

A systematic error appeared in our paper title “T Cell Responses Affected by Aminopeptidase N (CD13)-mediated Trimming of Major Histocompatibility Complex Class II-bound Peptides” (Larsen et al. July 1996, 184:183–189). During the computer conversion the symbol micro was erroneously replaced by an m. This affects measures of length and concentration. Thus, all measures in m should read micro except for the protease inhibitors PMSF, EDTA, and 1,10 phenanthroline, which are in millimolar; and the washing buffer for the fixed cells, where the L-glycine is also in millimolar.

To reiterate: (a) The beads used were 5.12 micrometer in diameter; (b) the concentrations of antibodies and peptides should all be in micromolar; (c) the concentrations of protease inhibitors were: PMSF 1 millimolar; EDTA 8 millimolar; 1,10 phenanthroline 1.2 millimolar; pepstatin A 69 micromolar; Na-*p*-tosyl-L-lysine chloromethyl ketone (TLCK) 128 micromolar; Na-*p*-tosyl-L-phenylalanine chloromethyl ketone (TLPK) 135 micromolar; and N-ethylmaleimide (NEM) 1 millimolar.

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