

Identification of a *Crassostrea gigas* ...

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ARTICLE INFO

Keywords:  
 Pfi  
*Crassostrea gigas*  
 I  
 S  
 P

ABSTRACT

T. Pfi (*Crassostrea gigas*) ...  
 fi . R  
 C. gigas  
 A 29  
 ff  
 R  
 R  
 T  
 S  
 I  
 T. fi  
 P fi

1. Introduction

T. M  
 ff  
 T. fi  
 R fi  
 P fi (*Crassostrea gigas*)  
 F  
 P fi /  
 (N , 2001). I K , P fi  
 20%  
 (K , 2013). O , P fi  
 P fi

M P fi  
 (B , 2004;  
 E , 2009; H , 2006; W , 2017; X ,  
 2017). H  
 P fi . A P fi  
 ff  
 fl ff  
 (G , 2015). N  
 T. P fi  
 ff  
 (B  
 , 2004; T S , 1961). H  
 fi  
 (G , 2014;  
 H , 2006). U  
 B -QTL

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*C. gigas* (Wang et al., 2018). B (2018) RNA (RNA), F RNA TCONS 00951105 U RNA, tyrosinase fi peroxidase (Fang et al., 2019). I, 36 ff P fi ffi

2. Material and methods

2.1. Parental source

A, B, S, C, T (Xu et al., 2017). S (Xu et al., 2018). B O (Fang et al., 2019), (B) (Wang et al., 2018).

2.2. Mating design and rearing

F<sub>1</sub> (W♀ B♂), BW (B♀ W♂), BB (B♀ B♂) WW (W♀ W♂) WB (W♀ B♂) A 36, 15 WB, 15 BW, 3 BB 3 WW. T O G

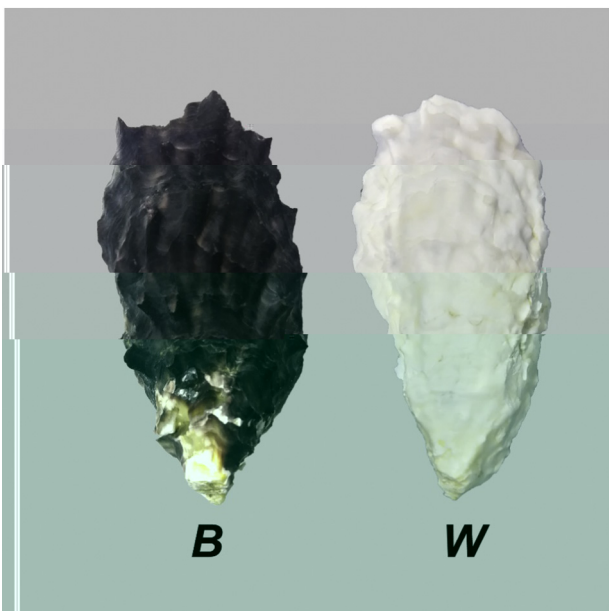


Fig. 1. R *C. gigas*. B = B ; W = W.

E, T, D- 70-L S 5 L<sup>-1</sup>, T G (2015). B fl, 18–22 Isochrysis galbana Nitzschia closterium. H fi S A S P T S B S P

2.3. Shell pigmentation record and statistical analysis

Off 2018. A 180 A T S (1961), fi F 2, T

- 0, ;
- 1, ;
- 2, ;
- 3, ;
- 4, ;

T A A (S ) A (S ). T, fi C. -fi. F

3. Results

A 29, 6 ff 1 ff 29 T T 1. O A ff ( 31) ( 35). T A fi (S ) (S ). A T 2, T ( 1–2, 7–12, 14, 18, 20–22, 26–27, 30, 33, 35)

Table 1

P	ff				C. gigas.					
	F	P	M	T	4	3	2	1	0	T
1	W♀1	0	B♂1	4	0	26	22	40	0	88
2	W♀2	0	B♂1	4	0	60	45	113	0	218
3	W♀3	0	B♂2	4	0	18	11	28	53	110
4	W♀4	1	B♂2	4	0	48	50	60	43	201
5	W♀5	0	B♂3	4	0	11	11	43	67	132
6	W♀6	0	B♂3	4	-	-	-	-	-	-
7	W♀7	0	B♂4	4	0	49	15	65	0	129
8	W♀8	0	B♂4	4	0	46	36	88	0	170
9	W♀9	0	B♂5	4	0	13	17	147	0	177
10	W♀10	0	B♂5	4	0	61	21	113	0	195
11	W♀11	0	B♂6	4	0	49	48	77	0	174
12	W♀12	0	B♂6	4	0	81	79	79	0	239
13	B♀13	4	B♂7	4	81	0	0	0	0	81
14	W♀13	0	B♂7	4	0	32	13	51	0	96
15	B♀14	4	B♂8	4	99	0	0	0	0	99
16	W♀14	0	B♂8	4	-	-	-	-	-	-
17	B♀15	4	B♂9	4	112	0	0	0	0	112
18	W♀15	1	B♂9	4	0	45	21	52	0	118
19	B♀1	4	W♂1	0	-	-	-	-	-	-
20	B♀2	4	W♂1	0	43	33	1	4	0	81
21	B♀3	4	W♂2	0	0	30	28	54	0	112
22	B♀4	4	W♂2	0	0	39	47	77	0	163
23	B♀5	4	W♂3	1	0	10	38	63	0	111
24	B♀6	4	W♂3	0	-	-	-	-	-	-
25	B♀7	4	W♂4	0	-	-	-	-	-	-
26	B♀8	4	W♂4	0	0	43	45	73	0	161
27	B♀9	4	W♂5	0	0	32	29	35	0	96
28	B♀9	4	W♂5	0	-	-	-	-	-	-
29	B♀11	4	W♂6	0	0	22	32	45	88	187
30	B♀12	4	W♂6	0	0	2	2	69	0	73
31	B♀13	4	W♂7	0	0	0	0	47	51	98
32	W♀13	0	W♂7	0	0	0	0	0	70	70
33	B♀14	4	W♂8	0	54	39	2	6	0	101
34	W♀14	0	W♂8	0	-	-	-	-	-	-
35	B♀15	4	W♂9	0	52	51	0	0	0	103
36	W♀15	1	W♂9	0	0	8	3	23	41	75

W: W; B: B

... S (13, 15, 17) ... (32) ... F (3, 5, 29, 31, 36, ...) ... (4), ... 3:1. O ... 29 ... T ... 3, ... F ... 2, P ... (20, 33, 35), ... 1:1 ... A ... (13, 15, 17). A ... Off ... (1, 7, 8, 26, 30). T.

... T ... 4. T ... fi ... 3:1 ... R ... T ... 4. T ... fi ... 1, ... 8 ... 27. T ... 4. Discussion ... M ... Mytilus edulis (I ... H ..., 1977), ... Argopecten irradians (A ... C ..., 1988), ... Chlamys nobilis ( ..., 2013). H ... P ... T ... (1995) ... M ... ff ... K - ... (2016) ... Littorina obtusata,

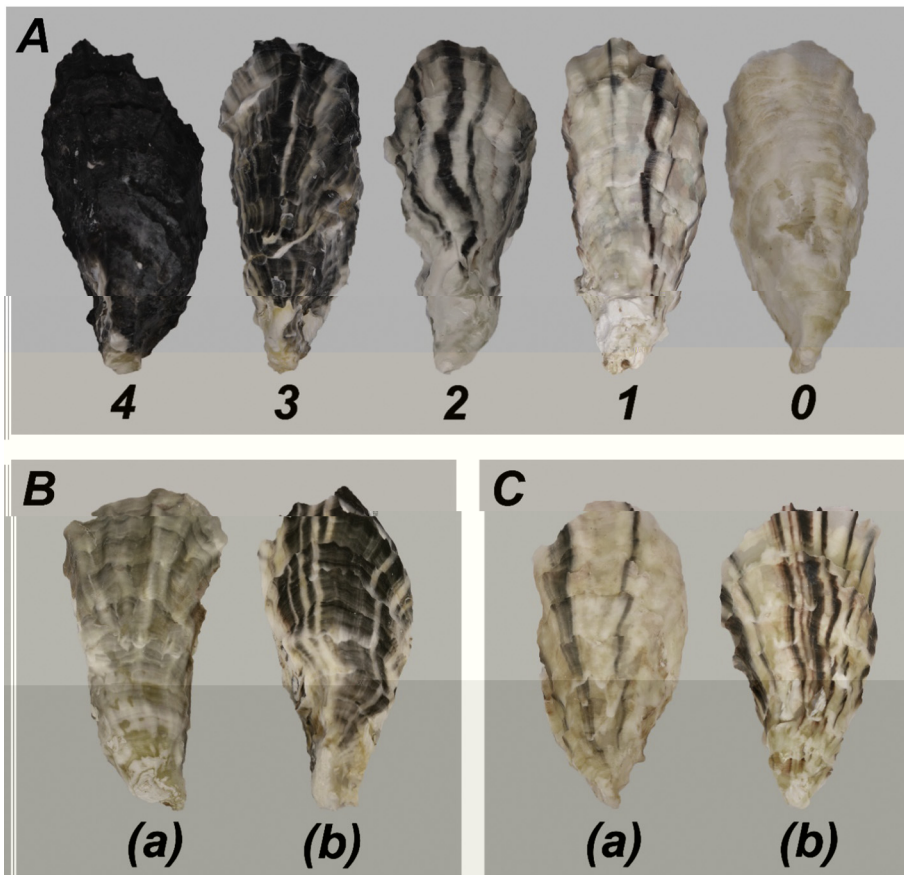


Fig. 2. Rostellum morphology of oysters. A: Offspring of the parental pair (F<sub>1</sub>) with rostellum scores 0–4; B: Offspring of the parental pair (F<sub>2</sub>) with rostellum scores 0–4; C: Offspring of the parental pair (F<sub>3</sub>) with rostellum scores 0–4. (F<sub>1</sub>, F<sub>2</sub>, F<sub>3</sub> represent the first, second, and third generations of offspring, respectively.)

Table 2

Linkage disequilibrium (LD) between the rostellum score and the shell shape

F <sub>1</sub> parental pair	Parental pair		Linkage disequilibrium (LD)		Parental pair				X <sup>2</sup> (P-value)
	F	M	F	M	S	S	T	E	
1	S	S		AA	88	0	88	1:0	NA
2	S	S		AA	218	0	218	1:0	NA
3	S	S		A	57	53	110	1:1	0.787
4	S	S	A	A	152	49	201	3:1	0.908
5	S	S		A	65	67	132	1:1	0.902
7	S	S		AA	129	0	129	1:0	NA
8	S	S		AA	170	0	170	1:0	NA
9	S	S		AA	177	0	177	1:0	NA
10	S	S		AA	195	0	195	1:0	NA
11	S	S		AA	174	0	174	1:0	NA
12	S	S		AA	239	0	239	1:0	NA
13	S	S	A	AA	81	0	81	1:0	NA
14	S	S		AA	96	0	96	1:0	NA
15	S	S	AA	AA - A	99	0	99	1:0	NA
17	S	S	AA	AA	112	0	112	1:0	NA
18	S	S	A	AA	118	0	118	1:0	NA
20	S	S	AA		81	0	81	1:0	NA
21	S	S	AA		112	0	112	1:0	NA
22	S	S	AA		163	0	163	1:0	NA
23	S	S	AA	A	111	0	111	1:0	NA
26	S	S	AA		161	0	161	1:0	NA
27	S	S	AA		96	0	96	1:0	NA
29	S	S	A		99	88	187	1:1	0.569
30	S	S	AA		73	0	73	1:0	NA
31	S	S	A		47	51	98	1:1	0.775
32	S	S			0	70	70	0:1	NA
33	S	S	AA		101	0	101	1:0	NA
35	S	S	AA		103	0	103	1:0	NA
36	S	S	A		34	41	75	1:1	0.566

S : S, shell shape; A : A, rostellum score; NA, not applicable.

L : L, linkage disequilibrium; A : A, linkage disequilibrium; S : S, linkage disequilibrium.

Table 3

F	P		L B		P					X <sup>2</sup> (P)
	F	M	F	M	S	S	U	T	E	
1	U	S	BB		0	88	0	88	0:1	NA
2	U	S	BB		0	218	0	218	0:1	NA
3	U	S	BB		0	57	53	110	0:1	NA
4	S	S	BB		0	158	43	201	0:1	NA
5	U	S	BB		0	65	67	132	0:1	NA
7	U	S	BB		0	129	0	129	0:1	NA
8	U	S	BB		0	170	0	170	0:1	NA
9	U	S	BB		0	177	0	177	0:1	NA
10	U	S	BB		0	195	0	195	0:1	NA
11	U	S	BB		0	174	0	174	0:1	NA
12	U	S	BB		0	239	0	239	0:1	NA
13	S	S			81	0	0	81	1:0	NA
14	U	S	BB		0	96	0	96	0:1	NA
15	S	S			99	0	0	99	1:0	NA
17	S	S			112	0	0	112	1:0	NA
18	S	S	BB		0	118	0	118	0:1	NA
20	S	U		B	43	38	0	81	1:1	0.693
21	S	U		BB	0	112	0	112	0:1	NA
22	S	U		BB	0	163	0	163	0:1	NA
23	S	S		BB	0	111	0	111	0:1	NA
26	S	U		BB	0	161	0	161	0:1	NA
27	S	U		BB	0	96	0	96	0:1	NA
29	S	U		BB	0	99	88	187	0:1	NA
30	S	U		BB	0	73	0	73	0:1	NA
31	S	U		BB	0	47	51	98	0:1	NA
32	U	U	A	A	0	0	70	70	-:-	NA
33	S	U		B	54	47	0	96	1:1	0.621
35	S	U		B	52	51	0	101	1:1	0.944
36	S	U	BB	B	0	34	41	75	0:1	NA

L: L; B: L; : ; B: ; S ; ; S

Table 4

F	P		L C		P	P				E	X <sup>2</sup>	P
	F	M	F	M		3	2	1	T			
1	S	S	C	C	S	19	10	35	64	3:1	0.732	0.002
						7	12	5	24			
7	S	S	C	C	S	34	9	55	98	3:1	0.885	0.050
						15	6	10	31			
8	S	S	C	C	S	31	23	76	130	3:1	0.8	0.009
						15	13	12	40			
26	S	S	C	C	S	23	37	63	123	3:1	0.795	0.000
						20	8	10	38			
30	S	S	C	C	S	2	2	50	53	3:1	0.706	0.457
						0	0	20	20			
T	-	-	-	-	S	109	80	279	468	-	-	-
						57	40	56	153			

S : S ; S : S ; L : C ; L : C ; C : S ; S

S, I, T, fi, fl, (H, 1975; L, P, 1990; S, B, 2000). I, N, I, T, ff, H, L, P, 1990; S, B, 2000). I, N, I, T, ff

T. ...  
 I. ...  
 T. fi ...  
 S. P fi ...  
 I. ...  
 F. ...  
 A. ...  
 I. ...

Acknowledgments

T. ... N N S  
 F. C. (31772843), S. P. (2017L GC009),  
 F. R. F. C. U.  
 (201762014), T. S. S. P. S.

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