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Change in Age Pattern of Persons with Dengue, Northeastern Brazil

To the Editor: Approximately 40% of the world's population is at risk for dengue (1). Epidemiologic characteristics of dengue differ by region, and disease incidence varies by patient age. In Southeast Asia, incidence of dengue fever (DF) and dengue hemorrhagic fever (DHF) is highest among children (2,3); in the Western Hemisphere, incidence of these diseases is higher among adults.

In Brazil, which has $\approx 80\%$ of dengue cases in the Western Hemisphere, adults are at risk for dengue virus (DENV) infection (3,4). However, in 2007, a total of 53% of persons in Brazil hospitalized with DHF were children <15 years of age; this proportion was highest (65.4%) in children in northeastern Brazil (5).

In Ceará, a state in northeastern Brazil, DENV-1 epidemics have occurred since 1987. DHF cases have been reported since 1994 when DENV-2 was identified. In 2003, a severe DENV-3 epidemic occurred, and DHF incidence was high among adults (6). However, since 2007, incidence of DENV infection has been highest among children (7). To better understand factors that could affect this change in risk by age group, we studied the temporal progression of age distribution of persons with dengue during 1998–2008 in Ceará.

We used data for Ceará from the National System of Notifiable Diseases (DF and DHF cases), the Hospital Admission Data System (dengue hospitalizations) (8), and the Central Public Health Laboratory (virus isolation). For each age group (<10, 10-19, 20–59, and ≥ 60 years), we calculated incidence of DF and hospitalization rate for DHF. We also calculated proportions of dengue serotypes per year (2001–2008). Medians for continuous variables were compared by using the Kruskal-Wallis test. Analyses were performed by using Epi Info version 6.0 software (Centers for Disease Control and Prevention, Atlanta, GA, USA).

From 1998 (10.8 cases/100,000 persons) through 2007 (236.7 cases/100,000 persons), DF incidence was lowest among persons <10 years of age. However, the incidence was highest (599.4 cases/100,000 persons) for this age group in 2008. In 2007, incidence among persons <10 years of age (236.7 cases/100,000 persons) was similar to that among persons 10–19, 20-59, and >60 years of age (305.6, 331.5, and 249.9 cases/100,000 persons, respectively).

Since 2007, the incidence of DHF among children was already higher (4,884 cases/100,000 persons) than among the other age groups (3,261, 3,387 and 2,789 cases/100,000 persons, respectively). In 2008, incidence of DHF among children was 8,992 cases/100,000 persons, which was 2× that among persons 10–19 and 20–59 years of age and $>3\times$ that among persons ≥ 60 years of age. Median age of persons with DHF decreased from 38 years in 2001 to 18 years in 2008 (p<0.0001). Children <10 years age, who in 2001 accounted for 5% of all cases, accounted for 33% of cases in 2008.

The hospitalization rate for dengue among children in Ceará followed a pattern similar to that for DHF and increased for children ≤ 10 years of age. In 2008, this pattern was greater for this age group (1.449/1,000 hospitalizations) than in any other age group. DENV-2 (52.3%) and DENV-1 (47.7%) were co-circulating in 2002. DENV-3 was isolated in 2003 and represented >40% of isolations. At this time, DENV-2 and DENV-1 represented 7.4% and 48.5% of isolations, respectively. DENV-3 then predominated in Ceará until 2006 when DENV-2 reemerged (1.4%). DENV-2 became the predominant serotype in 2007 (84%) and 2008 (76.1%) (Table).

The increase in DHF incidence among children in Ceará during 2007–2008 was greater than the overall increase in Brazil (4,5). Because the predominant serotype in Ceará in 2007–2008 was DENV-2, two hypotheses may explain this phenomenon.

First, a more virulent DENV-2 may have been introduced. Genetic sequencing of DENV-2 circulating in another state in Brazil during the 2008 epidemic, compared with the 1990 and 1998 epidemics, showed that all isolates had the same genotype (American/Asiatic); only a 2% had a phylogenetic change (9). Such a small difference cannot explain this change in the age group affected by dengue.

Second, the time when 3 serotypes circulated in Ceará may not have favored development of antibodies against DENV-3 in children <10 years of age, although they were susceptible to DENV-2. DENV-1 and DENV-2 were circulating in Ceará before 2002 and caused DF epidemics and a few DHF cases. However, these diseases occurred predominantly in adults. Conversely, children had little likelihood of being infected with DENV-2 because the incidence of dengue before 2002 was low and the 2000 birth cohort had little contact with DENV-2, which was no longer circulating. Thus, most persons susceptible to DENV-2 were children.

DENV-3 was circulating during 2003–2006 and affected persons of all ages. Thus, when DENV-2 reemerged in 2006, many adults in Ceará already had antibodies against it. However, children had no antibodies against DENV-2, although some had antibodies against DENV-2, although some had antibodies against DENV-3. This immunologic difference may have caused the higher incidence of dengue among children in Ceará during 2007–2008, particularly in view of severity of the 2 epidemics, increased risk for DHF (*10*), and number of hospitalizations for dengue during this period.

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| Table. Confirmed ca | ases of DF a | nd DHF, C | Ceará, Braz | zil, 1998–2 | 008* | | | | | | |
|----------------------|-----------------------|-------------|-------------|--------------|--------|--------|-------|--------|--------|--------|--------|
| Characteristic | 1998 | 1999 | 2000 | 2001 | 2002 | 2003 | 2004 | 2005 | 2006 | 2007 | 2008 |
| No. DF cases | 3,581 | 9,757 | 13,645 | 34,390 | 16,465 | 23,796 | 3,094 | 22,817 | 25,569 | 25,026 | 44,508 |
| No. DHF cases | 4 | 3 | 4 | 82 | 72 | 292 | 14 | 199 | 173 | 280 | 408 |
| Serotype of DENV i | solated, % | | | | | | | | | | |
| 1 | _ | _ | _ | 47.7 | 48.5 | 1.9 | 0 | 2.5 | 0 | 0 | 0 |
| 2 | - | _ | _ | 52.3 | 7.4 | 1.9 | 0 | 0.0 | 1.4 | 84.0 | 76.1 |
| 3 | _ | _ | _ | 0 | 44.1 | 96.2 | 100.0 | 97.5 | 98.6 | 16.0 | 23.9 |
| Incidence of DF and | DHF by ag | e, y† | | | | | | | | | |
| <10 | 10.8 | 37.8 | 65.4 | 174.7 | 78.5 | 128.0 | 14.3 | 126.5 | 116.0 | 236.7 | 599.4 |
| 10–19 | 26.7 | 95.4 | 129.3 | 321.4 | 160.6 | 250.4 | 34.3 | 198.2 | 247.9 | 305.6 | 574.4 |
| 20–59 | 68.3 | 198.7 | 263.6 | 659.3 | 304.3 | 416.9 | 53.6 | 365.2 | 412.6 | 331.5 | 521.9 |
| <u>></u> 60 | 143.1 | 210.0 | 194.8 | 423.4 | 223.3 | 313.0 | 39.1 | 441.5 | 422.2 | 249.9 | 301.0 |
| Incidence of DHF by | y age, y † | | | | | | | | | | |
| <10 | 0.000 | 0.000 | 0.000 | 0.236 | 0.114 | 1.423 | 0.068 | 1.011 | 0.890 | 4.884 | 8.992 |
| 10–19 | 0.062 | 0.000 | 0.059 | 0.796 | 0.449 | 3.601 | 0.065 | 2.512 | 1.771 | 3.261 | 4.968 |
| 20–59 | 0.064 | 0.094 | 0.058 | 1.361 | 1.474 | 4.655 | 0.318 | 3.074 | 3.053 | 3.387 | 4.358 |
| <u>></u> 60 | 0.174 | 0.000 | 0.152 | 1.890 | 1.289 | 5.134 | 0.000 | 2.650 | 0.832 | 2.789 | 2.868 |
| Hospitalizations for | dengue (cla | ssic and he | emorrhagio | c) by age, y | y‡ | | | | | | |
| <10 | 0.008 | 0.014 | 0.015 | 0.050 | 0.145 | 0.352 | 0.073 | 0.395 | 0.266 | 0.667 | 1.449 |
| 10–19 | 0.006 | 0.010 | 0.020 | 0.131 | 0.255 | 0.632 | 0.156 | 0.646 | 0.497 | 0.750 | 1.128 |
| 20–59 | 0.005 | 0.011 | 0.024 | 0.256 | 0.440 | 0.847 | 0.255 | 0.804 | 0.572 | 0.710 | 0.786 |
| <u>></u> 60 | 0.009 | 0.009 | 0.021 | 0.401 | 0.658 | 1.159 | 0.389 | 1.012 | 0.664 | 0.954 | 0.922 |

*Source: Hospitalization Information System, Notifiable Disease Information System, and Central Public Health Laboratory of Ceará. DF, dengue fever; DHF, dengue hemorrhagic fever, DENV, dengue virus; –, no information.

†Per 100,000 persons.

‡Per 1,000 persons.

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Apophysomyces variabilis Infections in Humans

To the Editor: The fungus *Apo-physomyces elegans* (order Mucorales) is a thermotolerant species that causes severe infections among humans. In contrast to other fungi that cause zy-gomycosis, which have a worldwide distribution and are rarely found in immunocompetent hosts, *A. elegans* has been reported mainly in areas with warm climates as an emerging pathogen that causes mostly cutaneous infections after injury to the skin (*1*). This fungus was discovered in 1979 (*2*) and until recently was considered the only species in the genus.

A polyphasic study of clinical and environmental strains of A. elegans, including analysis of several genes, showed that the genus contained 4 well-characterized species (3). Of 16 isolates tested in this study, only 2 from soil in India were A. elegans. Most of the isolates were A. variabi*lis.* The incidence of *A. variabilis* in humans is unknown and difficult to ascertain because most cases had isolates that were not properly preserved. These fungi usually cause necrotizing fasciitis, but rhino-orbito-cerebral or renal infections have also been reported (1). Whether these infections are produced by different Apophysomyces spp., have different responses to antifungal drugs, or have differences in virulence is unknown.

To assess incidence of *Apophy-somyces* spp. in a tertiary hospital (Government Medical College Hospital, Chandigarh, India), which usually receives patients with zygomycosis, a retrospective study was conducted during November 2001–April 2009. Nine patients were identified as having primary cutaneous zygomycosis. For 4 patients, fungal isolates were morphologically identified as *A. elegans*. A description of clinical findings, their management, and outcomes for these

9 patients has been reported (4). The 4 isolates were sent to the Universitat Rovira i Virgili (Reus, Spain) for molecular analysis.

The internal transcribed spacer region of these isolates was sequenced and compared with those of type strains of *Apophysomyces* spp. Fungi were identified by morphologic (Figure, panel A) and molecular analysis as *A. variabilis* (99.6%–99.7% sequence identity with sequence of type strain CBS 658.93 [FN556436]). GenBank accession nos. of the 4 isolates are FN813491, FN813490, FN556442, and FN813492.

Another patient was also infected with *A. variabilis* fungi. The patient was a 45-year-old woman with diabetes from Derabassi (Punjab), India, who was hospitalized because of swelling in her right breast and blackening of overlying skin. A diagnosis of right breast gangrene was made. Therefore, local debridement of the swelling was conducted, and tissue samples were tested by microbiologic culture and histopathologic analysis.

A KOH wet mount showed broad aseptate hyphae with right-angled branching. The fungal isolate was tentatively identified as A. elegans. Histopathologic analysis confirmed a diagnosis of zygomycosis. The patient was treated under local anesthesia by debridement of infected tissue and some of the healthy surrounding tissue (Figure, panel B). However, an antifungal regimen could not be given because she had disturbed renal function. Her condition deteriorated, septicemia was observed, and she died from sudden cardiac arrest on the sixth day after admission. The fungal isolate was also identified as A. variabilis (98.9% identity, GenBank accession no. FN556443).

Although most cases of infection with *A. variabilis* fungi have been reported in India (5), infections with this fungus may have a wider distribution. A recent study demonstrated that this species represented 0.5% of fungi