

Title	Nutrition analysis by nanoparticle-assisted laser desorption/ionisation mass spectrometry
Author(s)	Sahashi, Yuko; Osaka, Issey; Taira, Shu
Citation	Food Chemistry, 123(3): 865-871
Issue Date	2010
Type	Journal Article
Text version	author
URL	http://hdl.handle.net/10119/9049
Rights	NOTICE: This is the author's version of a work accepted for publication by Elsevier. Yuko Sahashi, Issey Osaka and Shu Taira, Food Chemistry, 123(3), 2010, 865-871, http://dx.doi.org/10.1016/j.foodchem.2010.05.008
Description	

Accepted Manuscript

Nutrition analysis by nano-particle assisted laser desorption/ionization mass spectrometry

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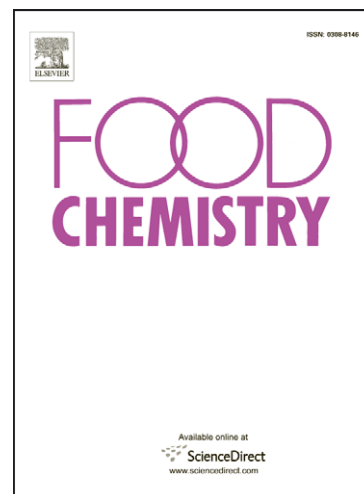
PII: S0308-8146(10)00568-6
DOI: [10.1016/j.foodchem.2010.05.008](https://doi.org/10.1016/j.foodchem.2010.05.008)
Reference: FOCH 9615

To appear in: *Food Chemistry*

Received Date: 19 November 2009
Revised Date: 25 March 2010
Accepted Date: 1 May 2010

Please cite this article as: Sahashi, Y., Osaka, I., Taira, S., Nutrition analysis by nano-particle assisted laser desorption/ionization mass spectrometry, *Food Chemistry* (2010), doi: [10.1016/j.foodchem.2010.05.008](https://doi.org/10.1016/j.foodchem.2010.05.008)

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1 Nutrition analysis by nano-particle assisted laser desorption/ionization mass
2 spectrometry.

3

4

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15

16 Abstract

17 We analyzed the bioactive compounds in *Panax ginseng* C.A. Meyer by using
18 nanoparticle-assisted laser desorption/ionization (nano-PALDI) mass spectrometry (MS).

19 To this end, we prepared manganese oxide nanoparticles ($d = 5.4$ nm) and developed a
20 nano-PALDI MS method to analyze the standard ginsenosides and identify these
21 ginsenosides in an extract of *Panax ginseng*. The nanoparticles served as an
22 ionization-assisting reagent in MS. The mass spectra did not show any background
23 interference in the low- m/z range. Our pilot study showed that the nanoparticles could
24 ionize the standard ginsenosides and also respective lipid and ginsenosides in the extract
25 without the aid of chemical and liquid matrices used in conventional MS methods.
26 Analysis of the post-source decay spectra obtained using nano-PALDI MS will yield
27 information regarding the chemical structure of the analyte.

28 Introduction

29 Herbal products have been used in traditional Chinese medicine (TCM) for a
30 long time and have recently gained attention as complementary and alternative
31 medicines (Hijikata, Miyamae, Takatsu & Sentoh, 2007; Sun et al., 2009; Xu & Xu,
32 2009). *Panax ginseng* C.A. Meyer is one of the most famous oriental herbs used in
33 TCM, and it contains many bioactive compounds, including triterpene glycosides called
34 ginsenosides. Although ginsenosides have been thought to be the main bioactive
35 components in *Panax ginseng* (Metori, Furutsu & Takahashi, 1997; Newman et al.,
36 1992; Wu et al., 1992), their role in the efficacy of *Panax ginseng* has not been
37 completely elucidated. Recent articles have reported that multiple components in *Panax*
38 *ginseng*, such as lipids, polysaccharides, peptides, and amino acids act synergistically
39 (Spelman, Burns, Nichols, Winters, Ottersberg & Tenborg, 2006; Zeng, Liang, Jiang,
40 Chau & Wang, 2008). Therefore, to elucidate the efficacy of this herb, simultaneous
41 analysis of the secondary-metabolite complexes of relatively small molecules like
42 ginsenosides is very important.

43 Mass spectrometry (MS) is a powerful technique used to analyze metabolites in
44 biological samples and tissues. It can be used to directly and simultaneously detect
45 multiple components in crude samples such as TCM products. Although MS combined

46 with liquid chromatography (LC) and gas chromatography (GC) can be used to analyze
47 ginsenosides (Cui, Song, Liu & Liu, 2001; Fuzzati, Gabetta, Jayakar, Pace & Peterlongo,
48 1999; Li, Mazza, Cottrell & Gao, 1996; Tawab, Bahr, Karas, Wurglics &
49 Schubert-Zsilavec, 2003; Wang et al., 2008), these techniques are time consuming
50 (analysis time, ~30 min.). Matrix-assisted laser desorption/ionization (MALDI) MS is a
51 soft and sensitive ionization technique that uses chemical matrices such as
52 4-hydroxy- α -cinnamic acid (CHCA), 2,5-dihydroxybenzoic acid (DHB), and sinapic
53 acid (SA) to facilitate ionization of the analyte. However, one of the main problems of
54 MALDI-MS is the overlapping of the matrix peaks and fragment peaks in the low-mass
55 region ($m/z \sim 800$). These shortcomings significantly complicate the application of these
56 techniques in the multiple-component analysis of samples such as TCM products and
57 their metabolites. It is our approach that these shortcomings can be overcome by using
58 advanced analytical techniques developed through interdisciplinary collaboration.

59 Nanomaterials have shown great potential in facilitating the development of new
60 technologies (Chithrani & Chan, 2007; Moritake et al., 2007; Taira, Hatanaka, Moritake,
61 Kai, Ichiyangi & Setou, 2007). Nanoparticles (NPs) have been used in the
62 development of solar cells (Kitada, Kikuchi, Ohono, Aramaki & Maenosono, 2009),
63 sensors (Ai, Zhang & Lu, 2009; Kalogianni, Koraki, Christopoulos & Ioannou, 2006),

64 catalysts (Mitsudome, Noujima, Mizugaki, Jitsukawa & Kaneda, 2009), drug delivery
65 systems (Moritake et al., 2007), and imaging techniques (Taira, Sugiura, Moritake,
66 Shimma, Ichiyanagi & Setou, 2008). However, there have been very few studies on the
67 use of NPs in food chemistry (Ravindranath, Mauer, Deb-Roy & Irudayaraj, 2009; Yang,
68 Kostov, Bruck & Rasooly, 2009). Previous studies used photospectroscopy with NPs to
69 detect analytes; however, this technique afforded limited detection of the multiple
70 components in food. In our previous reports, to obtain ionization-assisting agents that
71 could be used to perform nanoparticle-assisted laser desorption/ionization
72 (nano-PALDI) MS without significantly increasing the background signals, we prepared
73 metal oxide nanoparticles surrounded by amorphous SiO₂ and an amino group (Figure 1
74 c and d) (Moritake, Taira, Sugiura, Setou & Ichiyanagi, 2009; Taira, Kitajima,
75 Katayanahi, Ichiishi & Ichiyanagi, 2009; Taira et al., 2008). Here, we investigated the
76 suitability of nano-PALDI MS for analyzing lipid and ginsenosides in *Panax ginseng*
77 extracts. We assessed the ionization of several standard ginsenosides by using
78 nano-PALDI MS as an analysis marker for crude samples. Further, we used the
79 nano-PALDI MS technique to separate and evaluate the original ingredients in the
80 complicated MS spectrum for ginseng extract.

81 **Method**

82 Materials

83 Standard ginsenosides (Rb₁, Rb₂, Rc, Rd, Re, Rf, and Rg₁ with purity >98 %, >94%,
84 >99%, >99%, >99%, >99%, and >99%, respectively) were purchased from Funakoshi
85 (Tokyo, Japan). The extract was obtained from tissue-cultured *Panax ginseng* (TCPG)
86 (Nitto Denko, Osaka, Japan). The TCPG powder was extracted using hot water (80°C)
87 for 2 h, dried, and re-extracted with 70% (v/v) methanol. The extract thus obtained was
88 applied to a small column (Sep-Pak cartridge C18 and NH₂; Waters, Milford, USA) to
89 concentrate the ginsenosides.

90

91 Preparation of nanoparticles

92 Manganese oxide-based nanoparticles were prepared by mixing aqueous solutions
93 of MnCl₂·4H₂O (5 ml, 100 mM; WAKO Pure Chemicals, Japan) and
94 3-aminopropyltriethoxysilane (5 ml; γ-APTES; Shinetsu, Kagaku, Japan). After stirring
95 at room temperature for 1 h, the resulting precipitate was washed several times with
96 ultrapure water and dried at 55°C in an incubator. The dried samples were pulverized in
97 a porcelain mortar. The morphology and diameter distribution of the nanoparticles were
98 investigated using a transmission electron microscope (TEM; H-7100; Hitachi, Japan).

99

100 Nano-PALDI mass spectrometry

101 The utility of the nanoparticles as ionization-assisting materials in mass spectrometry
102 was confirmed in a MALDI-TOF-type instrument (TOF = time of flight;
103 Voyager-DE-RP; Applied Biosystems, Germany) by using a N₂ laser with an emission
104 wavelength of 337 nm. Samples of standard ginsenosides samples such as G-Rb₁, G-Rb₂,
105 G-Rc, G-Rd, G-Re, G-Rf, and G-Rg₁ were chosen for the analysis. The nanoparticles (1
106 mg) were dispersed in 1 mL of methanol or in 1 mL of a 10 mM methanolic solution of
107 sodium acetate. Each sample was independently dissolved in distilled water at a
108 concentration of 100 pmol/μL. Each analyte solution (1 μL) was pipetted on to the
109 surface of the nanoparticle-coated target plates. The peptides used for external
110 calibration were deposited on the plate to minimize the mass shift. The analyte surface
111 was irradiated with 100 laser shots in the positive mode.

112

113 Results and discussion

114 Ability of the nanoparticles to assist ionization of pure sample analytes

115 We used the standard ginsenosides G-Rb₁ [exact mass (Me): 1108.6], G-Rb₂ (Me:
116 1078.6), G-Rc (Me: 1078.6), G-Rd (Me: 947.2), G-Re (Me: 947.2), G-Rf (Me: 800.5),
117 and G-Rg₁ (Me: 800.5) to evaluate the usefulness of employing nanoparticles as laser

118 desorption/ionization materials. The mass spectra of the standard ginsenosides were
119 obtained in the presence of nanoparticles with sodium ions, thereby ensuring that the
120 precursor ions were obtained in the form of $[M + Na]^+$ ions (Taira et al. 2008). In this
121 study, the standard ginsenosides formed sodium-adduct ions and yielded high-intensity
122 signals. Further, to characterize a variety of ginsenosides, we performed post-source
123 decay (PSD) MS for structural analysis.

124 For G-Rb₁, we obtained a precursor $[M + Na]^+$ ion at m/z 1132.1. This ion yielded
125 fragment ions $[z1 + Na]^+$ (at m/z 789.3), which corresponded to the combination of the
126 agriconc moiety and disaccharide moiety of R₁ or R₂, and $[y1 + Na]^+$ (at m/z 364.7),
127 which corresponded to the disaccharide moiety of R₁ or R₂ (Figure 2a).

128 Similarly, the PSD spectra of G-Rb₂ and R_c showed a precursor ion at m/z 1102.1 and
129 the 2 derivative ions, namely, $[y1 + Na]^+$ at m/z 335.2, which corresponded to the
130 disaccharide moiety of R₁, and $[z1 + Na]^+$ at m/z 789.8 (Figure 2b) and 789.6 (Figure 2c),
131 which corresponded to the combination of the agriconc and disaccharide moieties of R₂.

132 For G-Rd, we obtained an $[M + Na]^+$ ion at m/z 970.1 and 3 fragment ions at m/z
133 789.3, 365.2, and 203.1. The fragment ion $[z1 + Na]^+$ at m/z 789.3 corresponds to an R₁
134 molecule without the glucose moiety. The m/z values for the smaller fragment ions $[y2$
135 $+ Na]^+$ and $[y1 + Na]^+$ (365.2 and 203.1, respectively) were consistent with the

136 molecular masses of sodium-adducted disaccharide and glucose moieties from R_2 and R_1 ,
137 respectively (Figure 2d).

138 For G-Re (m/z of the precursor ion, 970.1), we obtained 2 fragment ions $[z1 + Na]^+$
139 and $[y1 + Na]^+$ at m/z values 789.2 and 203.1, respectively. The mass difference
140 between the precursor ion and the fragment ion at m/z 789.2 was 180.9, which indicated
141 the loss of a glucose molecule. The fragment ion at m/z 203.1 indicated a
142 sodium-adducted glucose moiety from R_1 (Figure 2e). The same exact mass of R_f and
143 R_{g1} showed difference in the PSD spectra. The PSD of G-Rf showed only 1 fragment
144 ion $[z1 + Na]^+$ at m/z 365.2, which corresponded to the disaccharide moiety from R_3
145 (Figure 2f). In contrast, the PSD of G-R $_{g1}$ showed 2 fragment ions, namely, $[z1 + Na]^+$
146 and $[y1 + Na]^+$ at m/z values 643.8 and 202.9, respectively, which corresponded to the
147 agricone and glucose moieties of R_1 or R_3 and the divided glucose moiety from R_1 or R_3 ,
148 respectively (Figure 2g). We distinguished the molecules with the same exact mass on
149 the basis of the differences in the composition of disaccharides (G-Rf) and
150 monosaccharides (G-R $_{g1}$). These nano-PALDI PSD fragmentation patterns were in
151 good agreement with the MALDI PSD fragmentation patterns of standard ginsenosides
152 (data not shown). This finding indicated that our technique could also yield accurate
153 results under mild ionization conditions without unnecessary degradation of the

154 bioactive molecule. These results could be utilized for analyzing the index of raw
155 sample like plant extracts.

156 We observed a number of high-intensity signals for ginsenosides in the extract of
157 *Panax ginseng* t mass spectra obtained with nanoparticles in the absence (Figure 3a) or
158 presence (Figure 3b) of sodium ions. Although there were few MS signals with
159 intensity greater than m/z 600, we could confirm the signals that corresponded to
160 ginsenosides. In the case of the G-Rb₁ ions (m/z 1131.6 [M + Na]⁺; 1147.6 [M + K]⁺),
161 both sodium- and potassium-adduct ions were observed in the absence of sodium
162 acetate (Figure 3a inset), because the extract originally included salt ions, especially
163 sodium and potassium salts. In the MS spectrum, such salt ions preferentially appeared
164 in their adducted form, rather than the protonated form. However, the related signals
165 showed a convergence only in the case of the sodium-adducted form (m/z 1131.2).
166 Interestingly, the correlation between the signals of the sodium-adducted form of
167 G-Rg₁ (m/z 823.1) and G-Rb₂ or R_c (m/z 1102.1) appeared only in the presence of
168 sodium acetate (10 mM) (Figure 3b inset). In addition, the background noise in the
169 presence of sodium acetate (Figure 3b inset), was lower than in the absence of sodium
170 acetate (Figure 3a inset). The sodium-adducted forms of G-Rg₁ and G-Rb₂ or R_c were
171 more easily ionized than other ion-adducted forms, such as the proton- or

172 potassium-adducted form. This result indicated that the ginsenosides had optimal
173 ionization forms. Moreover, in the low molecular range (m/z 200–400), the signal
174 intensities in the presence of sodium ions (Figure 3b), were lower than that in the
175 absence of these ions (Figure 3a); thus, the signals in this region indicated a
176 preferential ionization to the protonated form. This technique can be used for accurate
177 and simple analysis of complex mixtures such as foods and nutrients; however, the
178 differences in the ionization characters of these samples must be carefully considered
179 while performing these analyses.

180 To perform structural analysis using post-source decay (PSD) nano-PALDI mass
181 spectrometry, we deduced that the 4 signals at m/z 551.5, 823.1, 1102.4, and 1132.1
182 were obtained from the extract of *Panax ginseng* in the presence of sodium ion and
183 determined that these signals originated from lysophosphatidylcholine (LPC)-(1-acyl
184 20:1) ($[M + H]^+$ ion), G-Rg₁ ($[M + Na]^+$ ion), G-Rb₂ or G-Rc ($[M + Na]^+$ ion), and
185 G-Rb₁ ($[M + Na]^+$ ion). For the precursor $[M + H]^+$ ion of LPC-(1-acyl 20:1) at m/z
186 551.1, the typical fragment ions $[y1]^+$ and $[z1 + H]^+$ were detected at m/z 85.9 and 298.5,
187 respectively; this finding provided information on the trimethylamine moiety and the
188 fatty acid (1-acyl 20:1) in the sequence. The PSD fragment patterns indicated that the
189 promptly obtained lipid fragment ions did not originate from the observed molecular

190 ions, because the prompt fragmentation occurred immediately after the formation of
191 highly unstable protonated precursor ions (Figure 4a) (Al-Saad, Zabrouskov, Siems,
192 Knowles, Hannan & Hill, 2003).

193 Similarly, the PSD spectra of ginsenosides showed fragment ions similar to those of
194 the standard ginsenosides G-Rg₁, G-Rb₂ or G-Rc, and G-Rb₁. The PSD spectrum of
195 G-Rg₁ showed 2 derivative ions that corresponded to the glucose ions (m/z 202.9; [M +
196 Na]⁺) of R₁ or R₂ and the agricones moieties (m/z 643.8; [M + Na]⁺) (Figure 4b).

197 We detected a precursor ion at m/z 1102.1 and 2 derivative sodium-adduct ions at m/z
198 336.3 and m/z 789.0, which corresponded to the disaccharide moiety of R₁ or R₃ and the
199 combination of the disaccharide and agricones moieties of R₁ or R₃, respectively. The
200 difference between G-Rb₂ and G-Rc can be attributed to the arabinose conformation
201 (arabinopyranose for G-Rb₂ and arabinofuranose for G-Rc) within the disaccharide
202 moiety of R₂; this conformation can complicate the distinction between G-Rb₂ and G-Rc
203 using the PSD MS technique (Figure 4c). The corresponding PSD spectrum of G-Rb₁ is
204 shown in Figure 4d. We detected an [M + Na]⁺ precursor ion at m/z 1132.1 and 2
205 fragment ions, namely, [M + Na]⁺ at m/z 788.4 and [M + Na]⁺ at m/z 365.0. These
206 fragment ions could be considered as the z1 and y1 ions, which are characteristic of the
207 cleavage of the glycosidic bonds at R₁ or R₂.

208 These fragment patterns were in good agreement with the PSD spectra of standard
209 ginsenosides (Figure 2 a, b, c, and g). We could identify the bioactive components such
210 as ginsenosides and lipids from the extract by using the nano-PALDI MS technique.

211

212 **4. Conclusions**

213 Nano-PALDI MS allowed ionization and background-free analysis of the small
214 molecules in a *Panax ginseng* extract. The nanoparticles could ionize the standard
215 ginsenosides in the presence of external sodium ions. The obtained signals corresponded
216 to those of sodium-adduct ions. Although conventional matrices do not ionize the
217 analyte in the presence of external salt ions, this technique can facilitate the analysis of
218 crude samples like plant extracts. Using this technique, we detected lipids and
219 ginsenosides in the *Panax ginseng* extract and identified the optimal ion forms of these
220 compounds. We mainly focused on using nano-PALDI MS to investigate the role of
221 ginsenosides as the active components of *Panax ginseng*. However, the contributions of
222 other compounds, such as saccharides, peptides, and proteins, should be investigated.
223 The nano-PALDI MS technique is a good substitute for MALDI and has great potential
224 for rapid screening of bioactive ingredients such as ginsenosides in plant extracts;
225 however, further studies are required to establish their traceability in foods and nutrient

226 product.

227 In addition, the nanoparticles may be utilized in the mass spectrometric analyses of
228 biomedical tissues (Taira et al., 2008) and in cellular analysis (Moritake et al., 2009).
229 The nanoparticle-based approach used in this study can be employed for simple and
230 efficient identification of various ingredients of foods and herbal products used in TCM.

231

232 **Acknowledgments**

233 We thank member of Takagi and Takamura laboratory in JAIST, particularly Prof. M.
234 Takagi and Prof. Y. Takamura, Ms. A. Makino and Ms T. Taniho for providing technical
235 assistance and advice. This research was supported by a WAKATE-B grant from the
236 Japan Society for the Promotion of Science to S. T. and a Grant-in-Aid to S. T. from
237 JAIST and the resource of coordinated research program to Nitto Denko and a A-STEP
238 to S. T. and Y. S. from Japan Science and Tech. Agency

239

240 **Figure legends**

241 Figure 1

242 A schematic illustration of nanoparticle-assisted laser desorption/ionization
243 (nano-PALDI) mass spectrometry (a). Transmission electron microscopy (TEM) image
244 of the nanoparticles (b). When reserpine (100 pmol) was used as a model drug with the
245 nanoparticles, the nano-PALDI mass spectra (c) did not show any background

246 interference in the low-mass region. In contrast, the mass spectra of reserpine with
247 4-hydroxy- α -cinnamic acid (CHCA) showed background noise in the low-mass region
248 (d).

249

250 Figure 2

251 The post-source decay nanoparticle-assisted laser desorption/ionization (nano-PALDI)
252 mass spectra of the standard ginsenosides G-Rb₁ (a), G-Rb₂ (b), G-Rc (c), G-Rd (d),
253 G-Re (e), G-Rf (f), and G-Rg₁ (g). The abbreviations for the sugar moieties are glc
254 (β -D-glucose), arap (α -L-arabinose; pyranose), araf (α -L-arabinose; furanose), and rha
255 (α -L-rhamnose).

256

257 Figure 3

258 Mass spectra of the extract with nanoparticles (NPs) alone (a) and with NPs in the
259 presence of sodium acetate (NaAc: 10 mM) (b). The superimposed spectra of
260 tissue-cultured *Panax ginseng* (TCPG) extract with NPs in the absence (upper) and
261 presence of (lower) additional NaAc.

262

263 Figure 4

264 The post-source decay nanoparticle-assisted laser desorption/ionization (nano-PALDI)
265 mass spectra of lysophosphatidylcholine (LPC)-(1-acyl 20:1) (a), ginsenoside (G)-Rg₁
266 (b), G-Rb₂ or G-Rc (c), and (G)-Rb₁ (d). The abbreviations for the sugar moieties are the
267 same as those used in Figure 2.

268

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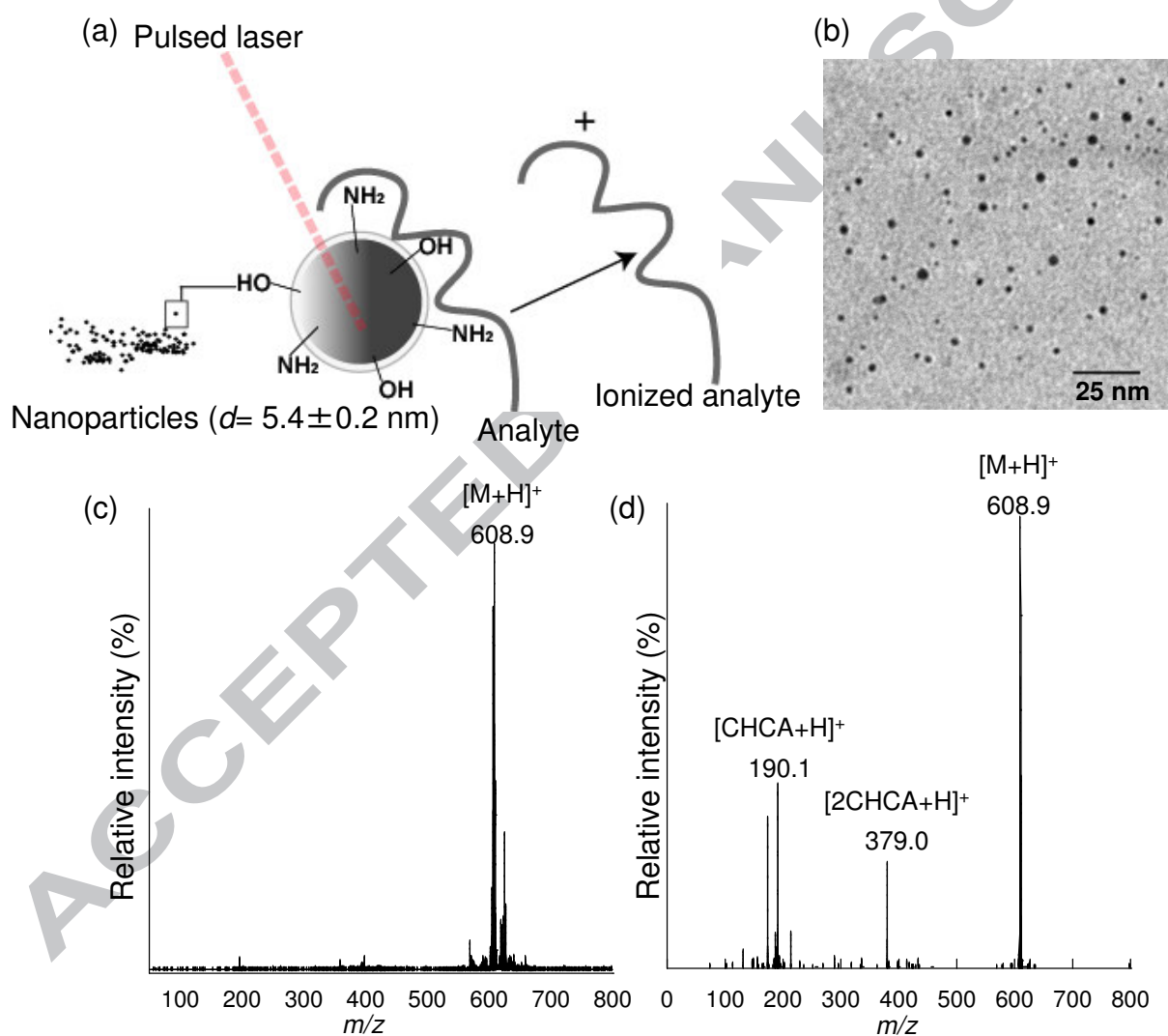


Figure 1 Taira et al.

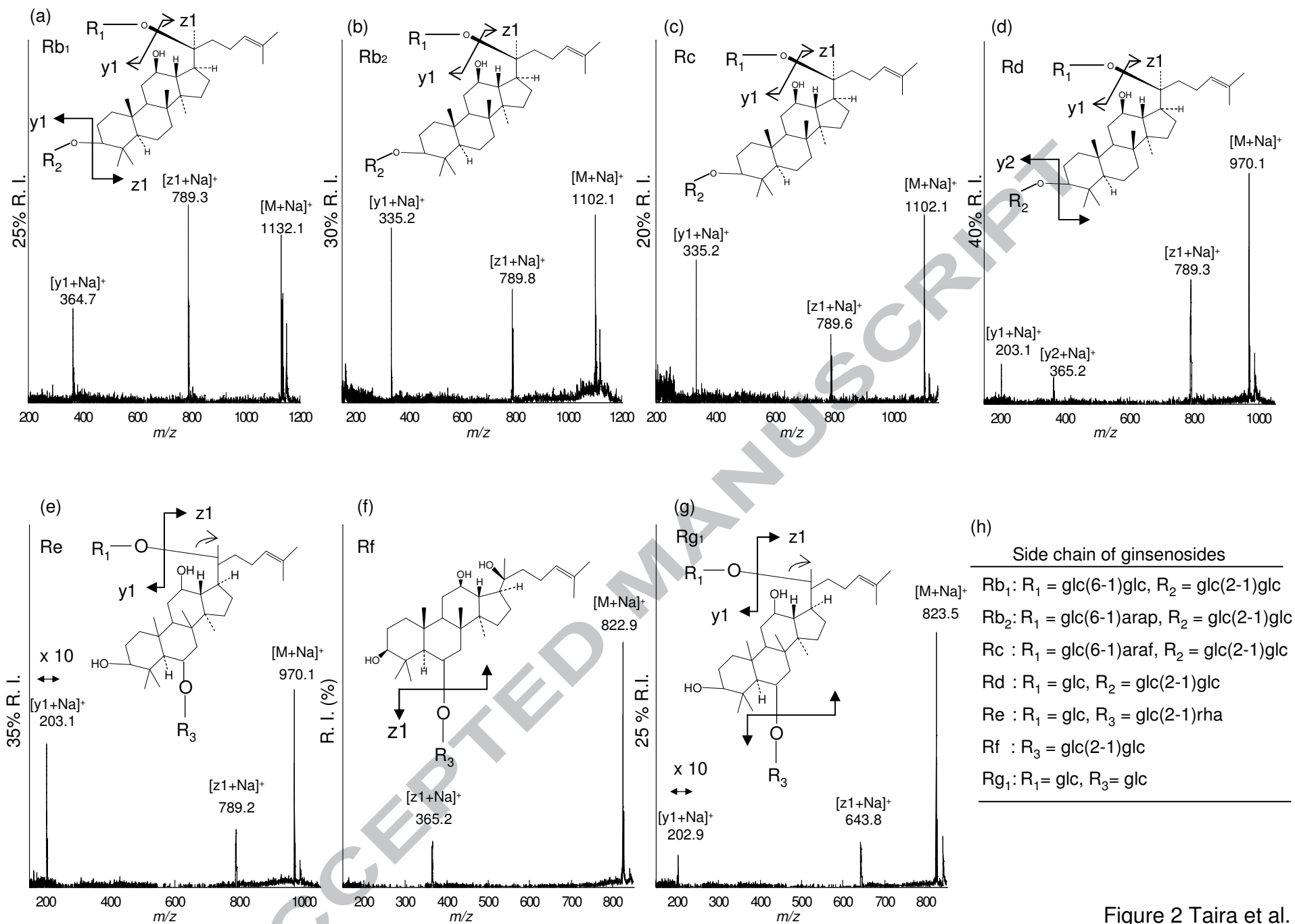


Figure 2 Taira et al.

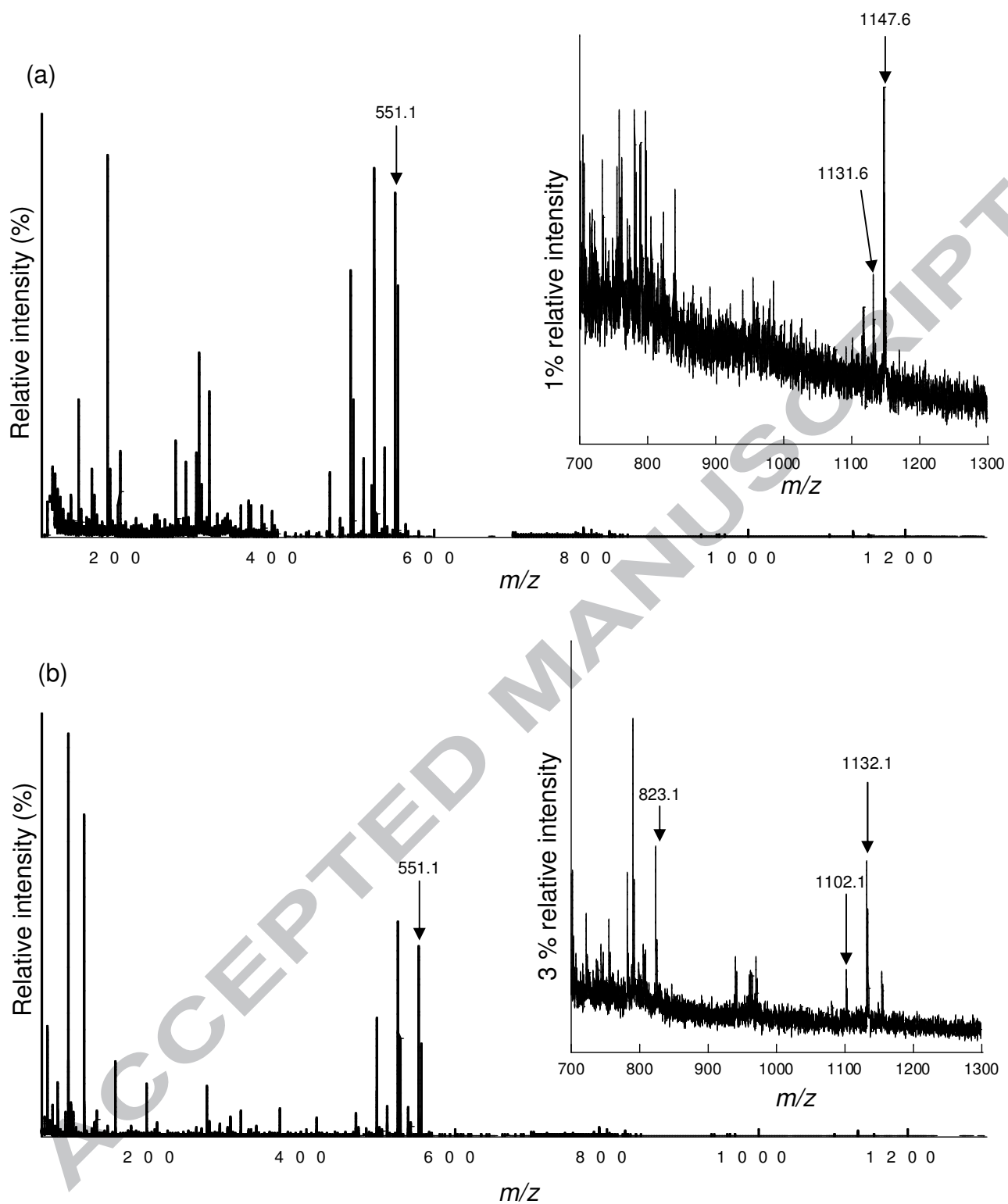


Figure 3 Taira et al.

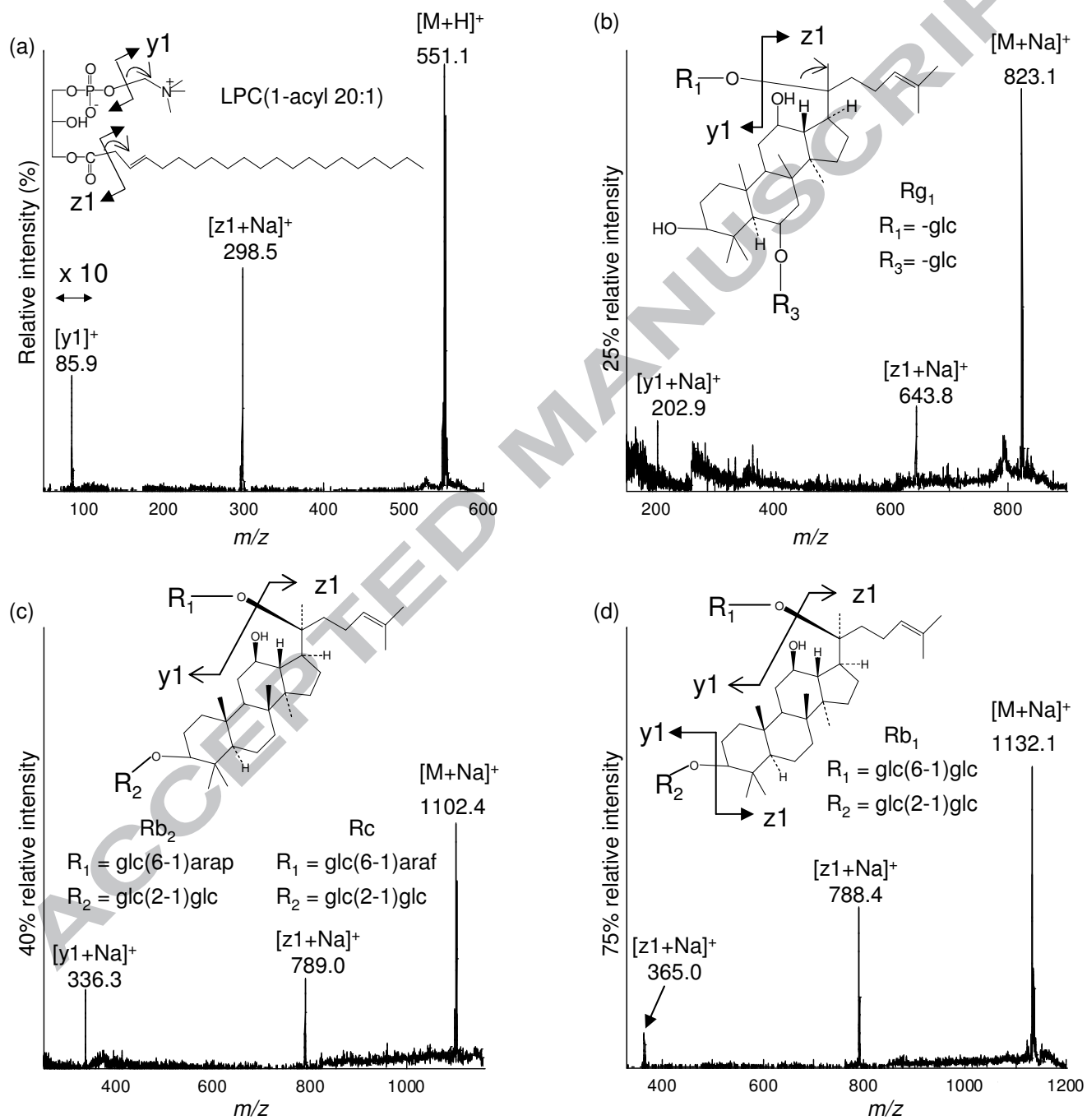


Figure 4 Taira et al.