



Characterization of novel EST-SNP markers and their association analysis with growth-related traits in the Pacific oyster *Crassostrea gigas*

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Abstract *Crassostrea gigas*

10 () () 10 0.011 0.01 0.01 (P 0.0001). 0 0.01 (P 0.01). 0. % *C. gigas*. *C. gigas*.

Keywords *Crassostrea gigas*

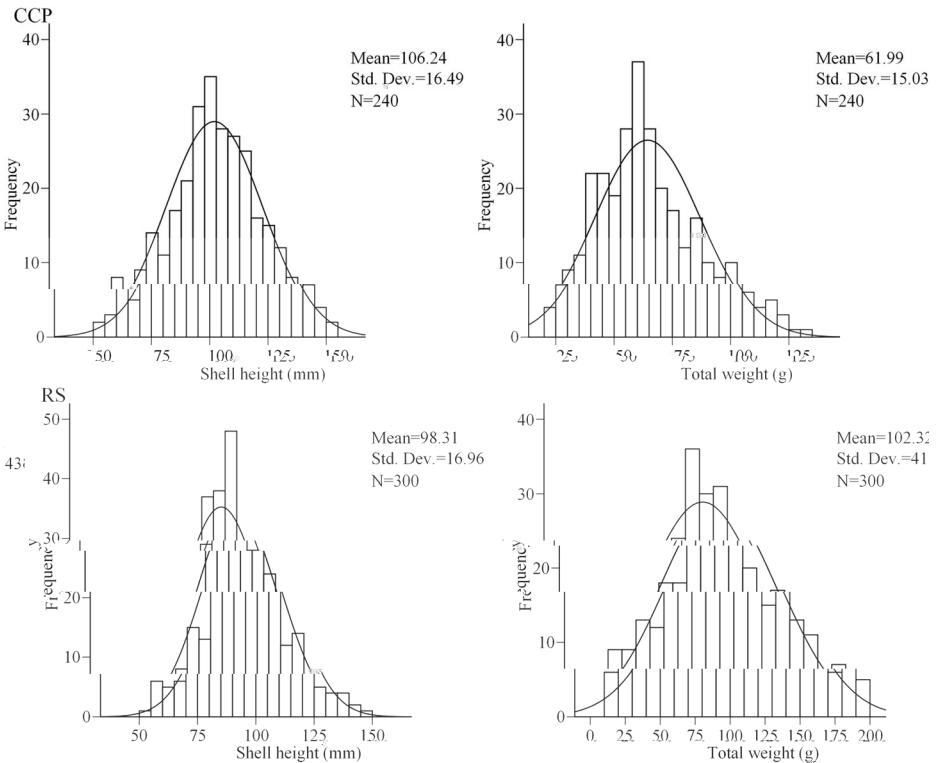


Fig. 1 T

10 μ^{-1} — 0

c — c () .

EST-SNP mining and validation

,1 1 T c c c G c

δ () .

c c c cc

T c c I , 0

c c c (I I , ,)

.0

δ c δ 0~1 0 c G 0~0%.

c

Amplification and genotyping

G c c c c 0

$\chi^2 = \frac{(n_{11} - E_{11})^2}{E_{11}} + \frac{(n_{12} - E_{12})^2}{E_{12}} + \frac{(n_{21} - E_{21})^2}{E_{21}} + \frac{(n_{22} - E_{22})^2}{E_{22}}$
 $= \frac{(10 - 10)^2}{10} + \frac{(0 - 0)^2}{0} + \frac{(0 - 0)^2}{0} + \frac{(10 - 10)^2}{10} = 0$
 $P > 0.10$

Genetic analysis and function prediction

$\chi^2 = \frac{(n_{11} - E_{11})^2}{E_{11}} + \frac{(n_{12} - E_{12})^2}{E_{12}} + \frac{(n_{21} - E_{21})^2}{E_{21}} + \frac{(n_{22} - E_{22})^2}{E_{22}}$
 $= \frac{(10 - 10)^2}{10} + \frac{(0 - 0)^2}{0} + \frac{(0 - 0)^2}{0} + \frac{(10 - 10)^2}{10} = 0$
 $P > 0.10$

Marker-QTL association analysis

$\chi^2 = \frac{(n_{11} - E_{11})^2}{E_{11}} + \frac{(n_{12} - E_{12})^2}{E_{12}} + \frac{(n_{21} - E_{21})^2}{E_{21}} + \frac{(n_{22} - E_{22})^2}{E_{22}}$
 $= \frac{(10 - 10)^2}{10} + \frac{(0 - 0)^2}{0} + \frac{(0 - 0)^2}{0} + \frac{(10 - 10)^2}{10} = 0$
 $P > 0.10$

$$\chi = (\hat{p}_1 - \hat{p}) / \sqrt{p}$$

$\hat{p}_1 = \frac{n_{11}}{n_{11} + n_{12}}$
 $\hat{p} = \frac{n_{11} + n_{21}}{n_{11} + n_{12} + n_{21} + n_{22}}$

$$p = \hat{p} (1 - \hat{p}) [1 - (1 - 1/N)(1 - 1/Ns)]$$

$N = \frac{n_{11} + n_{12} + n_{21} + n_{22}}{N}$
 (Ns)

Validating the candidate SNPs associated with growth

$\chi^2 = \frac{(n_{11} - E_{11})^2}{E_{11}} + \frac{(n_{12} - E_{12})^2}{E_{12}} + \frac{(n_{21} - E_{21})^2}{E_{21}} + \frac{(n_{22} - E_{22})^2}{E_{22}}$
 $= \frac{(10 - 10)^2}{10} + \frac{(0 - 0)^2}{0} + \frac{(0 - 0)^2}{0} + \frac{(10 - 10)^2}{10} = 0$
 $P > 0.10$

$$\Gamma = \frac{c}{(c - 1)^2} \left(\frac{c}{c - 1} - 1 \right)$$

$$pq(\alpha + \delta(q - p)) / V_A$$

$$p - q, \delta, c, \alpha$$

Results

SNP development and characterization

0 (1.0%), 10 (10.0%), 100 (100.0%)

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

0.0 0.1 0.2 0.3 0.4 0.5 0.6 0.7 0.8 0.9 1.0

Description of phenotypic variability in the experiment populations

1.0 0.9 0.8 0.7 0.6 0.5 0.4 0.3 0.2 0.1 0.0

71 71.5 72 72.5 73 73.5 74 74.5 75 75.5 76 76.5 77 77.5 78 78.5 79 79.5 80 80.5 81 81.5

(P < 0.01)

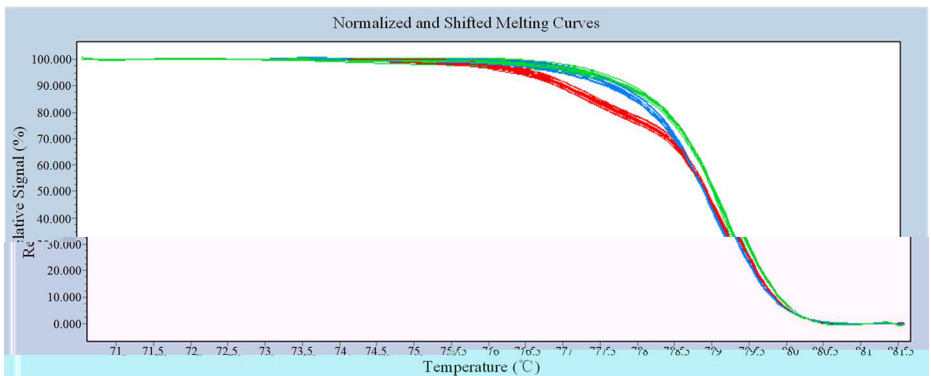


Fig. 2 Normalized and shifted melting curves for the SNP development and characterization. Green curves represent the 0.0, blue curves represent the 0.1, and red curves represent the 0.2.

Table 3 Γ c c c c

Δ	\hat{p}_1	\hat{p}	$-p_1$	χ
00 / Γ	0. 0	0.0 0	0.001	. 2 *
0 \hat{p} / G	0.	0.0 1	0.001	. 2 *
0 0 /	0. 0	0.0	0.000	. 11 *
1 / Γ	0.	0. 2	0.00 0	. 2 *
\hat{p} / G	0. 0	0.1 2	0.00	. 2 *
\hat{p} 1 / G	0.	0. 0	0.00	. 2 *
/	0.	0.	0.00 2	.01 *
/	0. 2	0. 0	0.00	.0 1 *
1 /	0. 2	0.1	0.00	.1 1 *

*P 0.0001, c c c

$G(\cdot)$ 0, 1 -c Γ c

c c Γ c 0 c

Discussion

c c Γ c c c

c c c Γ

0 cc 10 (. %) Γ C. gigas

00 01) c 0-11 Γ c

(00) c c c

c c c Γ c

c c c c c Γ c

c c c c c Γ c

c c c c c Γ c

Table 4 Γ c c () c c

	(%)	χ	(%)	χ
0 G	. 0	0.	. *	. 2 *
0 0 Γ	. 1	1. 1	11.0 *	. 0. 1 . *
\hat{p} 1 Γ	. 1	0.	1 . *	. 2 . 0 *

*P 0.01, c c c

$\frac{c}{c}$. I c c G
 (.) (. 1 3 , 1) .
 c T T
 cc (0 0 1) - G ()
 G c c
 (c c c G 1 001) .
 (. 1) . T c
 c c *C. gigas* c
 T c 0
 I 10 T T
 T c c
C. gigas c c c

Acknowledgements

(1) c c (0 1 0 0) - Q
 T c (0 1 0) 5 5 5

References

. (0 0) c () c
 (1) T c 1 1 - 1 1 Q T Q T
 I c c T
 (1) Q T 0 cc I () c
 c c 1 - 1 (1 0) c
 G (1 3) c T G 1 c (Q T)
 G (1) c T G 1 1 - 1 0 1
 (0 1) c 1 *Crassostrea gigas* c
 Q (0 0) G T c
 G (0 0) G T G 1 1 0 - 1 0

	(00)	c	c	c	
	0				
	(1 1)	c	1	3	3
	(1 1)	c			
	G	c	1	1	1
	(00)	c	0	c	c
			G		
	(01)				
	1	0	0		
	(00)	c			
	(00)	c	c	c	
	G				
	(00)	c	c	c	
	Q	3	3	3	(00)
	1	1	1	1	
	(010)				
	(Crassostrea gigas)				
	Q				
	(011)	c			
	(Crassostrea gigas)				
	Q				
	(01)	c	3	3	c
	Meretrix meretrix				
	(00)	c	1	1	1
	G	c	10		
	(1)	3			
	G	c	1	1	1
	(00)				
	1	1	1	1	
	Q				
	(01)	c			
	Meretrix meretrix				
	(01)	c			
	(00)	c			
	Coffea				
	G				
	(00)	c			
	(1)	c			
	(00)	c			
	(00)	c			
	Crassostrea gigas				
	0	1			
	(010)	Q			
	(Crassostrea gigas)				
	1	0			
	(01)	c			
	Crassostrea gigas				
	(1)	c			
	G	1	1	1	
	(00)				
	G	c	1	1	0
	(1 0)				
	(Zea mays)				
	(1)	c			
	Q				
	(00)	c			
	1	0			

$Q, Q, \dots, (01)T$ c, c, \dots (*Crassostrea gigas*)
 $T, G, G, \dots, G, \dots, (00)$ c
 $X, X, \dots, G, G, \dots, G(01)G$ c, c, \dots
 $c, c, \dots, (00)$ c, \dots, δ c, c, c, \dots
 $1-0$ $T(1)G$ $c, c, c, \dots, c, \dots, 1$
 $X, Q, \dots, (01)$ c
 $c, c, c, \dots, Crassostrea gigas, \dots, c$
 $X, Q, G, \dots, (01)QT$ c, c, \dots
 $c, c, Crassostrea gigas, \dots, c, I, 1, -1$