Validation of a urine metabolome fingerprint in dog for phenotypic classification

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In the paper entitled: “Validation of a urine metabolome fingerprint in dog for phenotypic classification” published in the December 2007 issue; 3(4): 453–463, the authors erroneously stated that the dogs were all neutered. Records

have shown that two of the dogs were entire (MS2 & MS3, samples 23–32). This error has made no impact on the interpretation of the data and re-analysis of the data provides no more meaningful interpretation. PCA plots indicate that these two dogs are not outliers, which suggests that “sex status” is not a major driver of variance in this study.

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