Corrigendum

Corrigendum to “Molecular Characterization of a Fully Human Chimeric T-Cell Antigen Receptor for Tumor-Associated Antigen EpCAM”

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Received 25 November 2014; Accepted 30 November 2014

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In the paper entitled “Molecular Characterization of a Fully Human Chimeric T-Cell Antigen Receptor for Tumor-Associated Antigen EpCAM,” there was an error in the amino acid/nucleotide sequences in Figure 2. Actually, the sequences were not for anti-EpCAM scFv but for anti-CEA scFv, which has been cloned by our laboratory in the previous study. Here we provide the correct version of Figure 2.

FIGURE 2: Schematic representations of CAR57-28ζ. The nucleotide and amino acid sequences of anti-EpCAM scFv from M13-57 hybridoma were analyzed using the IMGT/V-QUEST and IMGT/JunctionAnalysis online tools (<http://www.imgt.org/>), and each complementarity determining region (CDR) of the cloned scFv gene was determined according to Chothia's definition [15]. V_H, variable region of heavy chain; V_L, variable region of light chain; TM, transmembrane domain; Cyto, cytoplasmic domain.