

Errata

for A Reliable Sequence Alignment Method Based on
Probabilities of Residue Correspondences

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(December 13, 1999)

1. 6-8th lines in the section of “Probability alignment” at the left-hand side in page 1001.

(ii) Calculate a site pair (a_i, b_j) such that

$$p(a_i, b_j) = \max_{i_1 \leq k \leq i_2, j_1 \leq l \leq j_2} (p(a_k, b_l) \mid p(a_k, b_l) \geq p(a_k, \phi) \text{ and } p(a_k, b_l) \geq p(\phi, b_l)).$$