Dear Editor.

While in the process of submitting our study “Peptide clustering enhances large-scale analyses and reveals proteolytic signatures in mass spectrometry data” to *Nature Communications*, our corresponding author for that manuscript, Erik Hartman, was invited by Pratibha Kantekar to submit an accompanying data descriptor. We see this as an excellent opportunity to increase the visibility and accessibility of our data and are hereby submitting our data descriptor “Mass spectrometry peptidomics data from infected and uninfected porcine wounds” for your consideration.

I have no suggestions for people to include in the review process, nor any I wish to exclude.

This data descriptor has not been published or sent for consideration elsewhere. However, the related manuscript “Peptide clustering enhances large-scale analyses and reveals proteolytic signatures in mass spectrometry data” is now published in *Nature Communications*.

Thank you for your time and your consideration of this submission.

Best regards,

Fredrik Forsberg

Faculty of Medicine, Lund University