PHCOG MAG

Peptides from two sanguinovorous leeches analyzed by ultra-performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometric detector

Ling Xiao^{1,2}, Jing Nie², Danping Li², Keli Chen¹

¹College of Pharmacy, Hubei University of Chinese Medicine, ²Department of TCM, Hubei Provincial Institute for Food and Drug Control, Wuhan, China

Submitted: 10-02-2014

Revised: 25-04-2014

Published: 21-01-2015

ABSTRACT

Background: Hirudo nipponica Whitman and Poecilobdella manillensis Lesson fall into the family of Hirudinidae Whitman, both of them are sanguinovorous leeches and used a anticoagulant medicines in China. Their medicinal parts are the dried bodies. However, the peptides in the dried body of the two leeches have not been very clear up to now. Objective: To analyze the peptides from two sanguinovorous leeches, H. nipponica and P. manillensis. Materials and Methods: In this article it is reported that the peptides were obtained from anticoagulant active extracted parts of dried bodies of the two leeches and their molecular weights were analyzed by ultra-performance liquid chromatography with electrospray ionization quadrupole time-of-flight mass spectrometry mass spectrometric detector online. Results: Three peptide components were identified from H. nipponica with their molecular weight separately 14998, 15988, and 15956, six peptide components were identified from P. manillensis with molecular weight 9590, 13642, 14998, 17631, 15988, and 16567. Two of peptides from P. manillensis have the same molecular weight 14998 and 15988 as that in H. nipponica. Conclusion: And the two peptides are the main peaks in the base peak ion chromatogram because they occupied a large ratio of total base peak area. Hence the composition of the extracted active part of the two leeches are very close, difference is in that the extract of P. manillensis has more small peptide peaks, but the extract of *H. nipponica* has not. Furthermore, the tryptic digestion hydrolysates of the extracted active part of each sample were analyzed and the results showed that there were four peaks which only exist in P. manillensis, but not in Hirudo nipponia. They may be the identified peak between the two leeches. This work support the viewpoint that P. manillensis can be used as a medicinal leech as H. nipponia and these peptide components of dried bodies of the two species leeches are a basis for their chemical identification and further investigations in active action.

Key words: *Hirudo nipponia* Whitman, leech, peptide, *Poecilobdella manillensis* Lesson, ultra performance liquid chromatography-electrospray ionization quadrupole time-of-flight

INTRODUCTION

Hirudo nipponia Whitman has been one important medicine in Traditional Chinese Medicine, which effects is promoting blood circulation and removing blood stasis. Now it is mainly used in cerebral thrombosis and cerebral apoplexy in clinical treating and recorded in China Pharmacopoeia. *Poecilobdella manillensis* Lesson we is also used as a medicinal leech in the folk with the same function as *H. nipponia* Whitman, but

Address for correspondence: Prof. Keli Chen, Hubei University of Chinese Medicine, Wuhan - 430 065, China. E-mail: kelichen@126.com it has not been recorded in China Pharmacopoeia. Both of them fall into the famliy of *Hirudinidae* Whitman and are sanguinovorous leeches. Hirudin is a well-known and the strongest thrombin inhibitor in the world, which was isolated from saliva of *Hirudo nipponica* Whitman.^[1] Bufrudin was isolated from saliva of *P. manillensis* Lesson, which has the same effect and 50-60% same amino acid sequence as Hirudin.^[2] Baskova reported the proteins and peptides which was separated from the secretion of *Hirudo verbana*'s, *Hirudo medicinalis*'s and *Hirudo orientalis*'s salivary gland analyzed by SELDI mass spectrometry (MS) and two-dimensional electrophoresis in 2008.^[3] In China, the dried whole bodies of the two leeches are the medicinal parts in Traditional Chinese Medicine for many years, but up to now the



dried whole bodies of these leeches have yet seldom been analyzed to identify their biologically active part and peptide compounds. In this paper, we found an active part, the extract solution, and used the methods of ultra-performance liquid chromatography (UPLC) coupled with electrospray ionization quadrupole time-of-flight (ESI-Q-TOF) mass spectrometric detector to have determined the molecular weight of some peptide compounds in the extracted active part of the two leech species. No doubt, these peptides do not cover the whole compounds of the biologically active part of the dried bodies, but it can provide with new method and clue for the two sanguinovorous leeches' chemical identification and discovering new active compounds.

METERIALS AND METHODS

Instrument and reagent

Chromotographic analysis was conducted on a Waters series ultra-fast HPLC system Waters ACQUITY UPLC-Xevo G2-Q-TOF(ACQUIT UPLC Online Community). Chromatographic column, ACQUITY UPLC BEH C₁₈(1.7 μ m, 100 × 2.1 mm), was used for analysis. Milli-Q Water Purification System was used to purify the water for a test. Trypsin was from Promega Corporation. Sysmex CA-500 Coagulation analyzer was from Sysmex Corporation. Thrombocheck activated partial thromboplastin time (APTT) reagent and Thrombocheck prothrombin time (PT) reagent were from Sysmex Corporation. Acetonitrile and formic acid (chromatographic grade) were from TEDIA, USA.

Preparetion of sample

All dried samples of the bodies of *H. nipponia* Whitman and *P. manillensis* Lesson were get rid of the internal organs, and then were ground into powder. Transfer about 1-g sample powder to a flask, add 50 ml 0.9% sodium chloride and swirling for 1 h at 37°C, centrifuge for 5 min at 10,000 rpm, to obtain the extract solution from the supernatant liquor.

Anticoagulant effect in vitro determination

Human plasma 100 μ L were added to an eppendorf tube, and 100 μ L the extract solution of each sample or 0.9% sodium chloride, and then 100 μ L APTT or PT reagent were added into the tube. The value of APTT and PT of the mixture were detected by Sysmex CA-500 Coagulation analyzer. This operation was repeated 5 times and the mean value was taken as the final result.

Tryptic digestion of the sample

Samples were divided into two groups according to species, *P. manillensis* Lesson and *H. nipponia* Whitman, each group have four batches of samples. The extract method of each batch samples is as the same as the "preparation of sample".

Transfer 50 μ L of the extract solution to an eppendorf tube, diluted with 450 μ L 1% NH₄HCO₃, add 10 μ L 100 mM DTT in 1% NH₄HCO₃ and incubate the solution at 60°C for 15 min, add 50 μ L trypsin in 1% NH₄HCO₃ and leave the digested solution overnight at 37°C.

Liquid chromatography conditions

Acetonitrile: Water (2:98) with 0.2% formic acid was used as mobile phase A, and acetonitrile: Water (98:2) with 0.2% formic acid was used as mobile phase B, gradient elution, 0-60 min, 100% A \rightarrow 100% B. The column temperature was set at 65°C. The flow rate was set at 0.2 mL/min, and the injection volume was fixed at 1 µL.

Mass spectrometry conditions

The MS detection was performed on an ESI-Q-TOF MS System. The ionization mode was positive electrospray (ESI⁺). The desolvation temperature was fixed at 350°C with desolvation gas flow was fixed at 600 L/h, and the source temperature was fixed at 120°C. The cone voltage was fixed at 40 V, and the capillary voltage was fixed at 3 kV. The collision energies were fixed as 6 eV for low-energy scan, and 20-40 eV for high-energy scan. The LC-MS data acquisition mode was MS^E, which was controlled by MassLynx 4.0 Mass Spectrometry Software.(Waters SCN871).

Data processing

The extract solution and digest solution of the samples were injected into the UPLC at the above chromatography condition to acquire the MS data. Postacquisition data processing was performed by Mass Ent 1 and MarkerLynx XS, which are the "application manager" for MassLynx software. (Waters SCN871). Mass Ent 1 was used to transform multiple charges mass peak into a single charge mass peak. MarkerLnyx XS were used to identify the difference through the principal component analysis (PCA).

RESULTS AND DISCUSSION

The anticoagulant effect *in vitro* of the extracted solutions of samples

Prothrombin time APTT. Both of them are widely used in evaluating the anticoagulant effect. The results showed that the extracted solutions of two leech species have an anticoagulant effect. The active action of *P. manillensis* is stronger than the *H. nipponia*'s [Table 1].

Results of time-of-flight mass spectrometry analysis Electrospray ionization, a soft ionization technology, is an effective measure to analyze the protein and peptide,^[4-8] in which ESI source is coupled with LC online. When a molecule of protein or peptide is analyzed by ESI source, a series of multiple-charge icons peaks will be detected by TOF mass spectrometric detector. The molecular weight of every peak (M) can be calculated according to the mass-to-charge ratio (m/z) and the ion charges (e): m/z = (M + e)/e. MassEnt 1 can transfer the multiple charges mass peaks into a single charge mass peak and calibrate the calculated molecular weights.

Three peptide components named H1 ~ H3 were detected from *H. nipponia* and their molecular weight were 14998 (H1), 15988 (H2) and 15956 (H3), six peptide components named P1 ~ P6 weredetected from *P. manillensis* and their molecular weight were 9590 (P1), 13642 (P2), 14998 (P3), 17631(P4), 15988 (P5), and 16567 (P6). Figure 1 is the base peak ion chromatogram (BPI) of extracting solution of two leeches and 0.9% sodium chloride solution. Figure 2 is the multiple charges mass peak given by the ESI⁺ (left) and a single charge mass peak given by MassEnt 1 (right) of H1 ~ H3, P1 ~ P6. Table 2 shows the MS information of each peptide peak: The mass-to-charge ratio (m/z), the ion charges (e) of the multiple-charge ions peaks given by the ESI⁺, the calculated molecular weights and the weight given by MassEnt 1.

From Figure 1 it is noticed that H1 (detected from *H. nipponia*) and P3 (detected from *P. manillensis*) have the same molecular weight, and H2 (detected from *H. nipponia*) and P5 (detected from *P. manillensis*) also have the same molecular weight. And the two peptide peaks are the main peaks in the BPI of each species because they occupied a large ratio of total base peak area. Hence, the main composition of the extracted active part of the two leeches are very close or similar, difference is in that the extraction of *P. manillensis* has more small peptide peaks, but the extract of *H. nipponica* has not.

Analysis of tryptic digestion of samples

The results of TOF-MS analysis showed that *P. manillensis* and *H. nipponia* are close to each other in main peptide composition of dried bodies. Because of the complexity of composition, it is different to discriminate them in the dried bodies and the extraction only by vision observation.

The method of digestion was used to analyze the protein hydrolysates of the samples in order to identify them. The extracted solution of *P. manillensis* and *H. nipponia*

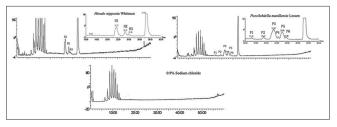


Figure 1: Base peak ion chromatograms of extracted solutions of two leeches and 0.9% sodium chloride solution

were digested with trypsase, then, the hydrolysates were analyzed by LC-MS. Figure 3 shows the BPIs obtained from hydrolysates of *P. manillensis* and *H. nipponia*. Because of the complexity of the sample, it is difficult to discriminate them only by visual observation of the two chromatograms. Hence, the data were processed by MarkerLnyx XS software. The software converts each data point into an exact mass/retention time (EMRT) pair and tabulates the results into a two-dimensional matrix, after a two-dimensional matrix is obtained, the extended statistics module PCA can be carried out. From the PCA plot, any two groups of samples can be paired for an orthogonal partial least squares data analysis (OPLS-DA).^[9] The data of samples

Table 1: The anticoagulant effect of samples *in vitro* ($X\pm S$, *n*=5)

Group	Name	APTT	PT
sampl e	Hirudo nipponi a Whitman	80.7±2.02*	28.4±2.12*
sampl e	Poecilo bdella manillensis Lesson	93.4±1.65*	38.5±1.78*
negative	0.9% sodium chloride solution	28.03±2.14	11.3 ± 1.82

Table 2: The peptide analysis of two species leech

Sample	Peak number	Retention time (min)	m/z (Mass-to-charge ratio)	e (Ion charges)	Calculated weights	Weights b Mass Ent
Hirudo nipponia Whitman	H1	21.72	790.3231 [M+19H] ¹⁹⁺ 834.1744 [M+18H] ¹⁸⁺ 883.2406 [M+17H] ¹⁷⁺ 938.3778 [M+16H] ¹⁶⁺	19 18 17 16	14997.14 14997.14 14998.09 14998.04	14998
			1000.8737[M+15H] ¹⁵⁺ 1072.3583[M+14H] ¹⁴⁺ 1154.8523[M+13H] ¹³⁺	15 14 13	14998.10 14999.01 15000.08	
	H2	23.35	842.4532 [M+19H] ¹⁹⁺ 889.2002 [M+18H] ¹⁸⁺	19 18	15987.62 15987.64	
			941.4465 [M+17H] ¹⁷⁺ 1000.2249 [M+16H] ¹⁶⁺ 1066.8977 [M+15H] ¹⁵⁺ 1142.9635 [M+14H] ¹⁴⁺	17 16 15 14	15987.59 15987.60 15988.46 15987.49	15988
			1230.8134 [M+13H] ¹³⁺	14	15987.57	
	H3 2.		840.7157[M+19H] ¹⁹⁺ 887.4274[M+18H] ¹⁸⁺	19 18	15954.60 15955.69	
		23.96	939.4463[M+17H] ¹⁷⁺ 998.2278[M+16H] ¹⁶⁺	17 16	15953.59 15955.64	15956
			1064.6343[M+15H] ¹⁵⁺ 1140.6901[M+14H] ¹⁴⁺ 1228.3045[M+13H] ¹³⁺	15 14 13	15954.51 15955.66 15954.96	
	P1		872.6291[M+11H] ¹¹⁺ 959.9008[M+10H] ¹⁰⁺	11 10	9587.92 9589.01	
		17.16	1066.4421[M+9H] ⁹⁺ 1199.8658[M+8H] ⁸⁺ 1370.9940[M+7H] ⁷⁺	9 8 7	9588.98 9590.93 9589.96	9590
	P2 1		910.5386[M+15H] ¹⁵⁺ 975.5080[M+14H] ¹⁴⁺	15 14	13643.08 13643.11	
		19.43	1050.4532[M+13H] ¹³⁺ 1137.9208[M+12H] ¹²⁺ 1241.1839[M+11H] ¹¹⁺	13 12 11	13642.89 13643.05 13642.02	13642
			1365.2053[M+10H] ¹⁰⁺ 1516.8870[M+9H] ⁹⁺	10 9	13642.05 13642.98	
Poecilobdella manillensis Lesson	P3 21.42		790.2769 [M+19H] ¹⁹⁺ 834.1862 [M+18H] ¹⁸⁺	19 18	14996.26 14997.35	
		21.42	883.2527 [M+17H] ¹⁷⁺ 938.3275[M+16H] ¹⁶⁺ 1000.8867[M+15H] ¹⁵⁺ 1072.3717[M+14H] ¹⁴⁺	17 16 15 14	14998.30 14997.24 14998.30 14999.20	14998
			1154.8862[M+13H] ¹³⁺	13	15000.52	
	P4 2	22.26	1176.4211[M+15H] ¹⁵⁺ 1260.3048[M+14H] ¹⁴⁺ 1357.2592[M+13H] ¹³⁺ 1470.2544[M+12H] ¹²⁺	15 14 13 12	17631.31 17630.27 17631.37 17631.05	17631
			1603.7512[M+11H] ¹¹⁺ 1764.1368[M+10H] ¹⁰⁺	11 10	17630.26 17631.37	
	P5 23.08		842.4561[M+19H] ¹⁹⁺ 889.1513 [M+18H] ¹⁸⁺ 041.4501 [M+17H] ¹⁷⁺	19 18 17	15987.67 15986.72 15987.80	
		23.08	$\begin{array}{l} 941.4591 [M+17H]^{17+} \\ 1000.1730 [M+16H]^{16+} \\ 1066.9244 [M+15H]^{15+} \\ 1142.9081 [M+14H]^{14+} \\ 1230.7559 [M+13H]^{13+} \end{array}$	17 16 15 14 13	15987.80 15986.77 15988.87 15986.71 15986.83	15988
		0.000	1275.3938[M+13H] ¹³⁺ 1381.5940[M+12H] ¹²⁺	13 12	16567.12 16567.13	
	P6	23.85	1507.1006[M+11H] ¹¹⁺ 1657.6050[M+10H] ¹⁰⁺ 1841.6814[M+9H] ⁹⁺	11 10 9	16567.11 16566.05 16566.13	16567

Xiao, et al.: Peptides from two leeches analyzed by UPLC-ESI-Q-TOF

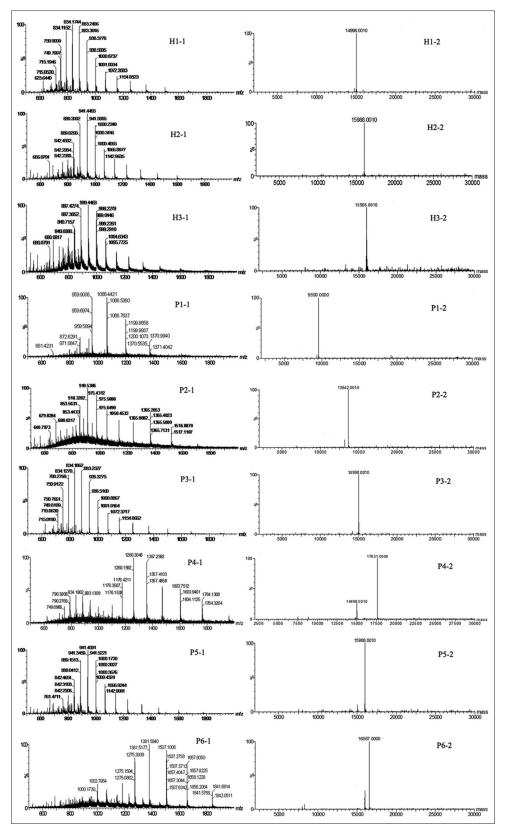


Figure 2: Multiple charges mass peak (left) and single charge mass peak (right) of H1~H3, P1~P6

were divided into two groups: Group 1 for *P. manillensis* and group 2 for *H. nipponia*. The resulting scatter plot (S-plot) from the OPLS-DA analysis can clearly display significant

markers that can differentiate the sample groups. Figure 4 shows the loading plot (S-plot) of the OPLS-DA result. The points on the Figure 4 are the EMRTs plotted by

covariance (X-axis) and correlation (Y-axis) values. The point the farther away from the X-axis is the greater the contribution to the difference between the two groups, while, the farther along the Y-axis is the higher the reliability of the analytical result. Every point is an EMRT pairs, which nearest the upper right corner of the S-plot are the leading contributing markers from the *P. manillensis* group, while, nearest the lower left corner of the S-plot are the leading contributing markers from the *H. nipponia* group [Figure 4].

Table 3 shows the combined list of the top 15 leading EMRT pairs obtained from the S-plot for both sample groups and the factor of change for each of the markers identified. The value of a factor of change indicates the intensity of difference between the peaks. No. 2-7 peaks are the markers which are highest in concentration in the P. manillensis group, and No. 10-15 peaks are the markers which are highest in concentration in the H. nipponia group. This table also can be displayed as a column plot, as shown in Figure 5. As a result, the markers tend to stretch out more toward the end of the S-plot and show the highest factor of change between the two groups. Hence, according to the highest factor of change and higher peak intensities, four EMRT pairs, which were detected only in P. manillensis and weren't in H. nipponia, can be regarded as useful for discriminating two species leeches: Peak 2 (24.74_492.2604), peak 3 (34.74_633.4192), peak 4 (22.30_711.8679), peak 5 (19.67_611.9691) [Table 3].

Trypsin digests the amino acid sequence in the peptide bond between lysine and arginine. Four peaks from the trypsin hydrolysate were detected in *P. manillensis* and weren't in *H. nipponia*, so that they can be taken as the discriminating peaks between the two species leeches.

Peak number	Retention Time	Mass	p[1]P	p(corr)[1]P	Factor of Change	peak intensities of H. nipponia	peak intensities of P. manillensis
1	3.01	120.081	0.11	0.99	2.9	97.6	279.8
2	24.74	492.2604	0.11	0.99	10000	1.63E-06	167
3	34.74	633.4192	0.09	0.87	10000	3.68E-06	139.3
4	22.30	711.8679	0.09	0.95	10000	7.47E-07	138.6
5	19.67	611.9691	0.08	0.94	10000	6.25E-07	114.3
6	24.73	766.0628	0.08	0.99	10000	1.40E-06	95.7
7	47.82	616.175	0.07	0.99	10000	1.90E-06	85.9
8	44.73	803.873	-0.06	-0.98	14.3	53.7	3.75
9	44.73	703.5149	-0.06	-0.99	17.9	54.8	3.06
10	24.91	492.2604	-0.06	-1.00	10000	62.8	1.44E-06
11	45.63	758.28	-0.06	-1.00	10000	49.2	2.77E-07
12	14.31	437.2334	-0.05	-1.00	10000	43.08	1.08E-07
13	45.6	884.4973	-0.05	-1.00	10000	41	3.07E-07
14	19.81	611.9694	-0.04	-1.00	10000	28.3	9.56E-07
15	44.8	703.515	-0.06	-1.00	10000	54.8	1.22E-06

Table 3: Top contributing markers from the two sample groups

CONCLUSION

Hirudotherapy is actively used in clinical practice in Russia, USA, Canada, etc., Hirudin is isolated in 1955

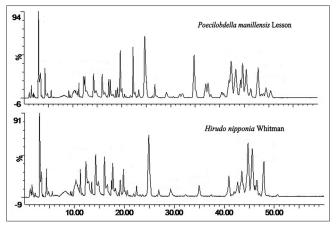


Figure 3: Base peak ion chromatogram from hydrolysates of *Poecilobdella manillensis* Lesson and *Hirudo nipponia* Whitman

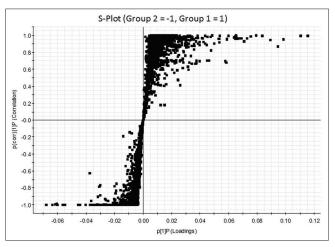


Figure 4: Scatter plot of Group 1 *Poecilobdella manillensis* versus Group 2 *Hirudo nipponia*

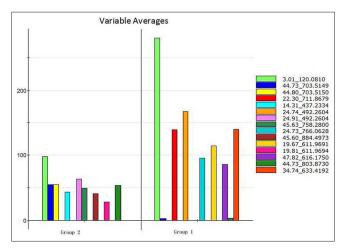


Figure 5: Column plot of the averages of Group 1 Poecilobdella anillensis versus Group 2 Hirudo nipponia

from the secretion of H. nipponia's salivary gland and is the main active compound in the humoral agent in hirudotherapy. In this article, anticoagulant active part was found in the dried body of the two species leeches, H. nipponia and P. manillensis, and they may have different anticoagulant action and effect from Hirudin. Furthermore, some peptide components were isolated from the anticoagulant active part of the dried body of the two species leeches. Among them, two of the peptides separately come from the two leeches have the same molecular weight, and the two peptide peaks occupied a very large ratio of total base peak area. Hence, the composition of the extracted active part of the two leeches is very close. This work shows that the dried body of the two species leeches have anticoagulant action, and P. manillensis may be used as a medicinal leech as H. nipponia. The two species leeches and their extraction can be discriminated by tryptic dehydrolytes of the extraction. The peptide components, isolated from the dried body of the two species leeches, are the basis for chemical identification and further investigations in active action.

ACKNOWLEDGEMENT

Technology major projecs of ——The quality and safety of Chinese medicine detection and risk control technology platform(NO. 2014ZX09304-307-02). Major drug discovery national science and technology major projects(2014ZX09304307001)

REFERENCES

- 1. Markwardt F. Untersuchunjen uber hirudin. Naturwissenschaften 1955;42:537-8.
- Electricwala A, Sawyer RT, Jones CP, Atkinson T. Isolation of thrombin inhibitor from the leech *Hirudinaria manillensis*. Blood Coagul Fibrinolysis 1991;2:83-9.
- Baskova IP, Kostrjukova ES, Vlasova MA, Kharitonova OV, Levitskiy SA, Zavalova LL, *et al.* Proteins and peptides of the salivary gland secretion of medicinal leeches *Hirudo verbana*, *H. medicinalis*, and *H. orientalis*. Biochemistry (Mosc) 2008;73:315-20.
- Deutsch EW, Lam H, Aebersold R. Data analysis and bioinformatics tools for tandem mass spectrometry in proteomics. Physiol Genomics 2008;33:18-25.
- Zhang L, Reilly JP. De novo sequencing of tryptic peptides derived from *Deinococcus radiodurans* ribosomal proteins using 157 nm photodissociation MALDI TOF/TOF mass spectrometry. J Proteome Res 2010;9:3025-34.
- 6. Pandey A, Mann M. Proteomics to study genes and genomes. Nature 2000;405:837-46.
- Chen J, Fu H, Chen Y, Zhao YF, Application of mass spectrometry on the sequencing of peptides and proteins. Chin J Org Chem 2002;22:81-90.
- 8. Cramer R. MALDI MS. Methods Mol Biol 2009;564:85-103.
- Yu K, Xin MT, Castro-Perez J, Fu XT, Chen YG, Guo HZ, et al. Using LC–oa-TOF MSE with a multivariate statistical sample profiling strategy to distinguish Chinese red ginseng from Korean red ginseng. Curr Trends Mass Spectrom Suppl 2009;10:93-100.

Cite this article as: Xiao L, Nie J, Li D, Chen K. Peptides from two sanguinovorous leeches analyzed by ultra-performance liquid chromatography coupled with electrospray ionization quadrupole time-of-flight mass spectrometric detector. Phcog Mag 2015;11:32-7.

Source of Support: Nil, Conflict of Interest: None declared.

Author Help: Online submission of the manuscripts

Articles can be submitted online from http://www.journalonweb.com. For online submission, the articles should be prepared in two files (first page file and article file). Images should be submitted separately.

1) First Page File:

Prepare the title page, covering letter, acknowledgement etc. using a word processor program. All information related to your identity should be included here. Use text/rtf/doc/pdf files. Do not zip the files.

2) Article File:

The main text of the article, beginning with the Abstract to References (including tables) should be in this file. Do not include any information (such as acknowledgement, your names in page headers etc.) in this file. Use text/rtf/doc/pdf files. Do not zip the files. Limit the file size to 1 MB. Do not incorporate images in the file. If file size is large, graphs can be submitted separately as images, without their being incorporated in the article file. This will reduce the size of the file.

3) Images:

Submit good quality color images. Each image should be less than 4096 kb (4 MB) in size. The size of the image can be reduced by decreasing the actual height and width of the images (keep up to about 6 inches and up to about 1800 x 1200 pixels). JPEG is the most suitable file format. The image quality should be good enough to judge the scientific value of the image. For the purpose of printing, always retain a good quality, high resolution image. This high resolution image should be sent to the editorial office at the time of sending a revised article.

4) Legends:

Legends for the figures/images should be included at the end of the article file.