

SUPPLEMENTARY MATERIAL

Table 1. Tryptase level, total IgE and specific IgE titers of the patient's serum from the day of anaphylaxis.

Tryptase level	49 µg/L
Total IgE	1106 kU/L
sIgE <i>Apis mellifera</i> (native)	0.24 kU/L
Api m 1	<0.1 kU/L
Api m 2	<0.1 kU/L
Api m 3	0.11 kU/L
Api m 5	<0.1 kU/L
Api m 10	<0.1 kU/L
sIgE <i>Vespula vulgaris</i> (native)	0.51 kU/L
Ves v 1	0.13 kU/L
Ves v 5	0.6 kU/L
CCD (HRP)	0.24 kU/L
CCD (MUXF3)	0.20 kU/L
sIgE snake venom mix (ELISA)	> 100 kU/L

Normal tryptase levels are < 11.4 µg/L, normal total IgE titers are < 150 kU/L, normal sIgE titer are < 0.1 kU/L. sIgE titer to snake venom was measured by using a standard curve of serial dilutions of a patient serum with a known sIgE titer to cat serum albumin.

Table 2. Identification of immuno-reactive spots from 2D analysis by mass spectrometry (MALDI-TOF-TOF; database SwissProt 2017_04.

Spot	ID	Accession number	Number of peptides identified
1	Basic phospholipase A2 CM-II OS= <i>Naja mossambica</i>	P00603	4
2	Basic phospholipase A2 CM-II OS= <i>Naja mossambica</i>	P00603	4
3	Acidic phospholipase A2 CM-I OS= <i>Naja mossambica</i>	P00602	5
4	Phospholipase A2 "basic" OS= <i>Najanigricollis</i> Basic phospholipase A2 homolog 2 OS= <i>Bothrops asper</i> Acidic phospholipase A2 CM-I OS= <i>Naja mossambica</i>	P00605 P24605 P00602	4 1 2
5	Phospholipase A2 "basic" OS= <i>Najanigricollis</i>	P00605	1

Figure 1.

P00605_NAJNG PLA2	1	NLYQFKNM I HCTV S RFW W HFA D YGC C YGRG G KGT F VDDI D RCC Q V H D NC E KAG K M G CWP Y L T I ----- LYK Y A C SQ G K 74
P00630_APIME PLA2	1	----- I I Y G T I W C G H G N KSSGP N EL G R<--- F K H I D A C R T H D M C P D V M S A G E S K H G I T N T A S H T R L S C D D K F 67
P00605_NAJNG PLA2	75	LTC S GG N SKCGAAVCN<--- C D I V A A N C F A G A R I D A N<----- Y N I N<----- E K R C Q 118
P00630_APIME PLA2	68	YD C L K N S A D T I SSY F V G K M Y F N L I D T R G Y K L E H P V T G C G E R T E G R C L H Y T V D K S K P K V Y Q W E D L R K Y 134

Alignment of the protein sequence of PLA2 from snake *Naja nigricollis*(Uniprot accession number P00605) and honey bee *Apis mellifera*, Api m 1(Uniprot accesión number P00630). Identical residues are highlighted in dark grey, similar residues are highlighted in light grey (Clustal W program). The box highlights the calcium binding site of snake PLA2 as described¹.

References:

1. Trento MVC, Sales TA, de Abreu TS, Braga MA, Cesar PHS, Marques TR, Marcussi S. Exploring the structural and functional aspects of the phospholipase A2 from *Naja* spp. Int J Biol Macromol. 2019;140:49-58.