

## PPDPF cDNA Cloning from Two *Bufo* Species and Tissue Expression in *B. gargarizans*

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**Abstract:** Cinobufacini injection was aqueous extract of toad (*Bufo*) dried skin, which has been widely used for cancer therapy in China. Recent studies indicated that the antitumor components included in cinobufacini are polypeptides. To identify these components, the skin plasmid cDNA library of Japanese toad (*B. japonicus formosus*) was screened. One of the transcripts is 864 base pair (bp) consisting of 144 bp 5'-untranslated region (UTR), 384 bp 3'-UTR and an open reading frame (ORF) of 336 bp encoding a polypeptide of 111 amino acid residues. Homology analysis indicated 86% homolog with pancreatic progenitor cell differentiation and proliferation factor (PPDPF) of *Xenopus laevis*, which implicated in pancreas development. PPDPF ORF of Chinese toad (*B. gargarizans*) was cloned from its skin first strand cDNA encoding completely the same polypeptide as that of Japanese toad. Real-time quantitative polymerase chain reaction (qPCR) indicated PPDPF expressed in all organs of Chinese toad tested so far, while the highest expression is detected from pancreas. Considering the previous reports on the relevance of abnormal expression of PPDPF with several cancers including pancreatic cancer, PPDPF might be one of the crucial antitumor components of *Bufo* skin, so the successful cloning of PPDPF from *Bufo* skin is likely to shed light on some new antitumor drug development in future.

**Keywords:** *Bufo japonicus formosus*; *Bufo gargarizans*; PPDPF cloning; qPCR

### 1. INTRODUCTION

Amphibian skin and its secretions are rich of large amounts of biologically active compounds having great potential in new drug discovery [1-3]. In China, *Bufo*-origin materials such as toad skin (Chan'pi), cortex (Chan'yi), secretions (Chan'su) and its whole dried body (Gan-chan) were widely used in many prescriptions of traditional Chinese medicine for clinical treatments of many kinds of diseases, especially tumor control [4-7]. Cinobufocini injection, aqueous extract of toad dried skin, is a such kind of clinical drug especially aimed at advanced cancer [8] including pancreatic cancer [9]. Recent studies indicated that peptides prepared from cinobufacini injection showed the same antitumor activity as that of the injection itself, which suggests that the polypeptides included in *Bufo* skin are the main active antitumor ingredients [10]. To analyze these polypeptides, the authors have been trying to clone cDNA from the skin cDNA plasmid library of Japanese toad (*B. japonicus formosus*) as well as from the skin first strand cDNA of Chinese toad (*B. gargarizans*) [11-13]. During these processes, the cDNA encoding pancreatic progenitor cell differentiation and proliferation factor (PPDPF) was obtained from both Japanese toad and Chinese toad.

PPDPF, also known as EXDPF (exocrine differentiation and proliferation factor) was first cloned from zebrafish, whose encoding protein has been reported as a decisive factor of the exocrine pancreas tissue development [14]. More interestingly, higher level of PPDPF ortholog has been detected in pancreatic cancer, breast cancer and kidney cancer [14], indicating that super expression of PPDPF is likely to be involved in the pathogenesis of malignancy. Here, PPDPF is cloned from two *Bufo* species, whose expression has been detected in the organs of Chinese toad by qPCR (real-time quantitative polymerase chain reaction), which will be reported here.

## 2. MATERIALS AND METHODS

### 2.1. Experimental Materials and Reagents

Japanese toad skin plasmid cDNA library held by the Japan Advanced Industrial Science and Technology (AIST, Tsukuba, Japan) was authorized Zhejiang Agricultural and Forestry University (ZAFU) for research as part of a Material Transfer Agreement. Concerning this library construction, as reported previously [12], pSD64TR (3250 base pairs) has been used as a vector, whose upstream primer is SP6 (5'-ATTTAGGTGACACTATAGAA-3') and the downstream one is S.D.A. (5'-TTATGTAGCTTAGAGACTC-3'), and *Eco*R I and *Xho* I as cloning sites, and the length of the cDNAs ranged from 500 - 2 000 base pairs (bp). The first strand cDNAs of Chinese toad skin were prepared from the individuals from the East lake Campus of ZAFU [11].

The RNA extraction kit was purchased from Shanghai Bocai Biotechnology Company; pGM-T vector, *Escherichia coli* competent cells (DH5 $\alpha$ ) and Real Master Mix (SYBR Green) for qRT-PCR from Beijing Tiangen Biotechnology Limited Company; PCR kit (PrimeStar Max DNA polymerase) and PrimeScript<sup>TM</sup> II 1<sup>st</sup> strand cDNA Synthesis kit from TaKaRa; and the primer synthesis and DNA sequencing were done by Shanghai Sang'ni Biotechnology Company.

### 2.2. *PPDPF* cDNA Cloning of *B. japonicus formosus* and *B. gargarizans*

cDNA cloning from Japanese toad was followed the method reported previously [12]. Briefly, Japanese toad skin plasmid cDNA library was transformed into *E. coli* (DH5 $\alpha$ ), and colony PCR was done using single colony suspension in LB medium as template, and *Xho*TT (5-AGATCTCTCGAGTTTTTTTTTTTTT-3, a self-designed primer complementary with the area compassing the connection point of cDNA polyA tail and the downstream cloning site of *Xho* I) and SP6 as primers. Following this process, the recombinant plasmids were extracted and sequenced.

For the cloning of *PPDPF* ORF from Chinese toad, its dorsal skin total RNA was extracted and its first strand cDNAs were synthesized according to the manufacturer's protocol [11, 12]. A set of primers was designed based on *PPDPF* cDNA sequence of Japanese toad (*PPDPF*-S: 5-CATCTTAAGTCCAAGCAAG-3; *PPDPF*-R: 5-ATGCATGGAGGGATTAGG-3), which was also used for qPCR. Then PCR product was ligated into a pGM-T vector and sequenced with its primers of SP6 and T7 (5'-TAATACGACTCACTATAGGG-3').

### 2.3. Sequence Analysis and the Phosphorylation Site Prediction

DNAstar/EditSeq was used to find out the ORF and deduce the amino acid sequence of the encoding protein. The potential phosphorylation sites of *PPDPF* were predicted via Net Phos 2.0 (<http://www.cbs.dtu.dk/services/NetPhos/>).

### 2.4. Homology Analysis of *PPDPF* Amino Acids

Eleven *PPDPF* amino acid sequences of other animals downloaded by NCBI blast program (<http://blast.ncbi.nlm.nih.gov/Blast.cgi>) and *PPDPF* of Japanese toad and Chinese toad were aligned by MEGA 5.2, which was also used to construct the phylogenetic tree of *PPDPF* (bootstrap with 1000 replications).

### 2.5. *PPDPF* Tissue Expression Analysis by qPCR

Total RNA was extracted from each organ of *B. gargarizans* including brain, heart, intestines, kidney, liver, lung, oviduct, pancreas, skin, spleen, stomach and testis, and the first strand cDNA was synthesized as mentioned above. qPCR was performed for analysis of *PPDPF* expression in Chinese toad by Mx3000P (Stratagene, Agilent Technologies Inc., La Jolla, CA, USA). Here  $\beta$ -*actin* was used as a reference gene (upstream primer: 5-TTGAGACCTTCAACACC-3, downstream primer: 5-CTTGATGTCACGCACA A-3). Three sets of the first strand cDNA of each organ from 3 individuals were used. Data was analyzed by MxPro Comparative Quantitation and MS excel.

## 3. RESULTS

### 3.1. *PPDPF* Cloning of *B. japonicus formosus* and *B. gargarizans*

Sequencing analysis indicated a transcript of 864 bp consisting of 144 bp 5'-untranslated region (UTR), 384 bp 3'-UTR and an ORF of 336 bp, whose encoding polypeptide is consisting of 111 amino acid

residues (Fig. 1) with high homology with PPDPF of other animals. This transcript has been deposited into GenBank and named as *B. japonicus formosus* PPDPF with an accession number of KJ128292.

From the skin first strand cDNAs of Chinese toad, 44 clones were obtained. Twenty-six out of them showed the same ORF of 336 bp with only one nucleotide difference from the ORF of Japanese toad PPDPF (Fig. 1).

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GTTTGTGCGTGGAGACAAGTCCCAGCTTCTCCTCC TCAAAAAGGGCCCCGTGCGCTAGCGTCTAACC CCCAGAGCCAGGCTC 80
CAGAGGACGTATACAGTCGGAATACAGGGTTTACATCATCTTAAGTCCAAGCAAGGTTTGA AACATGCGCAGCGATTCAT 160
                                     M A A I P
CCAGTGGCTCACTTGTGCGCAACACATGATTACTATCGTAGACGCCTGGGATCCACCTCTAGTAACAGCTCATGTGGGAGC 240
S S G S L V A T H D Y Y R R R L G S T S S N S S C G S
GTGGACTACTCTGGAGAGGTCATTCCTCACCACCCAGGCTCTCCAAAGTCAGATCCTGGTCACTGGTGGGCCAGCTTCTT 320
V D Y S G E V I P H H P G L P K S D P G H W W A S F F
TTTTGGTAAACCATCTCATCCC GTTATGACCACTGTTTCGGAATCCCCGGAGA AACTCAGGAAGCTTGCGCATGACCAATG 400
F G K P S H P V M T T V S E S P E N S G S L R M T N
GCC TTTTCCCCTGCGGCC TGGCTCAGGAGCCAGT GAGGAAGAACAGTCTCAATGAGTCC AAGACTGACTCCAGCACCTAA 480
G L F P C G L A Q E P V R K N S L N E S K T D S S T *
GCC TAATCCC TCCATGCA TCCCC TCC TACTGGAAAGACGCTGGGTGGCGGAGCCCAATACCCTCCCCATCCCCATCCC 560
CTACTCTTGACCCATAGGTTGCCCTTACTATAGCTTTAGTCACC AATCC TACCTTTTTGTTAATGTTGATTTGAAGAATA 640
ATGCTTTTTTACTACGTCAGTAACACTGAAACAT TCTTAATTGTTGTT TCCCTGCCTTACCATGTAATGTTTTTATTTTC 720
TAC TTCTGTATAAAGATGGGCAGCCTAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 800
AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 864
    
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**Figure 1** PPDPF cDNA and its deduced amino acid sequence of *Bufo japonicus formosus*

□: Start and stop codons; Nucleotide in shadow: Substituted by “T” in Chinese *Bufo*; -: Poly (A) tail; ==: Polyadenylation signal; ....: Primers of PPDPF-S and PPDPF-R.

Due to this nucleotide substitution did not result in the change of its encoding amino acid, the encoding polypeptide was completely the same as PPDPF of Japanese toad. The cDNA clone appeared in high frequency has been deposited into GenBank (accession number: KJ128291) and named as *B. gargarizans* PPDPF. Concerning the remaining 18 clones, they showed 1 or more nucleotide difference and were different from each other indicating the possible artificial introduction of nucleotide substitution during the experimental processes in vitro including PCR (data not shown).

### 3.2. Phosphorylation Site Prediction of *Bufo* PPDPF

Thirteen potential phosphorylation sites (Ser9, Ser23, Ser25, Ser28, Ser29, Ser36, Ser74, Ser101, Ser105, Ser109, Ser110, Tyr69, and Tyr70) were found in PPDPF of two *Bufo* species, which suggests that the function of PPDPF might be regulated by the upstream factors (Table 1).

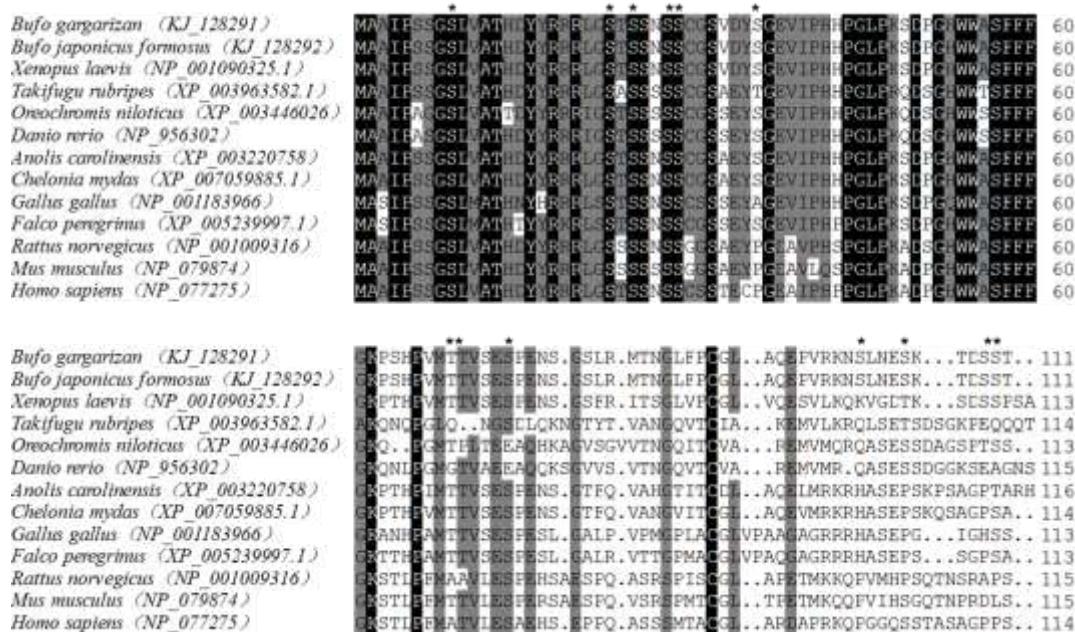
**Table1.** Potential phosphorylation sites (shade) of PPDPF two *Bufo* species

Position	Context	Score	Position	Context	Score
9	PSSG SAT	0.629	70	PIMTTVSES	0.789
23	RRLG STSSN	0.990	74	TVSE SPENS	0.998
25	LGST SSNSS	0.764	101	VRKNSL NES	0.995
28	TSSNS SCGS	0.970	105	SLNES KIDS	0.991
29	SSNS SCGSV	0.944	109	SKTDS ST	0.538
36	SVDYS GEVI	0.966	110	KTDS ST	0.690
69	HPIMTTVSE	0.898			

### 3.3. Homology Analysis of PPDPF

PPDPF homology analysis of two *Bufo* species indicated 86% homology with *Xenopus laevis*, 65% with *Homo sapiens*, and 59%-80% among 9 other animals (Fig. 2).

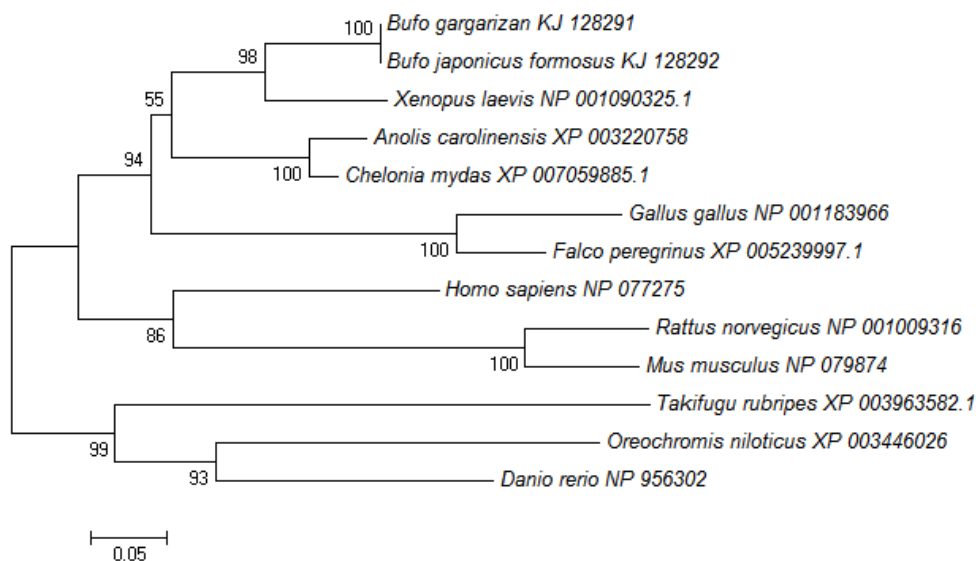




**Figure 2** Multiple amino acid sequence alignment of PPDPF of *Bufo* with that of 11 different animals

■: 100% homology; ■: Above 75% homology; □: Homology less than 75%. \*: Potential phosphorylation sites of *Bufo* PPDPF.

Based on the phylogenetic tree (Fig. 3), 3 fish (*Takifugu rubripes*, *Oreochromis niloticus*, *Danio rerio*), 3 amphibian species (*X. laevis* as well as two *Bufo* species), 2 reptiles (*Anolis carolinensis*, *Chelonia mydas*), 2 birds (*Gallus gallus*, *Falco peregrinus*), and 3 mammals (*Homo sapiens*, *Mus musculus*, *Rattus norvegicus*) gathered in one branch, which is consistent with the traditional animal taxonomy. Among 13 potential phosphorylation sites of *Bufo*, the first five were highly conserved among these 13 animals (Fig. 2).

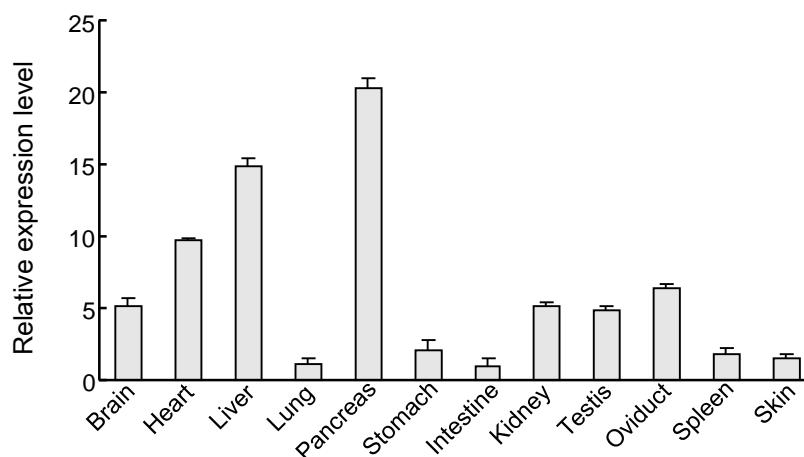


**Figure 3** Phylogenetic tree of PPDPF constructed by MEGA 5.2

The scale bar represents the number of changes per amino acid position.

### 3.4. PPDPF Expression Analysis in Different Organs of *Bufo gargarizans* by qPCR

qPCR analysis of *B. gargarizans* in the brain, heart, lung, liver, pancreas stomach, intestines, kidney, testis, oviduct, spleen and skin showed that *PPDPF* was expressed in all these organs, and the expression level was the highest in pancreas (Fig. 4).



**Figure 4** PPDPF expression in different organs of *Bufo gargarizans*

Abscissa represents the organs of *Bufo gargarizans* being tested, ordinate represents relative expression quantity relative to  $\beta$ -actin. Error bars show square deviation of each organ from 3 individuals.

#### 4. DISCUSSION

So far, there are many reports concerning on the pharmaceutical functions of *Bufo*-skin and its secretions [3, 15, 16]. In recent years, several reports clearly indicated the antitumor effects of the polypeptides extracted from cinobufacini injection [9, 17] which comes from toad skin. Therefore, it is both essential and urgent to identify these polypeptides. Authors have been screening/cloning the cDNA from the skin plasmid cDNA library of Japanese toad and the skin first strand cDNA of Chinese toad to contribute such studies [11-13], trying to seek out related factors.

In current study, PPDPF cDNA was successfully cloned from skin of Japanese toad and Chinese toad both encoding the same polypeptide consisting of 111 amino acid residues (Fig. 1) having high homology with that of other vertebrates (Fig. 2, 3). Many potential phosphorylation sites indicate PPDPF function might be upstream regulated (Table 1). qPCR showed PPDPF expressed in all the tested organs of Chinese toad (Fig. 4) indicating the therapeutic values of the whole body of *Bufo* but not limited to its skin.

Despite the fewer studies of PPDPF/EXDPF so far, it's certainly a decisive factor of the exocrine pancreas tissue development due to the PPDPF knocking down leading to cell cycle arrests through up-regulation of p21<sup>Cip</sup>, p27<sup>Kip</sup> and a severe reduction of the cell number of exocrine progenitor by 70%; however, the overexpression of PPDPF led to increased exocrine size indicating its involvement in the pancreatic development in *Danio rerio*<sup>[14]</sup>.

Pancreatic cancer is one of the leading causes of cancer deaths as it's often highly aggressive and resistant to treatments available at the time of diagnosis [18]. In addition, most malignant pancreatic tumors are derived from exocrine portion [19]. Interestingly, the EST (Expressed Sequence Tag) expression profile showed higher level of PPDPF ortholog has been detected in several cancers including pancreatic cancer, breast cancer and kidney cancer, indicating that super expression of PPDPF is likely to be involved in the pathogenesis of malignancy [14], suggesting the possible pivotal role towards cancer. In a word, the current study will lay foundation for exploring new antitumor drug to control cancer.

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