

Appendix 2-1

Chapter 2. Exploiting genomic data to identify proteins involved in abalone reproduction.

Omar Mendoza-Porras; Natasha A. Botwright; Sean M. McWilliams; Mathew T. Cook; James O. Harris; Gene Wijffels and Michelle L. Colgrave”

“Genetic information from several molluscan species (supplementary information 6) was used to build the customised database utilised in this study. Contigs, mRNA, SRA, ESTs sequences and transcriptome assemblies were downloaded from different public sources. SRA reads were assembled *de novo* into contigs using MIRA 3.4.0[€]. ESTs, mRNA or transcriptome assemblies were used directly to detect open reading frames (ORFs) using the getorf application. EMBOSS[¥] suite default parameters were applied to obtain the ORFs . For each sequence the longest ORF was kept and collated into the customised molluscan database”.

Table 3. Sources of genetic information used to assemble the molluscan database utilised in this study

Molluscan species	Source/reference
<i>Haliotis asinina</i>	Jackson et al. 2010 ^[1]
<i>Pinctada maxima</i>	Jackson et al. 2010 ^[1]
<i>Haliotis midae</i>	Franchini et al. 2011 ^[2]
<i>Littorina saxatilis</i>	Galindo et al. 2010 ^[3]
<i>Crepidula fornicate</i>	Henry et al. 2010 ^[4]
<i>Ilyanassa obsolete</i>	Lambert et al. 2010 ^[5]
<i>Lymnaea stagnalis</i>	Feng et al. 2009 ^[6]
<i>Littorina saxatilis</i>	EBI (http://www.ebi.ac.uk) accession (FR863689–FR864978)
<i>Biomphalaria glabrata</i>	http://www.snaildb.org/
<i>Aplysia californica</i>	http://hgdownload.cse.ucsc.edu/goldenPath/aplCall1/bigZips/
<i>Mytilus galloprovincialis</i>	http://mussel.cribi.unipd.it/
<i>Crassostrea gigas</i>	http://www.genoscope.cns.fr/spip/crassostrea-gigas-EST-library.html
<i>Lottia gigantea</i>	http://genome.jgi-psf.org/Lotgi1/Lotgi1.home.html
<i>Littorina saxatilis</i>	http://mbio-serv2.mbioekol.lu.se/Littorina/
<i>Dreissena polymorpha</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html

<i>Mytilus edulis</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html
<i>Crassostrea virginica</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html
<i>Argopecten irradians</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html
<i>Biomphalaria glabrata</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html
<i>Crassostrea, gigas</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html
<i>Lymnaea stagnalis</i>	http://www.nematodes.org/NeglectedGenomes/MOLLUSCA/wwwMolluscDB_blast.html

References

1. Jackson DJ, McDougall C, Woodcroft B, Moase P, Rose RA, Kube M, Reinhardt R, Rokhsar DS, Montagnani C, Joubert C, et al: **Parallel Evolution of Nacre Building Gene Sets in Molluscs.** *Molecular Biology and Evolution* 2010, **27**:591-608.
 2. Franchini P, van der Merwe M, Roodt-Wilding R: **Transcriptome characterization of the South African abalone *Haliotis midae* using sequencing-by-synthesis.** *BMC Res Notes* 2011, **4**:59.
 3. Galindo J, Grahame JW, Butlin RK: **An EST-based genome scan using 454 sequencing in the marine snail *Littorina saxatilis*.** *Journal of Evolutionary Biology* 2010, **23**:2004-2016.
 4. Henry JJ, Perry KJ, Fukui L, Alvi N: **Differential Localization of mRNAs During Early Development in the Mollusc, *Crepidula fornicata*.** *Integrative and Comparative Biology* 2010, **50**:720-733.
 5. Lambert JD, Chan XY, Spiecker B, Sweet HC: **Characterizing the Embryonic Transcriptome of the Snail *Ilyanassa*.** *Integrative and Comparative Biology* 2010, **50**:768-777.
 6. Feng ZP, Zhang Z, van Kesteren RE, Straub VA, van Nierop P, Jin K, Nejatbakhsh N, Goldberg JI, Spencer GE, Yeoman MS, et al: **Transcriptome analysis of the central nervous system of the mollusc *Lymnaea stagnalis*.** *Bmc Genomics* 2009, **10**.
 7. Jiang J-Z, Zhang W, Guo Z-X, Cai C-C, Su Y-L, Wang R-X, et al. **Functional annotation of an expressed sequence tag library from *Haliotis diversicolor* and analysis of its plant-like sequences.** *Marine Genomics.* 2011;4:189-96.
 8. Amparyup P, Klinbunga S, Jarayabhand P. **Identification and Expression Analysis of Sex-Specific Expression Markers of Thai Abalone *Haliotis asinina*, Linneaus, 1758.** *J Shellfish Res.* 2010;29:765-73.
- Tools for contigs and ORF files creation.
- €. <http://sourceforge.net/p/mira-assembler/wiki/Home/>
- ¥. <http://emboss.sourceforge.net/>