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PLECKSTRIN HOMOLOGY AND GREEN FLUORESCENT FUSION PROTEIN IN STARFISH BY USING BIOINFORMATICS

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Abstract

Different signaling and metabolic processes, including those that occur during fertilization, are tightly controlled by PLC isozymes. In this research, Bioinformatic databases will be utilized to fuse PLC's PH domain with GFP, which will then be used to study the starfish Patiria miniata's starfish chromosomes.

Keywords: Infertility, Phospholipase C, GFP, Patiria miniata

Introduction

In order to stimulate embryogenesis, cytoplasmic free Ca2+ is detected in the eggs. increase at fertilization [1] [2]. Anti-vertebrate protein inhibitors have been used in research in starfish eggs have been offered that this C2+ rise necessitates SFK (Src family kinase) in the egg that either activates PLCgamma directly or indirectly, resulting in IP3 production, it causes C2+ to be released from the endoplasmic reticulum of the egg (ER). Asterina miniata PLC-gamma was obtained from oocyte cDNA to study the endogenous measures in starfish eggs necessary for C2+ release during fertilization in greater detail. AmPLC gamma is a cDNA that encodes a protein that is 49 percent identical PLC-gamma in mammalians. Recombinant Src homology 2 (SH2) domains in AmPLC-gamma interacted with a 58kDa Src family kinase in a fertilization-responsive way [3] [4]. PLC from a sea urchin egg immunoprecipitates the PLC-gamma was shown to be phosphorylated in response to fertilization when it was tested with an antibody specific against AmPLC-gamma. Adding starfish eggs to the mix with AmPLC tandem gamma's SH2 domains (which block activation of PLC gamma) prevented release of Ca2+ at fertilization. These findings show that an endogenous starfish egg PLC-gamma interacts with an egg SFK and, via a PLC-gamma SH2-mediated mechanism, mediates Ca2+ release during fertilization [5] [6].Calcium signaling levels are maintained by the isoform PLCy, which assist to open a channel that allows for Ca2+ infusion over the plasma membrane and out of the endoplasmic reticulum, respectively [7]. PLC1 and PLC2 are two isoforms of the PLC class, growth factor stimulation of receptor and non-receptor (cytosolic) protein tyrosine kinase activation by polypeptide growth factor resulting in an increase in the activity of phospholipase, which can lead to angiogenesis, cell motility,



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ventricular contractility, among other things [8] [9]. GFP (Green fluorescent protein) is a widely recognized and transiently expressed fluorescent tag that can play a vital function in the localization of PLC γ . GFP will be fused to PLC's PH-domain. In this work, and the PH-GFP fusion protein will be utilized to investigate localisation in the starfish Patiria miniata [10]. Additional PLC family members have been demonstrated to influence Ca2+ signaling via previously undiscovered mechanisms, which suggests that this fusion protein may also Other non-membrane cytosolic proteins interact with and localize to compartments when exposed to sperm-egg interaction. Because the PH domain aids in marked protein-protein, protein-lipid interactions, and membrane binding it is used in this work for the development of fusion proteins that can bind to membranes [11] [12].

Materials and Methods

1-NCBI The NCBI houses a series of (computer files full of information) clearly connected with or related to (science that uses living things to improve the Earth) and natural communitydicine and is an important useful thing supply for bioinformatics tools and services [13]. Major computer files full of information include GenBank for DNA sequences and PubMed, a related to a list of references, computer file full of information for the study of how life and medicine work together. Other computer files full of information include the NCBI Epigenomics. All these computer files full of information are available online through the Entrez search engine [14].

2- Bioinformatics To construct a PH-GFP fusion protein, PLC PH domain of starfish PLC was amplified using bioinformatics to construct primers containing BsrG1 restriction sites the PJV53 – PAGFP plasmid [15] [16]. The NCBI database was used to retrieve the cDNA sequence in its entirety and the PH domain of AmPLC γ . PH domain was amplified by using NCBI's Primer-Blast to build the forward and reverse primers [17] [18] [19].

Result and Discussion

Asterina miniata phospholipase C-gamma mRNA, complete cds GenBank: AY486068.1 FASTA Graphics

Go to: 🖂	
LOCUS	AY486068 3816 bp mRNA linear INV 14-APR-2004
DEFINITION	Asterina miniata phospholipase C-gamma mRNA, complete ods.
ACCESSION	AY486068
VERSION	AY486068.1
KEYWORDS	 Alternative and the second seco
SOURCE	Patiria miniata (bat star)
ORGANISM	Patiria miniata
	Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Asterozoa;
	Asteroidea; Valvatacea; Valvatida; Asterinidae; Patiria.
REFERENCE	1 (bases 1 to 3816)
AUTHORS	Runft,L.L., Carroll,D.J., Gillett,J., Giusti,A.F., O'Neill,F.J. and Foltz,K.R.
TITLE	Identification of a starfish egg PLC-gamma that regulates Ca2+ release at fertilization
JOURNAL	Dev. Bicl. 269 (1), 220-236 (2004)
PUBMED	15081369
REFERENCE	2 (bases 1 to 3016)
AUTHORS	Gillett, J., Carroll, D.J., Runft, L.L., O'Neill, F.J., Giusti, A.F.,

Figure 1. Initial results page of the nucleotide search for Asterina miniata phospholipase C-gamma mRNA. https://www.ncbi.nlm.nih.gov/nuccore/40365362?log\$=activity



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phospholipase C-gamma [Patiria miniata]

GenBank: AAR85355.1

Identical Proteins FASTA Graphics

<u>Go to:</u> 🕑

LOCUS	AAR85355 1261 aa linear INV 14-APR-2004
DEFINITION	phospholipase C-gamma [Patiria miniata].
ACCESSION	AAR85355
VERSION	AAR85355.1
DBSOURCE	accession AY486068.1
KEYWORDS	
SOURCE	Patiria miniata (bat star)
ORGANISM	Patiria miniata
	Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Asterozoa;
	Asteroidea; Valvatacea; Valvatida; Asterinidae; Patiria.
REFERENCE	1 (residues 1 to 1261)
AUTHORS	Runft,L.L., Carroll,D.J., Gillett,J., Giusti,A.F., O'Neill,F.J. and
	Foltz,K.R.
TITLE	Identification of a starfish egg PLC-gamma that regulates Ca2+
	release at fertilization
JOURNAL	Dev. Biol. 269 (1), 220-236 (2004)
PUBMED	15081369
REFERENCE	2 (residues 1 to 1261)
AUTHORS	Gillett,J., Carroll,D.J., Runft,L.L., O'Neill,F.J., Giusti,A.F.,
	Jaffe,L.A. and Foltz,K.R.
TITLE	Direct Submission
JOURNAL	Submitted (24-NOV-2003) Biological Sciences, Florida Tech, 150 West
	University Blvd., Melbourne, FL 32901, USA
ORIGIN	
	ykkk ltpqevasvt km <mark>lkmgtvlt rfygkrrper rsfeicmetr qilwrrqtgr</mark>
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	glnw lvedhkissy p <mark>srlewwlrr efyamgktkn dtvslrdmks fmpyvnlkmn</mark>
181 tkdlke	yfne vdrwnkqeig fdgfvqlyhn lifqrevadr fkeyidernl vtvngmirfl

241 aqeqkdttan npiavkamme sfltdlgrpc qesdpkftvp efllylfspd neiwdkkfde

Figure 2. The Asterina miniata src cDNA was translated to Patiria miniata src protein. Amino acid sequence of the Patiria miniata Src family kinase protein, with the ph domain highlighted in brown from range(23-141).

https://www.ncbi.nlm.nih.gov/protein/40365363

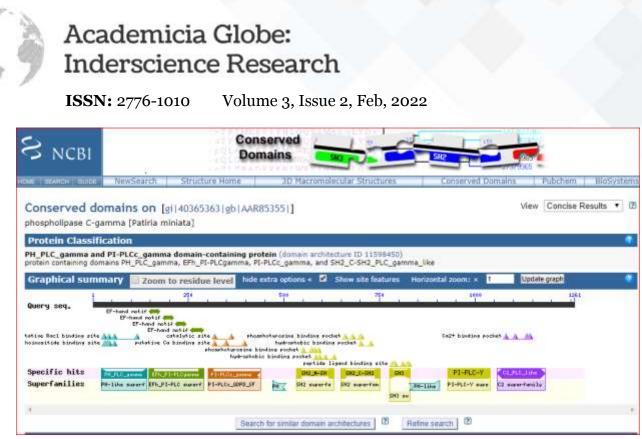


Figure 3. Conserved domain results for patiria minita . Conserved domain compares this protein sequence to the sequence for the same protein in other animals and identifies regions of high similarity (conserved regions). This shows that the ph domain is highly conserved in this protein. https://www.ncbi.nlm.nih.gov/Structure/cdd/wrpsb.cgi?INPUT_TYPE=live&SEQUENCE=AAR8535 5.1

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Primer pair 2									
	Sequence (5'->3')	Template strand	Length	Start	Stop	Tm	GC%	Self complementarity	Self 3' complementarity
Forward primer	AAGAAGAAGCTGACGCCCCA	Plus	20	29	48	61.77	55.00	4.00	2.00
Reverse primer	AGATTITATGGTCTTCCACTAGCCA	Minus	25	419	395	59.81	40.00	4.00	2.00
Product length	391								
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https://www.ncbi.nlm.nih.gov/tools/primer-

blast/primertool.cgi?ctg_time=1562583052&job_key=trxpXSXFKG0PV7hStTKcYM8pjVLiOpZP4w Figurre 4: primer-blast from NCBI are used to design forward primer and reverse primer , Next these primrs will cut out by using restriction enzyme BrsGI

ORIGIN						
1	cttcagaatg	gccaccaaca	gcctct <mark>acaa</mark>	gaagaagctg	acgccccagg	aggtggccag
61	cgtcaccaag	atgctgaaaa	tgggcaccgt	cctgacgcgc	ttctacggca	aacgacgacc
121	ggaaaggagg	tcgttcgaaa	tctgcatgga	gacgcggcag	atactgtgga	ggcgacagac
181	tgggcggaca	gacggagcag	ttaaaattcg	tgagataaaa	gagattcgtc	ccggtaagaa
241	ctcacgagac	ttcgagaggt	ggccggatga	agccaagaag	tatgatacct	cgctctgtct
301	tgtcatatgc	tacggtgccg	agttcagact	caagagcttg	tccgtcgttg	ccggcaatgc
361	cgatgaacga	cacaagtgga	tcgtcggcct	caac <mark>tggcta</mark>	gtggaa <mark>gacc</mark>	ataaaatctc
421	aagttaccca	agcagactag	aatggtggtt	acgacgggag	ttctacgcca	tggggaaaac
481	aaagaatgat	acggtgtcac	ttagggacat	gaagtcattc	atgccatacg	tcaacctgaa

Figure 5: the ph domain sequence from origin sequence ranging from nucleic acids(26 – 433) with primers are highlighted in yellow

https://www.ncbi.nlm.nih.gov/nuccore/40365362?log\$=activity



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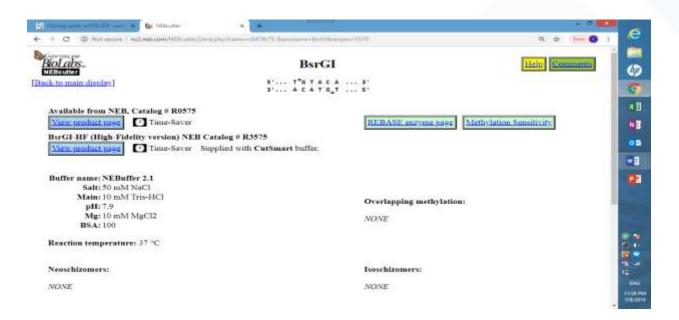


Figure 6: Picture of initial Nebcutter shows the detail of BsrGI restriction enzyme. http://nc2.neb.com/NEBcutter2/enz.php?name=c847fb75-&enzname=BsrGI&recpos=5570

Next, we will used the restriction enzyme BsrGI that cut out the primers match the sites chosen for the ph domain. so GFP – PH fusion protein can be made.

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Figure 7 : pjv53-GFP cloning vector DNA sequence found in NCBI.

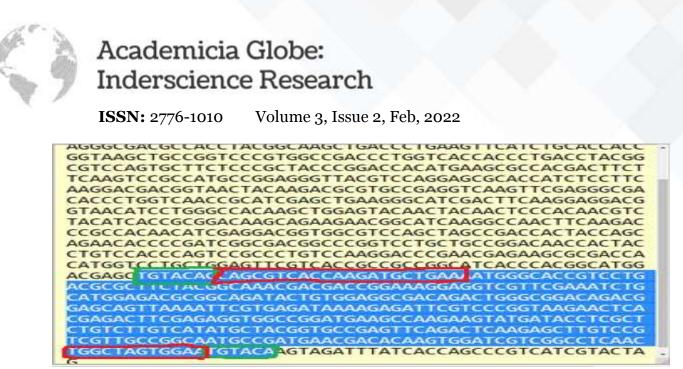


Figure 8: Combined PH domain with pjv53-GFP cloning vector and restriction enzyme highlight green . This sequence was copied and pasted into ORF Finder. https://www.ncbi.nlm.nih.gov/orffinder/

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Figure 9:BLAST results showing successful production of GFP-PH fusion protein. https://blast.ncbi.nlm.nih.gov/Blast.cgi

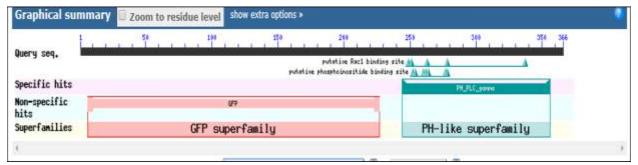


Figure 10: Zoomed-in version of successfully made GFP-PH fusion protein https://blast.ncbi.nlm.nih.gov/Blast.cgi



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The egg's Ca2+ levels grow in response to the sperm during fertilization, which is crucial in getting the egg to start developing at the very least, C2+ when IP3 levels rise, it is released from the endoplasmic reticulum is responsible for the rise in C2+ in echinoderm and vertebrate eggs [20] [21]. However, it has not been determined how IP3 is generated during fertilization [22] [23].

A phospholipase C enzyme is responsible for this (PLC), IP3 is generated from PIP2. The enzymes in this group contains δ , γ , and β isoforms. PLC β is activated by G proteins, whereas tyrosine kinases activate PLC [24]. Despite the fact that all three an increase in Ca2+ can trigger PLC isoforms the control of PLC δ remains a mystery, even if the enzymatic activity of all three PLC isoforms may be stimulated by an increase in Ca2+ [25] [26] [27]. One of these phospholipase C isoforms is activated most likely leads in the production of IP3 during fertilization.

Eggs contain PLC γ and PLC β pathway proteins. For example, expression of PLC pathway/ G protein -Dependent receptors such as the serotonin 2c or muscarinic m1 receptors allows for C2+ release in eggs when the appropriate antagonists are used [28] [29]. This implies the presence of functional PLC β and related G proteins. Exogenous tyrosine kinase/PLC γ receptors, such as those for PDGF or EGF, can be expressed in frog and starfish eggs to allow C2+ release as a result of exposure to these agonists. Ca2+ release is not caused by receptors with a single point mutation that don't activate PLC γ . A functional PLC is evident from these data. These studies have not been done on mammalian eggs, but immunoblotting has shown the presence of PLC γ .

Conclusion

Several prior research have looked into whether PLC γ or PLC β pathways are responsible for C2+ release during fertilization. Because of concerns about the selectivity of the pharmaceutical inhibitors used, the results of these trials have not been conclusive. To determine whether PLC γ - or PLC β -mediated Ca2+ release mechanisms are involved in fertilization, we used a recombinant PLC protein component to inject starfish eggs, and it inhibited PLC γ activation but not PLC β activation.

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