# Phylogeny of corallimorpharians (Anthozoa; Cnidaria) based on mitochondrial genomes

Mei-Fang Lin<sup>1,2,4</sup>, Marcelo Kitahara<sup>1,3</sup>, David Miller<sup>1,4</sup>, Chaolun Allen Chen<sup>2</sup>



<sup>1</sup>School of Pharmacy and Molecular Sciences, James Cook University, Townsville, Australia <sup>2</sup>Biodiversity Research Centre/ Taiwan International Graduate Program (TIGP)-Biodiversity, Academia Sinica, Taipei, Taiwan <sup>3</sup>Centro de Biologia Marinha, Universidade de São Paulo, São Sebastião, Brasil <sup>4</sup>ARC Centre of Excellence for Coral Reef Studies, James Cook University, Townsville, Australia



#### Objectives and approach

Order Corallimorpharia, Carlgren, 1940, is a small and poorly understood anthozoan group. Corallimorpharians resemble actiniarians in the absence of a calcareous skeleton, but have internal anatomies much more similar to those of scleractinians (den Hartog 1980). The phylogenetic position of corallimorpharians in relation to scleractinians is unclear and remains controversial (Fig. 1).



Corallimorpharia is classified into three or four families (Fig. 2). Systematic classification of this order is still controversial, mainly as a result of unclear original descriptions and the synonymisation of several genera and species (den Hartog 1980, Cha 2001; Table 1). This study is aimed at understanding corallimorpharian phylogeny by molecular phylogenetic analyses of whole mitochondrial (mt) genome sequences. Figure 1 Phylogenetic hypotheses of relationship between order Figure 2 Systematics of Corallimorpharia based on morphology and ecophysiology.



### Results

Phylogenetic relationships of corallimorpharians were determined by using whole mt genome sequences from a total of 12 taxa.

Three distinct types of gene organisation were found in the corallimorpharian study (Fig. 3):

(A) Most corallimorpharians were characterised by a common gene order;

(B) The shallow water azooxanthellate species Corynactis californica had a distinct mt genome organisation;

(C) Gene order in the deep-sea corallimorpharian Corallimorphus profundus was similar to the scleractinian pattern.

Phylogenetic trees were constructed by Maximum likelihood analysis and Bayesian inference (Fig. 4).

- 1. Three groups were resolved corresponding to the gene orders: 2. The validity of the family Ricordeidae:
  - (A) zooxanthellate corallimorpharians
  - (B) shallow water azooxanthellate corallimorpharian (C) deep-sea azooxanthellate corallimorpharian

Figure 4 Corallimorpharian mt genome phylogenetic tree

Table 1 Summary of the corallimorpharian classification schemes Carlgren 1949 (3 families, 10 ge Corall Corallir actis) (Coralli Nectactis, Sid Cha 2001 (3 f es, 11 p . 12 g Coral es, 11 ge Corall (Coral milies, 11 genera) Ricordeida

Sideractiidae Corallimo (Nectactis, Sideractis) (Corallimo

- 3. All genera in Discosomatidae are valid;

4. Corvnactis and Corallimorphus should be elevated to higher classification level; 5. Basal position of C. profundus.

Discosoma sp.1 (Gen size 20 908 bp: AT content 61 9% Figure 3 Corallimorpharian and scleractinian mt genome order 100/100 Discosoma sp.2 (20,912 bp; 61.9%) ue Amplevidie 100/100 Rhodactis mussoides (20.826 bp: 61.9% - Actinodiscus nummiformis (20.922 bp: 62.0%) Di Amplexidiscus fenestrafer (20,188 bp; 61.9%) Corallimorpharian Corvnactis californica 100/100 **Rhodactis indosinensis** (20,092 bp; 61.9%) 100/100 **Rhodactis** sp. (20,093 bp; 62.0%) R - Pseudocorynactis sp. (21,239 bp; 62.0%) - Ricordea yuma (22,015 bp; 63.1%) Ricordeidad 700xanthellate C - Ricordea florida (21,376 bp; 62.9% 100/10 Azooxanthellate (Shallow water) - Corynactis californica (20,632 bp; 61.0% Corallimorphida norphus profundus (20,488 bp; 61% (Deep-sea) D Acropora tenuis (18,338 bp; 62 strea radians (19,387 bp; 63.1%)

0.02

## Major conclusions

1. Gene order and phylogeny results support three corallimorpharian groups, revealing different evolutionary processes in corallimorpharians.

- 2. The similarity of gene order in C. profundus and scleractinians supports a close relationship between this corallimorpharian and scleractinians. 3. The family Discosomatidae contains most genera, the status of some of which is unclear. Results presented suggest that considerable diversity
- exists and that, in contradiction of den Hartog's view of lumping all Discosomatidae, most genera are likely to be valid.
- 4. In general, this study is consistent with there being 4-6 families and around 11 genera in the order Corallimorpharia.
- 5. Further analyses based on morphology and additional taxa (e.g. Sideractiidae) will be required to better understand relationships within the
- Corallimorpharia and between the various anthozoan lineages.

#### Acknowledgment

This study was supported by the Academia Sinica Thematic Grant (2006-2008) and grant from National Science Council, Taiwan (NSC-94-2621B001005) to CAC.

