

液滴萃取表面分析-质谱法用于新鲜与蒸制三七根切片中皂苷成分的快速鉴别

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摘要: 三七为五加科植物三七 (*Panax notoginseng*) 的干燥根及根茎, 为我国传统珍贵中药材, 具有止血散瘀、消肿止痛等功效。相比于新鲜三七, 蒸制三七对治疗肿瘤和心血管疾病等表现出更好的疗效。皂苷是三七主要的化学和药效成分, 本研究基于表面萃取结合芯片多通道纳喷质谱技术, 建立了液滴萃取表面分析-质谱方法 (LESA-MS), 能够直接、快速地对新鲜与蒸制三七根切片中木质部、韧皮部和形成层中皂苷成分进行鉴别。实验结果表明, 新鲜与蒸制三七根中皂苷成分及其含量具有一定的差异, 表现为在新鲜三七根切片中, 人参皂苷 Rg1、Rb1、Re、Rd, 三七皂苷 R1 及其丙二酰类成分较高。而在蒸制三七根切片中, 人参皂苷 Rg5、Rk1 等弱极性成分能被检测到, 大极性成分则相对含量较低。本方法具有快速、稳健且灵敏度高的优势, 且操作过程无需破碎、萃取、色谱分离等繁琐的步骤, 实现了对鲜三七与蒸三七根切片所含化学成分及二者差异的无损分析。

关键词: 液滴萃取表面分析; 质谱; 三七; 皂苷; 快速鉴别

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Rapid characterization of saponins in fresh and steamed notoginseng root slices by liquid extraction, surface analysis-mass spectrometry

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Abstract: Notoginseng (Sanqi), the root of *Panax notoginseng* (Burk.) F. H. Chen (Araliaceae), is one of the most valuable traditional Chinese medicines (TCM). It has been widely used in China with a long history for treatment of haemorrhage, edema, and cardiovascular disorders. Steamed *P. notoginseng* has been considered to have stronger therapeutic functions than raw *P. notoginseng* in the treatment of tumors, cardiovascular diseases, etc. Saponins are the principal chemical and pharmacological constituents in *P. notoginseng*. Thus, it is of great importance to determine the constituent saponins and determine any differences between fresh *P. notoginseng* and steamed *P. notoginseng*. We used a rapid and direct analytical method based on liquid extraction surface analysis combined with mass spectrometry (LESA-MS) to identify saponins in the xylem, phloem and cambium of fresh and steamed *P. notoginseng* root slices. The results revealed that ginsenosides Rg1, Rb1, Re, Rd, notoginsenoside R1 and their malonyl group versions were most abundant in fresh root slices, while in steamed slices ginsenosides Rg5, Rk1 and other minor polar components could be detected, and the relative content of large polar components

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was lower. The described method is fast, robust and sensitive and the process does not need traditional and cumbersome pretreatment such as crushing, extraction and separation. It is the first non-destructive study on the differences in saponins between fresh and steamed *P. notoginseng* root slices.

Key words: liquid extraction surface analysis; mass spectrometry; *Panax notoginseng*; saponin compound; rapid identification

三七为五加科人参属植物 *Panax notoginseng* (Burk.) F. H. Chen 的干燥根及根茎, 主产于中国云南文山^[1]。作为传统的珍稀药材, 三七有着悠久的延用历史。在清朝著作《本草纲目拾遗》中记载: “人参补气第一, 三七补血第一, 味同而功亦等, 故称人参三七, 为中药之最珍贵者”^[2]。现代研究表明, 三七外用可散瘀止血、消肿止痛, 内服可抗肿瘤、抗抑郁、改善心脑血管功能等^[3-6]。相比于新鲜三七, 有文献报道经高温高压蒸制的蒸三七对心血管功能障碍、肿瘤等疾病表现出更加积极的治疗效果^[7,8]。近年来, 对三七的化学组成已有较多的研究, 其主要含氨基酸、黄酮、三萜皂苷等成分, 其中三萜皂苷为三七主要的药效成分, 含量约为 12%^[9]。

三七中的皂苷成分主要为达玛烷型三萜皂苷。根据其糖基连接点在 C-3 位和 C-6 位的区别, 达玛烷型皂苷可以分为原人参二醇型 (protopanaxadiol, PPD) 和

原人参三醇型 (protopanaxatriol, PPT) (图 1A)。人参皂苷 Rg1、Rb1、Rd、Re 为三七中含量较高的 PPD 型皂苷, 而 PPT 型的主要有人参皂苷 Rg1 和三七皂苷 R1^[10]。当三七药材被蒸制后, 所含极性较大的皂苷成分会在高温高压下经过脱糖、脱水、羟基化等形式转化成极性较小的皂苷^[11]。目前针对三七、人参等植物药材的成分分析和质量标准研究大多需要繁琐的预处理过程, 例如对药材的清洗、粉碎、提取等。这些步骤不仅耗时费力, 而且消耗大量有机溶剂, 不符合绿色环保的发展理念^[12]。同时, 前处理过程中可能会损失部分活性成分, 对珍贵的三七药材资源造成过多的浪费^[13,14]。

本研究采用液滴萃取表面分析-质谱方法 (liquid extraction surface analysis-mass spectrometry, LESA-MS) 首次对三七根药材中的皂苷成分进行快速表征。该敞开式的方法结合了在线表面萃取与芯片多通道纳

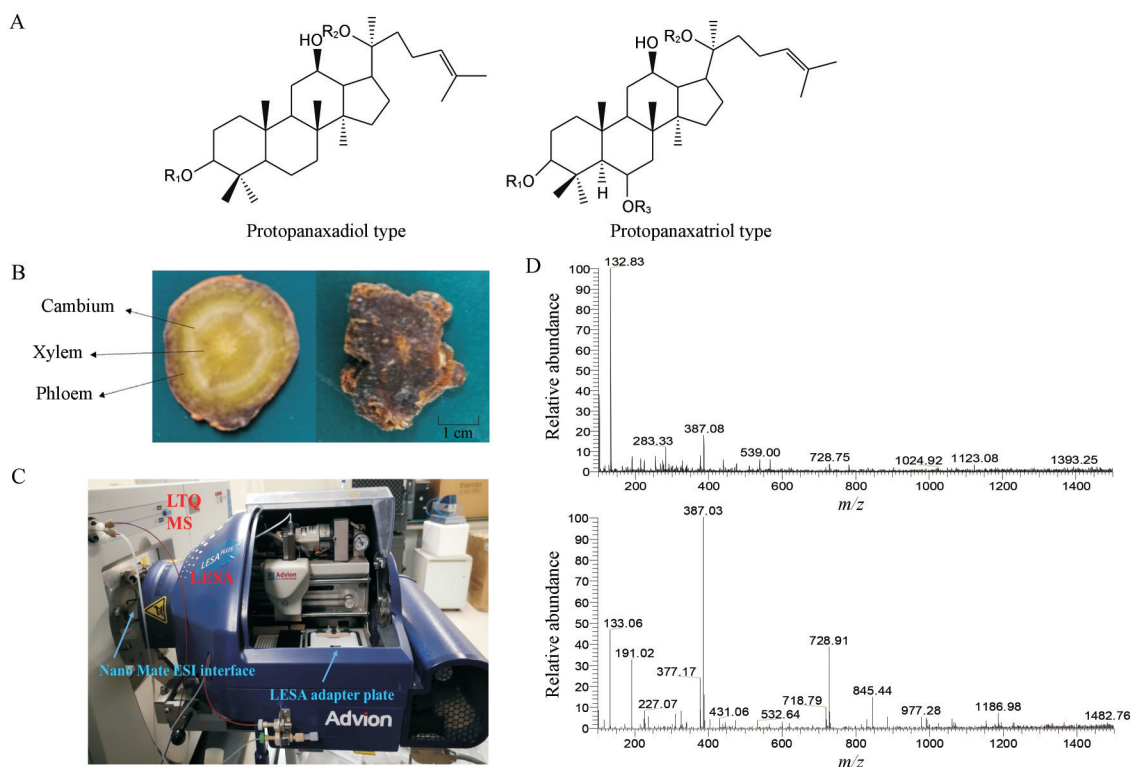


Figure 1 Graphical of experimental. A: PPD type and PPT type of dammarane saponins; B: Fresh *P. notoginseng* root slice (left) and steamed root slice (right); C: Liquid extraction surface analysis (LESA) device coupled with LTQ MS; D: Mass spectrum of steamed slice in phloem (upper) and mass spectrum of fresh slices in xylem (bottom)

喷技术, 无需繁琐冗长的样品前处理, 即可对中药材切片实现直接、快速、灵敏及高通量的分析^[15]。通过利用 LESA-MS 方法对新鲜三七根切片和在 125 °C 下蒸制 2 h 的三七根切片样品进行直接分析, 并初步探究新鲜与蒸制三七根木质部、韧皮部、形成层中皂苷成分的差异, 以期为新鲜三七和蒸制三七根切片的药效差异提供初步的理论与物质基础。

材料与方 法

实验仪器 LESA 液滴萃取表面分析系统 (美国 Advion 公司), 包括芯片多通道纳喷离子源, nano-ESI 芯片 (5 μm), LESA 适配板。ThermoFisher Scientific LTQ XL 线性离子阱质谱 (美国 ThermoFisher Scientific 公司)。高压灭菌锅 (日本 ALP 公司)。

试剂 实验用水由 Milli-Q 超纯水系统 (美国 Millipore 公司) 制得, 乙腈 (色谱纯, 美国 Fisher 公司), 甲酸 (色谱纯, 德国 Merck 公司)。

三七药材 三七根药材采于中国云南省文山种植基地, 药材经上海中医药大学吴立宏教授鉴定为五加科植物三七 *Panax notoginseng* (Burk.) F. H. Chen 的干燥根及根茎。蒸制三七采用高压灭菌锅在 125 °C 下蒸制 2 h。

LESA 方法 新鲜三七根和蒸制三七根切成表面平整的 3~5 mm 厚度的圆切片 (图 1B), 放置于通用的 LESA 适配板。采用负离子模式, 气压 (gas pressure): 0.5 psi, 应用电压 (voltage to apply): 1.70 kV。萃取溶剂为含 0.1% 甲酸的乙腈-水 (80:20), 每次吸头吸取的溶剂体积为 2.0 μL, 萃取溶剂在切片表面停留 10 s, 再通过带电枪头吸取后经 nano-ESI 芯片进行自动化的纳升级电喷雾进样。

MS 方法 采用负离子全扫描模式, m/z 扫描范围: 100~1 500, 毛细管温度为 190 °C。LESA-MS 装置如图 1C 所示。

数据处理 采用质谱数据处理软件 Xcalibur 2.2 导出色谱峰的响应值。采用多元统计分析软件 SIMCA 14.0 进行最小正交偏最小二乘法判别分析 (orthogonal partial least squares discriminant analysis, OPLS-DA)。

结果

1 新鲜与蒸制三七根切片所含皂苷成分的对 比分析

采用质谱数据处理软件提取图谱中的化合物峰 (图 1D), 根据文献所报道的分子量、质荷比及其来源, 共推测到 49 个化合物。其中皂苷类化合物有 45 个, 非皂苷类化合物有 4 个。以人参皂苷 Rg1 鉴定为例, 负离子模式下, 在质谱图中检测到 m/z 799.5 [M-H]⁻ 峰和 m/z

845.5 [M+HCOO]⁻ 峰, 推断出其分子式为 C₄₂H₇₂O₁₄, 推测为人参皂苷 Rg1 或人参皂苷 Rf。且在三七根中, 人参皂苷 Rg1 的含量远高于人参皂苷 Rf。因此, 根据峰的质谱信号强度, 可进一步推断出该化合物峰为人参皂苷 Rg1。在三七根中, 人参皂苷 Rg1 是含量最高的皂苷, 且响应信号容易确定。因此本实验以新鲜三七根木质部中的人参皂苷 Rg1 的信号响应为基准值, 其含量定义为 1, 鲜三七与蒸三七根切片各部位所含化合物的相对含量见表 1^[16-53]。表 1 的结果表明, 新鲜三七根切片中的皂苷成分和相对含量与蒸制三七根切片有明显差异。新鲜三七根切片中的不同部位皂苷的种类无明显差异, 其中木质部的含量最高, 韧皮部次之。而在蒸制三七根切片中, 韧皮部的皂苷含量最高, 木质部和形成层含量均较低。在新鲜三七根切片中, 人参皂苷 Rg1、Re、Rd、Rb1 及其丙二酰基形式, 以及三七皂苷 R1 相对于其他皂苷成分含量较高。对于蒸制三七根切片, 丙二酰人参皂苷 Rg1、Rd 和 Re 含量相对较低, 一些小极性皂苷化合物含量较高, 如三七皂苷 R10。相较新鲜三七, 蒸制三七中还可测到新出现的弱极性皂苷成分, 例如人参皂苷 Rg5、Rk1、Rs3、Rs7 和三七皂苷 Sft3、Sft4。根据不同皂苷分子结构及文献检索推测 (图 2), 当三七经过高温高压蒸制时, 大极性的皂苷成分会经过 C20 位脱糖转化为次生皂苷, 再经过水合作用或脱水作用在 C17 位侧链形成羟基化或双键, 从而转化为弱极性的皂苷化合物。

2 新鲜与蒸制三七根切片不同部位皂苷的多元统计 分析

采用有监督 OPLS-DA 模型对不同切片中皂苷化合物的质谱数据进行分析, 进一步揭示新鲜和蒸制三七根切片的木质部、韧皮部和形成层中皂苷类成分相对含量的差异。新鲜三七根各部位归为新鲜 (fresh) 类, 而蒸制则归为蒸制 (steamed) 类。从 OPLS-DA 的得分图 (图 3A) 可知, 新鲜与蒸制三七可显著分离, 其皂苷成分及含量有一定差异。该模型具有良好的解释能力和预测能力 (R^2X 、 R^2Y 和 Q^2 皆为 1)。此外, 在新鲜三七根切片中, 韧皮部和形成层皂苷成分和相对含量差异较小, 蒸制三七根切片中, 韧皮部与其他两个部位差异较明显。结合 S-Plots 和 VIP (variable importance on projection) 值 (图 3B 和 3C), 进一步筛选出 8 个新鲜与蒸制三七根中有较大差异的皂苷, 分别为人参皂苷 Rg1、Rd/Re、Rg2/F2、Rb1, 丙二酰-人参皂苷 Rg1, 三七皂苷 R1, 三七皂苷 R10 和丙二酰-越南参 R13。

选取上述 8 个皂苷化合物做堆积图比较新鲜与蒸制三七根主要皂苷的含量差异。图 4 结果表明, 新鲜三七根切片中皂苷化合物含量较高, 而蒸制三七根切

Table 1 Relative contents of compounds in different parts of fresh and steamed *P. notoginseng* root slices. Relative contents: The signal response of ginsenoside Rg1 in the xylem of fresh *P. notoginseng* was taken as the reference value, and its content was defined as 1

No.	Relative content						LESA-MS [M-Z] ⁻	Molecular formula	Compound
	Fresh root slice			Steamed root slice					
	Xylem	Phloem	Cambium	Xylem	Phloem	Cambium			
1	6.77	12.50	1.18	0.000 3	0.60	0.003 8	387.0 [M+HCOO] ⁻ , 341.3 [M-H] ⁻	C ₁₉ H ₁₈ O ₆ Methylophiopogonanone A ^[16]	
2	0.307	-	0.010 2	<0.000 1	0.06	0.000 5	431.3 [M-H] ⁻	C ₂₁ H ₂₀ O ₁₀ Kaempferol-3- <i>O</i> - α -L-rhamnoside ^[17]	
3	0.071 4	0.019	0.001	<0.000 1	0.024 3	0.000 3	447.2 [M-H] ⁻	C ₂₁ H ₂₀ O ₁₁ Kaempferol-3- <i>O</i> - β -D-galactoside ^[17]	
4	0.012 4	0.099 1	0.002 5	<0.000 1	0.135	0.001 7	475.2 [M-H] ⁻	C ₃₀ H ₅₂ O ₄ Protopanaxatriol ^[18]	
5	-	-	-	<0.000 1	0.024 6	0.000 3	478.4 [M-H] ⁻	C ₃₀ H ₅₄ O ₄ 25-OH-PPD ^[19]	
6	-	-	-	<0.000 1	<0.000 1	0.000 2	491.1 [M-H] ⁻	C ₃₁ H ₅₆ O ₄ 25-OCH ₃ -PPD ^[19]	
7	0.195	0.063 3	0.040 1	<0.000 1	0.017 7	0.000 5	599.1 [M+HCOO] ⁻ , 553.4 [M-H] ⁻	C ₃₀ H ₅₀ O ₉ Notoginsenoside R10 ^[20]	
8	0.017 1	0.025 5	0.002 3	<0.000 1	<0.000 1	0.000 2	621.5 [M-H] ⁻	C ₃₈ H ₆₂ O ₈ Ginsenoside Rh2 ^[21]	
9	0.010 3	0.008 2	-	<0.000 1	0.066 7	0.001 1	625.2 [M-H] ⁻	C ₂₇ H ₃₀ O ₁₇ Quercetin-3- <i>O</i> -sophoroside ^[22]	
10	-	-	-	<0.000 1	0.015 6	0.000 3	651.4 [M-H] ⁻	C ₃₆ H ₆₀ O ₁₀ Notoginsenoside T1 ^[23]	
11	0.038 3	0.008 7	0.003 7	-	0.020 6	0.000 1	653.4 [M-H] ⁻	C ₃₆ H ₆₂ O ₁₀ Notoginsenoside ST1/ notopanaxoside A ^[24,25]	
12	-	-	-	<0.000 1	0.025 4	<0.000 1	655.5 [M-H] ⁻	C ₃₆ H ₆₄ O ₁₀ Notoginsenoside SFt2 ^[26]	
13	-	-	-	-	0.020 4	<0.000 1	661.4 [M-H] ⁻	C ₃₈ H ₆₂ O ₉ Ginsenoside Rs7 ^[27]	
14	0.093 5	0.071 7	0.012 2	<0.000 1	0.025 2	0.000 3	683.4 [M+HCOO] ⁻ , 637.4 [M-H] ⁻	C ₃₆ H ₆₂ O ₉ Ginsenoside Rh1/F1 ^[21]	
15	0.009 2	0.007 2	0.016 4	<0.000 1	0.026 3	0.000 2	725.4 [M+HCOO] ⁻ , 679.4 [M-H] ⁻	C ₃₈ H ₆₄ O ₁₀ 6'- <i>O</i> -acetyl-ginsenoside Rh1 ^[28]	
16	0.007 8	0.002 8	0.002 5	<0.000 1	0.022 7	0.000 1	753.4 [M-H] ⁻	C ₄₁ H ₇₀ O ₁₂ Ginsenoside Mc ^[29,30]	
17	0.066 2	0.055 5	0.035 4	<0.000 1	0.11	0.001	781.0 [M-H] ⁻	C ₄₂ H ₇₀ O ₁₃ Ginsenoside Rh17 ^[31]	
18	-	-	-	<0.000 1	0.044 8	0.000 2	811.4 [M+HCOO] ⁻ , 811.7 [M+HCOO] ⁻	C ₄₂ H ₇₀ O ₁₂ Ginsenoside Rg5/Rk1 ^[32]	
19	0.111	0.041 7	0.026 1	<0.000 1	0.041 9	<0.000 1	815.5 [M+HCOO] ⁻ , 769.5 [M-H] ⁻	C ₄₁ H ₇₀ O ₁₃ Notoginsenoside R2 ^[28]	
20	0.009 6	0.013 5	-	<0.000 1	0.020 5	<0.000 1	831.5 [M+HCOO] ⁻ , 785.5 [M] ⁻	C ₄₂ H ₇₂ O ₁₃ Ginsenoside Rg3 ^[28]	
21	0.273	0.104	0.081	<0.000 1	0.012 2	0.000 2	829.5 [M+HCOO] ⁻	C ₄₂ H ₇₂ O ₁₃ Ginsenoside Rg2/F2 ^[33]	
22	1	0.524	0.319	<0.000 1	0.035 8	0.000 2	845.5 [M+HCOO] ⁻ , 799.5 [M-H] ⁻	C ₄₂ H ₇₂ O ₁₄ Ginsenoside Rg1 ^[28]	
23	-	-	-	<0.000 1	0.038 1	<0.000 1	881.4 [M+HCOO] ⁻ , 825.5 [M-H] ⁻	C ₄₄ H ₇₄ O ₁₄ Ginsenoside Rs3 ^[34]	
24	0.345	0.432	0.342	<0.000 1	0.018 9	<0.000 1	885.0 [M-H] ⁻	C ₄₅ H ₇₄ O ₁₇ Malonyl-ginsenoside Rg1 ^[35]	
25	-	-	-	<0.000 1	0.016 2	<0.000 1	897.5 [M-H] ⁻	C ₄₄ H ₇₈ O ₁₆ Notoginsenoside SFt3/SFt4 ^[36]	
26	0.028 8	-	0.002 1	-	-	<0.000 1	913.4 [M+HCOO] ⁻	C ₄₆ H ₇₆ O ₁₅ Koryoginsenoside R1 ^[37]	
27	0.034 2	0.018 9	0.011 9	<0.000 1	0.016 1	0.000 2	961.0 [M+HCOO] ⁻ , 915.5 [M-H] ⁻	C ₄₇ H ₈₀ O ₁₇ Notoginsenoside ST4 ^[38]	
28	0.351	0.155	0.075 8	<0.000 1	0.016 7	<0.000 1	977.5 [M+HCOO] ⁻ , 932.5 [M-H] ⁻	C ₄₇ H ₈₀ O ₁₈ Notoginsenoside R1 ^[39]	
29	-	-	-	-	0.017	0.000 1	977.7 [M+HCOO] ⁻ , 931.0 [M-H] ⁻	C ₄₇ H ₈₀ O ₁₈ Notoginsenoside Ft3 ^[40]	
30	0.008 5	0.007 3	0.006	<0.000 1	0.007 9	<0.000 1	987.5 [M-H] ⁻	C ₅₄ H ₈₀ O ₁₉ Acetyl-ginsenoside Re/Rd ^[41]	
31	0.323	0.274	0.139	<0.000 1	0.048 4	0.000 2	991.5 [M+HCOO] ⁻ , 945.5 [M-H] ⁻	C ₄₈ H ₈₂ O ₁₈ Ginsenoside Rd/Re ^[28]	
32	0.033 6	0.037	0.007 4	<0.000 1	0.034 8	0.000 2	993.5 [M+HCOO] ⁻	C ₄₇ H ₈₀ O ₁₉ Notoginsenoside H ^[39]	
33	0.022 9	0.017 8	0.010 1	-	0.032 7	<0.000 1	1 007.4 [M+HCOO] ⁻	C ₄₈ H ₈₂ O ₁₉ Notoginsenoside M/N ^[42]	
34	0.024 5	0.14	0.045 8	-	0.033	<0.000 1	1 031.3 [M-H] ⁻	C ₅₁ H ₈₄ O ₂₁ Mal-ginsenoside Re/Rd ^[43]	
35	0.173	0.062 5	0.064 6	-	0.022 9	<0.000 1	1 065.5 [M-H] ⁻	C ₅₁ H ₈₆ O ₂₃ Mal-vinaginsenoside R13 ^[44]	
36	0.018 1	0.009 5	0.005 7	<0.000 1	0.025 2	0.000 1	1 093.4 [M+HCOO] ⁻ , 1 047.5 [M-H] ⁻	C ₅₂ H ₈₈ O ₂₁ Notoginsenoside O/P ^[45]	
37	0.023 9	0.017 4	-	<0.000 1	0.016 2	0.096 2	1 123.5 [M+HCOO] ⁻ , 1 077.2 [M-H] ⁻	C ₅₃ H ₉₀ O ₂₂ Ginsenoside Rb2/Rb3/Rc ^[39]	
38	0.035 5	0.014 1	0.003 1	-	0.014	<0.000 1	1 139.6 [M+HCOO] ⁻	C ₅₃ H ₉₀ O ₂₃ Yesanchinoside H ^[46]	
	0.018 1	-	0.005 7	-	0.022 7	0.000 2	1 093.6 [M-H] ⁻		
39	0.226	0.041 7	0.081 6	<0.000 1	0.057 4	0.000 2	1 153.6 [M+HCOO] ⁻ , 1 107.6 [M-H] ⁻	C ₅₄ H ₉₂ O ₂₃ Ginsenoside Rb1 ^[28]	
40	0.024 9	0.019 2	0.006 1	-	-	-	1 149.5 [M-H] ⁻	C ₆₀ H ₉₀ O ₂₄ Acetyl-ginsenoside Rb1 ^[41]	
41	0.024 9	0.014 9	0.008	-	0.043 8	<0.000 1	1 209.6 [M+HCOO] ⁻ , 1 163.5 [M-H] ⁻	C ₅₆ H ₉₂ O ₂₅ Mal-ginsenoside Rb2/Rb3/Rc ^[43]	
42	0.037 3	0.004 8	0.004 2	<0.000 1	0.009 2	<0.000 1	1 165.5 [M+HCOO] ⁻	C ₅₅ H ₉₂ O ₂₃ Ginsenoside Rs1 ^[47]	
43	0.053 2	0.026 3	0.017 4	<0.000 1	0.010 3	<0.000 1	1 239.5 [M+HCOO] ⁻ , 1 193.6 [M-H] ⁻	C ₅₇ H ₉₄ O ₂₆ Mal-ginsenoside Rb1 ^[43]	
44	0.079 6	0.012 1	0.010 1	-	-	-	1 255.6 [M+HCOO] ⁻ , 1 209.6 [M-H] ⁻	C ₅₈ H ₉₈ O ₂₆ Ginsenoside Ra2 ^[48]	
45	0.053 2	0.026 3	0.017 4	<0.000 1	0.011 3	<0.000 1	1 285.4 [M+HCOO] ⁻ , 1 239.5 [M-H] ⁻	C ₅₉ H ₁₀₀ O ₂₇ Notoginsenoside R4 ^[49] / ginsenoside Ra3 ^[28]	
46	0.027 6	0.008 4	-	<0.000 1	0.015	0.000 1	1 297.8 [M+HCOO] ⁻ , 1 251.4 [M-H] ⁻	C ₆₀ H ₉₈ O ₂₇ Ginsenoside Ra5 ^[50]	
47	0.078 3	0.011 8	0.003 2	-	0.007 6	0.000 5	1 316.0 [M+HCOO] ⁻	C ₆₀ H ₁₀₂ O ₂₈ Ginsenoside Ra0 ^[51,52]	
48	0.044 7	0.011 1	0.014 4	-	0.033	0.000 3	1 371.5 [M+HCOO] ⁻ , 1 325.5 [M-H] ⁻	C ₆₂ H ₁₀₂ O ₃₀ Mal-notoginsenoside R4/ Mal-ginsenoside Ra3 ^[53]	
49	0.026 4	0.019 5	0.009 1	-	0.012 4	0.000 6	1 387.4 [M+HCOO] ⁻ , 1 341.6 [M-H] ⁻	C ₆₂ H ₁₀₆ O ₃₀ Notoginsenoside Q/S ^[45]	

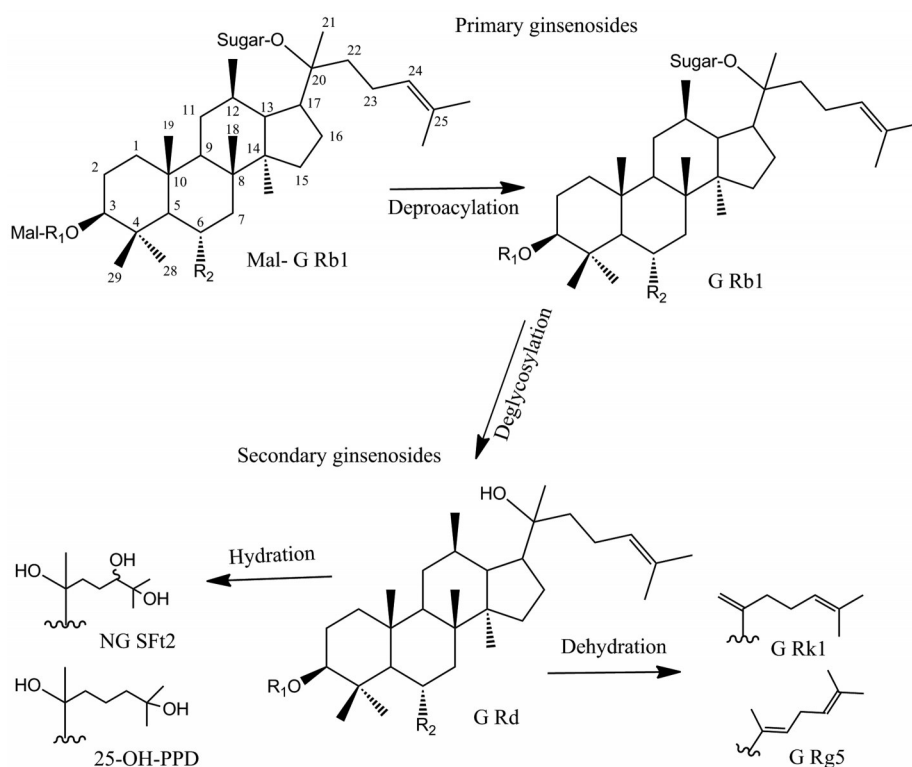


Figure 2 Proposed transformation of saponins in the process of steaming of *P. notoginseng* (take ginsenoside Rb1 for example). Mal-G Rb1: Malonyl-ginsenoside Rb1; G Rb1: Ginsenoside Rb1; G Rd: Ginsenoside Rd; NG SFt2: Notoginsenoside SFt2; G Rk1: Ginsenoside Rk1; G Rg5: Ginsenoside Rg5

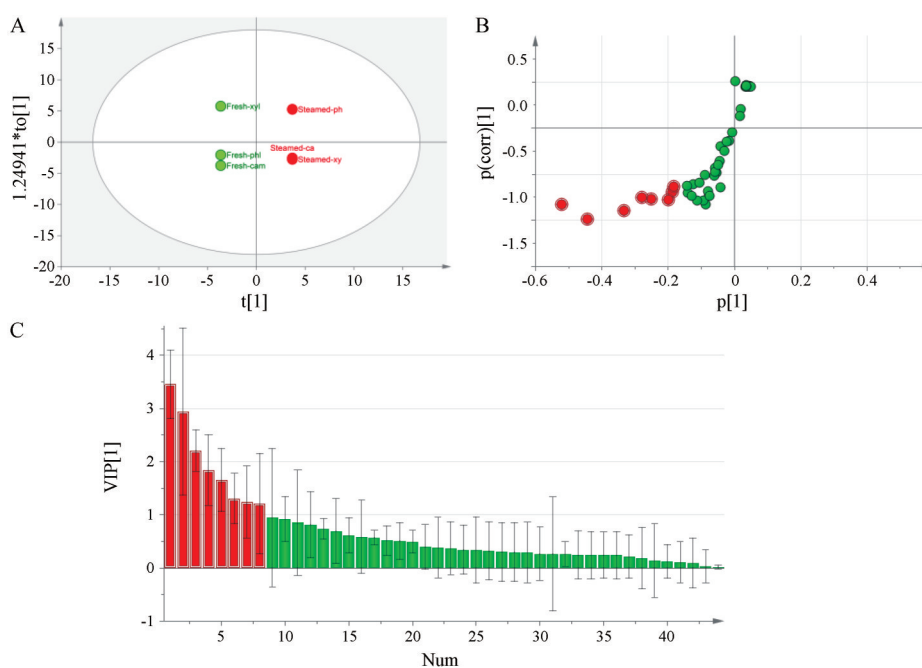


Figure 3 OPLS-DA results of different parts and saponin compounds in fresh and steamed root slices. A: OPLS-DA score chart; B: OPLS-DA S-Plots loading chart; C: OPLS-DA VIP chart

片中丙二酰类皂苷化合物相对较少, 与直观分析结果相吻合。这些皂苷成分和相对含量的差异均可在一定程度上区别新鲜三七根和蒸制三七根的不同部位。

讨论

本研究基于新鲜三七与蒸制三七的药效差异对新鲜与蒸制三七进行化学成分的差异性比较。达玛烷型

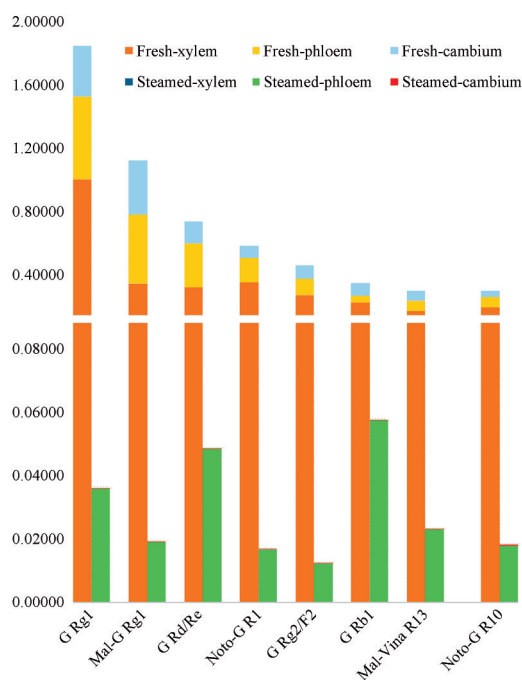


Figure 4 Accumulation diagram of relative contents of major saponins in the fresh and steamed roots slices were detected by LESA-MS. G Rg1: Ginsenoside Rg1; Mal-G Rg1: Malonyl-ginsenoside Rg1; G Rd/Re: Ginsenoside Rd/Re; Noto-G R1: Notoginsenoside R1; G Rg2/F2: Ginsenoside Rg2/F2; Mal-Vina R13: Malonyl-vinaginsenoside R13; Noto-G R10: Notoginsenoside R10

三萜皂苷是三七中主要的化学成分和药效成分, 遂以该类皂苷作新鲜三七与蒸制三七的差异对比。目前, 对三七中药材的基础成分研究大多采用繁琐的化学和液相方法, 该研究首次建立了一种稳健、快速、节约溶剂的三七根切片 LESA-MS 测定方法, 只需将三七根切成表面平整的切片形式, 只需 2.0 μL 提取溶剂, 即可在线操作对切片中化合物进行快速灵敏的分析。在整个分析过程中, 不需要破碎、萃取、分离等繁琐的传统工艺, 能够大大节省物力、人力与时间。该过程操作简便, 可以快速的表征新鲜三七根切片和蒸制三七根切片中皂苷的成分, 从而为中药的快速检测和测定提供了一种新的参考方法。本研究所用的仪器为离子阱质谱, 可提供多级碎片离子信息, 今后如将 LESA 离子源与高分辨质谱技术联用, 可大大提升对于未知化合物结构快速鉴别能力。

本研究通过快速表征皂苷成分, 共鉴定出 49 个化合物, 其中 45 个为皂苷类化合物。对比新鲜与蒸制三七根切片中皂苷成分可知, 人参皂苷 Rg1 为含量最高的皂苷, 在新鲜与蒸制三七的木质部、韧皮部和形成层部位分布广泛。新鲜三七中的丙二酰类皂苷及大极性皂苷类化合物成分较多, 而在蒸制三七中, 弱极性皂苷

化合物则有较多显现, 表明新鲜三七与蒸制三七的皂苷成分和含量均具有较大的差异。根据结构式推论, 新鲜三七中强极性成分可能经过高温高压蒸制脱糖脱水而转化成小极性皂苷成分, 与其他文献研究结果一致^[54]。同时, 本研究也采用多元统计方法 OPLS-DA 对新鲜与蒸制三七进行差异性对比, 所得模型为直观分析的结果提供良好的解释能力与贡献能力。因此, 通过直观解析与数理统计学一体化分析可以直接快速的找出具有较大差异性的皂苷化合物, 为新鲜与蒸制三七皂苷成分的差异性提供解释。

本研究采用快速且无需色谱分离的 LESA-MS 方法, 能够对三七切片中含量低, 难分离的皂苷化合物进行高通量的快速表征, 从而为三七中皂苷成分的鉴定提供更简便、快速的途径。目前, 相较于新鲜三七, 蒸制三七在抗肿瘤、抗心血管系统疾病、抗氧化等方面表现出更加显著的疗效。因此, 对新鲜三七和蒸制三七根的皂苷成分差异进行无损性的快速辨析将为探究三七中药材生熟药效的差异提供物质基础。

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