Letter to the Editor

Sir.

We refer to our paper entitled Epidemiological study in a new focus of cutaneous leishmaniasis in the Islamic Republic of Iran published in the Eastern Mediterranean Health Journal, 2003, Vol. 9 No. 4, pages 816–826.

In this paper we reported that in Jovein district, Sabzevar county in the north-east of the Islamic Republic of Iran, 7 of 45 (15.6%) *R. opimus* were found to be infected with *Leishmania* and these isolates were identified as *L. major*. These were identified by Professor K.P. Chang (University of Health Science/Chicago Medical School, United Sates of America) using a restriction fragment length polymorphism (RFLP)–PCR technique with *NAGT* gene.

However, subsequent to publication of the above-mentioned paper, characterization was made of one further isolate from an eighth case of *Rhombomys opimus* found to be infected with *Leishmaia* in the same area. This isolate was identified as *L. turanica*.

Thus of a total of 8 of 45 (17.8%) R. opimus found to be infected with Leishmania, 7 were infected with L major and 1 with L. turanica.

It has, until recently, been assumed that Leishmania parasites circulating in the R. opimus population in the Islamic Republic of Iran are L. major. This finding is the first report of the isolation and characterization of L. turanica from R. opimus from an area where there are many active colonies of the great gerbils. It is most probable that this is a continuation of the distribution of L. turanica in Turkmenistan which has now extended to the Islamic Republic of Iran.

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