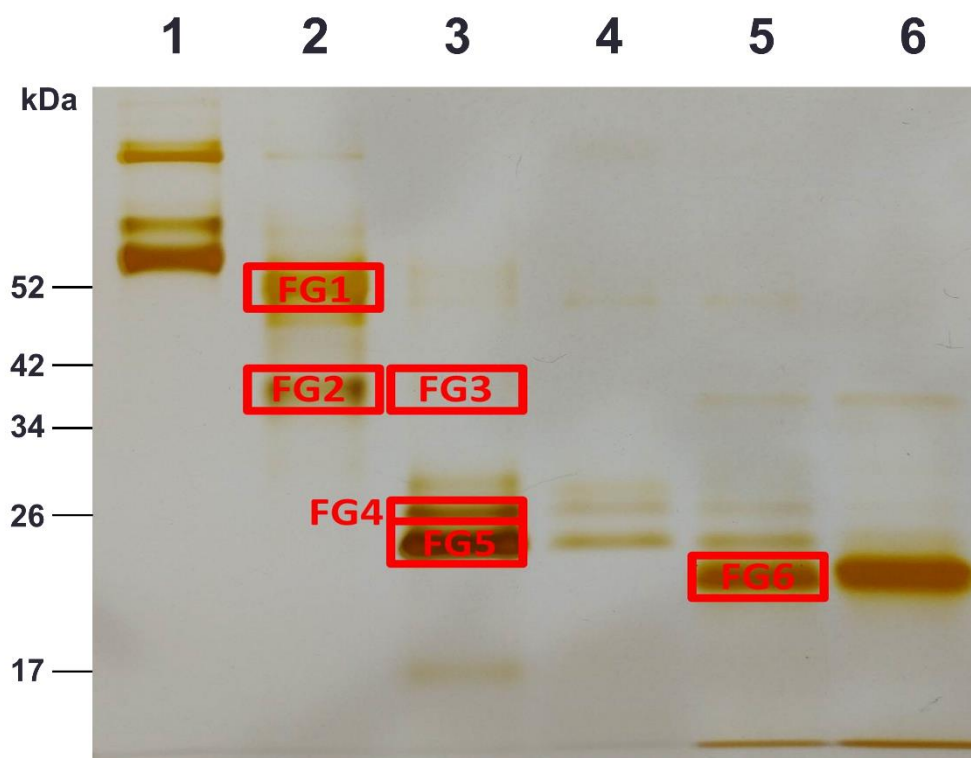


Comparing Traditional and Toxin-Oriented Approaches towards Antivenom Production against *Bitis arietans* Snake Venom

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Supplementary Figure S1. Selection of protein bands from peaks 2, 3 and 5 from fractionating of *B. arietans* venom by size exclusion chromatography to protein identification on LC-MS/MS analysis. 5 µg of each peak were submitted to 12.5% SDS-PAGE, in non-reducing conditions. Selected bands (FG1 to FG6) were cut and submitted to in-gel trypsin digestion.

Supplementary Table S1. Identification of protein bands content by LC-MS/MS from fractions obtained from *B. arietans* venom after size exclusion chromatography.

Sample	Score	Accession	Description	OS
FG1	78	SLRB_BITHR	Snaclec rhinocetin subunit beta	<i>Bitis rhinoceros</i>
	49	A0A140DC05_BITAR	C-type lectin 1b	<i>Bitis arietans</i>

	50	SLA_BITAR	Snaclec bitiscetin subunit alpha	<i>Bitis arietans</i>
	32	A0A0B6VJU7_PROFL	Beta-actin	<i>Probothrops flavoviridis</i>
	32	A0A0F7Z857_CROAD	Poly(a)-specific ribonuclease PARN	<i>Crotalus adamanteus</i>
	28	A0A1I9KNN4_VIPAA	Serine proteinase SP-3	<i>Vipera ammodytes ammodytes</i>
	28	A0A0F7Z8B1_CROAD	E3 ubiquitin-protein ligase	<i>Crotalus adamanteus</i>
	18	A0A6B2FD68_9SAUR	C-type lectin domain-containing protein	<i>Bothriechis nubestris</i>
	17	A0A0K8RXM4_CROHD	Cyclin-dependent kinase 1-like	<i>Crotalus horridus</i>
	131	A0A194AM96_9SAUR	Serine proteinase 17	<i>Agkistrodon piscivorus</i>
	42	A0A194APC3_9SAUR	Serine proteinase 14b	<i>Agkistrodon piscivorus</i>
	58	A0A8A8C7M3_BOTJA	Thrombin-like enzyme	<i>Bothrops jararaca</i>
	56	VSPY_MACLB	Chymotrypsin-like protease VLCTLP	<i>Macrovipera lebetina</i>
	66	E9JG14_ECHCS	Serine protease (fragment)	<i>Echis carinatus sochureki</i>
	61	D5KRX9_ECHOC	Snake venom serine protease (fragment)	<i>Echis ocellatus</i>
	116	A0A6B7FPJ0_VIPAA	Serine proteinase SP-8	<i>Vipera ammodytes ammodytes</i>
	81	J3RZT5_CROAD	Keratin, type I cytoskeletal 19	<i>Crotalus adamanteus</i>
	58	SLRB_BITRH	Snaclec rhinocetin subunit beta	<i>Bitis rhinoceros</i>
	50	SLA_BITAR	Snaclec bitiscetin subunit alpha	<i>Bitis arietans</i>
	57	A0A2H4Z2W1_DABSI	RVV-X heavy chain	<i>Dabola siamensis</i>
	52	A0A0F7ZB82_CROAD	Ankyrin repeat and BTB/POZ domain-containing protein 2	<i>Crotalus adamanteus</i>
	48	A0A1I9KNL5_VIPAA	Serine proteinase SP-2	<i>Vipera ammodytes ammodytes</i>

FG2	45	A0A0B8RV71_CROHD	Keratin, type II cytoskeletal cochleal-like	<i>Crotalus horridus</i>
	44	OXLA_DABRR	L-amino-acid oxidase	<i>Daboia russelii</i>
	43	A0A0F7ZAB1_CROAD	Keratin, type I cytoskeletal 18-like protein	<i>Crotalus adamanteus</i>
	36	A0A1W7RD94_AGKCO	Sister chromatid cohesion protein PDS5	<i>Agkistrodon contortrix contortrix</i>
	34	A0A0F7ZBJ1_CROAD	FAS-associated factor 1	<i>Crotalus adamanteus</i>
	33	A0A1W7REY6_AGKCO	Sister chromatid cohesion protein PDS5 B	<i>Agkistrodon contortrix contortrix</i>
	33	A0A0F7Z8R3_CROAD	Interferon-inducible GTPase 5-like	<i>Crotalus adamanteus</i>
	32	A0A0F7Z8B1_CROAD	E3 ubiquitin-protein ligase	<i>Crotalus adamanteus</i>
	29	A0A194APM7_9SAUR	C-type lectin 9d	<i>Agkistrodon piscivorus</i>
	28	A0A0F7ZEI3_CROAD	Protein c20orf29	<i>Crotalus adamanteus</i>
	27	A0A0F7Z857_CROAD	Poly(A)-specific ribonuclease PARN	<i>Crotalus adamanteus</i>
	26	A0A0F7ZE88_CROAD	Dermatan-sulfate epimerase	<i>Crotalus adamanteus</i>
	26	A0A1W7RF54_AGKCO	NudC domain-containing protein 2	<i>Agkistrodon contortrix contortrix</i>
	25	A0A0F7Z7G6_CROAD	PAT1 1-like protein	<i>Crotalus adamanteus</i>
	23	J3SFD0_CROAD	SURF1-like protein	<i>Crotalus adamanteus</i>
	20	A0A0F7Z8B2_CROAD	Exosome component 10 isoform 1	<i>Crotalus adamanteus</i>
	19	A0A0F7Z5N5_CROAD	Polycomb protein SCMH1 isoform X2	<i>Crotalus adamanteus</i>
	19	A0A0F7ZCD4_CROAD	Oxidation resistance protein 1 isoform X3	<i>Crotalus adamanteus</i>
	19	A0A0F7Z415_CROAD	Serine/threonine-protein kinase TBK1-like protein	<i>Crotalus adamanteus</i>
	18	A0A0F7ZEN3_CROAD	Ankyrin repeat domain-containing protein 12-like	<i>Crotalus adamanteus</i>

	16	A0A0F7Z988_CROAD	Creatine kinase	<i>Crotalus adamanteus</i>
	16	A0A1W7RHL5_AGKCO	Endoplasmic reticulum-Golgi intermediate compartment protein 2-like protein	<i>Agkistrodon contortrix</i> <i>contortrix</i>
	14	A0A0F7Z6K8_CROAD	Dynamin_N domain-containing protein	<i>Crotalus adamanteus</i>
	13	T1E5M6_CROHD	Acetyl-CoA C_myristoyltransferase	<i>Crotalus horridus</i>
FG3	73	VSPC_TRIGA	Snake venom serine protease 2C	<i>Trimeresurus gramineus</i>
	68	J3RZT5_CROAD	Keratin, type I cytoskeletal 19	<i>Crotalus adamanteus</i>
	61	A0A194AM96_9SAUR	Serine proteinase 17	<i>Agkistrodon piscivorus</i>
	54	D5KRX9_ECHOC	Snake venom serine protease (fragment)	<i>Echis ocellatus</i>
	52	D8MIA2_BITRH	Snake venom serine protease (fragment)	<i>Bitis rhinoceros</i>
	47	A0A6B7FPJ0_VIPAA	Serine proteinase SP-8	<i>Vipera ammodytes</i> <i>ammodytes</i>
	42	A0A194APM7_SAUR	C-type lectin 9d	<i>Agkistrodon piscivorus</i>
	33	A0A0F7ZAB1_CROAD	Keratin, type I cytoskeletal 18-like protein	<i>Crotalus adamanteus</i>
	29	A0A5A4WNG2_BITAR	Bitiscetin-3 subunit alpha	<i>Bitis arietans</i>
	24	A0A1W7RC08_AGKCO	Ubiquitin-like conjugating enzyme ATG3-like protein	<i>Agkistrodon contortrix</i> <i>contortrix</i>
	23	A0A0K8S0L2_CROHD	Iron-sulfur cluster assembly 1, mitochondrial-like protein	<i>Crotalus horridus</i>
	23	A0A0F7Z8B1_CROAD	E3 ubiquitin-protein ligase	<i>Crotalus adamanteus</i>
	22	A0A0F7ZEI3_CROAD	Protein C20orf29	<i>Crotalus adamanteus</i>
	22	A0A0F7Z7A4_CROAD	Protein phosphatase 1 regulatory subunit 12B-like	<i>Crotalus adamanteus</i>
	22	A0A1W7RF54_AGKCO	NudC domain-containing protein 2	<i>Agkistrodon contortrix</i> <i>contortrix</i>

	21	A0A0F7ZBJ1_CROAD	FAS-associated factor 1	<i>Crotalus adamanteus</i>
	21	A0A0F7Z958_CROAD	Guanine nucleotide-binding protein subunit alpha-11	<i>Crotalus adamanteus</i>
	20	A0A0K8RWZ3_CROHD	Proteasome-associated protein ECM29-like	<i>Crotalus horridus</i>
	19	A0A0F7Z3D0_CROAD	Ubiquitin C	<i>Crotalus adamanteus</i>
	19	A0A0F7Z434_CROAD	RAD21-like protein	<i>Crotalus adamanteus</i>
	19	A0A2H4Z2W1_DABSI	RVV-X heavy chain	<i>Daboia siamensis</i>
	18	A0A6B2FB67_BOTNI	U6 snRNA-associated Sm-like protein LSm8	<i>Bothriechis nigroviridis</i>
FG4	37	I7JA31_BITRH	C-type lectin-like protein 4	<i>Bitis rhinoceros</i>
	33	A0A0C5DKL1_MACLB	C-type lectin-like protein 3A	<i>Macrovipera lebetina</i>
	33	A0A140DC06_BITAR	C-type lectin 2	<i>Bitis arietans</i>
	33	A0A1B3AXS3_BITAR	C-type lectin-like protein	<i>Bitis arietans</i>
	33	A0A6B7FMP6_VIPAA	C-type lectin-like Snaclec-9	<i>Vipera ammodytes ammodytes</i>
	33	SLAE_MACLB	Snaclec A-14	<i>Macrovipera lebetina</i>
	33	SLAF_MACLB	Snaclec a15	<i>Macrovipera lebetina</i>
	33	SLA_DABPA	Snaclec VP12	<i>Daboia palaestinae</i>
	33	SLA2_BITGA	Snaclec 2	<i>Bitis gabonica</i>
	33	I7JX23_BITRH	C-type lectin-like protein 3	<i>Bitis rhinoceros</i>
	33	K9JBV0_DABSI	P68 alpha subunit	<i>Daboia siamensis</i>
	33	K9JDF2_DABRR	P68 alpha subunit	<i>Daboia russelii limitis</i>
	27	A0A0C5DKK6_MACLB	C-type lectin-like protein 2A	<i>Macrovipera lebetina</i>
	27	SLMA_MACLB	Snaclec macrovipectin subunit alpha	<i>Macrovipera lebetina</i>

FG5	54	SLA_BITAR	Snaclec bitiscetin subunit alpha	<i>Bitis arietans</i>
	52	SLB6_MACLB	Snaclec B6 (fragment)	<i>Macrovipera lebetina</i>
	49	A0A0C5DKL1_MACLB	C-type lectin-like protein 3A	<i>Macrovipera lebetina</i>
	46	J3RZT5_CROAD	Keratin, type I cytoskeletal 19	<i>Crotalus adamanteus</i>
	40	A0A1W7RE48_AGKCO	Ribonuclease P protein subunit p38-like protein	<i>Agkistrodon contortrix contortrix</i>
	38	A0A0B8RV71_CROHD	Keratin, type II cytoskeletal cochlear-like	<i>Crotalus horridus</i>
	34	A0A0F7ZAB1_CROAD	Keratin, type I cytoskeletal 18-like protein	<i>Crotalus adamanteus</i>
	31	A0A0F7Z805_CROAD	MBT domain-containing protein 1 isoform	<i>Crotalus adamanteus</i>
	29	SLB1_MACLB	Snaclec B1	<i>Macrovipera lebetina</i>
	29	A0A0F7Z8B1_CROAD	E3 ubiquitin-protein ligase	<i>Crotalus adamanteus</i>
	29	A0A194APM7_9SAUR	C-type lectin 9d	<i>Agkistrodon piscivorus</i>
	28	A0A1W7RD94_AGKCO	Sister chromatid cohesion protein PDS5	<i>Agkistrodon contortrix contortrix</i>
	26	A0A1W7REY6_AGKCO	Sister chromatid cohesion protein PDS5 B	<i>Agkistrodon contortrix contortrix</i>
	26	SL114_ECHCA	Snaclec carinactivase-1 regulatory subunit 14 kDa chain (fragment)	<i>Echis carinatus</i>
	25	A0A0C5DKK6_MACBL	C-type lectin-like protein 2A	<i>Macrovipera lebetina</i>
	24	A0A1W7RC08_AGKCO	Ubiquitin-like-conjugating enzyme ATG3-like protein	<i>Agkistrodon contortrix contortrix</i>
	23	A0A0F7Z6K1_CROAD	Phosphatidate phosphatase	<i>Crotalus adamanteus</i>
	22	A0A0K8S0L2_CROHD	Iron-sulfur cluster assembly 1, mitochondrial-like protein	<i>Crotalus horridus</i>
	20	A0A0F7ZC99_CROAD	Charged multivesicular body protein	<i>Crotalus adamanteus</i>

	18	A0A1W7RDB4_AGKCO	Sialic acid synthase-like protein	<i>Agkistrodon</i> <i>contortrix</i> <i>contortrix</i>
	17	A0A1W7RF54_AGKCO	NudC domain-containing protein 2	<i>Agkistrodon</i> <i>contortrix</i> <i>contortrix</i>
	16	A0A1W7RHL5_AGKCO	Endoplasmic reticulum-Golgi intermediate compartment protein-2 like protein	<i>Agkistrodon</i> <i>contortrix</i> <i>contortrix</i>
	15	T1E4Y0_CROHD	Secernin-3-like protein	<i>Crotalus horridus</i>
	15	A0A0B8RW51_CROHD	Cytochrome P450 2j5-like protein	<i>Crotalus horridus</i>
	15	A0A0K8RXU5_CROHD	Tubulin-glutamate carboxypeptidase	<i>Crotalus horridus</i>
	13	A0A194AQ76_SISMB	C-type lectin 15	<i>Sistrurus</i> <i>miliarius</i> <i>barbouri</i>
FG6	54	J3RZT5_CROAD	Keratin, type I cytoskeletal 19	<i>Crotalus</i> <i>adamanteus</i>
	52	A0A194APM7_9SAUR	C-type lectin 9d	<i>Agkistrodon</i> <i>piscivorus</i>
	49	A0A0B8RV71_CROHD	Keratin, type II cytoskeletal cochleal-like	<i>Crotalus horridus</i>
	35	A0A1W7REY6_AGKCO	Sister chromatid cohesion protein PDS5 B	<i>Agkistrodon</i> <i>contortrix</i> <i>contortrix</i>
	32	A0A0F72BJ1_CROAD	FAS-associated factor 1	<i>Crotalus</i> <i>adamanteus</i>
	30	A0A0F7ZAB1_CROAD	Keratin, type I cytoskeletal 18-like protein	<i>Crotalus</i> <i>adamanteus</i>
	29	VM2_BITAR	Disintegrin bitistatin	<i>Bitis arietans</i>

FG1 to FG6 were cut from the gel and subjected to in-gel digestion with trypsin. Analysis was performed using an Easy-nLC Proxeon nanoHPLC system coupled to an LTQ-Orbitrap Velos. The results obtained were compared with the “*Bitis*” and “*Snakes*” database downloaded from Uniprot (Taxid: 8570).