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# Molecular characterization and expression profiles of *cyclin A* and *cyclin B* during ovarian development of the giant tiger shrimp *Penaeus monodon*

# Virak Visudtiphole<sup>a</sup>, Sirawut Klinbunga<sup>a,b,\*</sup>, Kanyawim Kirtikara<sup>a</sup>

<sup>a</sup> Aquatic Molecular Genetics and Biotechnology Laboratory, National Center for Genetic Engineering and Biotechnology (BIOTEC), National Science and Technology Development Agency, 113 Paholyothin Road, Klong 1, Klong Luang, Pathumthani 12120, Thailand

<sup>b</sup> Center of Excellence for Marine Biotechnology, Faculty of Science, Chulalongkorn University, Bangkok 10330, Thailand

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### ABSTRACT

The meiotic maturation of oocytes is regulated by maturation promoting factor (MPF), a complex of cdc2 (Cdk1) and cyclin B and other Cdk/cyclin complexes. To better understand molecular aspects governing reproductive maturation of the giant tiger shrimp (*Penaeus monodon*), the full length cDNAs and genomic organization of *cyclins A* and *B* (*PMCyA* and *PMCyB*) were characterized. A single form of *PMCyA* contained an open reading frame (ORF) of 1326 bp corresponding to a deduced protein of 441 amino acids. Its genomic sequence contained 5 exons, 4 introns and untranslated regions (UTRs) spanning 2586 bp in length. In contrast, *PMCyB* possessed three isoforms with an identical ORF of 1206 bp (401 amino acids) but three different 3' UTR lengths of 416, 543 and 1117 bp, respectively. Their respective genomic sequences were composed of 8 exons, 7 introns and UTRs covering 4181, 4307 and 4940 bp. Expression levels of brodycA and *PMCyB* in ovaries of broodstock were much greater than those of juveniles (*P*<0.05). During ovarian development and after spawning of normal shrimp broodstock, *PMCyA* was not differentially expressed (*P*<0.05) whereas the level of *PMCyB* (*P*>0.05) but resulted in a lower expression of *PMCyA* at stage IV of ovarian development of this economically important species (*P*<0.05).

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## 1. Introduction

During oogenesis, immature oocytes resume meiosis and fertilization occurs at a particular stage of oocyte maturation depending on species (Kobayashi et al., 1991; Okano-Uchida et al., 1998). Progression through the cell cycle is regulated by association between cyclindependent kinases (Cdks) and their cyclin partners (e.g. cyclin A and cyclin B) governing at different points in the cell divisions (Westendorf et al., 1989; Knoblich and Lehner, 1993).

Cyclins play important roles in the reproductive development of eukaryotes. In all animals, oocyte maturation is regulated by the complex of cdc2 (Cdk1) and cyclin B (called maturation promoting factor, MPF). In various species, MPF has been elucidated to be involved on molecular resumption of arrested oocytes (Kobayashi et al., 1991; Kishimoto, 1999, 2003).

Both cyclins A and B bind to Cdk1 and synergistically allow cells to proceed through the G2-M checkpoint (Minshull et al., 1990; Lees and Harlow, 1993; Li et al., 2004). Nevertheless, they essentially confer different intrinsic roles during the M phase (Knoblich and Lehner, 1993). For instance, cyclin A2 regulates breakdown of the nuclear envelope and the

accumulation of cyclin B1 (Jackman et al., 2003) but cyclin B1 implicates centrosome separation during the cell separation event in both mitosis and meiosis (Gong et al., 2007; Voronina et al., 2003). In addition, cyclin A also forms a complex with Cdk2 to promote the G1-S phase transition for the chromosome replication at the end of meiosis (Pagano et al., 1992) and to potentiate the transcriptional activity of genes encoding estrogen and progesterone receptors (Rogatsky et al., 1999; Narayanan et al., 2005).

Reduced reproductive maturation of female giant tiger shrimp (*Penaeus monodon*) in captivity prevents effective domestication and selective breeding programs for genetic improvement in this species (Preechaphol et al., 2007). In the present study, the full length cDNAs and genomic organization of *P. monodon cyclin A* and *cyclin B* (*PMCyA* and *PMCyB*) were isolated and characterized. The former was reported for the first time in crustaceans. Expression patterns of *PMCyA* and *PMCyB* during ovarian development of normal and eyestalk-ablated *P. monodon* were examined by quantitative real-time PCR.

#### 2. Materials and methods

#### 2.1. Experimental animals

Juvenile shrimp (*P. monodon*; 4 month-old) were purchased from a commercial farm in Chachoengsao (eastern Thailand, N=5). Female

<sup>\*</sup> Corresponding author. Tel.: +66 2 6448150; fax: +66 2 6448190. *E-mail address:* sirawut@biotec.or.th (S. Klinbunga).

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A

152

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-543

536

B

A ACGCGGGCGGGAACTATCCTCATGTGGCGTCAATAGTGTGCGGGTTCTGGTCTTCGTTGG\_60 TGCCTTCCACCCACATATTCAAGGACCACGAGAACCGACTCCCCGCCGCCCACCGGAGAGA 180 A S T H I F K D O E N R V P A A O R R G GCA AGAGA GA GGA CCCGAGGGCGAG CA GTG TG CTT GT TAA CA GA CGGCCA GT GCT CGGTG 240 K R E D P R A S S V L V N R <mark>R P V L G A</mark> CTATCAACCCCAATTTGCCGAAACAACCAGCTAGAGCAGCCAAACAGGGTGTCAGTTATG 3.00 I N P N L R K Q P A R A A K Q G V S Y E AAACGGGTATTCCACAGCAAGCTGATGAAAATGCCTGCCCTACACAGAAATCTTTCTCGA 360 T G I P Q Q A D E N A C P T Q K S F S T CAAGT CAACC CTC TTTTACTATTCATGAAGAT GCATG CTTATCCATGGGAAC CACCAGTA 420 S Q P S F T I H E D A C L S M G T T S N ACAGCCAGAGTGCATCTAGCAATTCCACAGGACCTAACCCATTAAGTGAAAGGTTAAGTA 480 S O S A S S N S T G P N P L S E R L S K AACCATTACAAAGTGAAGAAGAATCTATTGAAGATATTGACCCATCTGTAACAGAATTGC 540 P L O S E E E S I E D I D P S V T E L O

AGAGGTCTCTATCCATCGAACATTCAGGATCTTCAGATGTAATGGAGATGAGTGTATGTG 600 R S L S I E H S G S S D V M E M S V C E AAGATGACCTTATGGTGGTAGAAACCACTCCCCGTGAAGATGTGCTTCATAGTAGAAATG 660 D D L M V V E T T P R E D V L H S R N D ACGACATATTTGATGTCCCAGAATATGCAGCAGACATTTACCAGTACCTGAGAGAAGCAG 720 D I F D V P E Y A A D I Y O Y L R E A E V C H K P R A N Y M S K Q T D I T A S M 

RWILVDWLVEVAEEYSLHTE AAACTTTA TA TTTGGCTGTTTCATA TA TTGACCGGTTTTTATCGCACATGTCTGTTAAAA 900 T L Y L A V S Y I D R F L S H M S V K R GGGATAAACTTCAGCTGGTTGGGACCACAGCTATGTTTATTGCTGCqtaaqtcttcaact 960

DKLQLVGTTAMFIAA gtttattgtggtgttgctatttatggaatattctaaatgtttatatttaataccggtgtc 1020 tgtgctaattacaatctttccacagGAAATATGAGGAGATCTACCCTCCTGACGTAGGAC 1080 K Y E E I Y P P D V G Q

AGTTTGCTTATATAACCGACAATACCTACAGAGTTGGGCAGATCTTGAGAATGGAGCATC 1140 FAYITDNTYRVGQILRMEHL 

I L K V L S F D M A V P T T H L F V N K AGTTTGCCAGATTGTGCAAATGTTCAGAAGAAACCTTGCATTTAGCCTTGgtaggtatta 1260 FARLCKCSEETLHLAL

tgactataacacctaaactttcatttcagTTCTTGGCAGAGGTTACTATGCTAGAATGTG 1380

FLAEVTMLECD ATCCTTTCCTGAGATTCCTGCCTTCTGTGATTGCAGCATCAGCTGTTTCATTAGCAAATC 1440 P F L R F L P S V I A A S A V S L A N H

ATACACAGgttagttttatataattccttacagtatgaactttaaacttcagaggtcaac 1500

taatagtaagcettgtataattaatetggacagatagttgacettgaetatacataeett 1560 ggettgttagettatttaatatttettttacagGGTCACACTGCCTGGCCATCACACATG 1620 GHTAWPSHM

GTAGA ATCCA CAGGA TA CAGTT TAGAA CACCT GAGAGAATGC TATGT AAA CCTACAT CGT 1680 V E S T G Y S L E H L R E C Y V N L H R GTTTTCTCCAGAGTTCATGAACCTCAGCAACACGCTATTAGAGATAAATATAGAGGCACA 1740 V F S R V H E P Q Q H A I R D K Y R G T AAgtaagtteteataatetaaatatteeaagttttgaagattatatgatggtatatattg 1800

aagaccatacaaagcatggacaatttgaaatccacattttaatccaatgctttagattat 1860 acaaggtctgacttggtttctaatatttgcagATGGCACGGAGTGTCCCGCTTAACTCCA 1920

WHGVSRLTP AGAAGCCACTTCCCCTGGTGATGAAACAAAAGTGTAATTAGGGTATGCTGATAATTGTGT 1980

RSHFPW\* GTGAAGATGTTTATGTAAATTTGTGTGGTGGTTGTTTTTCCCCCAAGACTGAATGGAATGTTAG 2040 CACTTTTTATACAGATAAGACAACTCGGTGATCATGAGTGATATACTCGGCTTTTTCCAA 2100 GCCTCTGGTAGAGTAAGGAAGTTTTATCCTTCACAGGATGTCTTATCAGTTGACAGGTGT 2220 AATTTTTTTTTAATAACCTTTTTATATTTTAATAGTGCAAGAAATCACCAGCTTTAA 2280 AAGGCCTGTGTGTAGCAGATCTCTTTTAGCGTTTGAACTGTCATCAGCTTACCAGTCACTGCA 2340 TCGTGGTAACCTTTGCTGCAGCCAAAACATATCAAGTGTTCAGAAAGACATGTACAAATG 2460 GCTGTATATATGTAAACTAGAATCCTTGAAGGATGAAGGATTTTTATCCTTCGATTTTTTA 2520 AAAAAA 2586

GGGAGTGTCGAGTCGCGTCGCTTGAGAGAGGACCTGTTTTGAAGGAGATTTCGCAGTGA 60 AAAAAGCCACACATTCTCGAATATCCGGTTGATTTTTATCCATCATGTCTTTGAGAACCA 120 MSLRTT

 ${\tt CCACGCATCTCAACAGT} g {\tt taagttccaagacgggaagacggcagaaagggcgagaaagg 180}$ THLNS

ttetattteetetageactatategecateggtatttagaaceteteeagtgeteggett 360 cccqtaqcqqqatcqqqaatatttattattaqqaatatttqaattqqaqaattctqtqqc 420 ggttatttatttgtttgttttcgtatgttttttttactttctatgtctgttatacttcca 480 tcqtttttaatqqtttqatqtctqtaattqacqccttcaaccttttacqtqaaqaatatt 540  $ccaattgctttgtaatttgaatatttaaaatgacttgggatatgcattgttgataggcct\ 600$ acttggtttgaatctcctgtgaatacggaatgatggttggggtctatttaaaggtgtgat 660 tgcatcccctaagttttattaagtggtagggaaacctgttttttccttcttttcttgcct 720 aattgaagtccattttttaaattgtttaattacactggttttgttccaaatgcatgtgaa 840 gcttatctagtctgtgatagaaagcggcgtcaagcagtctctgggacctcggattttgac 900 agtagattcagtcaaaagttatgacttgattcttccttgtaataaattgctagggaggta 960 tataagctcgtgatgagttgttcccttcaccagtgtagaggctacgtgtatgacagggtt 1020 agagaaatgggtgttttgtatttctgaaatcatcagacactttgagtatttggtttcaca 1080 AACCTGGGGCACGACCTGAACAATCCCCGCCAAAGTAGAGGCCCAAAATGATCCAGGGGCCCA 1200 N L G H D L N N P R K V E A K M I Q G P

GTCACCCGTCGTGCGTTTGTGGATGTTGGCAACCGTGCCATTCCTGTGCAAGGGCCCCAAA 1260 V T R R A F V D V G N R A I P V Q G P K CCTCCCCTCAAGCCTGGGGGAGATCTCCCCGAAATGAGTCCGTGAAGCTGCAGAAGCCCAAA 1320 P P L K P G E I S R N E S V K L Q K P K

GCCGGCCTCTCTGGGCTGCTCGCCAGgtgggacttttttcttcttcttgggggtaagggta 1380 AGLSGLLAR

gtaagggtatgcgaggagtgtgattccgaaatggtattccgtgatgcagctgaatgtctt 1560 tttttgattgtgattgctcaggaaatggttcgatattatcgtatcggattattaattttt 1620 ttgtattgtcagATCCCGCCAAAGAGAATGTGAAGCCCCTGAAGGAAGTGGTAGAGCATG 1680

SGKENVKPLKEVVEHV TGGAGCAGATGGATGTGGAGGAGGAGGAAGCCAAGGTGGAAGAGCTGGCTATTGCTTTCTCTA 1740 EOMDVEEEAKVEELAIAFST

CCCAGAGACTAGATGTTGAAGATATTGATGCCCAAGACAGTGATAATCCTCAGCTTGTAT 1800 Q R L D V E D I D A Q D S D N P Q L V S

CTGAATATGTGAATGATATCTACAAGTACCTGCGAGAGCTGGAGgtacagacctgttgat 1860 EYVNDIYKYLRELE 

ttttatgccacggcaagctaatcatgttggcaacactgaagtcggttgtcaaccagcttg 1980 cgtggaaatattaaatattcagacttcagggccggttgtcactaagagtgtcatattggt 2040 agtataaqtctaqaaacaaaccaatccatatqtattttaqatqcttaaatattqtqctqt 2100 gtggtgcaatcttgcaccgccagtggatggtaaaccccagccgttccttgcactcgtggg 2160 gtagtttggaagtacaataaacacagccagttttttccctctctaacagGATGCCAACAA 2220 DANK

AGT CA AGC CCAGA TA CCTAG AA GGC CA AGT AA TTA CA GGA AA GAT GA GAGCA ATT TT GAT 2280 V K P R Y L E G O V I T G K M R A I L I

TGACTGGCTTGTCCAAGTACACCTCCGCTTCACTCTGCTTCAAGAAACACTGTATCTGAC 2340 D W L V Q V H L R F T L L Q E T L Y L T

TGTTGCTATCATTGACAGATTTCTCCAGgtaactagatgttattacttatcttatttaa 2400 VAIIDRFLQ

tttttatttatttctgtttttgtaafaacttttttcctttttacattaatttttaaatgg 2460 tttcttaccacttccagACTCAGAGGAATATACCACGTAACAAGCTACAGTTAGTTGGTG 2520

TQRNIPRNKLQLVGV TGACTGCCATGTTCATTGCTAGCAAATATGAAGAAATGTATTGCCCAGAAATCGGGGGACT 2580 T A M F I A S K Y E E M Y C P E I G D F

TCGCATACATCACAGACAAAGCCTACTCAAAGGCAGAGATTCGTAAAATGGAGGTGACCA 2640 ΑΥΙΤΟΚΑΥSΚΑΕΙRΚΜΕνΤΜ

TGCTGAATGAGCTGGGCTTCAATGTATCCTATCCCCTTCCCTTACACTTCCTGCGAAGAA 2700 LNELGFNVSYPLPLHFLRRN

ACAGCAAAGCTGGCTCTgtgagtactgagtgattatggtatgatgatgcaatattactcc 2760 SKAGS

tataatataaaatactgatatgcttctgatttgtagtaaaactagtaattctgtgtcaac 2820 V D A S O H T L A K Y L M E L C L P E Y

ACAGCATGTGCCATTACAAGTCGTCAATGATTGCTGCATCTGCTCTCCCCTTTCACTTA 2940 S M C H Y K S S M I A A S A L C L S L K Agtatgagatttaagattttaactaggattacttcaagtgacattaaattttactcgctt 3000 cctagattgcagttaatattgtgctttaataattgctacttatgttttaatccagtcctc 3060 atggttttattgttagtgcttttcttaagttgaaatggatgttgctaatatttgtccatc 3120 

Fig. 1. Nucleotide sequences illustrating organization of PMCvA (A) and PMCvB (B) genes. Coding nucleotides and deduced amino acids of each exon are capitalized. Introns are italicized and illustrated with lower letters. Start and stop codons (asterisks) are illustrated in boldface. The N-terminal cyclin destruction box (RXALGXIXN) and the C-terminal cyclin domains are boxed and highlighted, respectively. The length of 3' UTRs of PMCyB-m and PMCyB-l extended from that of PMCyB-s and PMCyB-m is single and double underlined, respectively. Cytoplasmic polyadenylation elements (CPU, U/AUUUUAU/A) are illustrated in boldface and underlined. Polyadenylation signals (AATAAA) are boldfaced and italicized.

attaaaattcctccccacaccttcagtaattttttatattttccttttagGTTGTTGGAT 3300

GGCAATAACTGGAGTGATACATTGACTTTCTATTCTCGCTACACTGAACAACAGCTCATG 3360 G N N W S D T L T F Y S R Y T E Q Q L M

CCAGTCATGTGCAAAATGGCATCAGTTGTAGTAAAGAGCAGTAGTGCCAAGCAACAGgta 3420 P V M C K M A S V V V K S S S A K Q Q

atttaaaatatgaatttagactgtttttacttggttcaaatgtgtacattttgaacttca 3540 tcatgcttatacagtttgcttacatcttgcaatggagatgcaaagcttgccacttgctca 3600

ttttcatctcccctgtattaggtctgatccctagttttatgcatgttttgtgtaaagaag 3660  $tcttgttaattagtattttacttgcag \texttt{GCTGTAAGACAGAAGTACAAA\texttt{GCCAGCAAGTTG}\ 3\,72\,0$ 

AVRQKYKASKL M K I S E I P O L K S K L I N S L A E K AGTGCGTCTTATGCATGAGGTGGTTGCCATTTATAAAGTAAATATTGTACATGTTGAATG 3840 SASYA \*

AATGGACTGGTTTTTGTACATATTGCTTTAAGATGCCAGTTTTTCTTTTTCATAAGCTTT	3900
CAGATTTA TA AAA TA CT GACTA CCAGTATT TA AT <b>TTTTA T</b> TTTA CA AGA CT AGT TC TC T	3960
GGGTT AGCATGTA CAAC CCAGAGTGAC TGGTTTTGGCATCTATCCATTTATCATGGT CTT	4020
CATTGTATTAATATAATTTTAGACAGGACAAAAATGGCTATTTGATGAAAGGAACTTGCA	4080
AAGAAGTATATGCTATAATGTCTTGTAATGTATATATGATCTGTGATAAACTTTTAGTTA	4140
TGTTCTGTATGTATACCTATATAAAAACCAGTTTTATCCAAGATCTGTATTCCACTATT	4200
<b>TTTTAT</b> GCCTTTAGTTATACATTGCAACCATAAGTGTGGATTGACAGTAAATAAGTTGAT	4260
TTA TA GAT TA TTG TT CA TAA CCATT CC TTT CC TAT AG TTT TC AGCAT <u>CAC CA AAG AA <b>AAT</b></u>	4320
AAA TA TGCATGCTTTACTTTATAAA TG CAA AA GAA CAGGCTTGTG TG CCAAA TATTG TAC	4380
ACATTTTATTCTTTATTTTAGAACTCGATGTTGATTTTATGTGCAGGATGCATATTCCTT	4440
TATGCAGA TTAGGTG CTTTCCCAAA TG TG <b>A TT TTA A</b> CGTC TG CACTT TTA AA ATA TT CCG	4500
TAAATTGTCACAGGTATATTTTTTGTATGGTAATGCAGTTTCGGTAATTTAGGATTAATG	4560
AACCAGGGAATCCTGTTTTCCATAATTTTTGTTGTACCTTTTTGATCAGCAGTTGCTTTG	4620
GAT GG TAG TG GTC CC CA AAT TT CTG GG GACAA AGG TG GAC GT TG TAT TT A GC GG CTG TAA	4680
AATAGGATAGAAGTGGGGGAGAAAAAAAAAAATGGAAAACTATTTTTTCTGGTATTGCATCC	4740
TAGAAAAGATCTAGTTCTTGACCAAAGGATGTATTAAATTGTTCTCATGTGGATTTTACT	4800
AGA CTAAA CTAGGTG CTATATT CTTAC TATGTATA CTTCC AG TTGAA AGA AA TGTAA ATG	4860
~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	4920

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