Globalization of Infectious Diseases: The Impact of Migration

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With up to 2% of the world’s population living outside of their country of birth, the potential impact of population mobility on health and on use of health services of migrant host nations is increasing in its importance. The drivers of mobility, the process of the international movement, and the back-and-forth transitioning between differential risk environments has significance for the management of infectious diseases in migrant receiving areas. The management issues are broad, high-level, and cross-cutting, including policy decisions on managing the migration process for skilled-labor requirements, population demographic and biometric characteristics, and family reunification; to program issues encompassing health care professional education, training, and maintenance of competence; communication of global events of public health significance; development of management guidelines, particularly for nonendemic diseases; access to diagnostic and therapeutic interventions for exotic or rare clinical presentations; and monitoring of health service use and health outcomes in both the migrant and local populations.

Current estimates are that ∼2% of the world’s population resides in a nation different than the one in which they were born. Although small in terms of percentages, the total number of the foreign-born individuals living abroad is significant. Worldwide, the number is >200 million people, which is equivalent to the size of many large nations. These mobile, migrant populations are composed of several groups, including immigrants, migrant and seasonal workers, refugees, asylum seekers, international students, and others. Table 1 lists the numerical impacts of immigration on the major immigrant-receiving developed nations.

Historical patterns of immigration, seasonal work, and population flows coupled with the presence of national policies that encourage or support immigration as a source of population growth combine to ensure that the global distribution of these migrant populations is not uniform. Foreign-born residents make up significantly different percentages of national populations. Long-standing immigration policies in nations such as Australia, Canada, Israel, and the United States produce diverse national populations that reflect both recent and longer-term population flows, which in turn have an impact on health outcomes [6]. Other nations, particularly those in Europe, have foreign-born populations that are reflective of refugee movements, asylum seeking, or temporary worker flows.

In each of these cases, the health characteristics of the foreign-born or migrant cohorts of the populations are influenced by the health environments and situations at their place of original residence, the environments through which they made the transition, and their new destination [7]. For many migrants, the transition period is inconsequential in terms of health because of the short duration of transportation. For other migrants, such as those transitioning through refugee camps [8] or those experiencing trafficking or smuggling [9] as a means of arrival, the transition period can greatly affect health. If the health and disease parameters that influence the prevalence of infections at a migrant’s point of origin and transition periods are different from those at the destination, the process of migration and travel can bridge the difference in disease prevalence and function as a method of transfer between regions.

The population-based downstream health impact and consequences of the epidemiological bridging produced by migration is directly related to 2 basic factors. The first is the degree of difference between origin and the destination [10], and the
second is the size of the mobile population that moves between the different disease prevalence patterns. In 2002, >700 million international arrivals were recorded [11]. Specific individual health status and outcomes are influenced by several other factors, such as the amount time spent in the location, complex social and economic factors, access and availability to affordable health care, educational attainments, and cultural health practices.

In terms of infectious disease epidemiology, the association between the introduction of disease and migration has long been recognized. Border health and quarantine medicine practices were developed out of attempts to control the importation of epidemic infections. Over the past 40 years, both the process and scope of migration have undergone several major shifts, and many of those changes have altered the nature of migration-associated infectious disease.

Since the 1960s, the immigration patterns in many immigration-receiving patterns have changed. Regions supplying immigrants to Australia, Canada, and the United States, which were historically western and central European population movements, have predominately shifted to Latin American, African, and Asian sources. Source regions are now more likely to encompass less developed regions of the world, many in tropical and subtropical environments. During the same period, the conditions determining the overall health of the population—including institutional health services and public health control capacities in many immigrant source areas—have experienced significant detrimental challenges. Developed regions of the world have continued to make great strides in the control of infectious diseases of significance to public health. Many infections in the more developed regions are at historical levels of low prevalence or elimination of endemic transmission.

In most immigrant source regions, infectious diseases continue to represent major causes of death and morbidity due to respiratory infections, gastrointestinal and diarrheal diseases [12], tuberculosis [13], and HIV infection/AIDS [14]. The risks of infectious disease acquisition in those source nations are manifest in certain immigrant and migrant populations after they move to their new destination. As migration continues in an increasingly globalized world, health care providers at the primary care and specialist level can expect to be faced with the challenges of recognition, diagnosis, and management of diseases that are themselves the consequences of international factors.

THE INCREASING IMPORTANCE OF MIGRATION IN NATIONAL INFECTIOUS DISEASE EPIDEMIOLOGY

During the past 50 years, many national and some international disease control, immunization, and communicable disease prevention strategies have been successful. Incidence and prevalence rates of many previously serious infections have decreased dramatically. In several developed nations, local transmission of vaccine-preventable diseases has practically ceased. However, these successes have not been uniform, and infectious diseases continue to be major causes of mortality and morbidity in some regions of the world.

Immigrants and refugees originating from areas where infections persist can pose a significant challenge for national disease control and or elimination strategies. The factors that support and sustain the prevalence of diseases in the less developed world are beyond the control of national and state/provincial/municipal public health programs at the migrant’s ultimate destination. As a consequence, the continued disparities in prevalence levels of infectious diseases in a world of increasing travel and migration makes national disease control or elimination almost impossible. These same factors also increase the epidemiological likelihood that many infections that are now very well controlled in the developed world will be increasingly observed in migrant or foreign-born populations resident in host destinations.

Tuberculosis is an example of the situation described above. Beginning in the early part of the 20th century, extensive tuberculosis-control programs, coupled with the development of effective pharmacotherapy, resulted in the marked reduction of the prevalence of the disease in North America, Australia, and many parts of Europe. Over the past 50 years, tuberculosis in

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Table 1. Visible minority, ethnic, and foreign-born populations, by total population, for selected nations.

<table>
<thead>
<tr>
<th>Nation</th>
<th>Total population</th>
<th>Visible minority/ethnic population</th>
<th>Proportion of foreign born persons, %</th>
<th>Source</th>
</tr>
</thead>
<tbody>
<tr>
<td>United States</td>
<td>284,796,887</td>
<td>7,091,442 (24.9%)</td>
<td>11.1</td>
<td>US Census Bureau [1]</td>
</tr>
<tr>
<td>Australia</td>
<td>19,996,058</td>
<td>...*</td>
<td>25</td>
<td>Australia Bureau of Statistics [2]</td>
</tr>
<tr>
<td>Canada</td>
<td>29,639,035</td>
<td>3,983,845 (13.4%)</td>
<td>18.4</td>
<td>Statistics Canada [3]</td>
</tr>
<tr>
<td>Israel</td>
<td>6,500,000</td>
<td>1,500,000 (Arabs, 23%)</td>
<td>42.2 [4]</td>
<td>Government of Israel—Ministry of Immigrant Absorption [5]</td>
</tr>
</tbody>
</table>

* Calculation based on the arrival of 1 immigrant every 3 minutes and 51 s gives an estimate of 136,819 foreign-born persons arriving per year.
Table 2. Selected imported infectious diseases illustrating high to low prevalence impacts.

<table>
<thead>
<tr>
<th>Disease or syndrome</th>
<th>High-prevalence regions</th>
<th>Impacts in low-prevalence zones</th>
<th>Interventions</th>
</tr>
</thead>
<tbody>
<tr>
<td>Tuberculosis</td>
<td>Africa, Asia</td>
<td>Acute infections in migrants; secondary, local transmission; multiple-drug resistance patterns</td>
<td>Delay in regional control or elimination program goals</td>
</tr>
<tr>
<td>Severe acute respiratory syndrome</td>
<td>East and Southeast Asia</td>
<td>Outbreaks of atypical pneumonia and unexplained death</td>
<td>Realignment of public health prevention and diagnosis resources; diminished resources for other programs</td>
</tr>
<tr>
<td>Monkeypox</td>
<td>Sub-Saharan Africa</td>
<td>Outbreaks of animal to human transmission with vesicular, febrile illness; potential for death</td>
<td>Potential for introduction in susceptible wild animal reservoirs; establish zoonotic-human cycles</td>
</tr>
<tr>
<td>Leishmaniasis—viscerotropic and visceral</td>
<td>Middle East Asia, South Asia, Africa</td>
<td>Acute infections in migrants</td>
<td>Clinical, diagnostic service impacts; transfusion donor and product screening</td>
</tr>
<tr>
<td>Malaria—Plasmodium falciparum</td>
<td>Sub-Saharan Africa</td>
<td>Acute morbidity and mortality in migrants; potential for secondary transmission by mosquitoes or blood products</td>
<td>Introduction of temporary or sustained transmission outside existing endemic regions</td>
</tr>
<tr>
<td>Strongyloidiasis</td>
<td>Global tropical zones: Africa, Southeast Asia</td>
<td>Acute dissemination syndromes; chronic eosinophilia syndromes with or without pulmonary involvement; chronic urticaria</td>
<td>Late-onset dissemination syndromes</td>
</tr>
<tr>
<td>West Nile virus–associated syndromes</td>
<td>Middle East Asia, North Africa</td>
<td>Acute meningoencephalitis in immune naive population; established zoonotic cycles</td>
<td>Shifting in zoonotic host populations and densities; secondary environmental impacts; naive population herd immune response</td>
</tr>
</tbody>
</table>
of transfusion-related trypanosomiasis, historical screening of
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is American trypanosomiasis (Chagas disease), which is rarely
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grams developed in the 1980s and 1990s and the burden of
disease control programs and outcomes continue or expand,
impact of migration on infectious disease epidemiology in
low-prevalence regions of the world, including most of
North America, Australia, and Europe, will likewise increase.
Current examples to illustrate this include malaria, which has
been historically diagnosed in returning visitors and tourists
from zones of endemicity to zones of nonendemicity, or linked
to returning migrants or rare examples of transmission outside
of zones of endemicity [20]; Hansen disease (leprosy), a con-
dition in which migrants and foreign-born populations make up
the majority of national case loads in immigration-receiving
nations such as Canada [21]; and HIV infection/AIDS, asso-
ciated with a profound global disparity in its prevalence and
occurrence, combined with growing immigration and popu-
lation mobility patterns, resulting in high rates of HIV in-
fection in foreign-born populations. As HIV/AIDS prevention pro-
grams developed in the 1980s and 1990s and the burden of
disease fell to the developed world, new cases will be increas-
ingly prevalent in some migrant populations. Another example
is American trypanosomiasis (Chagas disease), which is rarely
imported from areas of endemicity, but high levels of immi-
gation from areas of the Western Hemisphere where this par-
asitic disease is endemic have resulted in national policy de-
cisions on transfusion blood product screening [22] and
donation practices [23] in migration-receiving nations (for ex-
ample, where blood product screening or pretreatment [24]
may not be effective approaches to reducing population risk
of transfusion-related trypanosomiasis, historical screening of
potential donors and establishing donor ineligibility criteria
may be done [25]).
Other examples include vaccine-preventable diseases. Mi-
gration and refugee movements frequently deliver individuals
from areas of the world where immunization programs may
less comprehensive to destinations where immunization has
proved very effective at disease control or elimination. As a
consequence, migrants and other mobile populations are in-
creasingly related to individual cases or small outbreaks of im-
ported vaccine-preventable disease with local transmission to
the developed world. Specific examples include polio [26], he-
patitis A [27] and hepatitis B [28], mumps [29], measles [30,
31], pertussis [32, 33], diphtheria [34], and rubella [35, 36]
(table 2).
The current and future influence of migration on the epi-
demiology of infection diseases in the developed world is not
limited to infections for which there are current or historic
control strategies. The nature of modern migration is that it is
a dynamic process, and immigrant source nations and regions
evolve in response to a complex series of political, social, and
international factors. Conflict, political upheaval, and social
unrest traditionally initiate the movements of refugees and
other displaced individuals. These unorganized movements are
often followed by subsequent generations of migrants in or-
ganized and regular migration movements. These sustained
population flows are associated with family reunification move-
ments and nonfamilial but intracultural linkages that support
secondary arrivals.
Just as the refugee movements from Southeast Asia and Af-
rica starting in the 1960s brought new infectious disease chal-
enges in terms of malaria, hepatitis B, and Hansen disease to
the host nations, more recent movements from the Balkans and
Middle East can be expected to result in future infectious dis-
ese events. Common infections, such as hantavirus infection
in the Balkans [37] or leishmaniasis in the Middle East [38],
can be expected to be represented in migrant communities
originating from those areas. As the populations of new arrivals
from these regions increases over the next decade, the incidence
of infectious diseases common to those areas will correspond-
ingly increase in destination countries related to primary, per-
manent movements or secondary movements associated with
temporary stays [39].

CURRENT AND FUTURE IMPLICATIONS
FOR THE MANAGEMENT OF INFECTIOUS
DISEASES IN MIGRATION-RECEIVING NATIONS

The impact of migration-associated changes in global infectious
disease epidemiology will have immediate and direct impli-
cations for those who provide clinical infectious disease and lab-
atory diagnostic services in ways not traditionally considered.
The first is an increased need for international public health information. Effectively managing infectious diseases in situations where many of the influences originate beyond national borders will require an evolving reference base of accurate, verifiable baseline and outbreak data linked to dynamic population health determinants, such as demography, biometrics, and mobility, which are also influenced by health services availability, access, and affordability; and socioeconomic, biological and genetic, environmental, and behavioral factors.

In combination, these factors have immediate and pressing consequences for expanding educational programs at all levels of clinical and administrative health services. As national populations become more diverse, infectious disease physicians will need knowledge and information previously associated with international public health providers. For example, information regarding the extensive transmission of hepatitis C during schistosomiasis control programs in parts of northern Africa [40] has been important to those working in public health in the region. However, that same knowledge has important implications for those practitioners involved in the care of migrants after their relocation.

The long latency phase of some infections ensures that the majority of the chronic disease sequelae and complications will only present clinically long after arrival. The long-term consequences of infections acquired before immigration occurs will become more commonly encountered by health care practitioners. Examples include strongyloidiasis [41], for which it is necessary to appreciate the risks of antecedent infections when foreign-born individuals [42, 43]—even those who have resided at their new home for many years—are treated with agents that suppress their immunity, such as cancer chemotherapy and systemic corticosteroids. Incidents of chemotherapy-induced reactivation of latent infections are not infrequent in patients of this type, and clinicians will increasingly need to be aware of the global patterns of disease to anticipate and recognize what historically have been clinically rare events [44].

Both the quantity and quality of the international public health information required by clinical and diagnostic practitioners in low-prevalence zones will be required to change. The past movements and population demographic shifts with current and the projected movements of migrants need to be linked with the evolving international epidemiology of both chronic prevalence gap conditions (e.g., tuberculosis, syphilis, HIV/AIDS, and strongyloidiasis) and new, emerging conditions of public health significance (e.g., hantavirus, West Nile virus, severe acute respiratory syndrome, monkeypox, avian influenza). These emerging conditions will continue to pose particular challenges to the existing communication and educational frameworks.

Second, there will be an increased need to understand the immigration and migration process. As the pattern of exotic imported infections becomes more commonly associated with the migration process and migrants on the move become significant risk groups for diseases of otherwise low incidence in immigration-receiving nations, clinicians will need better understanding of the variations in the migration process itself as a risk determinative factor. Traditionally, the foreign-born have been associated into one group for epidemiological purposes. Like all complex populations, there are profound differences in foreign-born migrant populations, such as immigrants, refugees, asylum seekers, and migrant workers, that affect health determination and outcomes. The unifying factor may be foreign birth and travel to a new destination, but beneath these characteristics are several components that affect the clinical management of infectious diseases. The awareness, appreciation, and clinical expertise related to the importance of those factors are currently limited to those clinicians who have substantial work experience in the area of immigration health. As the foreign-born populations expand, that knowledge will become increasingly important for all clinicians and for the services that support diagnostic and therapeutic interventions.

The management of illness and disease in foreign-born populations can be affected by the status of the individual themselves. Illegal or irregular migrants may not present for care, attend follow-up assessments, or fill prescriptions out of fear of resulting immigration enforcement actions. This can affect compliance in cases of long-term treatment and may delay initial presentation, complicating public health follow-up for communicable diseases.

Practitioners not accustomed to dealing with migrants may not be aware of the extent, intent, or outcome of routine immigration screening relevant to infectious diseases. Routine immigration screening practiced by major immigration-screening nations is limited to a small number of infections and usually only to those conditions that could pose a threat to public health through transmission or create an excessive resource burden on existing health services.

**SUMMARY**

The increasing foreign-born component in the populations of immigration-receiving regions, including the United States, Australia, Canada, and western Europe, coupled with the sustained prevalence of infectious diseases in immigration-source nations, will have a growing effect on the nature and practice of clinical infectious diseases. Global factors beyond the influence of national infectious disease control strategies will affect national disease-control programs. Migrants will become increasingly important risk groups and index cases for infections of low incidence in the immigration-receiving nations. As the demography of immigration source nations evolves, classical infectious disease concerns in migrants may become less rel-
evant [45], and new challenges can be expected as immigrants and refugees arrive from new regions of the world with different background infectious disease epidemiology.

For many infections, particularly those that have been reduced to very low prevalence or eliminated in some nations, management and control will increasingly become a process of dealing with global rather than local or national epidemiology. More and more clinical infectious disease practitioners will be confronted by situations that represent the consequences of existing infection-control practices and public health systems that operate beyond their own borders. Meeting these challenges will require the acquisition and sharing of information more commonly associated with international public health. As the process of globalization continues to reduce geographical limits to disease, wider appreciation of the importance of global and international factors will be required.

Strategies to meet those challenges include increasing the networking of clinicians; specific migration-focused surveillance, detection, and interpretation systems similar to those used for disease outbreaks among international travelers; and the rapid availability and accessibility to public health and infectious disease information relevant to migrants and mobile populations, including diagnostic, clinical evaluation, management, and control expertise.

What is becoming increasingly evident is that globalization and migration have ensured that infectious disease “hoof beats” in migrant populations, and the host populations within the transmission envelope of the disease may actually be the zebras that many clinicians would only consider after more common causes of the presentations were ruled out or would not consider at all. Education and training in the use of the tools of global epidemiology in infectious diseases requires a commitment from clinicians, educational institutions, health care facilities, national and international authorities in migration [46], and health to act on this emerging area of impact on international public health as a result of globalization.

Acknowledgments

Conflict of interest. All authors: no conflict.

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