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Addition/Correction

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Correction to Effects of the Oncogenic $V_{664}E$ Mutation on Membrane Insertion, Structure, and Sequence-Dependent Interactions of the Neu Transmembrane Domain in Micelles and Model Membranes: An Integrated Biophysical and Simulation Study

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Biochemistry, **2012**, *51*, 12, 2558−2568 DOI: 10.1021/bi201269w

Page 2558. The correct version of Figure 1 and its caption are given below.

Neu: RASW*VTFIIATVVGVLLFLILVVVVGILIKRRR
Neu*: RASW*VTFIIATV**E**GVLLFLILVVVVGILIKRRR
NeuQM: RASW*VTF**A**I**L**T**L**V**L**VLLFLILVVVVGILIKRRR
Neu*DM: RASW*VTF**A**IAT**L**EGVLLFLILVVVVGILIKRRR

Figure 1. Sequences of peptides used in this study, including the transmembrane domain of the wild-type (proto-oncogenic) rat Neu protein (Neu) and the transmembrane domain containing the oncogenic $V_{664}E$ mutation (Neu*). Also shown are mutants that lead to significant weakening of helix−helix interactions in *Escherichia coli* membranes for wild-type TM (specifically $I_{659}A/A_{661}L/V_{663}L/G_{665}L$, NeuQM) and the oncogenic TM ($I_{659}A/V_{663}L$, Neu*DM). For all four peptides, the native Pro655 residue was changed to Trp (W* in sequence) to aid in detection of the peptide in analytical ultracentrifugation experiments.

Received: June 26, 2012
Published: July 25, 2012