Differentiation of Acanthamoeba Strains from Infected Corneas and the Environment by Using Restriction Endonuclease Digestion of Whole-Cell DNA

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Restriction endonuclease digestion of Acanthamoeba whole-cell DNA was used to study the relationship between 33 morphologically identical strains from keratitis cases (30 strains), contact lens storage containers (2 strains), and soil (1 strain). Samples digested with BgllI, EcoRI, or Hindlll and separated by agarose gel electrophoresis contained detectable mitochondrial DNA restriction fragment length polymorphisms (RFLPs). By comparing RFLPs, the strains could be assigned to seven multiple-strain and three single-strain groups. The largest of these contained nine strains, eight of which were isolated in keratitis cases in various locations worldwide and may indicate a group particularly associated with keratitis. Restriction endonuclease analysis of whole-cell DNA is proposed as ^a valuable technique for detecting mitochondrial DNA RFLPs in the differentiation of morphologically identical Acanthamoeba strains and may therefore be useful in resolving the complex taxonomy of the genus, which has hitherto been founded on subjective morphological criteria.

Acanthamoebae are small free-living amoebae with a life cycle characterized by a trophozoite stage and a cyst stage (25, 26). These organisms are common in the environment and have been isolated from soil, natural and man-made aquatic sites, and the atmosphere (16, 25, 27). Acanthamoebae are increasingly being recognized as a cause of severe sight-threatening keratitis (12, 28). Since the disease was first recognized (13), several hundred cases have been reported, mostly from the United States (29). Contact lens wearers appear most at risk from infection (20, 28), accounting for approximately 85% of cases (29). Poor hygiene practices, notably the preparation of homemade saline rinsing solutions, are recognized risk factors among contact lens wearers (20, 28).

Although Acanthamoeba keratitis may be routinely diagnosed by the culture or microscopic examination of clinical material (32), the identification of species involved in the disease is unclear. The taxonomic classification of the members of the genus Acanthamoeba is derived from morphological observations of the trophozoite and cyst forms of these organisms (25, 26). The most detailed study is that of Pussard and Pons, who recognized 18 species that were assigned to three morphological groups (26). While this classification clearly defines the genus, the variation in cyst morphology seen even within cloned strains makes the identification of many described species a subjective and unreliable process (11, 31, 33). This has prompted the investigation of other methods for the differentiation of acanthamoebae. Isoenzyme electrophoresis (8, 11) and restriction endonuclease digestion of mitochondrial DNA (mtDNA) or whole-cell DNA (3, 4, 7, 19) have proved most useful in this respect. These methods have all demonstrated wide interstrain variation within species and also similarities between strains of separate species (7, 11).

While such studies have highlighted the inadequacies of the present taxonomic classification of the acanthamoebae,

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MATERIALS AND METHODS

Amoebae. The Acanthamoeba strains examined in this study are listed in Table 1. The majority of strains were from cases of keratitis not reported in the literature and were identified only by the names of the patients from whom they were isolated. Where possible, a reference to a strain is cited in Table 1. The cyst forms of all the strains appeared morphologically similar when viewed by light microscopy. Strains Ac/PHL/7a through Ac/PHL/7c were three consecutive isolates from the same patient obtained over a 7-month period. By cyst morphology, strains Ac/PHL/6 (Bitzer), Ac/PHL/8 (CCAP 1501/3d), Ac/PHL/10 (Garcia), and Ac/ PHL/23 (SHI) have previously been identified as Acanthamoeba polyphaga (13, 14, 22) and Ac/PHL/29 (CCAP 1501/la, Neff Strain) has been identified as Acanthamoeba castellanii (23). The two contact lens storage case isolates, Ac/PHL/30 and Ac/PHL/31, were isolated from asymptomatic patients and found to be cytopathogenic for Vero tissue culture cells at 32°C (14). Trophozoites were adapted to axenic culture in Chang's serum-casein-glucose-yeast extract medium (1), which was modified in this study by the inclusion of 0.1% filter-sterilized Panmede liver digest

the strains examined have mainly been of environmental origin and unknown pathogenicity. The relationship between 33 morphologically identical Acanthamoeba strains from patients with keratitis (30 strains), contact lens storage containers (2 strains), and soil (1 strain) was therefore investigated by restriction endonuclease digestion of wholecell DNA. The detection of mtDNA restriction fragment length polymorphisms (RFLPs) that enabled the strains to be assigned to seven multiple-strain and three single-strain groups has prompted this report.

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TABLE 1. Acanthamoeba strains examined in this study

Isolate	Country of origin		Reference	
Ac/PHL/1 (Ac 112)	Australia	AK.		
Ac/PHL/2 (Ac 116)	Australia	AK	$\frac{2}{2}$	
Ac/PHL/3	England	AК		
Ac/PHL/4	England	AК		
Ac/PHL/5	The Netherlands	AK		
Ac/PHL/6 (Bitzer)	United States	АK	13	
Ac/PHL/7a	England	AК	30	
Ac/PHL/7b	England	AК	30	
Ac/PHL/7c	England	AK	30	
Ac/PHL/8 (CCAP 1501/3d)	England	AK	22	
Ac/PHL/10 (Garcia)	United States	AК	13	
Ac/PHL/12	England	AK		
Ac/PHL/13	England	AK		
Ac/PHL/14	England	AK		
Ac/PHL/15	Germany	AK		
Ac/PHL/16	France	AK		
Ac/PHL/17	England	AK		
Ac/PHL/18	The Netherlands	AK		
Ac/PHL/19	England	AK		
Ac/PHL/20	England	AK		
Ac/PHL/21	France	AK		
Ac/PHL/22	England	AK		
Ac/PHL/23 (SHI)	England	AK	14	
Ac/PHL/24	The Netherlands	AK		
Ac/PHL/25	England	AK		
Ac/PHL/26	France	AK		
Ac/PHL/28	England	AК		
Ac/PHL/29 (CCAP 1501/	United States	Soil	23	
1a, Neff strain)				
Ac/PHL/30	England	CLC	14	
Ac/PHL/31	England	CLC	14	
Ac/PHL/32	Australia	AK		
Ac/PHL/33	England	AK	15	
Ac/PHL/34	England	AК	1	

^a AK, Acanthamoeba keratitis; CLC, contact lens container.

(Paines & Byrne Ltd., Greenford, England) and maintained in tissue culture flasks (GIBCO, Uxbridge, Middlesex, England) at 32°C.

DNA isolation. Approximately 5×10^7 trophozoites of each strain were used for DNA isolation. Late-log-phase cultures were concentrated by centrifugation at 500 \times g for 10 min at room temperature and suspended in 5 ml of cell lysis buffer (100 mM disodium EDTA, ¹⁰⁰ mM NaCI, ¹⁰ mM Tris hydrochloride [pH 8.0]). Proteinase K (20 mg/ml) was added to a final concentration of $300 \mu g/ml$, and the cells were lysed in 1% Sarkosyl (20%, wt/vol). After gentle mixing by inversion, the lysates were incubated at 56°C for a period ranging from 4 h to overnight. The samples were then chilled on ice for 5 min and extracted twice by gentle mixing for 5 min with equal volumes of phenol-chloroform-isoamyl alcohol (25:24:1) and once with chloroform-isoamyl alcohol (24:1). After each extraction, the aqueous and organic phases were separated by centrifugation at 2,000 \times g for 15 min at room temperature, and the upper aqueous phase was retained. Nucleic acids were precipitated by the addition of an equal volume of isopropanol and holding at -20° C for a period ranging from ¹ h to overnight (18). The precipitate was pelleted by centrifugation at $2,000 \times g$ for 10 min at room temperature and dissolved in ⁵ ml of TE buffer (10 mM Tris hydrochloride, ¹ mM disodium EDTA [pH 7.5]). RNase (5 mg/ml) was added up to a final concentration of 50 μ g/ml, and the mixture was incubated at 37°C for ¹ h. After extraction once each with phenol-chloroform-isoamyl alco-

FIG. 1. Agarose gel electrophoresis of EcoRI digests of Acanthamoeba whole-cell DNA. Lanes: 1, Ac/PHL/28; 2, Ac/PHL/24; 3, Ac/PHL/23 (SHI); 4, Ac/PHL/22; 5, Ac/PHL/20; 6, Ac/PHL/19; 7, Ac/PHL/18; 8, Ac/PHL/15; 9, Ac/PHL/12; 10, Ac/PHL/10; 11, Ac/PHL/b; 12, Ac/PHL/6 (Bitzer); 13, Ac/PHL/5; 14, Ac/PHL/4; 15, Ac/PHL/3; 16, Ac/PHL/8 (CCAP 1501/3d); 17, Ac/PHL/30; 18, Ac/PHL/29 (CCAP 1501/la, Neff strain); 19, DNA fragments from lambda HindIII- ϕ X 174 RF HaeIII digestion.

hol (25:24:1) and chloroform-isoamyl alcohol (24:1), the DNA was precipitated by adding 0.1 volume of ³ M sodium acetate (pH 5.2) and an equal volume of isopropanol and holding at -20° C for a period ranging from 1 h to overnight. The DNA was pelleted by centrifugation, washed twice with 70% ethanol, and dried briefly under ^a vacuum. The DNA was dissolved in 300 μ l of double-distilled H₂O and stored at -20°C until use.

Restriction endonuclease analysis. DNA samples (2 to 3 μ g in 18 μ I) were digested at 37°C for 2 h with 5 to 10 U of the restriction endonuclease BamHI, BglII, EcoRI, HindIII, KpnI, or PstI by using appropriate reaction buffers provided with the enzymes (Northumbria Biologicals Ltd., Cramlington, Northumberland, England). The samples were loaded

FIG. 2. Agarose gel electrophoresis of HindIlI digests of Acanthamoeba whole-cell DNA. Lanes: ¹ and 18, DNA fragments from lambda HindIII- ϕ X 174 RF HaeIII digestion; 2, Ac/PHL/28; 3, Ac/PHL/24; 4, Ac/PHL/23 (SHI); 5, Ac/PHL/22; 6, Ac/PHL/20; 7, Ac/PHL/19; 8, Ac/PHL/15; 9, Ac/PHL/12; 10, Ac/PHL/10 (Garcia); 11, Ac/PHL/5; 12, Ac/PHL/7b; 13, Ac/PHL/4; 14, Ac/PHL/3; 15, Ac/PHL/8 (CCAP 1501/3d); 16, Ac/PHL/30; 17, Ac/PHL/29 (CCAP 1501/la, Neff strain).

FIG. 3. Differentiation of *Acanthamoeba* strains by $EcoRI$ summarized in Table 2. RFLP groups (see Table ³

onto horizontal 0.7% agarose gels (20 by 20 by 0.5 cm) prepared in TBE buffer (89 mM Tris, 89 mM boric acid, 2.5 mM disodium EDTA [pH 8.2]) for electrophoresis in TBE strains tested. buffer at 1.5 V/cm overnight (18). DNA standards of lambda HindIII and ϕ X 174 RF HaeIII digests (Pharmacia LKB Ltd., Milton Keynes, England) were included as size markers. Gels were stained with 1 μ g of ethidium bromide per ml $Bgl1I, EcoRI,$ and HindIII are shown in Table 3. in distilled $H₂O$ for 1 h, destained in distilled water for 20 min, and photographed under shortwave UV transillumination with Polaroid 665 film and a Kodak Wratten 23A orange filter.

Analysis of results. The sizes of the RFLPs in kilobases were estimated with the GELPROG computer program written by K. Merrifield at the Institute of Cancer Research, Chester Beatty Laboratories, London, England.

RESULTS

Digestion of Acanthamoeba whole-cell DNA with the restriction endonucleases BamHI, KpnI, and PstI resulted in only ^a smearing of DNA after agarose gel electrophoresis and staining with ethidium bromide (result not shown). Increasing the restriction endonuclease concentration and incubation time did not alter this effect. In contrast, samples digested with BgII, EcoRI, and Hindlll produced discrete bands of DNA that enabled specific RFLPs to be detected (Fig. ¹ and 2). This permitted the strains to be assigned to seven multiple-strain and three single-strain RFLP groups that were consistent for each restriction endonuclease used (Fig. ³ to 5). Of the 33 strains examined, only Ac/PHL/22, Ac/PHL/25, and Ac/PHL/29 (CCAP 1501/la, Neff strain)

FIG. 4. Differentiation of Acanthamoeba strains by HindIII RFLP groups (see Table 3).

 $\frac{D}{C}$ $\stackrel{f}{\leftarrow}$ $\stackrel{g}{\leftarrow}$ $\stackrel{g}{\leftarrow}$ $\stackrel{g}{\leftarrow}$ $\stackrel{g}{\leftarrow}$ showed unique RFLPs. With EcoRI, the RFLPs of strains showed unique KFLFs. With ECONI, the KFLFs of strains
assigned to groups A and C appeared closely related, but
- used. The most common RFLP profile belonged to the nine
strains assigned to group A, of which eight were from they were clearly separated when HindIII and BglII were used. The most common RFLP profile belonged to the nine strains assigned to group A, of which eight were from keratitis cases in Australia (Ac/PHL/1 and Ac/PHL/2), the United States (Ac/PHL/6 [Bitzer]), England (Ac/PHL/17, $\equiv \equiv \equiv \equiv -$ strains assigned to group A, of which eight were from
 $\equiv \equiv \equiv \equiv$ - keratitis cases in Australia (Ac/PHL/1 and Ac/PHL/2), the
 $\equiv \equiv \equiv$ - United States (Ac/PHL/6 [Bitzer]), England (Ac/PHL/17,
 \equiv - - Ac/PHL and the Netherlands (Ac/PHL/24). This was followed by group F (five strains), group G (four), groups B and C (three each), groups D and E (two each, assuming Ac/PHL/7a, Ac/PHL/7b, and Ac/PHL/7c of group D are the same strain), and groups H, I, and J (one each). These findings are

> The four strains identified as A. polyphaga by cyst morphology were separated into groups A (Ac/PHL/6 [Bitzer] and Ac/PHL/23 [SHI]), B (Ac/PHL/8 [CCAP 1501/3d]), and E (Ac/PHL/10 [Garcia]). A. castellanii (Ac/PHL/29 [CCAP 1501/1a, Neff strain]) showed unique RFLPs from all the strains tested.

> The sums of the sizes (in kilobases) of the RFLPs detected in strains Ac/PHL/7a, Ac/PHL/23 (SHI), Ac/PHL/28, and $Ac/PHL/29$ (CCAP 1501/la, Neff strain) by digestion with Bg/II , Ec/ORI , and $HindIII$ are shown in Table 3.

DISCUSSION

By either cyst morphology or biochemical analysis, A. castellanii, A. hatchetti, A. Iugdunensis, A. polyphaga, and A. rhysodes of morphological group 11 (26) and A. culbertsoni of morphological group III have been described as causing keratitis. The 33 strains examined in this study formed cysts typical of group II Acanthamoeba spp. and resembled A. polyphaga or possibly A. castellanii (25, 26), as the differentiation of these species by this means is considered subjective (11, 31, 33). Using restriction endonuclease digestion of Acanthamoeba whole-cell DNA, we were able to detect RFLPs that differentiated these strains into seven multiple-strain and three single-strain groups labeled A through ^J directly by agarose gel electrophoresis. Of these, group A contained the largest number of strains (nine), of which eight were isolated from keratitis cases in various locations worldwide. This group may therefore indicate the type most frequently associated with keratitis through either a greater prevalence in the environment or increased virulence for corneas.

				Although <i>Acanthamoeba</i> keratitis has been successfully						
Kb23.1 $9.4 -$ 6.6 4.4										
$2.3 - 2.0 -$										
$1.35 -$ 1.1										

FIG. 5. Differentiation of Acanthamoeba strains by BgIII RFLP groups (see Table 3).

TABLE 2. RFLP group assignment of Acanthamoeba strains

RFLP group	Acanthamoeba strain
	A Ac/PHL/1 (Ac 112), Ac/PHL/2 (Ac 116), Ac/PHL/ 6 (Bitzer), Ac/PHL/17, Ac/PHL/15, Ac/PHL/24, Ac/PHL/23 (SHI), Ac/PHL/30, Ac/PHL/34
	B Ac/PHL/5, Ac/PHL/8 (CCAP 1501/3d), Ac/PHL/12
	C Ac/PHL/3, Ac/PHL/4, Ac/PHL/20
	DAc/PHL/7a, Ac/PHL/7b, Ac/PHL/7c, Ac/PHL/18
	E Ac/PHL/10 (Garcia), Ac/PHL/31
	F Ac/PHL/13, Ac/PHL/14, Ac/PHL/16, Ac/PHL/19, Ac/PHL/21
	G Ac/PHL/26, Ac/PHL/28, Ac/PHL/32, Ac/PHL/33
	H $Ac/PHL/22$
	I Ac/PHL/25
	JAc/PHL/29 (CCAP 1501/1a, Neff strain)

treated with propamidine isethionate (35), the disease frequently necessitates surgical intervention (6, 34). Reports on the susceptibility of Acanthamoeba strains to chemotherapeutic agents and contact lens disinfectants often give conflicting results (5, 6, 9, 17, 21, 35). Strain Ac/PHL/33 of group G is inherently resistant to propamidine isethionate (15), while strain Ac/PHL/23 (SHI) of group A is susceptible (unpublished observations). This suggests that innate differences in susceptibility to antimicrobial agents occur between Acanthamoeba strains. Whether such variation correlates with RFLP group assignments of strains merits further investigation. Acquired drug resistance can also complicate the treatment of Acanthamoeba keratitis. Strains Ac/PHL/ 7a, Ac/PHL/7b, and Ac/PHL/7c represent three consecutive isolates from the same patient over a 7-month period. Ac/PHL/7c is resistant to propamidine isethionate, while Ac/PHL/7a and Ac/PHL7b are susceptible (34). Since all three showed identical RFLPs, this result indicates that resistance was acquired by the original infecting strain during the course of treatment.

RFLP profiles may also be useful in identifying pathogenic Acanthamoeba strains. The virulence of acanthamoebae has been shown to attenuate during axenic culture (30) and may account for the description of both pathogenic and nonpathogenic strains within species (10). With the exception of group J, which contains only an environmental strain, Ac/ PHL/29 (CCAP 1501/1a, Neff strain), the RFLP profiles defined here are those of pathogenic strains associated with keratitis. Therefore, environmental strains with RFLPs of groups A through ^I could be assumed to be potential human pathogens.

Byers et al. (4) previously studied the relationship of 13 Acanthamoeba strains, including Ac/PHL/29 (CCAP 1501/ la, Neff strain), by restriction endonuclease digestion of purified mtDNA. The number and positions of the EcoRI RFLPs for this strain are identical to those obtained in this study. Bogler et al. (3) estimated the size of the mtDNA genome for this strain to be 40.6 kb, which is in accord with the value of 40.1 kb found in our study. It is therefore concluded that restriction endonuclease digestion of Acan-

TABLE 3. Total sizes of Acanthamoeba RFLPs

Acanthamoeba strain	Size of RFLPs (kb) with indicated restriction endonuclease					
		Sum				
	Belll	EcoRI	HindIII	Mean \pm SD		
Ac/PHL/7a	37.5	43.3	38.1	39.6 ± 2.6		
Ac/PHL/23 (SHI)	42.6	39.6	39.2	40.5 ± 1.5		
Ac/PHL/28	42.2	37.1	39.5	39.5 ± 2.6		
Ac/PHL/29 (CCAP 1501/1a, Neff strain)	43.3	40.8	36.1	40.1 ± 3.7		

thamoeba whole-cell DNA gives rise to RFLPs originating from mtDNA. The reason for the failure of the restriction endonucleases BamHI, KpnI, and PstI to cleave the mtDNA in the Acanthamoeba whole-cell DNA samples is not known. This failure may be due to the absence of endonuclease sequence recognition sites on the mtDNA or due to methylation of the sites, which would render them resistant to digestion (24).

In conclusion, restriction endonuclease digestion of whole-cell DNA is ^a potent technique for differentiating morphologically identical Acanthamoeba strains by the detection of mtDNA RFLPs. Although this study was not intended to be a taxonomic investigation of the acanthamoebae, four strains of A. polyphaga and one of A. castellanii that had been identified by cyst morphology were included. Of these, only two of the four strains of A. polyphaga showed identical RFLPs, while the one strain of A. castellanii was found to have a unique RFLP profile. It is therefore unclear whether Acanthamoeba mtDNA RFLPs indicate intra- or interspecies differences. Future studies will address this problem with a larger number of strains of the presently described species of the genus.

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