## Erratum: Development of potent monoclonal antibody auristatin conjugates for cancer therapy

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In the legend to Figure 4d on page 781, cIgG Ag<sup>-</sup> should be cAC10 Ag<sup>-</sup>.

## Corrigendum: A model of molecular interactions on short oligonucleotide microarrays

Li Zhang, Michael F Miles & Kenneth D Aldape Nat. Biotechnol. 21, 818-821 (2003)

In the legend to Figure 1 on page 819, text in parts b and c was transposed. The legend should have read as follows: (b) Weight factors. (c) Nearest-neighbor stacking energy. These stacking energies weakly correlated (r = 0.6) with that found in aqueous solution<sup>8</sup>, and are smaller in magnitude.

In the legend to Figure 2 on page 820, figure parts were referred to incorrectly. The legend should have read as follows: Accuracy test. Known concentrations of 14 'spike-in' genes are compared with those obtained from (a) PDNN, (b) MAS5.0 and (c) dChip. Each line represents a gene in 14 samples. The microarray raw data were obtained from the '1532 series' human data (see Methods for URL). For genes other than the 'spike-ins', standard deviations (s.d.) versus the averages of the log-transformed expression levels are shown in d, e and f as determined using PDNN, MAS5.0 and dChip<sup>7</sup>, respectively. Each of these figures contains 12,474 genes; top half shown in red.

