

Annex 6 – Proteins identified by MaxQuant in the venom from *Vipera ammodytes montandoni*

Accession	Description	Family
JAG65874.1	Xaa-Pro aminopeptidase 2 [Boiga irregularis]	Aminopeptidase
KAG8144505.1	putative Xaa-Pro aminopeptidase 2 protein [Naja naja]	Aminopeptidase
XP_007437861.2	endoplasmic reticulum aminopeptidase 1 [Python bivittatus]	Aminopeptidase
XP_015676063.1	xaa-Pro aminopeptidase 2 [Protobothrops mucrosquamatus]	Aminopeptidase
XP_026530513.1	xaa-Pro aminopeptidase 2 [Notechis scutatus]	Aminopeptidase
XP_039186994.1	xaa-Pro aminopeptidase 2 isoform X2 [Crotalus tigris]	Aminopeptidase
ACE73566.1	cysteine-rich secretory protein Cv-CRP, partial [Crotalus viridis]	CRISP
AMB36337.1	cysteine-rich secretory protein CRISP-1 [Vipera ammodytes ammodytes]	CRISP
AWX67630.1	cysteine-rich secretory protein, partial [Boiga irregularis]	CRISP
B7FDI0.1	Cysteine-rich venom protein [Vipera berus nikolskii]	CRISP
P0DMT4.1	Cysteine-rich venom protein [Echis coloratus]	CRISP
Q2XXP4.1	Cysteine-rich venom protein TRI1 [Trimorphodon biscutatus]	CRISP
Q8JGT9.1	Cysteine-rich venom protein tigrin [Rhabdophis tigrinus tigrinus]	CRISP
Q8JI40.1	Cysteine-rich venom protein ablomin [Gloydus blomhoffii]	CRISP
P0C6A5.1	Disintegrin VA6 [Vipera ammodytes ammodytes]	DIS
P0C6B0.1	Disintegrin VLO5B [Macrovipera lebetina obtusa]	DIS
QHR82751.1	disintegrin, partial [Vipera anatolica senliki]	DIS
ACF70483.1	secreted L-amino acid oxidase precursor [Daboia russelii]	LAAO
B5U6Y8.1	L-amino-acid oxidase [Echis ocellatus]	LAAO
J7H670.1	L-amino acid oxidase Lm29 [Lachesis muta]	LAAO
JAA74732.1	LAO-Den-1 [Denisonia devisi]	LAAO
JAS04522.1	L-amino acid oxidase [Agkistrodon conanti]	LAAO
P0DI84.1	L-amino-acid oxidase [Vipera ammodytes ammodytes]	LAAO
QBF53419.1	L-amino acid oxidase LAAO-II [Vipera ammodytes ammodytes]	LAAO
QHR82782.1	nerve growth factor 1 [Vipera anatolica senliki]	NGF
A0A2I4HXH5.1	Snake venom 5'-nucleotidase [Naja atra]	Nucleotidase
AHJ80886.1	5'-nucleotidase, partial [Macrovipera lebetina]	Nucleotidase
KAG5858140.1	snake venom 5' nucleotidase precursor [Bothrops jararaca]	Nucleotidase
1708179C	phospholipase A2 II [Vipera aspis]	PLA
1708179D	phospholipase A2 III [Vipera aspis]	PLA
1AOK	Chain B, VIPOXIN COMPLEX [Vipera ammodytes meridionalis]	PLA
1Q5T	Chain B, Phospholipase A2 inhibitor [Vipera ammodytes meridionalis]	PLA
1RGB	Chain A, Phospholipase A2 [Vipera ammodytes meridionalis]	PLA
AAB36471.1	phospholipase A2 acidic subunit, PLA2-I subunit A [Vipera aspis]	PLA
ABD24039.1	phospholipase A2-IV [Daboia russelii russelii]	PLA
AEO99168.1	phospholipase A2 [Ovophis monticola]	PLA
F8QN51.1	Acidic phospholipase A2 Vur-PL3 [Vipera renardi]	PLA
F8QN52.1	Acidic phospholipase A2 PL1 [Vipera renardi]	PLA
F8QN53.1	Acidic phospholipase A2 Vur-PL2B [Vipera renardi]	PLA
O93233.1	Phospholipase A2 inhibitor Gloydus brevicaudus siniticus[]	PLA
P04084.3	Acidic phospholipase A2 homolog vipoxin A chain [Vipera ammodytes meridionalis]	PLA
P0DKR3.1	Acidic phospholipase A2 Cblalpha [Pseudocerastes fieldi]	PLA
P24294.1	Acidic phospholipase A2 PLA-2 [Eristicophis macmahoni]	PLA
BAN82155.1	phospholipase b [Ovophis okinavensis]	PLB
ETE59578.1	Phospholipase B-like 1, partial [Ophiophagus hannah]	PLB

QB53421.1	phospholipase B [Vipera ammodytes ammodytes]	PLB
AJO70723.1	C-type lectin-like protein 3B [Macrovipera lebetina]	SNACLEC
AJO70726.1	C-type lectin-like protein 3A [Macrovipera lebetina]	SNACLEC
APB93443.1	snaclec-6 [Vipera ammodytes ammodytes]	SNACLEC
APB93445.1	snaclec-8 [Vipera ammodytes ammodytes]	SNACLEC
B4XS20.1	Snaclec A15 [Macrovipera lebetina]	SNACLEC
JAC96618.1	C-type lectin I [Echis coloratus]	SNACLEC
Q6T7B7.1	C-type lectin 1 [Bitis gabonica]	SNACLEC
QB53411.1	C-type lectin-like Snaclec-9 [Vipera ammodytes ammodytes]	SNACLEC
AAF01040.1	metalloproteinase precursor, partial [Atractaspis engaddensis]	svMP
ADI47619.1	metalloproteinase [Echis coloratus]	svMP
ADI47620.1	metalloproteinase [Echis coloratus]	svMP
ADW54345.1	group III snake venom metalloproteinase [Echis ocellatus]	svMP
ADW54352.1	group III snake venom metalloproteinase, partial [Echis ocellatus]	svMP
AGL45259.1	H3 metalloproteinase precursor 1 [Vipera ammodytes ammodytes]	svMP
AHB62069.1	metalloproteinase H4-A [Vipera ammodytes ammodytes]	svMP
AJC52543.1	metalloproteinase F1 precursor [Vipera ammodytes ammodytes]	svMP
AMB36349.1	metalloproteinase of class P-II MP11-1 [Vipera ammodytes ammodytes]	svMP
AMB36350.1	metalloproteinase of class P-II MP11-2 [Vipera ammodytes ammodytes]	svMP
AMB36351.1	metalloproteinase of class P-II MP11-3 [Vipera ammodytes ammodytes]	svMP
AMB36352.1	metalloproteinase of class P-III MP11-1 [Vipera ammodytes ammodytes]	svMP
ANT80539.1	venom metalloprotease PIII-SVMP EOC00089 [Echis ocellatus]	svMP
B8K1W0.1	Zinc metalloproteinase-disintegrin-like daborhagin-K [Daboia russelii]	svMP
JAS05432.1	Metalloproteinase type II 5a [Sistrurus miliarius barbouri]	svMP
JAV01848.1	BATXSVMPIII16 [Bothrops atrox]	svMP
JAV01853.1	BATXSVMPIII11 [Bothrops atrox]	svMP
JAV01854.1	BATXSVMPIII10 [Bothrops atrox]	svMP
KAG5858154.1	snake venom metalloprotease precursor [Bothrops jararaca]	svMP
Q2UXR0.1	Zinc metalloproteinase-disintegrin-like Eoc1 [Echis ocellatus]	svMP
Q7T046.1	Coagulation factor X-activating enzyme heavy chain [Macrovipera lebetina]	svMP
Q98995.1	Zinc metalloproteinase/disintegrin [Macrovipera lebetina]	svMP
QB53408.1	MPi-3 [Vipera ammodytes ammodytes]	svMP
QB53415.1	metalloproteinase of class P-III MP11-2 [Vipera ammodytes ammodytes]	svMP
QB53416.1	metalloproteinase-like protein of class P-III MP11-3 [Vipera ammodytes ammodytes]	svMP
QB53417.1	metalloproteinase of class P-III MP11-4 [Vipera ammodytes ammodytes]	svMP
ALA20853.1	phosphodiesterase, partial [Pseudagkistrodon rudis]	svPDE
P0DQQ4.1	Venom phosphodiesterase CdcPDE [Crotalus durissus collilineatus]	svPDE
W8E7D1.1	Venom phosphodiesterase [Macrovipera lebetina]	svPDE
XP_007430581.1	ectonucleotide pyrophosphatase/phosphodiesterase family member 3, partial [Python bivittatus]	svPDE
ADI47572.1	serine protease, partial [Echis coloratus]	svSP
AMB36343.1	serine proteinase SP-2 [Vipera ammodytes ammodytes]	svSP
AMB36344.1	serine proteinase SP-3 [Vipera ammodytes ammodytes]	svSP
AMB36345.1	serine proteinase SP-4 [Vipera ammodytes ammodytes]	svSP
AMB36346.1	serine proteinase SP-5, partial [Vipera ammodytes ammodytes]	svSP
AUS82476.1	serine endopeptidase [Agkistrodon piscivorus]	svSP
AUS82539.1	serine endopeptidase [Crotalus scutulatus]	svSP
C0HK18.1	Thrombin-like enzyme collinein-4 [Crotalus durissus collilineatus]	svSP
E0Y418.1	Serine protease VLSP-1 [Macrovipera lebetina]	svSP
JAS04414.1	Serine proteinase 1 [Agkistrodon conanti]	svSP
JAV51413.1	Serine proteinase 9 [Agkistrodon contortrix contortrix]	svSP

O93421.2	Snake venom serine protease pallase [Gloydius halys]	svSP
P00991.2	Kunitz-type serine protease inhibitor 1 [Vipera ammodytes ammodytes]	svSP
P82981.1	Thrombin-like enzyme contortrixobin [Agkistrodon contortrix contortrix]	svSP
Q9PT41.1	Factor V activator [Macrovipera lebetina]	svSP
QBF53412.1	serine proteinase SP-6 [Vipera ammodytes ammodytes]	svSP
QBF53414.1	serine proteinase SP-8 [Vipera ammodytes ammodytes]	svSP
QHR82809.1	serine protease 2 [Vipera anatolica senliki]	svSP
QHR82827.1	serine protease inhibitor 3 [Vipera anatolica senliki]	svSP
QHR82828.1	serine protease inhibitor 5 [Vipera anatolica senliki]	svSP
XP_015671564.1	snake venom serine protease serpentokallikrein-1 [Protobothrops mucrosquamatus]	svSP
C0K3N5.1	Vascular endothelial growth factor A [Vipera ammodytes ammodytes]	VEGF
P67863.2	Snake venom vascular endothelial growth factor toxin vammin [Vipera ammodytes ammodytes]	VEGF
2E3X	Chain A, Coagulation factor X-activating enzyme heavy chain [Daboia siamensis]	-
A0A1I9KNP0.1	Vaa serine proteinase homolog 1 [Vipera ammodytes ammodytes]	-
AAN59983.1	ammodytin I2 [Vipera aspis aspis]	-
AAO62996.1	ophanin [Ophiophagus hannah]	-
AAX38181.1	VLAIP-A [Macrovipera lebetina]	-
ABI33937.1	hyaluronidase [Echis ocellatus]	-
ABU68556.1	PLA2(IIA)-Cau1 [Causus rhombeatus]	-
ABV89584.1	thrombin-like protein, partial [Deinagkistrodon acutus]	-
AFE84762.1	glutaminyl-peptide cyclotransferases [Daboia russelii]	-
AHJ80887.1	hyaluronidase, partial [Macrovipera lebetina]	-
AMH40734.1	Kunitz/BPTI inhibitor-2 [Vipera ammodytes ammodytes]	-
APB93441.1	SVMPi-1 [Vipera ammodytes ammodytes]	-
BAN89449.1	glutaminyl_cyclase, partial [Ovophis okinavensis]	-
CAA01524.1	ancrod-like protein [Calloselasma rhodostoma]	-
CAA01526.1	ancrod-like protein [Calloselasma rhodostoma]	-
CAE47105.1	vaspin acidic subunit (1) variant [Vipera aspis aspis]	-
CAE47176.1	ammodytin I1(A) variant [Vipera ammodytes ammodytes]	-
CAE47181.1	ammodytin I1(D) isoform [Vipera ammodytes montandoni]	-
CAJ55260.1	renin-like aspartic protease [Echis ocellatus]	-
CAK12655.1	MLD-containing dimeric disintegrin subunit ml-G2 precursor, partial [Macrovipera lebetina transmediterranea]	-
ETE57638.1	hypothetical protein L345_16645, partial [Ophiophagus hannah]	-
ETE59790.1	hypothetical protein L345_14477 [Ophiophagus hannah]	-
ETE59912.1	78 kDa glucose-regulated protein [Ophiophagus hannah]	-
ETE62249.1	Protein unc-79-like protein, partial [Ophiophagus hannah]	-
ETE63622.1	SWI/SNF complex subunit SMARCC1, partial [Ophiophagus hannah]	-
ETE63958.1	Rho GTPase-activating protein 10, partial [Ophiophagus hannah]	-
ETE68362.1	Neurobeachin-like protein 2, partial [Ophiophagus hannah]	-
ETE70241.1	hypothetical protein L345_03955, partial [Ophiophagus hannah]	-
ETE71055.1	hypothetical protein L345_03129 [Ophiophagus hannah]	-
ETE73956.1	Renin, partial [Ophiophagus hannah]	-
JAA96189.1	Peroxiredoxin 4 [Crotalus horridus]	-
JAC94969.1	Cathepsin D [Opheodrys aestivus]	-
JAC95083.1	Pre-mRNA-processing factor 39 [Hypsigena sp. JMG-2014]	-
JAV48710.1	tumor protein p53 binding protein, 2 [Agkistrodon contortrix contortrix]	-
KAG6539588.1	type-2 vomeronasal receptor [Crotalus adamanteus]	-
KAG8123366.1	hypothetical protein E2320_018771 [Naja naja]	-
KAG8125038.1	hypothetical protein E2320_020305 [Naja naja]	-
KAG8125449.1	hypothetical protein E2320_020711, partial [Naja naja]	-

KAG8131107.1	hypothetical protein E2320_017660 [Naja naja]	-
KAG8132840.1	hypothetical protein E2320_010673 [Naja naja]	-
KAG8142765.1	hypothetical protein E2320_005966, partial [Naja naja]	-
LAA18635.1	hypothetical protein, partial [Micrurus lemniscatus carvalhoi]	-
LAA19405.1	hypothetical protein, partial [Micrurus lemniscatus carvalhoi]	-
LAA23376.1	hypothetical protein, partial [Micrurus lemniscatus carvalhoi]	-
LAA26205.1	hypothetical protein, partial [Micrurus lemniscatus carvalhoi]	-
LAA30658.1	hypothetical protein, partial [Micrurus lemniscatus carvalhoi]	-
LAA43750.1	hypothetical protein, partial [Micrurus corallinus]	-
LAA45417.1	hypothetical protein [Micrurus corallinus]	-
LAA66478.1	hypothetical protein, partial [Micrurus corallinus]	-
LAA67683.1	hypothetical protein, partial [Micrurus lemniscatus lemniscatus]	-
LAA69240.1	hypothetical protein, partial [Micrurus lemniscatus lemniscatus]	-
LAA96095.1	hypothetical protein, partial [Micrurus lemniscatus lemniscatus]	-
LAA98218.1	hypothetical protein, partial [Micrurus lemniscatus lemniscatus]	-
LAB17776.1	hypothetical protein [Micrurus spixii]	-
LAB18218.1	hypothetical protein [Micrurus spixii]	-
LAB26629.1	hypothetical protein [Micrurus spixii]	-
LAB48077.1	hypothetical protein, partial [Micrurus surinamensis]	-
LAB59127.1	hypothetical protein, partial [Micrurus surinamensis]	-
LAB63721.1	hypothetical protein, partial [Micrurus surinamensis]	-
O13069.1	Thrombin-like enzyme KN-BJ 2 [Bothrops jararaca]	-
P00761	Trypsin	-
P02769	Albumin	-
P04264	Keratin, type II cytoskeletal 1	-
P13645	Keratin, type I cytoskeletal 10	-
P15636	Protease 1	-
P35527	Keratin, type I cytoskeletal 9	-
P35908	Keratin, type II cytoskeletal 2 epidermal	-
Q3TTY5	Keratin, type II cytoskeletal 2 epidermal	-
Q7Z794	Keratin, type II cytoskeletal 1b	-
Q86YZ3	Hornerin	-
QEE02313.1	periplakin, partial [Lycodon subcinctus]	-
QGV16220.1	recombination activating protein 1, partial [Smithophis atemporalis]	-
QHR82750.1	cystatin 4, partial [Vipera anatolica senliki]	-
XP_007426346.1	calpain-2 catalytic subunit [Python bivittatus]	-
XP_007426835.1	peptidyl-prolyl cis-trans isomerase B, partial [Python bivittatus]	-
XP_007427414.1	hexokinase-1, partial [Python bivittatus]	-
XP_007427945.1	aprataxin isoform X3 [Python bivittatus]	-
XP_007431360.1	zinc finger protein RFP-like [Python bivittatus]	-
XP_007436345.1	ELKS/Rab6-interacting/CAST family member 1-like isoform X2 [Python bivittatus]	-
XP_007437328.1	glutaminy-peptide cyclotransferase, partial [Python bivittatus]	-
XP_007445078.1	peptidyl-prolyl cis-trans isomerase F, mitochondrial-like, partial [Python bivittatus]	-
XP_013907759.1	PREDICTED: kinesin-like protein KIF28P [Thamnophis sirtalis]	-
XP_013910680.1	PREDICTED: solute carrier family 35 member G2 isoform X2 [Thamnophis sirtalis]	-
XP_013912549.1	PREDICTED: coiled-coil domain-containing protein 40 [Thamnophis sirtalis]	-
XP_013915889.1	PREDICTED: maltase-glucoamylase, intestinal-like [Thamnophis sirtalis]	-
XP_013927039.1	PREDICTED: adhesion G protein-coupled receptor B2 isoform X2 [Thamnophis sirtalis]	-
XP_013931521.1	PREDICTED: TNF receptor-associated factor 6 [Thamnophis sirtalis]	-
XP_015671453.1	digestive cysteine proteinase 2-like [Protobothrops mucrosquamatus]	-

XP_015672624.1	zinc finger protein 740 [Protobothrops mucrosquamatus]	-
XP_015679678.1	spermatogenesis-associated serine-rich protein 2-like [Protobothrops mucrosquamatus]	-
XP_015680317.1	junctional adhesion molecule A [Protobothrops mucrosquamatus]	-
XP_015683377.1	threonine--tRNA ligase, cytoplasmic-like [Protobothrops mucrosquamatus]	-
XP_015687762.1	DNA topoisomerase 2-alpha [Protobothrops mucrosquamatus]	-
XP_015744661.1	DNA repair protein complementing XP-G cells [Python bivittatus]	-
XP_025018835.1	CAP-Gly domain-containing linker protein 1 isoform X8 [Python bivittatus]	-
XP_025020245.1	kelch-like protein 6, partial [Python bivittatus]	-
XP_025027126.1	Golgi apparatus protein 1 [Python bivittatus]	-
XP_025027687.1	G2 and S phase-expressed protein 1 isoform X3 [Python bivittatus]	-
XP_025027842.1	caskin-1 [Python bivittatus]	-
XP_025028798.1	EF-hand calcium-binding domain-containing protein 4B isoform X4 [Python bivittatus]	-
XP_025029238.1	zinc finger and BTB domain-containing protein 38 [Python bivittatus]	-
XP_025029781.1	EF-hand calcium-binding domain-containing protein 3-like [Python bivittatus]	-
XP_025030431.1	RAS protein activator like-3 [Python bivittatus]	-
XP_025033114.1	low-density lipoprotein receptor-related protein 8-like, partial [Python bivittatus]	-
XP_026522209.1	leukemia inhibitory factor receptor isoform X2 [Notechis scutatus]	-
XP_026524175.1	uncharacterized protein LOC113412638 [Notechis scutatus]	-
XP_026531030.1	mitogen-activated protein kinase kinase kinase 4 isoform X3 [Notechis scutatus]	-
XP_026531079.1	centromere protein J-like [Notechis scutatus]	-
XP_026534175.1	PLIalpha-like protein [Notechis scutatus]	-
XP_026536026.1	echinoderm microtubule-associated protein-like 1 [Notechis scutatus]	-
XP_026545819.1	neuromedin-B-like, partial [Notechis scutatus]	-
XP_026549755.1	BPTI/Kunitz domain-containing protein-like, partial [Notechis scutatus]	-
XP_026572509.1	growth arrest and DNA damage-inducible proteins-interacting protein 1 [Pseudonaja textilis]	-
XP_026577188.1	alpha-2-macroglobulin-like [Pseudonaja textilis]	-
XP_026581458.1	lebercilin-like protein, partial [Pseudonaja textilis]	-
XP_029140249.1	E3 ubiquitin-protein ligase RNF123 [Protobothrops mucrosquamatus]	-
XP_029140997.1	drebrin-like protein A [Protobothrops mucrosquamatus]	-
XP_029141240.1	dystonin [Protobothrops mucrosquamatus]	-
XP_029141257.1	DENN domain-containing protein 1B [Protobothrops mucrosquamatus]	-
XP_029142797.1	zinc finger protein 300-like [Protobothrops mucrosquamatus]	-
XP_032064726.1	C-Jun-amino-terminal kinase-interacting protein 4 isoform X3 [Thamnophis elegans]	-
XP_032067965.1	small integral membrane protein 3 [Thamnophis elegans]	-
XP_032071327.1	reticulon-4 isoform X4 [Thamnophis elegans]	-
XP_032082288.1	protein Jade-3 isoform X2 [Thamnophis elegans]	-
XP_032083046.1	platelet-activating factor receptor isoform X2 [Thamnophis elegans]	-
XP_032083638.1	digestive cysteine proteinase 2-like [Thamnophis elegans]	-
XP_032088853.1	uncharacterized protein LOC116519199 [Thamnophis elegans]	-
XP_032089769.1	pleckstrin homology domain-containing family O member 1 isoform X2 [Thamnophis elegans]	-
XP_032091487.1	1-aminocyclopropane-1-carboxylate synthase-like protein 1 [Thamnophis elegans]	-
XP_032092252.1	putative olfactory receptor 13C6 [Thamnophis elegans]	-
XP_032095013.1	LOW QUALITY PROTEIN: zinc finger protein 91-like [Thamnophis elegans]	-
XP_034257482.1	probable tRNA(His) guanylyltransferase isoform X5 [Pantherophis guttatus]	-
XP_034257560.1	putative deoxyribonuclease TATDN2 [Pantherophis guttatus]	-
XP_034257685.1	tripartite motif-containing protein 55-like [Pantherophis guttatus]	-
XP_034257953.1	histone deacetylase 9 isoform X2 [Pantherophis guttatus]	-
XP_034260201.1	vomer nasal type-2 receptor 26-like [Pantherophis guttatus]	-
XP_034262134.1	LOW QUALITY PROTEIN: splicing factor ESS-2 homolog [Pantherophis guttatus]	-
XP_034263433.1	C3a anaphylatoxin chemotactic receptor-like [Pantherophis guttatus]	-

XP_034265777.1	adenylate cyclase type 10 isoform X8 [Pantherophis guttatus]	-
XP_034271514.1	suppressor of cytokine signaling 7 isoform X2 [Pantherophis guttatus]	-
XP_034272377.1	small ribosomal subunit protein uS2-like [Pantherophis guttatus]	-
XP_034279009.1	albumin-like [Pantherophis guttatus]	-
XP_034280079.1	kunitz/BPTI-like toxin [Pantherophis guttatus]	-
XP_034285082.1	adenylyl cyclase-associated protein 2 [Pantherophis guttatus]	-
XP_034290209.1	intercellular adhesion molecule 5-like [Pantherophis guttatus]	-
XP_034290552.1	ubiquitin carboxyl-terminal hydrolase 8 [Pantherophis guttatus]	-
XP_034292909.1	geranylgeranyl transferase type-2 subunit beta isoform X3 [Pantherophis guttatus]	-
XP_034295364.1	period circadian protein homolog 2 isoform X2 [Pantherophis guttatus]	-
XP_034297623.1	neuroligin-3 isoform X4 [Pantherophis guttatus]	-
XP_039178534.1	myomesin-2 [Crotalus tigris]	-
XP_039178879.1	probable ATP-dependent RNA helicase DDX31 [Crotalus tigris]	-
XP_039179757.1	centrosomal protein of 70 kDa isoform X2 [Crotalus tigris]	-
XP_039179918.1	pantetheinase-like [Crotalus tigris]	-
XP_039182639.1	zinc finger protein 85-like isoform X3 [Crotalus tigris]	-
XP_039185187.1	little elongation complex subunit 1 isoform X3 [Crotalus tigris]	-
XP_039188353.1	heparin cofactor 2 [Crotalus tigris]	-
XP_039188792.1	signal-transducing adaptor protein 2 isoform X2 [Crotalus tigris]	-
XP_039189813.1	maestro heat-like repeat family member 5 isoform X6 [Crotalus tigris]	-
XP_039189909.1	myosin regulatory light chain 10 isoform X1 [Crotalus tigris]	-
XP_039193803.1	A-kinase anchor protein 4 [Crotalus tigris]	-
XP_039196843.1	glutathione peroxidase 3 [Crotalus tigris]	-
XP_039212786.1	actin, alpha skeletal muscle B-like isoform X1 [Crotalus tigris]	-
XP_039212967.1	carboxypeptidase N catalytic chain [Crotalus tigris]	-
XP_039214639.1	IQ domain-containing protein J [Crotalus tigris]	-
XP_039219335.1	extracellular matrix protein 1 [Crotalus tigris]	-
XP_039221953.1	guanine nucleotide-binding protein-like 3-like protein isoform X3 [Crotalus tigris]	-
XP_039222233.1	cyclin-dependent kinase 1 [Crotalus tigris]	-
XP_039224524.1	pregnancy zone protein-like [Crotalus tigris]	-

Protein family	Count
Aminopeptidase	6
DIS	3
LAAO	7
NGF	1
Nucleotidase	3
PLA	15
PLB	3
SNACLEC	8
svMP	26
svPDE	4
svSP	21
VEGF	2
CRISP	8
-	168
TOTAL	275

svMP	Snake Venom Metalloprotease
svSP	Snake Venom Serine Protease
PLA	Phospholipase A
SNACLEC	Snake Venom Lectins
CRISP	Cysteine-rich secretory protein
LAAO	L-amino Acid Oxidase
svPDE	Snake Venom Phosphodiesterase
DIS	Disintegrin
PLB	Phospholipase B
VEGF	Snake venom vascular endothelial growth factors
NGF	Nerve growth factors