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Author Correction: Unrevealed genetic diversity of GII Norovirus in the swine population of North East Italy

L. Cavicchio, L. Tassoni, A. Laconi, G. Cunial, L. Gagliazzo, A. Milani, M. Campalto, G. Di Martino, M. Forzan, I. Monne & M. S. Beato

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This Article contains a formatting error in Figure 1 which makes the key unreadable. The correct Figure 1 appears below.

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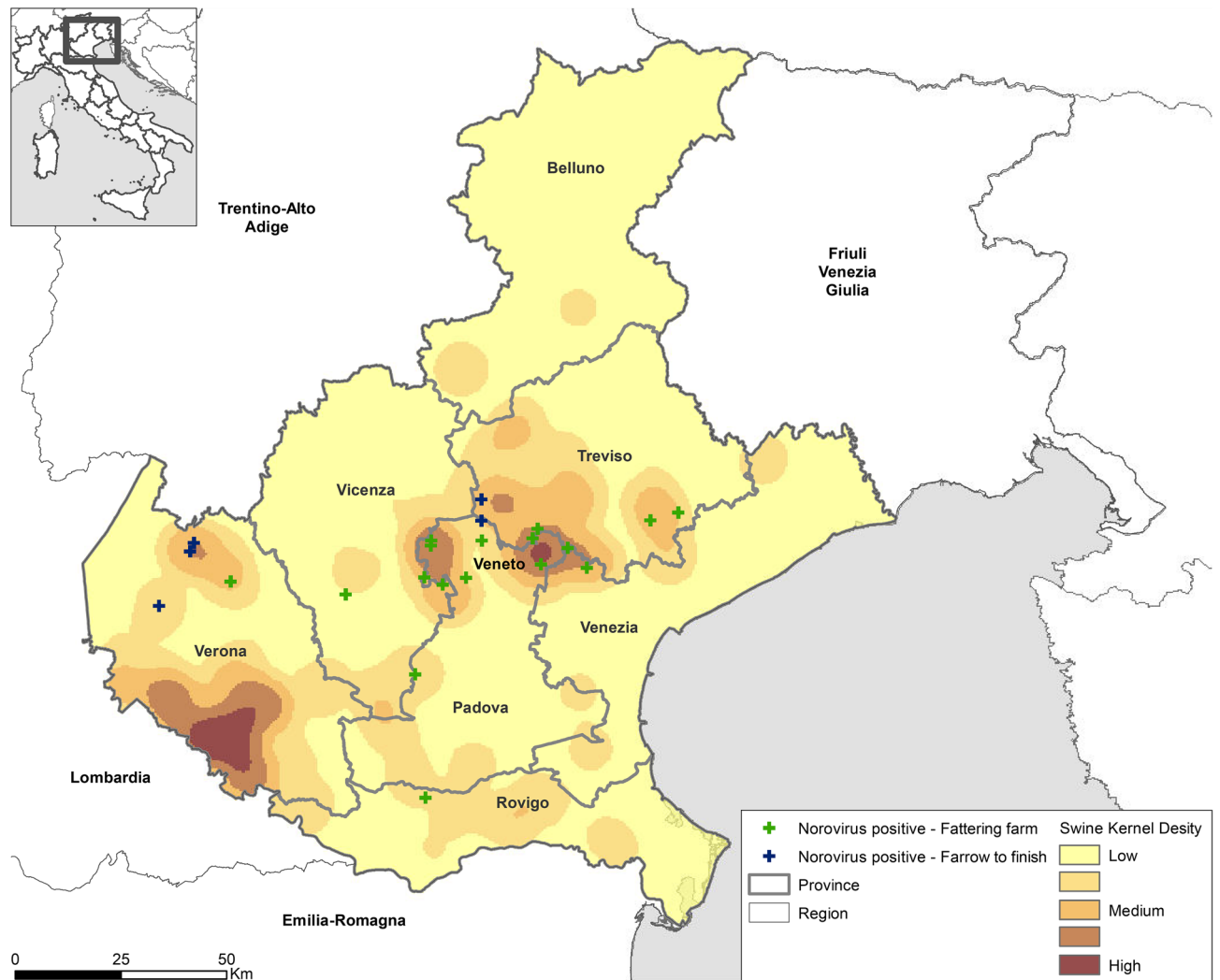


Figure 1. Density map (Kernel Density) of Veneto swine farms and distribution of NoV positive swine farms: 2018–2019. Green crosses identify the NoV positive fattening farms, dark blue crosses the NoV positive farrow to finish farms. The Kernel Density tool calculates the density of features in a neighbourhood around those features. The input data was the industrial swine farms in Veneto region and the density was calculated considering the potential capacity of each farm. The parameter used to calculate the density was a radius of 500 m and the raster cell of the analysis was 10 km².



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