

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 milh should read micro except for the protease inhibitors PMSF, EDTA, and 1.10 phenantroline, which are in millimolar; and the washing bugger 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 phenantroline 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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