

CORRECTION

Correction: Identification of Reproduction-Specific Genes Associated with Maturation and Estrogen Exposure in a Marine Bivalve *Mytilus edulis*

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[Fig 1](#) of the published article is incomplete. Please view the complete [Fig 1](#) and its legend here. There are errors in [Table 1](#) and [Table 2](#) of the published article. Please view the correct tables here.



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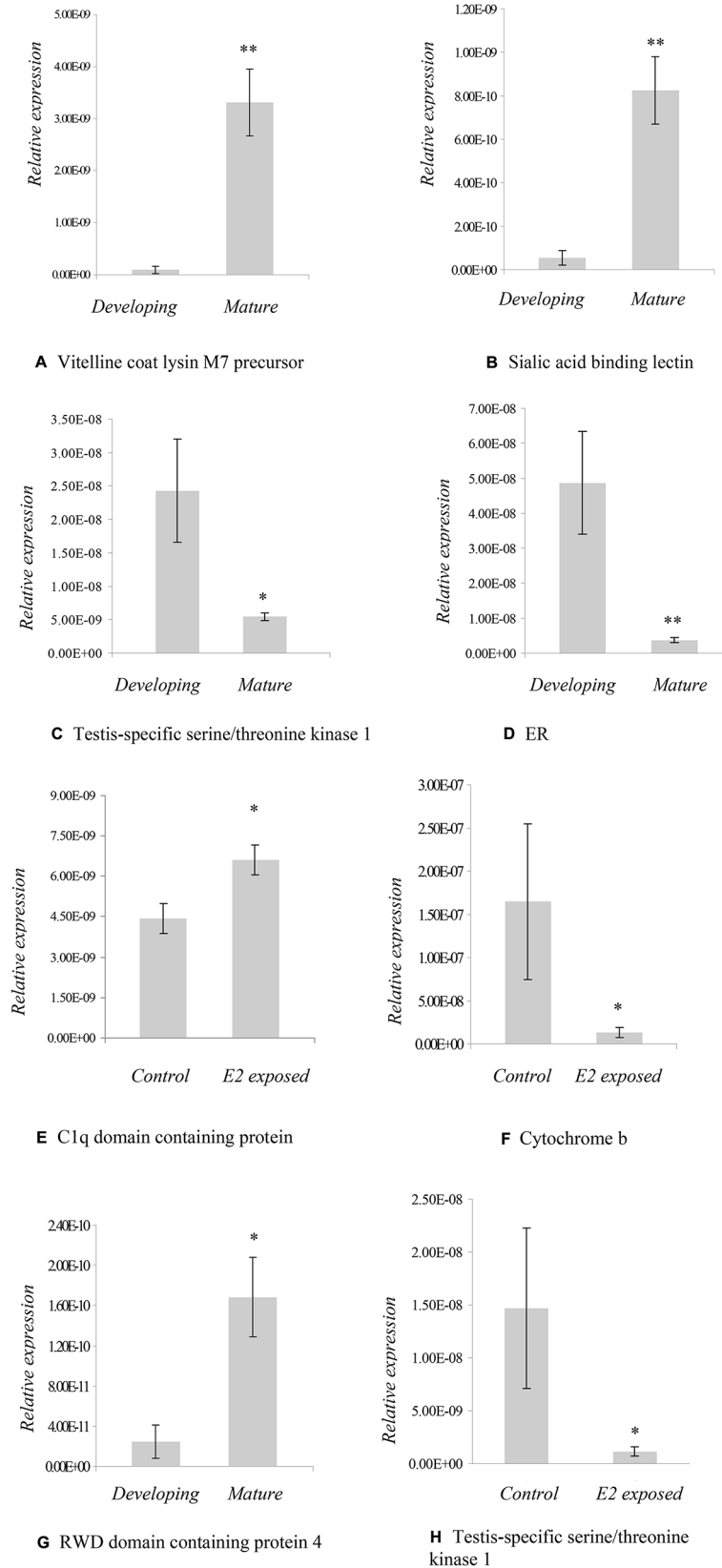


Fig 1. Real-time quantitative RT-PCR validation of differential screening results of *M. edulis* developing gonad versus mature gonad samples (1A–1E) and *M. edulis* experimentally-exposed to E2 (1F–1H). Data plotted as mean±SEM, n = 15 samples. * = $p < 0.05$; ** = $p < 0.01$.

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Table 1. Differentially expressed (subtracted) mRNAs identified in *M. edulis* testis at two stages of gonadal development. ^aDown-regulated in early developing testes relative to mature testes. ^bUp-regulated in early developing testes relative to mature testes.

Clone accession no.	Category & gene identity	Length (bp)	Homolog species / Accession no.	E-value
HQ690234	^a Senescence-associated protein	155	<i>Brugia malayi</i> XP_001900327.1	5.0E ⁻¹⁰
AJ492924.1	^a Histone 2A	301	<i>M. edulis</i> AJ492924.1	1.0E ⁻¹⁰⁸
AY484747.1	^a 16S ribosomal protein	931	<i>M. edulis</i> AY484747.1	0
FM995162.1	^a Vitelline coat lysin M7 precursor	635	<i>M. edulis</i> BAA03551.1	3.0E ⁻¹²³
HQ678182	^a Sialic acid binding lectin	397	<i>Helix pomatia</i> ABF00124.1	4.0E ⁻¹⁵
CAX33833	^a Putative vitelline envelope receptor for lysin (VERL)	663	<i>M. edulis</i> CAX33833	4.0E ⁻¹²⁵
HQ678180	^b Testis-specific serine/threonine kinase 1 (TSTK1)	815	<i>Strongylocentrotus purpuratus</i> XP_787865.1	4.0E ⁻⁷⁰
HQ678181	^b Testis-specific A-kinase-anchoring-protein	182	<i>Gallus gallus</i> XP_002162537.1	9.0E ⁻⁶
HQ67816	^b Histone H2A isoform 2	332	<i>Halotis discus discus</i> ACJ12611.1	1.0E ⁻⁵³
HQ678184	^b Beta-tubulin	511	<i>C. gigas</i> AAU93877.1	9.0E ⁻⁸¹
AB257133	^b ER	111	<i>M. edulis</i> BAF34366.2	1.0E ⁻¹²
HQ678183	^b Bindin precursor 5 repeat variant (acrosomal protein)	392	<i>C. gigas</i> ABQ18234.1	7.0E ⁻¹⁴
HQ678185	^b Phosphodiesterase 1	533	<i>S. purpuratus</i> NP_001091918.1	5.0E ⁻⁶²
AY130198.1	^b Cytochrome c oxidase subunit III	254	<i>M. edulis</i> AAV68300.1	2.0E ⁻³¹

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Table 2. Differentially expressed (subtracted) mRNAs identified in *M. edulis* testis following E2 exposure. ^aDown-regulated in control testes relative to E2-exposed testes. ^bUp-regulated in control testes relative to E2-exposed testes.

Clone Accession No.	Category & gene identity (BlastX)	Length (bp)	Homolog species/Accession no.	E-value
HQ664951	^a Complement C1q-like protein	111	<i>Ailuropoda melanoleuca</i> XP_002918680.1	7.0E ⁻⁵
HQ690237	^a Alpha tubulin	297	<i>C. gigas</i> BAD80736.1	2.0E ⁻⁵¹
HQ690238	^a Beta tubulin	501	<i>Rattus norvegicus</i> NP_954525.1	2.0E ⁻⁹⁴
HQ690235	^a Ribosomal protein L7	240	<i>C. gigas</i> AJ557884.1	3.0E ⁻³⁰
HQ690239	^a Bromodomain adjacent to zinc finger domain, 1A	369	<i>G. gallus</i> XP_426440.2	7.0E ⁻²⁴
HQ690240	^a Elongation factor 1 gamma	162	<i>Saccoglossus kowalevskii</i> NP_001171816.1	1.0E ⁻¹³
HQ664949	^a Vitelline envelope zona pellucida domain 9	900	<i>Halotis rufescens</i> ABE72949.1	1.0E ⁻²⁶
HQ664950	^a C1q domain containing protein	141	<i>Argopecten irradians</i> ADD17343	7.0E ⁻⁵
HQ664948	^a RWD domain containing protein 4A	240	<i>Caligus rogercresseyi</i> ACO11028.1	3.0E ⁻¹⁷
HQ664952	^a Hemagglutinin/amoebocyte aggregation factor precursor	240	<i>Salmo salar</i> ACI68653.1	1.0E ⁻¹⁰
YP_073337	^a Cytochrome c oxidase subunit II	448	<i>M. edulis</i> YP_073337.1	4.0E ⁻⁵⁶
YP_073338.1	^a NADH dehydrogenase subunit 1	647	<i>M. edulis</i> YP_073338.1	4.0E ⁻⁶¹
HQ690236	^a Triosephosphate isomerase TIM	156	<i>Metapenaeus ensis</i> AAP79983.1	3.0E ⁻¹³
CAX33833	^a Putative vitelline envelope receptor for lysin (VERL)	663	<i>M. edulis</i> CAX33833	4.0E ⁻¹²⁵
AAV68423	^b Cytochrome c oxidase subunit 1	534	<i>M. edulis</i> AAV68411.1	8.0E ⁻⁹⁰
AAV68416	^b Cytochrome b	302	<i>M. edulis</i> AAV68404.1	6.0E ⁻²⁸
HQ690243	^b Ferritin-like protein	492	<i>Pinctada fucata</i> AAQ12076.1	2.0E ⁻⁷⁸
HQ690241	^b Senescence-associated protein	318	<i>Trichoplax adhaerens</i> XP_002118266.1	6.0E ⁻⁴⁹
HQ690244	^b Spectrin beta chain	293	<i>Harpegnathos saltator</i> EFN75523.1	1.0E ⁻¹⁰

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Reference

1. Ciocan CM, Cubero-Leon E, Minier C, Rotchell JM (2011) Identification of Reproduction-Specific Genes Associated with Maturation and Estrogen Exposure in a Marine Bivalve *Mytilus edulis*. PLoS ONE 6(7): e22326. doi: [10.1371/journal.pone.0022326](https://doi.org/10.1371/journal.pone.0022326) PMID: [21818309](https://pubmed.ncbi.nlm.nih.gov/21818309/)