Dengue Virus, Nepal

To the Editor: Dengue virus belongs to the genus Flavivirus, family Flaviviridae. It has 4 serotypes: dengue virus type 1 (DENV-1), dengue virus type 2 (DENV-2), dengue virus type 3 (DENV-3), and dengue virus type 4 (DENV-4). Dengue virus is maintained in a cycle between humans and Aedes aegypti, domestic day-biting mosquitoes. Dengue virus induces clinical illness, which ranges from a nonspecific viral syndrome (dengue fever [DF]) to severe and fatal hemorrhagic disease (dengue hemorrhagic fever [DHF]). DF/DHF occurs primarily in tropical and subtropical areas of the world. Domestic dengue virus infection occurs in >100 countries; >2.5 billion persons live in these areas. Approximately 100 million cases of DF, 500,000 cases of DHF, and several thousand deaths occur annually worldwide (1). During the past decades, dengue virus has emerged in southern Asia; DF/DHF epidemics have occurred in Bhutan, India, Maldives, Bangladesh, and Pakistan (2–4).

From August through November 2006, the number of febrile patients increased in 4 major hospitals in the Terai region of Nepal: Nepalgunj Medical College, Bheri Zonal Hospital in Nepalgunj, Tribhuvan Hospital in Dang, and Narayani subregional hospital in Birgunj. Patients with severe symptoms were referred to Sukraraj Tropical and Infectious Disease Hospital, Kathmandu, for diagnosis and treatment. The clinical features in most patients were consistent with signs of DF, but some patients showed signs (high fever, rash, ecchymosis, epistaxis, positive tourniquet test, liver dysfunction, and thrombocytopenia [platelet count <100,000/mm³]) consistent with the World Health Organization (WHO) definition of DHF. Ascites and plural effusion developed in 2 patients. Blood specimens were collected from all patients at the time of admission to the local hospitals. Particle agglutination (PA) assay (Pentax Ltd, Tokyo, Japan) (5) and immunoglobulin (Ig) M–capture ELISA (Dengue/JE IgM Combo ELISA kit, Panbio Ltd, Brisbane, Queensland, Australia) were performed. Dengue virus–specific IgM was detected in 11 patients who had fever, headache, and rash (Table). Each of these patients had negative

Table. Clinical and laboratory data for 11 patients admitted to hospitals and diagnosed with dengue fever or dengue hemorrhagic fever, Nepal, 2006*

<table>
<thead>
<tr>
<th>Patient age, y/Sex</th>
<th>Month admitted</th>
<th>Location</th>
<th>Initial diagnosis</th>
<th>Travel history</th>
<th>Clinical signs and symptoms</th>
<th>Selected laboratory and other test results</th>
</tr>
</thead>
<tbody>
<tr>
<td>20/M</td>
<td>Sep</td>
<td>Kathmandu</td>
<td>DF</td>
<td>Yes</td>
<td>Fever, headache, nausea</td>
<td>Hb 15.4 g/dL; TLC 10,500/mm³; Pit 185,000/mm³; blood culture for salmonellae negative; ALT 38 IU/L</td>
</tr>
<tr>
<td>27/F</td>
<td>Sep</td>
<td>Bardiya</td>
<td>Viral fever</td>
<td>No</td>
<td>Fever, headache, vomiting</td>
<td>TLC 5,600/mm³; blood culture for salmonellae negative</td>
</tr>
<tr>
<td>3/M</td>
<td>Sep</td>
<td>Salayan</td>
<td>Encephalitis</td>
<td>No</td>
<td>Fever, vomiting, convulsions</td>
<td>Widal negative; TLC 4,700/mm³</td>
</tr>
<tr>
<td>13/M</td>
<td>Oct</td>
<td>Sindhuli</td>
<td>Typhoid fever</td>
<td>No</td>
<td>Fever, headache</td>
<td>Widal negative; TLC 4,500/mm³; blood culture for salmonellae negative; Brucella antigen negative; chest radiograph normal</td>
</tr>
<tr>
<td>22/M</td>
<td>Oct</td>
<td>Birgunj</td>
<td>DHF</td>
<td>No</td>
<td>Fever, headache, vomiting, ascites</td>
<td>Bil 0.8 mg/dL; ALT 80 IU/L; Pit 22,000/mm³; chest radiograph normal</td>
</tr>
<tr>
<td>55/F</td>
<td>Oct</td>
<td>Dang</td>
<td>DF</td>
<td>No</td>
<td>Fever, headache, muscular pain</td>
<td>Pit 51,000/mm³; TLC 7,600/mm³; MP negative; ESR 20 mm/h; Bil 0.7 mg/dL Brucella negative; Widal negative; TLC 5,600/mm³</td>
</tr>
<tr>
<td>22/F</td>
<td>Oct</td>
<td>Birgunj</td>
<td>Viral fever</td>
<td>No</td>
<td>Fever, headache, body ache</td>
<td>Pit 95,000/mm³; TLC 4,700/mm³; Hb 13.1 g/L; Bil 0.8 mg/dL; ALT 26 IU/L</td>
</tr>
<tr>
<td>13/M</td>
<td>Nov</td>
<td>Dang</td>
<td>DF</td>
<td>No</td>
<td>Fever, headache, rashes</td>
<td>Bil 0.81 mg/dL; Pit 31,000/mm³; PT 2 min 30 s (control 14)</td>
</tr>
<tr>
<td>35/F</td>
<td>Nov</td>
<td>Birgunj</td>
<td>DHF</td>
<td>No</td>
<td>Fever, headache, bruises, tourniquet: positive</td>
<td>ALT 127 IU/L; Pit 110,000 /mm³; PCV 38.8%; TLC 5,500/mm³; ultrasonography liver size, 16.8 cm</td>
</tr>
<tr>
<td>40/M</td>
<td>Nov</td>
<td>Birgunj</td>
<td>DF</td>
<td>No</td>
<td>Fever, headache, rashes</td>
<td>Bil 0.7 mg/dL; Widal test negative; TLC 6,800/mm³; Pit 164,000/mm³</td>
</tr>
<tr>
<td>42/M</td>
<td>Nov</td>
<td>Dang</td>
<td>DF</td>
<td>No</td>
<td>Fever, headache, rashes</td>
<td></td>
</tr>
</tbody>
</table>

*Blood specimens were collected at time of hospital admission. Diagnosis was confirmed by using immunoglobulin M–capture ELISA. DF, dengue fever; Hb, hemoglobin; TLC, total leukocyte count; Pit, platelets; ALT, alanine aminotransferase; DHF, dengue hemorrhagic fever; Bil, bilirubin; MP, malaria parasites; ESR, erythrocyte sedimentation rate; PT, prothrombin time; PCV, packed cell volume.
results for Japanese encephalitis virus—specific IgM. Of the 11 patients, 10 had no history of travel to India or other dengue-endemic countries. DF or DHF was initially diagnosed for 7 patients, and viral encephalitis, typhoid fever, or viral fever was diagnosed for others without serologic tests. Reverse transcription–PCR and virus isolation were performed at Institute of Tropical Medicine, Nagasaki University, Nagasaki, Japan, but the dengue virus genome was not detected, and no virus was isolated, likely because sample collection was delayed and the sample was transported to Japan in a deteriorated condition.

DF/DHF have been considered to be a possible public health threat to Nepal because DF/DHF epidemics have occurred recently in India and Pakistan, which reported several thousand cases and >100 deaths (6). The first DF case in Nepal was reported in 2004 (7). Further, the first DENV-2 strain of Nepal origin was isolated from a Japanese traveler who visited Nepal and in which DF developed after the patient returned to Japan. The isolated DENV-2 (GenBank accession no. AB194882) was 98% homologous with DENV-2 isolated in India (8). The prevalence of dengue virus antibody was reported to be 10.4% in the southwestern region of Nepal (9). These reports suggest that dengue virus has been circulating in Nepal for several years. Thus, DF/DHF has likely been misdiagnosed and illness caused by dengue virus underestimated in Nepal. In contrast, Japanese encephalitis has been a public health problem in southwestern region of Nepal, and large epidemics have occurred almost every year since 1978 (10). Nepal has no dengue surveillance programs, and health professionals do not usually consider dengue as a differential diagnosis.

The emergence occurred in the lowland Terai belt region, which borders the state of Bihar, India. The Aedes mosquito is known to persist in this region. The emerging DENV-2 is likely to have been introduced into Nepal from India.

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Human Tuberculosis Caused by Mycobacterium bovis, Taiwan

To the Editor: Mycobacterium bovis is one of the causative agents of tuberculosis (TB) in humans and animals. Drinking unpasteurized milk, eating undercooked meat, and close contact with infected animals are the main sources of infection for humans. Currently, 119 M. bovis spoligotypes are contained in the fourth international spoligotyping database (SpolDB 4) and are categorized into 3 main sublineages corresponding to ST prototypes 482, 683, and 479 (1).

Although an M. bovis surveillance program for farm animals has been implemented by the Taiwan Council of Agriculture, no surveillance system exists for human TB cases caused by M. bovis. To monitor the epidemiology of M. bovis in domestic animals, a regular tuberculin skin test (TST) is compulsory for cattle and sheep and optional for deer in Taiwan (2).

In 2005, screening of Mycobacterium spp. infections by TST was performed for 111,412 cattle and 73,396 caprines and ovine herds, of which 188 (0.17%) and 148 (0.2%), respectively, were positive (2). We used spacer oligonucleotide typing (spoligotyping) and