Diversity and Distribution of Symbiodinium Associated with Seven Common Coral Species in the Chagos Archipelago, Central Indian Ocean

Sung-Yin Yang1*, Shashank Keshavmurthy1*, David Obura2,3, Charles R. C. Sheppard3, Shakil Visram1,2, Chaolun Allen Chen1,4*

1 Biodiversity Research Center, Academia Sinica, Nangang, Taipei, Taiwan, 2 Coastal Ocean Research and Development Indian Ocean (CORDIO), Mombasa, Kenya, 3 School of Life Sciences, University of Warwick, Coventry, United Kingdom, 4 Institute of Oceanography, National Taiwan University, Taipei, Taiwan

Abstract
The Chagos Archipelago designated as a no-take marine protected area in 2010, lying about 500 km south of the Maldives in the Indian Ocean, has a high conservation priority, particularly because of its fast recovery from the ocean-wide massive coral mortality following the 1998 coral bleaching event. The aims of this study were to examine Symbiodinium diversity and distribution associated with scleractinian corals in five atolls of the Chagos Archipelago, spread over 10,000 km². Symbiodinium clade diversity in 262 samples of seven common coral species, Acropora muricata, Isopora paliifera, Pocillopora damicornis, P. verrucosa, P. eydouxi, Seriatopora hystrix, and Stylophora pistillata were determined using PCR-SSCP of the ribosomal internal transcribed spacer 1 (ITS1), PCR-DDGE of ITS2, and phylogenetic analyses. The results indicated that Symbiodinium in clade C were the dominant symbiont group in the seven coral species. Our analysis revealed types of Symbiodinium clade C specific to coral species. Types C1 and C3 (with C3z and C3i variants) were dominant in Acroporidae and C1 and C1c were the dominant types in Pocilloporidae. We also found 2 novel ITS2 types in S. hystrix and 1 novel ITS2 type of Symbiodinium in A. muricata. Some colonies of A. muricata and I. paliifera were also associated with Symbiodinium A1. These results suggest that corals in the Chagos Archipelago host different assemblages of Symbiodinium types then their conspecifics from other locations in the Indian Ocean; and that future research will show whether these patterns in Symbiodinium genotypes may be due to local adaptation to specific conditions in the Chagos.

Introduction
Mutualistic symbiosis between scleractinian corals and dinoflagellates (genus Symbiodinium) contributes to high productivity in coral reef ecosystems, providing important resources and functions for consumers, including humans [1–3]. Corals and coral reefs have suffered as a result of several environmental and anthropogenic factors that have caused their destruction worldwide [2,4–10]. Most coral-algal symbioses are sensitive to increasing seawater temperature and high irradiance [2,11–16]. However, particular partner combinations may resist episodes of high thermal stress. Despite the general conclusion that coral-algal mutualistic symbioses are sensitive to changes in their environment, biogeographic studies indicate that different regional environments also significantly influence the ecology and evolution of this relationship [17–22]. These environmental factors include average annual temperatures, seasonal changes in water clarity and substantial seawater temperature variation across tropical and subtropical regions [23–24].

From various studies, the diversity of Symbiodinium is well established [1,5,25–33]. Molecular (small ribosomal subunit (SSU rDNA), larger subunit (LSU rRNA), chloroplast large subunit ribosomal DNA, internal-transcribed regions (ITS1 and ITS2)) and phylogenetic classification of Symbiodinium into functionally distinct evolutionary entities (using alpha-numeric designations equivalent to ‘species’) has shown them to belong to nine divergent clades [1,5,25–33]. LaJeunesse et al. [27,34] using internal transcribed spacer (ITS) region argued that the ‘clades’ of Symbiodinium required further resolution into operational taxonomic units, ‘subclade,’ and/or sub species levels, thereby providing a more complete picture of Symbiodinium diversity and systematics [34], also see [22] [35]. Furthermore, the results from ITS2 and other analysis now support the designation of the ITS2 sequence as individual Symbiodinium species and hence the term ITS2 types is being used to refer to the variation in the Symbiodinium diversity [32]. Ecophysiological function has been inferred for several of these clades based either on biogeographic / ecological distribution or physiological stress experiments
Symbiodinium are genetically and physiologically diverse and the presence of a particular Symbiodinium type influences a reef coral’s tolerance to thermal stress (See [3] for detailed analysis, Also see [20] for Symbiodinium evolution and symbiosis response to climate change). Studies have shown that corals associate with different Symbiodinium clades or types depending on the environmental conditions.

Overall composition of Symbiodinium clades in the seven coral species sampled from all four locations in the Chagos Archipelago revealed the presence of only two Symbiodinium clades; C and A (Fig. 2 and Fig. S1).

Among the divergent lineages of Symbiodinium, clade C is ecologically abundant and diverse among reef corals in both the Indo-Pacific and western Atlantic, whereas clade A is relatively common in shallow water scleractinians in the Caribbean and Indian Ocean [22,27,34] but is rare in the Pacific [reviewed in [5], [45], but see [46]]. Although surveys have shown that clade C is the dominant symbiont associated with scleractinian corals in the Pacific Ocean and West Indian Ocean [1,7,19,22,47–48], a recent study on the same sampling scale in the north-eastern Indian Ocean (Andaman Sea) showed a significant proportion of clade D and a diversity of Symbiodinium types in different coral hosts [22]. Although some regions of the Indian Ocean have been surveyed [7,22,49], there is no information of Symbiodinium diversity from the central Indian Ocean either at regional or local scale, which limits our understanding of the relationship between coral host and Symbiodinium on a broad biogeographical scale.

The Chagos Archipelago, designated as a no-take marine protected area in 2010 [50–51], is located in the central Indian Ocean on the Chagos-Laccadive Ridge that extends as far south as 20°S. It covers 550,000 km² with >60,000 km² shallow limestone platform and reefs [51]. Chagos is a valuable location for studying corals, as it contains >25% of the Indian Ocean reef area [51] in a condition and habitat that is largely unaffected by direct, local human impacts, although during the El Niño event of 1998, the Chagos suffered severe coral bleaching and mass mortality especially among large colonies of tabular Acropora [52]. However, significant and rapid recovery was observed afterwards [24]. Also, coral diseases are extremely low in the Chagos reefs [51]. The role of the Chagos Archipelago in the Indian Ocean is very important for a number of reasons, including conservation of commercial fish stocks [50] and supports densities of coral reef fishes one to two orders of magnitude greater than in other Indian Ocean locations [51]. It also provides a scientific reference point for several aspects of Indian Ocean research and for global studies of reef condition and reef restoration. Further, the Chagos archipelago appears to act as a marine biodiversity corridor between the north-eastern and western Indian Ocean [51], such that knowledge of its coral and coral reefs is very important for conservation efforts. This has led to its designation as the world’s largest marine protected area [50,53].

In the wake of global climate change and future sea surface warming predicted for this region and others [24], we conducted this study, 1. To document the diversity of Symbiodinium in seven

### Table 1. Information on sampling locations, coral species sampled at each location and associated Symbiodinium clades (RFLP).

| Site           | Coral species | 28S-RFLP |
|----------------|---------------|----------|
| Salomon Islands (90) | I. palifera (17) | C        |
| P. damicornis (24) |              | C        |
| P. eydouxi (10)  |              | C        |
| P. verrucosa (12) |              | C        |
| S. hystrix (4)   |              | C        |
| S. pistillata (23) |             | C        |
| Peros Banhos (23) | A. muricata (20) | C        |
| I. palifera (3)  |              | C        |
| Diego Gracia (78) | I. palifera (21) | C (4), C+A (7) |
| P. damicornis (11) |              | C        |
| S. hystrix (21)  |              | C        |
| S. pistillata (25) |             | C        |
| Chagos (71)       | A. muricata (22) | C (11), A (2), C+A (9) |
| I. palifera (12)  |              | C        |
| P. damicornis (8) |              | C        |
| S. hystrix (6)   |              | C        |
| S. pistillata (23) |             | C        |

The numbers in the brackets denote sample obtained from each site, for each coral species and number of clade C and A.

doi:10.1371/journal.pone.0035836.t001

Figure 1. Map showing the location of the Chagos Archipelago, including the four sampling sites: Salomon Islands, Diego Garcia, Chagos Bank and Peros Banhos. Areas shown in red are the actual locations from which the samples were obtained at each site. doi:10.1371/journal.pone.0035836.g001
common scleractinian species in the Chagos Archipelago and 2. To know whether such remote reefs contribute to some unique Symbiodinium clades/types from the extracted DNA, DNA extraction followed the protocol of [55] followed by amplification of DNA using the nuclear ribosomal internal transcribed spacer 1 (ITS1) regions and the nuclear ribosomal internal transcribed spacer 2 (ITS2) regions (see below). Before analyzing the samples using ITS1 and ITS2, DNA from all the samples were screened for Symbiodinium clades using large subunit ribosomal DNA (lssr-DNA) restriction fragment length polymorphism (lssr-DNA-RFLP) and further confirmed with small subunit ribosomal DNA-RFLP (ssl-DNA-RFLP) [42]. RFLP analysis was carried out following the protocols of [42].

**ITS1 and Single Strand Confirmation Polymorphism (SSCP)**

The ITS1 region was amplified using the primer set with a fluorescent Hex-labelled forward primer ‘SymITSFP (5’-CTCGAGCTCTGGAGCGTGTGTTGG-3’) and ‘SymITSRP’ (5’-TATCGCRCTTCRCTGCGCCT-3’). Single-stranded conformation polymorphism (SSCP) was performed following the protocol described in [56]. SSCP were preformed using Gelscan 3000 (Corbett Research). PCR products were first denatured at 95°C for 3 min and on ice immediately for 3 min. Then PCR products were loaded onto the 4% nondenaturing TBE-polyacrylamide gel (1.5 ml of 40% acrylamide / bis-acrylamide (37.5:1), 95 m of 25% APS and 30 u of TEMED), pulsed into gel for 25s and flushed. This was followed by a run in 1200 V, 22°C with 0.6x TBE buffer for 40 min.

**ITS2 and Denaturing gradient gel electrophoresis (DGGE)**

ITS2 region were amplified using primer set ‘TitSintfor2’ (5’-AAT TGC AGA ACT CCT CCG TGG-3’) and ‘TitS2 clamp’ (5’-CGC CCG CCG CCG CCC CCC GGC ATG CAT ATG CTT AAG TCC AGC GGG TCT-3’) from [57] and using touch-down PCR [27]. PCR products of ITS2 were denatured using 45–80% denaturing gradient gel for 16 h on CBS Scientifc system (Del Mar, CA, USA). Gels were stained with 1 x SYBR Gold (Life Technologies, Invitrogen, USA) for 20 min, and were photographed using a gel documentation system (Vilber Lourmat, France).

The method used for assigning the ITS2-DGGE fingerprint followed [22]. Prominent bands of each fingerprint were sent for direct sequencing then matched with the sequences from Genbank (Table S1).

**Direct sequencing**

For samples of Acroporidae for which ITS2-DGGE did not successfully detect the presence of clade A (this might be due to the low copy number of clade A), the region between ITS1 (primer

---

**Table 2.** SSCP and DGGE Symbiodinium types associated with each species from different locations.

| Coral species | Sample location | SSCP-ITS1 | DGGE-ITS2 |
|---------------|-----------------|-----------|-----------|
| *A. muricata* | Great Chagos Bank | A1        | A1        |
| Peros Banhos  | C               | C1        |
|               | Ca2             | C2        |
|               | Ca3             | C3        |
|               | Not detected    | C3kk      |
| *L. palifera*| Great Chagos Bank | A1        | A1        |
| Salomon Islands | C               | C1        |
| Diego Gracia | Ca              | C3        |
| Peros Banhos  | Ca2             | C2        |
|               | Ca3             | C3        |
|               | Ca4             | C40       |
|               | Ca5             | Not detected |
| *P. damicornis*| Great Chagos Bank | C        | C1        |
| Salomon Islands | C2              | C1        |
| Diego Gracia | Not detected    | C1        |
| *P. eydouxi* | Salomon Islands  | C        | C1        |
|               | C2              | C1c       |
| *P. verrucosa*| Salomon Islands  | C        | C1        |
|               | C2              | C1c       |
| *S. pistillata*| Great Chagos Bank | C        | C1        |
| Salomon Islands | Cs              | Not detected |
| Diego Gracia | Csh             | Not detected |
| *S. hystrix* | Great Chagos Bank | C        | C1        |
| Salomon Islands | Cs              | Not detected |
| Diego Gracia | Csh             | Not detected |

Common Scleractinian species in the Chagos Archipelago and 2. To know whether such remote reefs contribute to some unique Symbiodinium clades/types from the extracted DNA, DNA extraction followed the protocol of [55] followed by amplification of DNA using the nuclear ribosomal internal transcribed spacer 1 (ITS1) regions and the nuclear ribosomal internal transcribed spacer 2 (ITS2) regions (see below). Before analyzing the samples using ITS1 and ITS2, DNA from all the samples were screened for Symbiodinium clades using large subunit ribosomal DNA (lssr-DNA) restriction fragment length polymorphism (lssr-DNA-RFLP) and further confirmed with small subunit ribosomal DNA-RFLP (ssl-DNA-RFLP) [42]. RFLP analysis was carried out following the protocols of [42].

**Materials and Methods**

**Ethics Statement**

Coral tissues were collected and exported from British Indian Ocean Territory with permission granted by the Administrator of the Territory, Foreign and Commonwealth Office, London, United Kingdom.

---

**Symbiodinium in Corals of the Chagos Archipelago**

Seven common scleractinian coral species (Isopora palifera, Acropora muricata, Pocillopora damicornis, *P. eydouxi*, *P. verrucosa*, *Stylophora pistillata* and *Seriatopora hystrix*) were sampled at depths of 2 m to 7 m from five parts of the Chagos Archipelago in March 2006 (Fig 1). Sampled sites were Salomon, Peros Banhos and Diego Garcia atolls, and from Eagle Islands and Three Brothers from the Great Chagos Bank. In all 262 samples (Table 1) collected from 7 coral species were preserved in 95% ethanol for DNA extraction.

**Molecular analysis of Symbiodinium clade and types**

Use of combinations of genetic analyses better resolves the Symbiodinium identity in a given sample [54]. Thus, we used three techniques to analyze Symbiodinium clades/types from the extracted DNA. DNA extraction followed the protocol of [55] followed by amplification of DNA using the nuclear ribosomal internal transcribed spacer 1 (ITS1) regions and the nuclear ribosomal internal transcribed spacer 2 (ITS2) regions (see below). Before analyzing the samples using ITS1 and ITS2, DNA from all the samples were screened for Symbiodinium clades using large subunit ribosomal DNA (lssr-DNA) restriction fragment length polymorphism (lssr-DNA-RFLP) and further confirmed with small subunit ribosomal DNA-RFLP (ssl-DNA-RFLP) [42]. RFLP analysis was carried out following the protocols of [42].

**ITS1 and Single Strand Confirmation Polymorphism (SSCP)**

The ITS1 region was amplified using the primer set with a fluorescent Hex-labelled forward primer ‘SymITSFP (5’-CTCAGCTCTGGAGCGTGTGTTGG-3’) and ‘SymITSRP’ (5’-TATCGCRCTTCRCTGCGCCT-3’). Single-stranded conformation polymorphism (SSCP) was performed following the protocol described in [56]. SSCP were preformed using Gelscan 3000 (Corbett Research). PCR products were first denatured at 95°C for 3 min and on ice immediately for 3 min. Then PCR products were loaded onto the 4% nondenaturing TBE-polyacrylamide gel (1.5 ml of 40% acrylamide / bis-acrylamide (37.5:1), 11.445 ml of ddH2O, 0.3 ml of 100% glycerol, 0.9 ml 10x TBE, 50 μl of 25% APS and 50 μl of TEMED), pulsed into gel for 25s and flushed. This was followed by a run in 1200 V, 22°C with 0.6x TBE buffer for 40 min.

**ITS2 and Denaturing gradient gel electrophoresis (DGGE)**

ITS2 region were amplified using primer set ‘TitSintfor2’ (5’-AAT TGC AGA ACT CCT CCG TGG-3’) and ‘TitS2 clamp’ (5’-CGC CCG CCG CCG CCC CCC GGC ATG CAT ATG CTT AAG TCC AGC GGG TCT-3’) from [57] and using touch-down PCR [27]. PCR products of ITS2 were denatured using 45–80% denaturing gradient gels for 16 h on CBS Scientific system (Del Mar, CA, USA). Gels were stained with 1 x SYBR Gold (Life Technologies, Invitrogen, USA) for 20 min, and were photographed using a gel documentation system (Vilber Lourmat, France).

The method used for assigning the ITS2-DGGE fingerprint followed [22]. Prominent bands of each fingerprint were sent for direct sequencing then matched with the sequences from Genbank (Table S1).

**Direct sequencing**

For samples of Acroporidae for which ITS2-DGGE did not successfully detect the presence of clade A (this might be due to the low copy number of clade A), the region between ITS1 (primer
SymITSFP to lsurDNA (primer D1/D2 R) was amplified, cloned and three clones for each sample were sequenced. The molecular cloning and sequencing methods were followed as per [55].

Environmental Parameters
Monthly mean SST values (1948–2010) were acquired from NOAA Earth System Research Laboratory NCEP/NCAR data (http://www.esrl.noaa.gov/psd/cgi-bin/data/timeseries/time-series1.pl; downloaded 14 Mar. 2010), and chlorophyll a and coloured dissolved organic matter (CDOM) (2005-2010) records were acquired from the Giovanni online data system (http://gdata1.sci.gsfc.nasa.gov/daac-bin/G3/gui.cgi?instance_id=oce-an_month; downloaded 14 Mar. 2010), which was developed and is maintained by the NASA Goddard Earth Sciences (GES) Data and Information Services Center (DISC), MD, USA. All the data were acquired from 3 regions of the Indian Ocean; Central Indian Ocean (CIO) – Chagos Archipelago; West Indian Ocean (WIO) – Zanzibar; and East Indian Ocean (EIO) – Andaman Sea.

Results
Symbiodinium nomenclature depends on the method of molecular analysis used. For example, types C and Ca in SSCP is designated as C1 and C3 in DGGE. Since we have utilized both SSCP and DGGE for analysis, to avoid confusion, throughout the text we will follow the nomenclature used for DGGE (see Table 2 for more details).

PCR-SSCP of ITS1
Separate taxonomic units were identified within clade C; C1, C3, C3z, C40, Ca5, C1c, C2, Cs and Csh. SSCP results for A. muricata, P. damicornis, P. eydouxi, P. verrucosa, S. pistillata and S. hystrix are shown in Fig. 3A. Symbiodinium type diversity in Pocillopora damicornis, P. eydouxi, P. verrucosa, S. pistillata and S. hystrix revealed the presence of C1c and C2 in three Pocillopora species, Cs in S. pistillata and S. hystrix, and Csh in S. hystrix (Fig. 3A). The coral I. Palifera was found to be associated with the greatest number of symbionts including 5 taxa from clade C (C3, C3z, Ca5 and C3z, and C3i; Fig. 3A) and with Symbiodinium type A1 within clade A.
Figure 3. ITS1-SSCP and ITS2-DGGE fingerprints of Symbiodinium types from representative samples of seven corals species from the Chagos Archipelago. Information of DNA samples from different coral species run in each lane of ITS1-SSCP is shown next to the gel photo (A).
For ITS2-DGGE analysis, samples from all representative Symbiodinium types are shown on the gel (B). Sample number is shown on each lane gel with markers denoted by the letter M on each gel. Symbiodinium type name is shown at the bottom of the lane. Acropora muricata – Lane 1–7; Isopora palifera – Lane 1–17 (inside red box); Pocillopora damicornis – Lane 1–4; Pocillopora verrucosa – Lane 5–7; Pocillopora eudoxi – Lane 8–10; Seriatopora hystrix – Lane 1 and 2 and Stylophora pistillata – Lane 1–5. The bands pointed with black arrows were cut and sequenced to confirm the Symbiodinium types.

doi:10.1371/journal.pone.0035836.g003

Discussion

Several studies in the past few years have advanced our knowledge on the biogeographic distribution of Symbiodinium diversity in corals and have contributed to our understanding of the coral-symbiont association with respect to their specificity to regional environments [17–22]. Notably, [22] compared symbiosis patterns in the Indian Ocean (north-east and west Indian Ocean) with the Great Barrier Reef. This showed the occurrence of high incidences of clade D Symbiodinium in coral hosts in high temperature, turbid waters of the Andaman Sea (Thailand) compared with clade C in the Great Barrier Reef and Zanzibar. It was suggested that high temperature and turbidity might explain, in part, the ecological success of clade D in these areas [22].

Samples form the Chagos that we analyzed did not show any presence of clade D, even though the composition of Symbiodinium varied among reefs (Fig. 2). The absence of clade D in this survey could be because insufficient taxa were sampled. However, not one of the 262 specimens of seven species of major and abundant coral species in the Chagos showed the presence of Symbiodinium clade D. Our results indicate high incidence of Symbiodinium clade C types with occasional occurrence of Symbiodinium A1 in the samples analyzed, which is different form the previously investigated locations in the Indian Ocean.

The type of coral-Symbiodinium association in the Chagos observed through this study may be due to the fact that the central Indian Ocean atolls are bathed by water very low in dissolved organic matter (CDOM) with much less pelagic chlorophyll than the northeast Indian Ocean (Andaman Sea) and west Indian Ocean (Zanzibar) [Fig. 4 see also [22,58]]. Water quality not only affects the distribution of corals but also the Symbiodinium composition [22,59]. Also, the mass mortality in 1998 in the Chagos archipelago was triggered as least as much by high irradiance as by warming of seawater; the trade winds that year did not develop [60] leading to prolonged calm sea surface conditions conducive to greater light penetration [61]. This is illustrated by many examples of coral colonies whose shaded sides survived while top surfaces exposed to light did not [62]. The bleaching in the central Indian Ocean in recent years appears to be as much a consequence of very stable water layers and thermoclines given extended periods of low winds, leading to enhanced light penetration, as was to raised temperatures [61,63]. However, the average sea surface temperatures in the Chagos atolls are around 28°C, with the warmest being about 30°C [64], which is cooler than in the north-east Indian Ocean (Andaman Sea) where average sea surface temperatures exceed 31°C in summer (Fig. 4). All the above-described conditions can be conducive to the presence of Symbiodinium clad C and A in the corals of the Chagos Archipelago. But these correlations needs to be explored with caution since presence of corals and their association with a particular Symbiodinium types dose not necessarily correlate with their physiological adaptation to the prevailing environmental conditions in a given area.

Symbiodinium type diversity

Previous studies have shown that it is important to resolve the high diversity within Symbiodinium clade C [20,22,34,46,65]. Symbiodinium types C1 and C3 are common in the Pacific Ocean and Indian Ocean [20,22]. In this study we could detect at least 14 types of clade C; C1, C1a, C3, C3z, C3i, C40, C3kk, C132a, C132, C40, C41, C40, C41, C40, Cs and Cs1. *P. palifera* had most diverse Symbiodinium clade C types (C3, C3z, C3i, C40 and Ca3) (Fig. 3B). The variety of C types present in *P. palifera* may be due to vertical transmission of Symbiodinium from parental corals [19]. *A. muricata* hosted type C1 and C3, C3kk (novel Symbiodinium type) together with Symbiodinium A1 (29% of total samples analyzed), otherwise it was mainly associated with C1. The Presence of Symbiodinium A1 in *Acroporidae* in the Chagos is consistent with previous reports of these associations at various locations in the Pacific [46]. Similarly, two Symbiodinium types C40 and C3z found in the Acroporidae have been previously recorded in the west Pacific Ocean, eastern Indian Ocean off the coast of Australia and at several locations in Thailand respectively [19,22,66]. This indicates that Acroporidae may be flexible in its association with different Symbiodinium clades, however, whether such flexibility is related to local environmental conditions in the Chagos is a matter of future investigations. Symbiodinium type Cs was associated with both *S. hystrix* and *S. pistillata*, whereas Cs1, C132a, C132 were only found in *S. hystrix* (Fig. 3B). It has been observed that association of C1b-c in *Pocillopora* species is generally in the clear water reefs in the West Pacific Ocean [59]. This might explain the presence of type C1c (close related with C1b-c) in *Pocillopora* spp in the Chagos archipelago (Fig. 3B). C1c is a common Symbiodinium type that has been found in pocillloridae in the Pacific [19]. However, we could not detect the presence of Symbiodinium type *C1h that is associated with *Pocillopora* spp in Zanzibar [22].

What makes Symbiodinium clade C dominant in the corals of the Chagos Archipelago?

Patterns of Symbiodinium associated with corals sampled from Chagos can be explained by the different environmental conditions...
that prevail in different parts of the Indian Ocean (see Fig. 4). This ocean lacks equatorial upwelling due to the climatological winds, which tend to be westerly rather than easterly. As a result of this, a warm pool is found in the Eastern Indian Ocean rather than in the west, as is the case in the tropical Atlantic and Pacific. This is attributed to the presence of large-scale, land-sea contrasts between the Asian land mass and the Indian Ocean, resulting in the monsoonal winds and seasonal reversing of currents north of ~20°S. Also lack of winds over the surface of the ocean observed in the central Indian Ocean results in less chlorophyll and CDOM (Fig. 4). The average sea surface temperature in the Chagos is usually <30°C (Fig. 4, also see [67]) with temperatures exceeding 31°C on rare occasions, together with high PAR that results in coral bleaching [62,63]. This is conducive to the dominance of *Symbiodinium* clade C and presence of clade A to a certain extent in the Chagos corals. However, changing environmental conditions as a result of global climate change (see Fig. 4) shows an increasing trend in average sea surface temperature, which might also be working against such associations.

**Conclusion**

Time will reveal whether corals in the Chagos Archipelago adapt to the global climate change and increase in average sea surface temperatures. Presently there is no sign of high temperature resistant *Symbiodinium* clades (clade D) in the corals of the Chagos that we analyzed. The marine environment of the Chagos Archipelago is an exceptional place, which is a result of the absence of overfishing, pollution and minimal other human impacts. The reefs of the Chagos make up perhaps 50% of the total reef area in the Indian Ocean that remains in the least disturbed, low threat category. The view that reefs in the Chagos need strong protection is based not only on studies of the Chagos itself, but from knowledge and observation of the generally poor condition of reefs elsewhere in the Indian Ocean (from Madagascar to Kuwait, and from Sri Lanka westwards) and other oceans too. Now, since the Chagos Archipelago has been designated as a no-take MPA, it is suggested that more such areas be designated and that efforts be made to ensure that these areas are not disturbed by human activities. The Chagos Archipelago is the largest marine protected area in the central Indian Ocean.

**Future directions**

Our study has shown that most common coral species in the Chagos archipelago are associated with *Symbiodinium* clade C and A. Whether such coral-*Symbiodinium* associations present in the Chagos are physiologically adapted or acclimated to environmental conditions prevailing in the Chagos Archipelago and environmental tolerances of these *Symbiodinium* types needs to be confirmed through laboratory based physiological experiments.

**Supporting Information**

**Figure S1** RFLP (28S and 18S) pattern of *Symbiodinium* clades in corals from Chagos Archipelago. RFLP banding profiles shown for 28S rDNA (a) include: type A (lane 2), type A+C (lanes 3 and 4), type C+Csh (Csh; lane 5), type C+Cs (Cs; lanes 6 and 7) and type C (lanes 8–11), with the 100 bp DNA marker in lane 1. RFLP patterns of 18S rDNA (b) include: type A (lane 2), type A+C (lane 3), type C (lanes 4, 6 and 7) and type C2 (lane 8). The 100 bp DNA marker is in lane 1.

**Figure S2** Phylogenetic analysis of 28S-rDNA sequences. Maximum likelihood tree of all the samples from this study combined (a) and Neighbour-joining trees showing details of Clade C group (b) and clade A group (c). Samples from this study are represented by colored letters.

**Table S1** The biogeographic and host information for *Symbiodinium* used for phylogenetic analysis.

**Acknowledgments**

Many thanks go to members of the Coral Reef Evolutionary Ecology and Genetics Laboratory (CREEG), Biodiversity Research Center, Academia Sinica (BRCAS) for the assistance with DNA extraction, and two anonymous referees, and J.R. and T. L. for constructive comments. This is the Coral Reef Evolutionary Ecology and Genetics Group, BRCAS contribution no. 52.

**Author Contributions**

Conceived and designed the experiments: CAC CS DO. Performed the experiments: SY SK. Analyzed the data: SY SK. Contributed reagents/materials/analysis tools: CAC DO. Wrote the paper: SK SY CAC.

**References**

1. Rowan R (1998) Diversity and ecology of zooxanthellae on coral reefs. J Phycol 34: 407–417.
2. Hoegh-Guldberg O (1999) Climate change, coral bleaching and the future of the world’s coral reefs. Marine Freshwater Research 50: 839-866.
3. Stat M, Carter D, Hoegh-Guldberg O (2006) The evolutionary history of *Symbiodinium* and scleractinian host - Symbiosis, diversity, and the effect of climate change. PeerJ Ecol Evol Syst 2: 23–43.
4. Douglas AE. (2003) Coral bleaching – how and why? Mar Pollut Bull 46: 385–392.
5. Baker AC (2003a) Flexibility and specificity in coral-algal symbiosis: diversity, ecology, and biogeography of *Symbiodinium*. Annu Rev Ecol Syst 34: 661-689.
6. Baker AC (2003b) Symbiont diversity on coral reefs and its relationship to bleaching resistance and resilience. In: Rosenberg E, Loya Y (eds) Coral health and disease. Springer, Berlin, 177–191.
7. Baker AC, Starger CJ, McClanahan TR, Glynn PW (2004) Corals’ adaptive response to climate change. Nature 430: 741-742.
8. Hughes TE, Baird AH, Bellwood DR, Card M, Connolly SR, et al. (2003) Climate change, human impacts, and the resilience of coral reefs. Science 301: 929–933.
9. Bellwood DR, Hughes TP, Folke C, Nyström M (2004) Confronting the coral reef crisis. Nature 429: 927–933.
10. Hoegh-Guldberg O, Mumby PJ, Hooten AJ, Steneck RS, Greenfield P, et al. (2007) Coral reefs under rapid climate change and ocean acidification. Science 318: 1737–1742.
11. Muscatine L, Grossman D, Dosin J (1993) Release of symbiotic algae by tropical sea anemones and corals after cold shock. MEPS 77: 233–243.
12. Glynn PW (1996) Coral reef bleaching: facts, hypotheses and implications. Global Change Biology 2: 495–509.
13. Brown BE (1997) Adaptations of reef corals to physical environmental stress. Adv Mar Biol 31: 221–299.
14. Brown BE, Dunne RP, Scoffin TP, Le Tisser MDA (1994) Solar damage in intertidal corals. MEPS 105: 219-230.
15. Coles SL, Brown BE (2003) Coral bleaching-capacity for acclimatization and adaptation. Adv Mar Biol 46: 184–212.
16. van Woerden R (2009) Coral's prolonged struggle against unfavorable conditions. Galaxea 11: 53–58.
17. Leuck WKV, Loi T, Carter D, Hoegh-Guldberg O (2001) Genetic variability of the symbiotic dinoflagellates from the wide ranging coral species Seriatopora hystrix and Acropora tenuis in the Indo-West Pacific. MEPS 222: 97–107.
18. Chen CA, Lam KK, Nakano Y, Tsai WS (2003) A stable association of the stress-tolerant zooxanthellae, Symbiodinium clade D, with the low-temperature-tolerant coral, Oenothera fruticosa (Scleractinia: Fabricius) in subtropical non-reefal coral communities. Zool Stud 42: 540–550.
19. LaJeunesse TC, Bhagooli R, Hidaka M, DeVantier L, Done T, (et al.) (2004a) Closely related Symbiodinium spp. differ in relative dominance in coral reef host communities across environmental, latitudinal and biogeographic gradients. MEPS 284: 147–161.
20. LaJeunesse TC (2005) “Species” radiating of symbiotic dinoflagellates in the Atlantic and Indo-Pacific since the Miocene-Pliocene transition. Mol Ecol Evol 22: 570–581.
21. Lien YT, Nakano Y, Platlong S, Fukami H, Wang JT, (et al.) (2007) Occurrence of the putatively heat-tolerant Symbiodinium phylotype D in high-latitude outlying coral communities. Coral Reefs 26: 35–44.
22. LaJeunesse TC, DT, Sampayo EM, Possumpun N, Brown B, (et al.) (2010a) Long-standing environmental conditions, geographic isolation and host-symbiont specificity influence the relative ecological dominance and genetic diversification of coral endosymbionts in the Symbiodinium clade D. Jour Biol 375: 780–795.
23. Coles, S. L, Jokiel, P. L. (1979) Effects of temperature on photosynthesis and respiration in hermatypic corals. Mar Biol Arch 43: 209–216.
24. Sheppard CRC (2003) Predicted recurrents of mass coral mortality in the Indian Ocean. Nature 425: 294–297.
25. Rowan R, Powers DA (1991) Molecular genetic identification of symbiotic dinoflagellates via length heteroplasmy in domain V of chloroplast large subunit (cp23S)-ribosomal DNA sequences. Mar Biotechnol 5: 130–140.
26. Rowan R, Knowlton N (1995) Intraspecific diversity and ecological zonation in the reef coral Galaxea 11: 53–58.
27. Visram S, Wiedenmuller J, Douglas AE (2006) Molecular diversity of symbiotic algal of the genus Symbiodinium (Zoanthellales) in ciliartians of the Mediterranean Sea. J Mar Biol Ass UK 86: 1291–1293.
28. Coldewey-Hj, Curnick D, Harding S, Harrison LR, Golluch M (2010) Potential benefits to fisheries and biodiversity in the Chagos Archipelago/British Indian Ocean Territory as a no-take marine reserve. Mar Pollut Bull 60: 1906–1915.
29. Sheppard CRC, Rowe B, Carr P, Chen CA, Chabbe C, (et al.) (2012) Reef and islands of the Chagos Archipelago, Indian Ocean: Why it is the world's largest no-take marine protected area. Mar Freshwater Res DOI: 10.1071/AR12045.
30. Sheppard CRC (1999) Coral decline and weather patterns over 20 years in the Chagos Archipelago, Central Indian Ocean. Ambio 28: 472–478.
31. Mangi S, Hooper T, Rodwell L, Simon D, Snow D, (et al.) (2010) Establishing a marine protected area in the Chagos Archipelago: socio-economic considerations. Report of workshop held 7 January 2010, Royal Holloway, University of London, UK. 23 p.
32. Sampayo EM, Dove S, LaJeunesse TC (2009) Cohesive molecular genetic data delineate species diversity in the dinoflagellate genus Symbiodinium. Mol Ecol 18: 500–519.
33. Chen CA, Wallace CC, Yu JK, Wei NV (2000) Strategies for amplification by polymerase chain reaction of the complete sequence of the gene encoding nuclear large subunit ribosomal RNA in corals. Mar Biotechnol 2: 146–153.
34. Ulstrup KE, van Oppen MJH (2003) Geographic and habitat partitioning of genetically distinct zooxanthellae (Symbiodinium) in Acropora corals on the Great Barrier Reef. Mol Ecol 12: 3477–3484.
35. LaJeunesse TC, Trench RK (2000) The biogeography of two species of subclade D, with the low-temperature-tolerant clade C and the high-temperature-tolerant clade D, in high-latitude coral communities. Coral Reefs 19: 55–61.
36. Santos SR, Gutierrez RC, Coiffoth MA (2005a) Phylogenetic identification of symbiotic dinoflagellates via length heteroplasmy in domain V of chloroplast large subunit ribosomal DNA sequences. Mar Biotechnol 5: 130–140.
37. Tchernov D, Gorbunov MY, de Vargas C, Yadav SN, Milligan AJ, (et al.) (2004) Membrane lipids of symbiotic algae are diagnostic of sensitivity to thermal stress. Proc Royal Soc Lon B Biol Sci 275: 2273–2282.
38. Stat M, Loh WKW, Hoegh-Guldberg O, Carter D (2008a) Symbiont acquisition and host-symbiont interactions between algal endosymbionts and coral hosts define their bleaching response to heat and light stress. Proc Royal Soc Lon B Biol Sci 273: 2273–2282.
39. Chen CA, Wang JT, Fang LS, Yang YW (2005a) Clutactuating algal symbionts in Acropora palmate (Scleractinia: Acroporidae) from Taiwan. MEPS 295: 115–121.
40. Garren M, Walsh SM, Giaccone A, Knowlton N (2006) Patterns of association between symbionts and members of the Montastrea annularis species complex on spacial scales ranging from within colonies to between geographic regions. Coral Reefs 25: 503–512.
41. Oliver T, Palumbi S (2009) Distributions of stress-resistant coral symbionts match environmental patterns at local but not regional scales. MEPS 375: 93–103.
42. van Oppen MJH, Baker AC, Coiffoth MA, Willis BL (2009) Bleaching resistance and the role of algal endosymbionts. In: van Oppen MJH, Lough JM (eds) Coral Bleaching. Ecological Studies 203. Springer-Verlag, Berlin and Heidelberg 83–102.
43. LaJeunesse TC, Loi WKV, Trench RR (2009) Do introduced endosymbiotic dinoflagellates “take” to new hosts? Biol Invasions 11: 993–1003.
44. Chen CA, Yang YW, Wei NV, Tsai WS, Fang LS (2003b) Symbiont diversity in Scleractinian corals from tropical reefs and subropical non-reef communities in Taiwan. Coral Reefs 24: 11–22.
45. LaJeunesse TC, Loi WKV, Van Wees R, Hoegh-Guldberg O, Schmidt GW, et al. (2008) Low symbiont diversity in southern Great Barrier Reef corals, related to those of the Caribbean. Limned Oceanogr 48: 2056–2054.
46. Visram S, Wiedenmann J, Douglas AE (2006) Molecular diversity of symbiotic algae of the genus Symbiodinium (Zoanthellales) in ciliartians of the Mediterranean Sea. J Mar Biol Ass UK 86: 1291–1293.
47. Sheppard CRC (1999) Coral decline and weather patterns over 20 years in the Chagos Archipelago, Central Indian Ocean. Ambio 28: 472–478.
48. LaJeunesse TC, Smith R, Walther M, Pinzón J, Pettay DT, (et al.) (2010b) Host-symbiont recombination versus natural selection in the response of coral-dinoflagellate symbioses to environmental disturbance. Proc R Soc Lond, B Biol Sci 277: 2925–2934.
49. West JM, Slam RV (2003) Resistance and resilience to coral bleaching: Implications for coral reef conservation and management. Conserv Biol 17: 956–967.
50. Sheppard CRC (2006) Longer term impacts of climate change. In: Cote I, Reynolds J (eds) Coral Reef Conservation Cambridge University Press. pp 264–290.
51. Webster PJ, Moore AM, Louchmig JP, Lebron RR (1999) Coupled ocean-atmosphere dynamics in the Indian Ocean during 1997–98. Nature 401: 356–360.
52. Sheppard CRC, Obura D (2005) coral reefs of the Western Indian Ocean: an overview. In: McClanahan TR, Sheppard CRC, Obura D (eds) Coral reefs of the Indian Ocean and their conservation Oxford University Press. pp 3–32.
53. LaJeunesse TC, Smith R, Walther M, Pinzón J, Pettay DT, (et al.) (2010b) Host-