Connectivity of stony corals in Spermonde Archipelago, South Sulawesi, Indonesia

Sutrisno Puspitaningasih¹, Janne Timm² and Marc Kochzius¹

- ¹ Marine Biology, Vrije Universiteit Brussel, Pleinlaan 2, 1050 Brussels, Belgium E-mail: psutrisn@vub.ac.be
- Biotechnology and Molecular Genetics, University of Bremen, Leobener Strasse UFT, 28359 Bremen, Germany

Due to increasing changes of environmental conditions and over-exploitation of coastal habitats, many species inhabiting coral reefs suffer depletion of populations, which may lead to local extinctions. The change of species composition may have an influence on the resilience of the whole ecosystem to disturbances and changing environmental conditions. Important factors for an assessment of the status and chance for recovery and adaptation of exploited and disturbed populations are the genetic diversity and connectivity to other populations. This research aims to investigate the genetic diversity and genetic population structures of stony corals in Indonesia, especially in Spermonde, South Sulawesi. Several potential model taxa were sampled, but genetic analyses will focus on two species, depending on abundance at the different sample sites. In order to investigate the influence of early life history stages on connectivity, one species will be a broadcast spawner with pelagic eggs (Acropora tenuis, A. millepora), while the other species will have brooded eggs (Pocillopora damicornis, Seriatopora hystrix). Sampling was done on September 2012 from nine locations in Spermonde Archipelago, South Sulawesi, Indonesia. The genetic population structure will be analysed by utilising nuclear microsatellites, which will be amplified by PCR. An Analysis of Molecular Variances (AMOVA) will be done to test for population structures with the programme Arlequin (ver. 3.5, Excoffier & Lischer, 2010). In order to reveal detailed patterns of genetic divergences, pairwise F-statistics will be calculated with the same program, and a hierarchical AMOVA will be conducted to test spatial groupings of populations. In addition, a Bayesian analysis as implemented in the software STRUCTURE (ver. 2.3.3, Hubisz et al., 2009) will be performed, testing for different numbers of clusters in the dataset and giving the corresponding probabilities. The findings of the ongoing study will be provided for a synergetic assessment of the sustainable use of coral reef resources and their conservation.

References

- Beerli P. and M. Palczewski. 2010. Unified framework to evaluate panmixia and migration direction among multiple sampling locations. Genetics 185:313-326.
- Ehlers A., B. Worm and T.B.H. Reusch. 2008. Importance of genetic diversity in eelgrass *Zostera marina* for its resilience to global warming. Marine Ecology Progress Series 355:1-7.
- Excoffier L. and H.E.L. Lischer. 2010. Arlequin suite ver 3.5: a new series of programs to perform population genetics analyses under Linux and Windows. Molecular Ecology Resources 10:564-567
- Froukh T. and M. Kochzius. 2007. Genetic population structure of the endemic fourline wrasse (*Larabicus quadrilineatus*) suggests limited larval dispersal distances in the Red Sea. Molecular Ecology 16:1359-1367.
- Hubisz M.J., D. Falush, M. Stephens and K. Pritchard. 2009. Inferring weak population structure with the assistance of sample group information. Molecular Ecology Resources 9:1322-1332.
- Hughes T.P., A.H. Baird, D.R. Bellwood, M. Card, S.R. Connolly, C. Folke, R. Grosberg, O. Hoegh-Guldberg, J.B.C. Jackson, J. Kleypas, J.M. Lough, P. Marshall, M. Nyström, S.R. Palumbi, J.M. Pandolfi, B. Rosen and J. Roughgarden. 2003. Climate change, human impacts, and the resilience of coral reefs. Science 301:929-933.
- Hughes A.R. and J.J. Stachowicz. 2004. Genetic diversity enhances the resistance of a seagrass ecosystem to disturbances. PNAS 101:8998-9002.
- Jensen J.L., A.J. Bohonak and S.T. Kelley. 2005. Isolation by distance, web service. BMC Genetics 6:1-6.

- Knittweis L., W.E. Krämer, J. Timm and M. Kochzius. 2009. Genetic structure of *Heliofungia actiniformis* (Scleractinia: Fungiidae) populations in the Indo-Malay Archipelago: implications for live coral trade management efforts. Conservation Genetics 10:241-249.
- Kochzius M. and D. Blohm. 2005. Genetic population structure of the lionfish *Pterois miles* (Scorpaenidae, Pteroidae) in the Gulf of Agaba and the northern Red Sea. Gene 347:295-301.
- Kochzius M. and A. Nuryanto. 2008. Strong genetic population structure in the boring giant clam *Tridacna crocea* across the Indo-Malay Archipelago: implications related to evolutionary processes and connectivity. Molecular Ecology 17:3775-3787.
- Kochzius M., C. Seidel, J. Hauschild, S. Kirchhoff, P. Mester, I. Meyer-Wachsmuth, A. Nuryanto and J. Timm. 2009. Genetic population structures of the blue starfish *Linckia laevigata* and its gastropod ectoparasite *Thyca crystallina*. Marine Ecology Progress Series 396:211-219.
- Nuryanto A. and M. Kochzius. 2009. Highly restricted gene flow and deep evolutionary lineages in the giant clam *Tridacna maxima*. Coral Reefs 28:607-619.
- Starger C.J., S.S.R. Yeoh, C.-F. Dai, A.C. Baker and R. Desalle. 2008. Ten polymorphic STR loci in the cosmopolitan reef coral, *Pocillopora damicornis*. Molecular Ecology Resources 8:619-621.
- Timm J. and M. Kochzius. 2008. Geological history and oceanography of the Indo-Malay Archipelago shape the genetic population structure in the False Clown Anemonefish (*Amphiprion ocellaris*). Molecular Ecology 17:3999-4014.
- Underwood J.N., L.D. Smith, M.J.H. van Oppen and J.P. Gilmour. 2009. Ecologically relevant dispersal of corals on isolated reefs: implications for managing resilience. Ecological Applications 19:18-29.
- Van Oppen M.J.H., J.N. Underwood, A.N. Muirhead and L. Peplow. 2007. Ten microsatellite loci for the reef-building coral *Acropora millepora* (Cnidaria, Sclreactinia) from the Great Barrier Reef, Australia. Molecular Ecology Notes 7:436-438.
- Wang S., L. Zhang and M. Matz. 2009. Microsatellite characterization and marker development from public EST and WGS databases in the reef-builing coral *Acropora millepora* (Cnidaria, Anthozoa, Scleractinia). Journal of Heredity 100:329-337.
- Wild C., O. Hoegh-Guldberg, M.S. Naumann, F.M. Colombo-Pallotta, M. Ateweberhan, W.K. Fitt, R. Iglesias-Prieto, C. Palmer, J.C. Bythell, J.-C. Ortiz, Y. Loya and R. van Woesik. 2011. Climate change impedes scleractinian corals as primary reef ecosystem engineers. Marine and Freshwater Research 62:205-215.