



<http://dx.doi.org/10.11646/zootaxa.3893.4.9>

<http://zoobank.org/urn:lsid:zoobank.org:pub:46C298CB-D934-4300-8D20-53F71F1054D8>

## Absence of consistent genetic differentiation among several morphs of *Actinia* (Actiniaria: Actiniidae) occurring in the Portuguese coast

ANA M PEREIRA, CRISTIANA BRITO, JOANA SANCHES, CARLA SOUSA-SANTOS  
& JOANA I ROBALO

MARE – Marine and Environmental Sciences Centre and UIEE – Eco-Ethology Research Unit ISPA-IU. Rua Jardim do Tabaco, 34, 1149-041 Lisbon, Portugal. E-mail: ana\_pereira@ispa.pt

### Abstract

*Actinia equina*, the beadlet sea anemone, is a very labile species, displaying variable colour patterns, broad habitat choice and diverse modes of reproduction. Historically, studies using genetic markers such as allozymes and differences in habitat choice lead several authors to propose that different colour morphs could represent different species. One of the species defined was *A. fragacea*. In this paper, the relationships between brown, red and green colour morphs of *A. equina* and *A. fragacea* were studied, using two DNA fragments (one mitochondrial and one nuclear). Individuals were sampled from three different areas in Portugal separated by a maximum distance of 500 km. This is the first study applying direct sequencing of selected gene fragments to approach the validity of *Actinia* morphs as different genetic entities. The results show that, at least in the Portuguese coast, these colour morphs do not correspond to the two valid species recognized in the literature. The existence of cryptic species is discussed.

**Key words:** *Actinia equina*, *Actinia fragacea*, Portugal, Cnidaria, morphotypes

### Introduction

The beadlet sea anemone *Actinia equina* L. is a widespread species that lives in the rocky intertidal and subtidally to depths of up to 20m. This anemone is predominantly a scavenger, feeding unselectively on macrofaunal carrion (Davenport *et al.* 2011).

Historically, this species was considered to be distributed from North Russia and the Baltic Sea to tropical West African waters, including the archipelagos of Madeira, Azores and Canaries, South Africa, Mediterranean and Black sea, the Red Sea and Far East (Stephenson 1935; Manuel 1981; Cha *et al.* 2004). In its range, this species is considered very labile, showing variable colour patterns (individuals may display red, brown, orange, green and mottled column and tentacles), broad habitat choice, and diverse modes of reproduction (asexual and/or sexual). Their high level of morphological diversity, allied with its extensive distribution, lead different authors to propose that different morphs could represent different species (e.g., Carter & Thorpe 1981; Haylor *et al.* 1984; Monteiro *et al.* 1997; Schama *et al.* 2005).

The first morph of *A. equina* recognized as a species was *A. fragacea* Tugwell, which presents a red or dark red column and tentacles with green, yellow or blue spots in the column. According with previous authors (e.g. Haylor *et al.* 1984 and references therein), this species shows a different distribution and ecology from *A. equina*, lacking viviparity and presenting only sexual reproduction. The identity of this species was corroborated by the use of allozymes, which showed that this morphotype was consistently different from *A. equina* in areas where they occurred in sympatry in the English Channel, suggesting reproductive isolation between them (Carter & Thorpe 1981).

Later, in a study investigating the ecological and genetic relationships between the red, brown and green morphs of *A. equina* in Britain (Isle of Man), the distinct character of the green morph was evaluated (Haylor *et al.* 1984). In this study, the green morph occupied different tidal levels and microhabitats when compared to brown and red morphs (which were indistinct one from each other). Again, allozymes showed that this morphotype was

## References

- Carter, M.A. & Thorpe, J.P. (1981) Reproductive, genetic and ecological evidence that *Actinia equina* var. *mesembryanthemum* and var. *fragacea* are not conspecific. *Journal of the Marine Biological Association of the United Kingdom*, 61, 79–83.  
<http://dx.doi.org/10.1017/S0025315400045926>
- Cha, H.-R., Buddmeier, R.W., Fautin, D.G. & Sandhei, P. (2004) Distribution of sea anemones (Cnidaria, Actiniaria) in Korea analysed by environmental clustering. *Hydrobiologia*, 530/531, 497–502.  
<http://dx.doi.org/10.1007/s10750-004-2667-3>
- Chomsky, O., Douek, J., Chadwick, N.E., Dubinsky, Z. & Rinkevich, B. (2009) Biological and population-genetic aspects of the sea anemone *Actinia equina* (Cnidaria: Anthozoa) along the Mediterranean coast of Israel. *Journal of Experimental Marine Biology and Ecology*, 375, 16–20.  
<http://dx.doi.org/10.1016/j.jembe.2009.04.017>
- Clement, M., Posada, D. & Crandall, K.A. (2000) TCS: a computer program to estimate gene genealogies. *Molecular Ecology*, 9, 1657–1659.  
<http://dx.doi.org/10.1046/j.1365-294x.2000.01020.x>
- Davenport, J., Moloney, T.V. & Kelly, J. (2011) Common sea anemones *Actinia equina* are predominantly sessile intertidal scavengers. *Marine Ecology Progress Series*, 430, 147–155.  
<http://dx.doi.org/10.3354/meps08861>
- Douek, J., Barki, Y., Gateño, D. & Binkevich, B. (2002) Possible cryptic speciation within the sea anemone *Actinia equina* complex detected by AFLP markers. *Zoological Journal of the Linnean Society*, 136, 315–320.  
<http://dx.doi.org/10.1046/j.1096-3642.2002.00034.x>
- Excoffier, L.G.L. & Schneider, S. (2005) Arlequin ver. 3.0: an integrated software package for population genetics data analysis. *Evolutionary Bioinformatics Online*, 1, 47–50.
- Geller, J.B. & Walton, E.D. (2001) Breaking up and getting together: evolution of symbiosis and cloning by fission in sea anemones (genus *Anthopleura*). *Evolution*, 55, 1781–1794.  
<http://dx.doi.org/10.1111/j.0014-3820.2001.tb00827.x>
- Haylor, G.S., Thorpe, J.P. & Carter, M.A. (1984) Genetic and ecological differentiation between sympatric colour morphs of the common intertidal sea anemone. *Marine Ecology Progress Series*, 16, 281–289.  
<http://dx.doi.org/10.3354/meps016281>
- Manuel, R.L. (1981) *British Anthozoa*. Academic Press, London, 241 pp.
- Monteiro, F.A., Solé-Cava, A.M. & Thorpe, J.P. (1997) Extensive genetic divergence between populations of the common intertidal sea anemone *Actinia equina* from Britain, the Mediterranean and the Cape Verde Islands. *Marine Biology*, 129, 425–433.  
<http://dx.doi.org/10.1007/s002270050183>
- Rodríguez, E., Barbeitos, M.S., Brugler, M.R., Crowley, M.M., Grajales, A., Gusmão, L., Haussermann, A.R. & Daly, M. (2014) Hidden among Sea Anemones: the first comprehensive phylogenetic reconstruction of the Order Actiniaria (Cnidaria, Anthozoa, Hexacorallia) reveals a novel group of Hexacorals. *PLoS One*, 9, 1–17.  
<http://dx.doi.org/10.1371/journal.pone.0096998>
- Schama, R., Solé-Cava, A.M. & Thorpe, J.P. (2005) Genetic divergence between east and west Atlantic populations of *Actinia* spp. sea anemones (Cnidaria: Actiniidae). *Marine Biology*, 146, 435–443.  
<http://dx.doi.org/10.1007/s00227-004-1462-z>
- Schama, R., Mitchell, M. & Solé-Cava, A.M. (2011) *Actinia ebhayiensis* sp. nov., a new species of sea anemone (Anthozoa: Actiniaria: Actiniidae) from South Africa. *Journal of the Marine Biological Association of the United Kingdom*, 92 (5), 885–894.  
<http://dx.doi.org/10.1017/S0025315411001305>
- Shearer, T.L., Van Oppen, M.J.H., Romano, S.L. & Worheide, G. (2002) Slow mitochondrial DNA sequence evolution in the Anthozoa (Cnidaria). *Molecular Ecology*, 11, 2475–2487.  
<http://dx.doi.org/10.1046/j.1365-294x.2002.01652.x>
- Solé-Cava, A.M. & Thorpe, J.P. (1987) Further genetic evidence for the reproductive isolation of green sea anemone *Actinia prasina* Gosse from common intertidal beadlet anemone *Actinia equina* (L.). *Marine Ecology Progress Series*, 38, 225–229.  
<http://dx.doi.org/10.3354/meps038225>
- Sonnenberg, R., Nolte, A.W. & Tautz, D. (2007) An evaluation of LSU rDNA D1-D2 sequences for their use in species identification. *Frontiers in Zoology*, 4, 1–6.  
<http://dx.doi.org/10.1186/1742-9994-4-6>
- Sousa-Santos, C., Robalo, J.I., Collares-Pereira, M.J. & Almada, V.C. (2005) Heterozygous indels as useful tools in the reconstruction of DNA sequences and in the assessment of ploidy level and genomic constitution of hybrid organism. *DNA Sequence*, 16, 462–467.  
<http://dx.doi.org/10.1080/10425170500356065>
- Stephenson, T.A. (1935) *The British sea anemones*. Ray Society, London, 148 pp.

- Swofford, D.L. (1998) PAUP\*. *Phylogenetic Analysis Using Parsimony (\* and Other Methods)*, version 4. Sinauer Associates, Sunderland, Massachusetts.
- Thompson, J.D., Gibson, T.J., Plewniak, F., Jeanmougin, F. & Higgins, D.G. (1997) The ClustalX windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *Nucleic Acids Research*, 24, 4876–4882.  
<http://dx.doi.org/10.1093/nar/25.24.4876>
- Watts, P.C., Allcock, A.L., Lynch, S.M. & Thorpe, J.P. (2000) An analysis of the nematocysts of the beadlet anemone *Actinia equina* and the green sea anemone *Actinia prasina*. *Journal of the Marine Biological Association of the United Kingdom*, 80, 719–724.  
<http://dx.doi.org/10.1017/S002531540000254X>