Correction


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Received: 8 August 2010 / Published: 24 August 2010

The authors wish to make the following corrections to this paper [1]:

**Results and Discussion:** The sequence in Figure 5 (b) is wrong. The figure should be replaced with the one below.

![Image of the corrected Figure 5 (b)](image_url)

**Section 3.2.1:** The sentence “This library has $1.8 \times 10^{18}$ kinds of sequence variation.” should be read as: “This library has $1.15 \times 10^{18}$ kinds of sequence variation.”

**Section 3.2.2:** The sentence “The membranes were then blocked with 100 ng/mL of BSA in TBSTE,
and the aptamers were incubated with the proteins in TBSTE.” should be read as: “The membranes were then blocked with 100 mg/mL of BSA in TBSTE, and the aptamers were incubated with the proteins in TBSTE.

**Section 3.2.3:** The sentence “The structures of the aptamers were analyzed by circular dichroism (CD) spectroscopy using a JASCO (J-725) spectropolarimeter.” should be read as: “The structures of the aptamers were analyzed by circular dichroism (CD) spectroscopy using a JASCO (J-720) spectropolarimeter.”

**Section 3.2.4:** The sentence “Next, VEGF (400 U) dissolved into 10 mM acetate buffer (pH 6.0) was immobilized on the sensor chip as stipulated in the manufacturer’s manual.” should read as: “Next, VEGF (200 pmol) dissolved into 10 mM acetate buffer (pH 6.0) was immobilized on the sensor chip as stipulated in the manufacturer’s manual.”

**Conclusions:** The sentence “V7t1, the truncated mutant of Vap7, has a short sequence (29-mer) and a strong affinity for VEGF” should be read as: “V7t1, the truncated mutant of Vap7, has a short sequence (25-mer) and a strong affinity for VEGF.”

**Reference**


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