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

Svetlana O Doronina, Brian E Toki, Michael Y Torgov, Brian A Mendelsohn, Charles G Cerveny, Dana F Chace, Ron L DeBlanc, R Patrick Gearing, Tim D Bovee, Clay B Siegall, Joseph A Francisco, Alan F Wahl, Damon L Meyer & Peter D Senter
Nat. Biotechnol. **21**, 778–784 (2003)

In the legend to **Figure 4d** on page 781, cIgG Ag⁻ should be cAC10 Ag⁻.

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[8], 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[7], respectively. Each of these figures contains 12,474 genes; top half shown in red.