

Sympson et al. Vol. 125, No. 3, May 1994. Pages 681–693.

The following is a correction for the above-mentioned article.

We have found by sequence analysis that, owing to an inadvertent error, the mutated rat stromelysin-1 cDNA sequence designated M2 (Pro⁹³ → Val⁹³) was in fact the same as the sequence M1 (Pro⁹² → Gly⁹²). Therefore, all the transgenic mice designated as M2 actually have the M1 autoactivating mutation. Since the phenotypes of both sets of M1 and M2 transgenic animals are the same, this error does not alter the results or interpretation of the data.