>Smar-ACP  
MKWIAVYLLLTIIVLTIVAPVEGQVTFSRDWTPAGKRGMDCGFVKTKLLRDIAVLLQVKYHFAELCMLTLAWLMLDGGQFTEILWTSCDGAFP

>Smar-AKH  
MTKFTWLSMTLLVLMVFITVDVNGQINFSPGWGQGKRSLSDDKPVNGYSDCSETMIEVYRLLK

>Smar-Allatostatin A  
MTFAVTWWCLLLTAPTLLMSEYIYDISSESDSNDQEKRGLNTPWKLPEGYVYLRKVDPSTGEYQIGKKDNQYRRRFSFGLGKRFHSDVLLENNEDEVGKRGSQRNRHFGLGKRQPVDYLQGRIGRYNMGLGKRSVDSREAEEVAIEEMKRGASKFNFGLGKRTKPYSFGLGKRWDDRGVEENLIEEYKRAKTYGFGLGKRDEEEMEEEKKNRPYQFGLGKRDRSYSFGLGKRMEEEKKT

>Smar-Allatostatin B  
MLLSWTSSVTIVLVIASVLAASASEDKRAWSDLNGAWGKRNWDQLRGVWGKRGAGQLPNSVWGKREDAPSDWNAFRGSWGKRNNWNKLQGVWGKRDSSDWNKLQGLWGKRASWTHST

>Smar-Allatostatin C  
MASSGKFCILIFALVLTLSHVTSKSIGEHEHPNFNTDLSLVDDDGSMDTALINYLFARQMIKRLQSSMDVTDLQRKRSYWKQCAFNAVSCFGKK

>Smar-Allatostatin CC  
MYSLILVFCVCMLTLPYVSCQIEMNSLKNMAKTFHLSSSDSNYFQHPVKRSTMLLDRLVTALQKAFKQETQEVTGMELQRRRPNGRVYWRCYFNAVSCFRRKK

>Smar-Allatotropin  
MKPVCLVILLFGLLVSATSSSTDEPANRVRQTRGFKNSALATARGFGKRTMLNDLVDSTDRAIMSNEQLADLMSRNPQFAQQILTKFVDTDGDGVLSFRE

>Smar-Bursicon alpha  
MVTLVFMVAAMMCIFSSRLATADECHITPVIHVLKYPGCNQKPIPSFACQGRCTSYVSGSKMWQMERSCMCCQEMGVREANVTLHCPHARPGEPKFRKVTTRAPVDCMCRPCTSVEKHLVQPQESAPWLSDPNFNDAILSV

>Smar-Bursicon beta  
MTNTWSAFAFFTIITFVLFITTKSLRAFPESTCETLPSFIHIIKEEYDSRTKLVRTCEGDVAVNKCEGTCTSQMQPSVTTSTGFLKECYCCRESYLQEREVILQRCFNFDGETLAGDMSLMKIRLKEPAECQCYKCGE

>Smar-Corazonin  
MGFQKTKLLLIVASILVFIICTSGQTFQYSKGWEPGRKRAVDRSYQVRDWDAGRKRENIRGLDSSTAWILGLKRAAFN

>Smar-CCAP  
MQYVTIHGFVTLLLIVFSCTIGCAAQKRRPFCNAFTGCGRKRSELPASNVNDLETDLLLDKLSHQILGLVHVLEALHMRIETSRQNQQPLPMIETDSRIPNYILDRKRRSTKI

>Smar-CCHamide 1-1  
MHGCRCNTALFILLVLSLLVSSATGRRGCLNYGHSCLGAHGKRSSNRQAMRRDIANFLPLHKTAAEYNTISEENLLRDESDTRKWNDKWRDMVISALEKDD

>Smar-CCHamide 1-2  
MSTLKFIFALLTLAVYVYQVQGLRGCTNYGHSCFGAHGKRTPKNDEKDQTSFLDSTENTKHVNQITTNDVPGVKNGLNSAFLRKWMTALRQSGNDEILQ

>Smar-DH31  
...DKRNLDLGFSRGFSGSQAAKHLMGLAAANFAGGPGRRKRSEE

>Smar-DH44 (CRF-like)  
MLLRLFSFIVCVCAIACVGARSLSRLQECTDCTLVPADDNRYSIQDDFNNKGFILKARRIPKWSLFANSPDEVSSERMIHGFSMTRLDGTKKRNDGTNLSIVNPVEVIRQRMMIDAARERQNQIDANAEMLREIGKRQPKAWRSDWH

>Smar-EFLamide  
MKLLCIDPSLFIFTYLVLSLIPSTNLVSSQEIERVPESYGMDSSVLKRSARISQEDVYRMFVLLNKIKDRGGIKRIGSEFIGKRYSQAEDIEKKLGSEFLGKRGIGSEFLGKRSQRSQAEN

>Smar-EH  
MRQSESQRAAVSTVVILLLLHLDRASSRSINLCIQNCAQCKKMFGPYFEGQLCAETCIELRAKMTPDCADANSISPFLNKFER

>Smar-EH-like  
MACLLSAACVLASVACISEESSLGVCIRNCGQCKRMYGDFFLGQHCAEECLQTEGRGDLPDCNNPKSLYRFLGKT

>Smar-Elevenin  
MKIVIRSVIPLLICLILLLVFVSHSEQVDCRIYVFAPKCRGISAKRGIDPPGRQQQPNQKLDINADPYAPYEYNPTDYSWDERASDSNYGSRISSDFVAPIFDSSSPPSRFKNSDDRDQGRFLRALVKMYFDQRQDSED

>Smar-ETH  
MTTFTSSFFVPTQMYILVLFAVFLLQTEAQFFAKTSKNLPRIGRRVDHQQEVSEIIPPSFKHLLAFVRKFDSDANGCLSPEELIEIPLFQMAFENEDFTPLEIAADVVEEYKSEVEGNLKDIVAKVLAASYAEKK

>Smar-FMRFamide  
MQQVLLSFLVVTIPALAIPVSARCFLQPSLGGVPNENSSPLLCTLMSSRESADEVSEDDDKLMDSDALEPINNELTSEENSMDDDQKDALLRSLRQEPGHKFLRFGRASSNHNFLRFGRDPEHKFLRFGRDQHKFLRFGRSVANGEENRLRLRENQSKMTVLPMFMRLGRGPEHVFMRFGRQGSNSEGHKFMRFGRTQNDTPVQDENTQQEKA

>Smar-FMRFamide-like  
..TEQKFARFDRSNFDENGDPQENWLDSENRKLLSLRSARLIRELGRKRGSLEKNFLRFGRSLENGNDINTKRDTSSLDDSHMSYLKSDQLNQNKLVPKRNKLENNFLRFGRWKTENLNEQYE

>Smar-ILP  
MSSHLRTAAAVLILLSVVLSIAAQESAYFQELFTPHEIKIRAKQKYCGRNLVEVLQLVCARNVLNNLEAEEISDLLGNDFARSLMGIKHKIQTRGITDECCRKGCSFNELKSYCAEEP

>Smar-Inotocin  
MKSTHFVNIFIYSVFIFIMADGCYITNCPPGGKRSGNEKSGRGVRQCTPCGPGGIGRCYGPDICCGANVGCFVGTRESAICRLENLYSLPCQNEGRACGTDGTCSADGFCCSTDQCKADESCRGKVHHTNNLQRVLDGEIDLNDVMGPQR

>Smar-ITP  
MNHTLFFRVFVVLSAIIASTCLVSARSLNLEDVSGVHRINKRSFHTLGCLGDYDTAGFSRLDRLCEDCYDMYRDSQVRAMCRSSCFTTDTFKKCAEALLVNMEEEKLGDVVNRLYGRD

>Myosuppressin  
MKLLTLYKILLVISVVIIPSIFSHPPPQCDTDDPLPPRLLRICNALRTISEYTELMEDYLDEEVMHTLAVSDMKRDERDTGHVFMRFG

>Smar-NPF  
MSLSISTRSTLALICVIVVLYVFCAPAQATSGPDQAVSMTEALKYLQALDKYYSQVARPRFGRSLPMRYPSKDMSVEAEANRLLEQRRR

>Smar-NPLP1  
MQGRTCLRMLLLAIVFSHLTQGALQDTGPEATTISGRLTELSGQQERIKPPVKRYVGSVARAGGLPPFFHGKRHEIESTDDEDENNEIIKRYLGSIVRQGIFSHNKRQDEEEMLDDAVEKRHLGSVLRAGDSRLVGRDLFSSLQDKRFMGSLARAGELGPGGRMSGKKRYLGSVARVDGIPFRAKRTPQDTESPDWESEENLDDNEEDVGNEEEWEEDDLLIPFKRNIASLARNGWLPHTRSLRRHDAPTYSVSEEAPKRNIYYRPASASGRSRLLGWLRREEARLKEEAEARNGQGKRTIAALARSGELPVRYRFSRSADRLPAPPFMRMATYRGGGGGGYSIRSPHAFASLHEGGWNRFKRSFEQLDRMQMHLDALDDLCDSWDLERCGPHKGEKKSMERHSSESRKDEGGEMSMTV

>Smar-Proctolin  
MTVKCAVVFALLMTLMYCWSAQCRYLPSRADNTRAEEIREILRE...

>Smar-Pyrokinin  
MWVSLCVWFGIFCSGFGLSLFEVDKRQGLIPFPRIGRALPVDNSILLSSQEIRDALYQGLFRSNRLKRDVEVEGGDDWAEITNDGFQGSPDEKKTVEKTSKLAPRLGRSRYRIGASPFQPRLGRAYVSFGPRLGRSKLPPRLGKRLNRN

>Smar-RYamide  
MFSSKQSTLFYLLTILGMCALFWRVESQQFYPNGRYGRSDKMPALSDVRGTREMTVSFFGDGTVQCTYTGYPDFYRCK...

>Smar-SIFamide  
MASKTTIILLVVAIVAICLVVDVTSANYRKPPFNGSIFGKRAPEDSTAEKLFAMCAIATDACSQWFPASEAK

>Smar-sNPF  
MHSIITCSFLLFATLFILTFPLENASPAPYTDYDNIRELYELLLRNEALNDRTGHQVVRKGSRDPSLRLRFGRRSDPAWQEGREPSLRLRFGRSADDVRGYSKQLRFGRSDDTAWQHDVASSENDVIEN

>Smar-Sulfakinin  
MNCTVIFLVHYLVLVCTFVSSSNGSPSVARSSKHDSRLANIMAPYLYLKLHDQAARRPVDSESESVKDEVESEFDDFFESFDKTKRNFDDYGHMRFGKREFDDYGHPRYGRSA

>Smar-Tachykinin  
MMNFRGIWSLRLSIFIIFFFGPIFGQQVQSLNDKIKNMQLDEDNLLRKSRNVFMGMRGKKMSMDQDRDLQRAATSIAEIKRINGFMAMRGNKINSIHNIFDEVENPYIFPGDKRAKGFLGMRGKKQPSGSEMNRWSNSKFFAMRGKRD

>Smar-Trunk (PTTH?)  
MMKRKKVSFLFGFVLGFYWLAGGQGNATGENVMTNFLETMGKATSKTSKTTETNDRLWTCEWSEHWLDLGNDYFPRYIRTAKCTTEKCWFNFFKCEGRAFTVKVLRKRQDECISHVNNKTVILLEDWVFEERAVNFCCVCVPVWKKKKATLIFFFFSS