ERRATUM

Erratum to: Spatially explicit uncertainty modeling of zoonotic pathogen distribution: a case of *Listeria monocytogenes* in New York State, USA

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The original version of this article unfortunately contained mistakes.

The *Listeria monocytogenes* study used as input for modeling in this article was obtained from Cornell University. We did not provide acknowledgment of the source and we neglected to cite a previous paper that describes some analysis of these data. We apologize for this oversight and publish the following correction to our Materials and Methods, Acknowledgments, and References section.

Materials and methods (page 3; first paragraph)

The initial survey for *Listeria monocytogenes* and other *Listeria* spp. was conducted as described in Chapin *et al.* (2014). We replicated the random forest analysis of environmental distribution by independently obtaining and processing spatial and meteorological GIS predictor data (Table 2). The results obtained in this article using the mean decrease in Gini index criterion were comparable to those in Fig. 2A of Chapin *et al.* (2014). We used the replicated random

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forest analysis results for the presented habitat suitability and spatial uncertainty modeling reported in this article.

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References

- Chapin TK, Nightingale KK, Worobo RW, Wiedmann M, Strawn LK (2014) Geographical and meteorological factors associated with isolation of *Listeria* species in New York State produce production and natural environments. *J Food Prot* 77:1919–1928