Corrigendum: Biased signaling of protease-activated receptors

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Figure 3 shows a snake diagram of the N-terminal amino acid sequence of human protease-activated receptor (PAR2). It indicates the sites at which different proteases cleave PAR2. These sites are correct. The

error is that we accidentally repeated a sequence of amino acid residues (11–16, GAAILL) in positions 17–22 of the snake diagram. We wish to replace the snake diagram with the correct version.

This minor correction does not affect our discussion or conclusions.

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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