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## Connectivity of the coral *Acropora tenuis* in Spermonde Archipelago

### INTRODUCTION

Coral reefs have the highest species biodiversity of all marine ecosystems.

However, since they are declining globally due to anthropogenic impacts, conservation is essential.

• Marine Protected Areas (MPA) are an important conservation tool.

MPA's should be arranged in networks, because coral reefs are scattered/patchy habitats. This ensures connectivity and increases resilience.

Hermatypic corals are ecosystem engineers that build the three-dimensional structure of the reef. **Insights of population dynamics help to develop management strategies for conservation.** 

Exploring the Indonesian region in Sulawesi, study of a hermatypic, mass-spawning coral: *Acropora tenuis* in Spermonde Archipelago.

#### Research objectives

A. population structure on a local scale:

Code Sites in Spermonde #

Kodingareng Lompo

Kodingareng Keke

Barang Lompo

Bone Tambung

SL | Sarapo Lompo

Lanyukang

Kapoposang

**Sample Total** 

What is the composition of genetic diversity?

(allelic richness + heterozygosity)

Hypothesis: Genetic diversity is shaped by local sea surface currents.

B. gene flow and connectivity on a global scale:

Is there migration and/or dispersal across the Indian Ocean?

(F-statistics + Mantel test + AMOVA + Bayesian cluster analysis)

Hypothesis: There are two distinct genotype clusters: an African and an Indonesian cluster.

**Fig 1:** Adult colonies of *Acropora tenuis*. (Picture: Puspita Sutrisno, 2013) 
(B) the coast of Tanzania. (Veron, 2000)

#### MATERIAL & METHODS

Acropora tenuis coral branch: 80 samples from 8 locations.

- Genomic DNA extraction and purification
- Microsatellite analysis (SSR length polymorphism)
  - 1. 7 markers from (Underwood et al., 2009) + 9 markers from literature search (NEW)
  - 2. Primer validation + Multiplex PCR with fluorescent labeled primers
- DNA capillary sequencer
  - 1. Fragment Length Analysis of PCR products
  - 2. Scoring of fragment lengths

#### Genotypes = length polymorphisms of microsatellites

- Statistical analysis of the length polymorphism Fstat (F-statistics), GenAlEx (AMOVA + Mantel test), STRUCTURE (Bayesian cluster analysis):
- Data comparison with samples from the Tanzanian coast
  Dataset from a previous study (Rosa van der Ven, 2012)
  Measure of gene flow and connectivity

#### RESULTS

- Successful primer validation + Multiplex PCR for the 7 markers (Underwood et al., 2009)
   Successful fragment length analysis → scoring process ongoing.
- Primer validation for the 9 NEW markers ongoing.

# LA BA BT BL KK KK

**Spermonde Archipelago** 

SAMPLING

South Sulawesi

**Fig 3:** Samples collected in Spermonde, SE-Sulawesi. **TOP RIGHT**– Table of sampling sites: site code, site name and number of samples collected there. **DOWN**– Spatial distribution of the collection sites.

#### References:

- Underwood J N, Smith H L D, Van Oppen M J H, Gilmour J P (2009) Ecologically relevant dispersal of corals on isolated reefs: implications for managing resilience. Ecological Applications 19(1):18-29.

   Veron 1 (2000) Corals of the world. Australian Institute of marine science, p360-361 and Veron 1 (2013) Internet (http://coral aims gov au/)
- Veron, J (2000). Corals of the world. Australian Institute of marine science. p360-361 and Veron, J (2013). Internet, (http://coral.aims.gov.au/).
  The map of South-Sulawesi (http://en.wikipedia.org/wiki/File:Sulawesi\_blank\_map.png) is under the terms of the GNU Free Documentation License (last accessed: 28 February 2014).
- The map of the Spermonde Archipelago was taken with Google earth V 7.1.2. Spermonde Archipelago, Sulawesi, ID. 4°55′11.51″S, 119°13′19.93″O, Eye alt 121.28km [image:2013-04-10]. (accessed: 2014-02-26).