



**Figure 5.** MALDI mass spectra of a mixture of peptides and proteins. The mixture of peptides and proteins (50 fmol/ $\mu$ L each) and the matrix solution were mixed together in equal volumes. The matrix solution was a 5:1 mixture of the CHCA solution with (a) deionized water; (b) Tf solution (0.10  $\mu$ g/ $\mu$ L); and (c) BSA solution (0.10  $\mu$ g/ $\mu$ L). Signal 1, [bovine insulin] $^+$  (5730 Da); 2, [human insulin] $^+$  (5808 Da); 3, [cytochrome C] $^{2+}$ ; 4, [IGF-I] $^+$  (7649 Da); 5, [apomyoglobin] $^{2+}$ ; 6, [cytochrome C] $^+$  (12362 Da); 7, [apomyoglobin] $^+$  (16952 Da); 8, [Tf] $^{7+}$ ; 9, [Tf] $^{6+}$ ; 10, [Tf] $^{5+}$ ; 11, [Tf] $^{4+}$ ; 12, [BSA] $^{7+}$ ; 13, [BSA] $^{6+}$ ; 14, [BSA] $^{5+}$ ; and 15, [BSA] $^{4+}$ .

the effect should not be observed when using liquid matrices.<sup>26,27</sup>

The present results suggest that the enhancement brought about by either Tf or BSA could be applicable to the improvement of sensitivity in the detection of proteins by MALDI-TOFMS in general. However, when Tf or BSA was used as an enhancer in a MALDI-TOFMS system, signals from Tf and BSA were also detected, which sometimes interfered with the analysis of the target proteins. Therefore, neither Tf nor BSA appears to be the best possible enhancer. Further studies are currently underway in order to discover the best macromolecule as an enhancer.

## CONCLUSIONS

We have demonstrated that the signal intensities of insulin and of several peptides and proteins were enhanced in

CHCA premixed with Tf or other peptides or proteins. The characteristics of this type of enhancement are as follows: (1) Tf (80 kDa) and BSA (66 kDa) led to better signal enhancement than did small peptides and proteins (<20 kDa) or IgG (150 kDa); (2) the optimum S/N value was observed when the added amount of peptide or protein was within the range 0.26–0.62 pmol; and (3) the signals of peptides of high molecular weight (>3000 Da) were enhanced by the addition of Tf or BSA to CHCA, although the signals of small peptides (<2500 Da) were not enhanced. This type of enhancement may be useful for the improvement of protein analyses with MALDI-TOFMS.

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