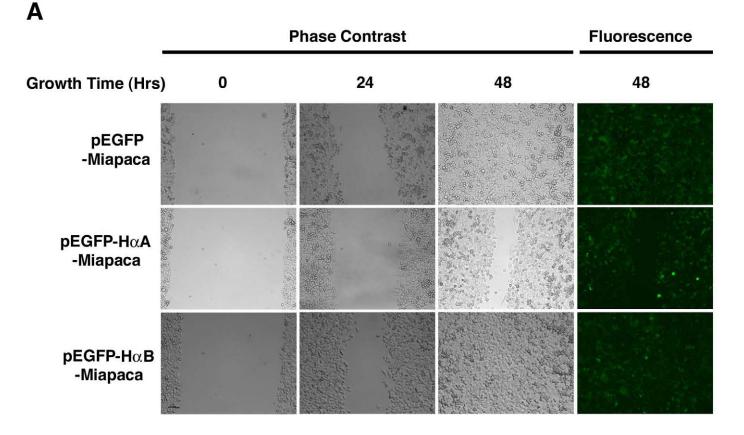
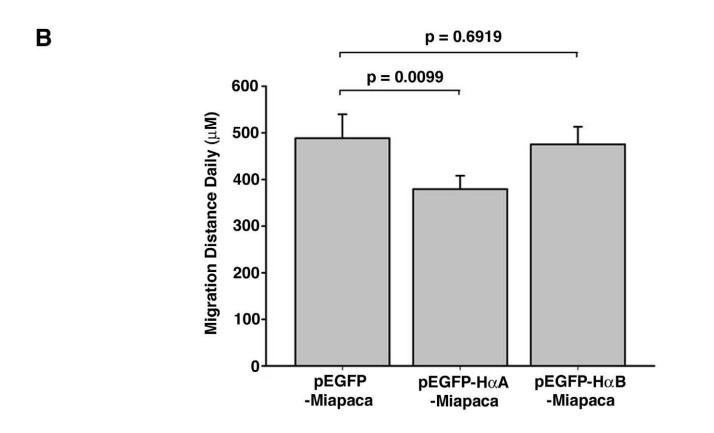
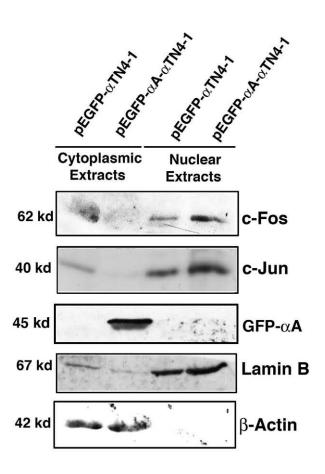


Fig.9 Deng et al.





Supplementary Fig. 1 Deng et al.



Supplementary Fig. 2

Deng et al.

AP-1 Complex **■**