

Coleoptera Genome and Transcriptome Sequences reveal numerous Differences in Neuropeptide Signaling between Species

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Hypothenemus hampei

Tribolium Natalisin-R	MSTTESWKEEEGNGS---FQGFYDGLLPVSSQTPVVVDSENGFMPVWVKVLWT	VIFMVM	57				
Sequence-1	-----M		1				
Sequence-3	-----	ILYAGM	6				
Tribolium Tachykinin-R	MNFTQEFLYTGTYSIMDNSSEYDYTTNIS---YNDTEEGNQFILPVWRQVLWS	ILYAGM	57				
Tribolium Natalisin-R	ILIATGGNCIVIWIVTAHRRMRTVTNYFLVNLSLADLLLTFNCIFNFSYMIQRDWPFGS		117				
Sequence-1	VIVATGGNLIVIIYIVLAHKRMRTVTNYFL		30				
Sequence-2	---ATGGNLIVIIYIVLAHKRMRTVTNYFLRELKIA		32				
Sequence-3	VIVATGGNLIVIIYIVLAHKRMRTVTNY		33				
Tribolium Tachykinin-R	VIVATGGNLIVIWIVFSHKRMRTVTNYFLNLSVADTMVSTLNVTNFVYMLNSHWPFGE		117				
Tribolium Natalisin-R	LYCIISNFIANATVAASVFTLTGISCDRYLAIVHPLQPRMSKRASLITITFIWLASMTVA		177				
Tribolium Tachykinin-R	LYCKISQFI AVL SVCASFSLMSISIDRYMAIMTPLRPMGRVTVLLAVTTWLLGVIIG		177				
Tribolium Natalisin-R	FPCLLYSTTITNKYK-GVERTGCILIPDGKVVGSFDFAYQMFFLIITYVIPVTL	MSFS	236				
Sequence-5	-----MTYT		4				
Tribolium Tachykinin-R	SPSLMFFRTYTMPYKDGEERVICYPEWPDGTTNESMMEYAYNVGFLFVTYVPIGSM	MTYT	237				
Tribolium Natalisin-R	YTIMGKELWGSRSIGEMTQRQIDSIIRSKRKVVKMFIFVVFIFAICWLPYHGYFLVYYDT		296				
Sequence-5	YARIGIELWGSQSIGECTQRQMDNIKSKR		33				
Tribolium Tachykinin-R	YARIGIELWGSQSIGECTQRQMEINIRSKRRVVKMMVVVVIIFAVCWLPYHLYFIVISYFP		297				
Tribolium Natalisin-R	DIIFS	KYTHVYLA	FYWFAM	SNAMVN	PLIYYWMNARYVKT	-----	336
Sequence-4	---	SAYIQETFLAIYWLAMSNSMYPNPIIYCWMNARY					33
Tribolium Tachykinin-R	EITNSTYIQETYLA	IYWLAMSNSMYPNPIIYCWMNAR	FRRGFKQFFSCLPFIHVS	PGALTR			357
Tribolium Natalisin-R	-----						336
Tribolium Tachykinin-R	REVLTSRRRSYSGSPDHNRIRNG						381

Hycleus phaleratus

Tribolium_Natalisin-R	MSTTESWKEEEGNGS---FQGFYDGLLPVSSQTPVVVD	SGENFMPVWVKVLWTVIFMVM	57			
Sequence-1	-----	GNQFILPIWRQALWSILYAGM	21			
Tribolium_Tachykinin-R	MNFTQEFLYTGTGYSIMDNSSEYDYTTNIS--YNDTEEG	GNQFILPVWRQVLWSILYAGM	57			
Tribolium_Natalisin-R	ILIATGGNCIVIWIVTAHRRMRTVTNYFLVNLSLADLLLTFNCIFNFSYMIQRDWPFGS		117			
Sequence-1	VIVATGGNLIVIWIVLAHKRMRTVTNYFL-----		50			
Sequence-4	-----	NLSIADTMVSTLNVTNFNVYMLNSHWPFGE	30			
Tribolium_Tachykinin-R	VIVATGGNLIVIWIVFSHKRMRTVTNYFLNLSVADTMVSTLNVTNFNVYMLNSHWPFGE		117			
Tribolium_Natalisin-R	LYCIISNFIANATVAASVFTLTGISCDRYLAIVHPLQPRMSKRASLITITFIWLASMTVA		177			
Sequence-4	LYCKITQFIAVLSVCASF-----		50			
Tribolium_Tachykinin-R	LYCKISQFIAVLSVCASFSLMSISIDRYMAIMTPLRPMGRVTVLLAVTTWLLGVIIG		177			
Tribolium_Natalisin-R	FPCLLYSTTITNKYK-GVERTGCILIPDGKVVGSFDFAY	QMFFLIITYVIPVTLMSFS	236			
Sequence-2	-----	YNVAFLFVTYVVPIGSMTYT	20			
Tribolium_Tachykinin-R	SPSLMFFRTYTMPYKDGEERVICYPEWPDGTTNESMMEYAY	NVGFLFVTYVVPIGSMTYT	237			
Tribolium_Natalisin-R	YTIMGKELWGSRSIGEMTQRQIDSIIRSKRKVVKMFIFVVFIFAICWLPYHGYFLVYYDYT		296			
Sequence-2	YARIGLELWGSQSIGECTQRQIENIKSKR-----		49			
Sequence-3	-----	WLPYHLYFIVTSYFP	15			
Tribolium_Tachykinin-R	YARIGIELWGSQSIGECTQRQMENIRSKRRVVKMMVVVVIIFAVCWLPYHLYFIVISYFP		297			
Tribolium_Natalisin-R	DIIFS	KYTHVYLA	FYWFAM	SNAMVN	PLIYYWMNARYVKT-----	336
Sequence-3	EITNSPYIQETYLA	IYWLAMSNSMYPNPIIYCWMNA-----		50		
Tribolium_Tachykinin-R	EITNSTYIQETYLA	IYWLAMSNSMYPNPIIYCWMNARFRRGFKQFFSCLPFIHVS	PGALTR		357	

Figure S1. When the *Tribolium* natalisin receptor is used as query in a tblastn search of genomic short reads from either *Hypothenemus* or *Hycleus*, the predicted protein sequences after translation of the reads with the smallest e-values are more similar to the *Tribolium* tachykinin receptor. Yellow highlighting indicates identical amino acid residues between the short sequences and the two *Tribolium* receptors, red highlighting indicates different residues. Black highlighting indicates conceptual translation of residues that are part of an intron. Note that in all cases the short sequences are more similar to the tachykinin receptor. Hence, one can conclude that neither of these species has a natalisin receptor.

1

Pogonus	MHRRW	AVVCVYAASLATA	LPVTLLEDAKAAEAHLE	NKVKRSHMDE	SLNLNAE	SNSGAPVAYFSN	PSASKRA
Ignelater	---	MRSRLIWLGVVSYLT	TLVSSLP	TSLVEDVKANEIK	DNKVKRAHSDT	EENPRISTAYYNN	PTAIKRG
Photinus	---	MPSKLAWLGVISYLT	FASTLPT	TSLVEDVKANSIK	DSKVKRARS	EENPHLPATYYKN	PSAIKRG
Aquatica	---	MRSNFAWLSTIISYLT	FANALPT	TSLVEDVKANSIK	DNKVKRAHSDS	QESSHMPVYYKN	PTAIKRG
Nicrophorus	MKHTLRALP	FLLYFAALALALP	PASVVEDVKSSDIK	NSKVKRAPVNAE	---	S HGESR	PTAVKRG
Aleochara	MHPAHTGRT	LAVLLTVIVVSSAIP	PASVLEDLISHDVA	NKKVKRAQVAVSK	ENPQPAQ	KELFY	PSAVKRS
Oryctes	MILSRHLPL	LLTVFSCILLISGLPT	TSLVEDVKDNEVR	NIKVKRAHDTLSR	---	SEKAPHDSTPYLSN	PTAIKRG
Coccinella	---	MGVOTRLFLFLIFCSYGC	SAIPSSMIDQIKAREMR	PNKVKRVSTAAEE	---	PEESLIGNHYGN	PTAIKRG
Harmonia	---	MGLQSRLLILIFCLYD	ASATPSSMIEEIKARELR	PTKVKRVSMIAEG	---	PEESMKIGNHYGD	PTAIKRG
Dendroctonus	---	MAKWAFSSLLGAYL	VAVSLALPSSILVQEIKNSEMK	SNKVKRAQSD	---	SNEPEENIGKMSYFTNR	RPTSAIKRG
Hypothenemus	MNSKW	SEFNCLLVASMI	VLITWALPSSLIEELKNNVNS	GHKVKRGQPD	SGTSGP	SAEENMINTNYFDK	PTSIAIKRG
Anoplophora	MGLRHVILD	FLIVASSIVLVIGIPT	TSLVDEIKSSELR	DNKVKRAHPSDLL	---	TSEENGDDVPYYNK	PSAIKRG
Leptinotarsa	MGLRQVYLD	LLIVATLVASAAAIPT	TSLVEEIKSSELR	NNKVKRAHPPNLA	---	SSEENGEIIPYYTK	PSAIKRG
Aethina	MD	SHFGARLFVLTSC	LALVCSIPASLVEEIKSSELR	DSKVKRTPS	---	NGEONSEIQYFTK	PSAAKRG
Hycleus	MEVIWL	CFKWIVWTAYL	TTVLAIPGSLVDEIKANELEANNKVKRAHSI	---	---	VEETVKDVPYYGK	PTAIKRG
Tenebrio	MELRC	SLKWATLASCMVLT	LAIPASLIEEIKASELR	NNKVKRAHPPM	N	NVEERSRDVPYYSK	PTAIKRG
Tribolium	MELRW	SIRWATLASCLALS	FAIPASLVEEIKTNELR	NNKVKRAHPQL	N	VGEHGREVPYYSK	PTAIKRG

78

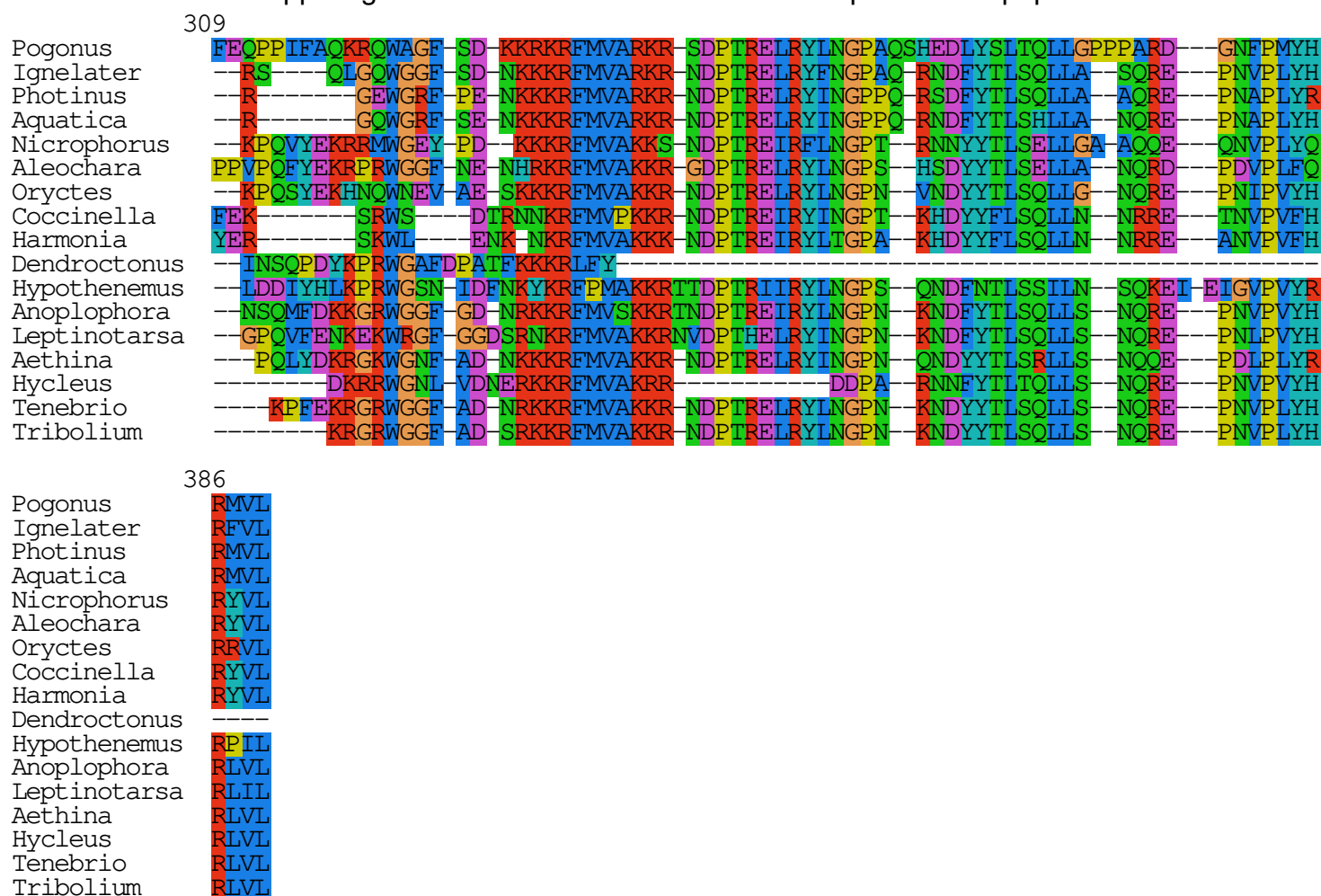
Pogonus	SNLNTDV	SDEATNEWAE	OOOOOOOOOO	PKIDPLK	MIGYDDNPFE	DKTIAEYEGKFHYGTINKAKLDEQ	MENA	
Ignelater	TSQ	TLGGWEDQS		KESLANLAPS	LYSSDDGQFD	DKSLLDYEKAYRYGTINKEKLDEA	LENA	
Photinus	TSQ	PFDNWEDLN		KEFTINLAPS	LYSN-DGKID	DKSIVDYEKAYQYGTINKEKLDEA	LENA	
Aquatica	TNQ	PMENWEDLN		KESLLNIAPS	LYSTYDGKFD	DKSLLDYDKGYGTINCKDLDEA	LENA	
Nicrophorus	INLNKGQ	LPLLTTEWDQORFRPS	EEEEFAASP	LYTSED	ELGN	DKTVAEYEGKFYGVNKKDLDEA	LENA	
Aleochara	EGNRKLS	SQLPLLTTEWDQORYR	EQAERVSPS	LYSSF		DDSPLEYAKALRYASNRDQ	LEKQNA	
Oryctes	INFND	QIGGWDDQ	SIFONT	DIPSS	LYDS	QI-DDKSTIAEYEGKRYGTINCKDLDEA	LENA	
Coccinella	TEMKNEAADDESSDDNDWSP	SEQAIYQVND	DFQST	VYNDE	SSQSANDMDDYDKGLQYGN	REKFDEA	IENA	
Harmonia	TEMKNEVPEDESSSDTDWVQSQ	GIYQVDN	DFQST	SYNDE	SSQSAKSVDYDKGFQYGENREKFDEA	IENA		
Dendroctonus	INLKEAD	MGEWNRDQ	YGGRDFSLDNIQSG	LYDFPNLPMD	DKTIAEYEGKYGTINKEKLDAALQ	NA		
Hypothenemus	TNFQDKY	LGDDLLDFG	N	GDDIQSG	LFN	SPLD	DKSLNEYEGKYRYGTGKEKLDEA	LENA
Anoplophora	TINLNKSP	LPDQOSLNEWEREC	SIFRNOAVLNPISPS	LYNSPDDSYD	DKTIAEYEGKFYGTINCKIDEA	LENA		
Leptinotarsa	ANIDLKSLP	DLDWEREC	ADFINRNTILNPISSS	LFQYEPENNFD	DKTIAEYEGKYRYGTINKEKLDEA	LENA		
Aethina	AN	LKNANAEQOSLSDWTQDC	VLT	DSLSSIQSS	LYNNPNQ	YD	DKTVEEYEGKFQYGASKDKLDEA	LENA
Hycleus	TNQ	LPLIILSP	LDWEHEQOTFYENPENLANIQSS	LYNSD	NFD	DKTIAEYEGKYHYGTINKEKLDEA	LENA	
Tenebrio	TN	SLKNPTDQOSLNEWEQEC	SLYQSP	EGLANIQSG	LYNNADAPFD	DKSVAEYEGKFYGTINKEKLDEA	LENA	
Tribolium	AN	NLNKNSP	EQOSLSDWEQEC	SLYQNPDSLADIQSS	LYN	AENPFD	DKTIAEYEGKFHYGTINKEKLDEA	LENA

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Pogonus	ILKSEMYGDPAST	NOYRYYGGSDRRRRKRSN	LKRSASFANRYKREVELSPEDILTLLALWDEDRLR
Ignelater	VLKSELYGDPGAL	NOYRYFDGSSDERRR KR	RAIKNLRLNSRYKREIELTPEEVLTLTLWENERRR
Photinus	VLKSELYGSPGSA	NOYRYFEGGNDEKRR KR	AAF KRLNSRYKRDIELTPEEFLTLTLWENERRH
Aquatica	VLKTELYGNPASF	NOYRYFDADER KR	GT YKLKDRFKRDIELTPEEFLTLTLWENERRL
Nicrophorus	ILKSELYGEPAAV	NOYRYYGMEGD KKRKR	RNAQKTRMDNRYKREVDLSPEDILAILTLWENERQH
Aleochara	ILKSEMYDLPATL	PNQYQYYGNDDRRRRKRRSTNGKLAKKAARVQKARSITNRLKRSADMSPEQILAILTLWENERMN	
Oryctes	VLKSELYGDPVSI	NOYRYYGGVADEKRRRRR	SNAQKMRFDTRIKRDVDLSPEDILTLLSLMENERQR
Coccinella	VLKSEFYGNLEGD	KKKRKK	RNSRQYSYGN AASDDLSPEEVLDTLLTYESERQK
Harmonia	VLKSEFYGNLDND	KKKRKK	RSSRLNGYGN IASDDLSPEEVLLELNLTYEGERQK
Dendroctonus	ILKSELYGDNV	GPSIWPPYEDRRRR KR	ETVSNKRNLTLYKRNVDLTPEVLRLLQFYEKNORE
Hypothenemus	ILKSDLLIDPY	RRYN	NRSKRTVDLTPEGMAALLSLYEKNHR
Anoplophora	VLKSELYGEPGAI	NOYRYYGADD KKR KR	REVKKLRFDGRIKREVDLTPEEILALLSLYESNRQR
Leptinotarsa	VLKSEIY EPTSI	NOYRYFGAEAA KKR KR	RDACKLRLESRMKREVDLSPEEILTLLSLYENNHR
Aethina	VLKSEYYGEPGAV	NOYRFYGGDD RKR KR	RSACKLRLSQRLKRDVELTPEEILTLLNLYEKSRQN
Hycleus	VLKSELYGEPGAI	NOYRYYGNDDK KRRRRR	RDVQKSSRTNRLKRDITLTPEEILSILTYENEHQN
Tenebrio	VLKSELYGDPAPL	NOYRYYENDDR RRR RR	RDARKIRPDNRKREVDLTPEEILTILTYENERQS
Tribolium	VLKSELYGDPAPL	NOYRYYGNDDQ RRR KR	RDARKIRLDSRMKREVDLTPEEIFTLTLTYENER

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Pogonus	Q	---	PAGYRINWPRYNPDIDLDNDREE	---	---	---	NEIDEPEESWLESPV	---	YPGIRNP	---	VHAS	---	PF	---	ARE	---	YAHNOVQ
Ignelater	A	---	ODDYRPTWSRYEPNLDLDSSSEN	---	---	---	D-VEQEDENWLDTPV	---	YPVYPHATNR	---	INQLSGEFVRGNPNYIYEDK	---	---	---	---	---	---
Photinus	T	---	SDDYRPTWSRYEPNVDLDRNEN	---	---	---	E-LDEENWLDTPV	---	YPVYPHATNR	---	---	---	---	---	---	---	ASPNMYIEDK
Aquatica	S	---	ODDYRPTWSRYEPNVDLDRNEN	---	---	---	D-VDQDEENWLDTPV	---	YPVYPHATNR	---	---	---	---	---	---	---	GNPNYIYEDK
Nicrophorus	GPP	---	ANFHPSWQRF	---	---	---	NFMDSTISGDKNEPEIEQNDWLDSPV	---	YPHISEH	---	LSPM	---	---	---	---	---	NPNNYY-DS
Aleochara	K	---	DATPWLNYDEDGYGD	---	---	---	MSSTDQEDWLEAPV	---	HPHATDE	---	H	---	---	---	---	---	MMQSPNNYYDVSPQA
Oryctes	P	---	SNYRNWQRYDSNLDLDKDNQD	---	---	---	SYEEDEDEGTWLDTPV	---	YPHVGP	---	INSM	---	---	---	---	---	PSSN-YYYE
Coccinella	P	---	---	---	---	---	SNLEQDNIWLDVPV	---	YPGATLPSTSNF	---	---	---	---	---	---	---	GPAYHLDRSPSN
Harmonia	P	---	---	---	---	---	GNFQDHAVWLDVPV	---	YPGPVVPNTSNL	---	---	---	---	---	---	---	GPSYHLDRSPLD
Dendroctonus	LK	---	---	---	---	---	PVEDNDEIWFNEPMRYG	---	SV-D	---	RP	---	---	---	---	---	ANNRFHQ
Hypothenemus	-S	---	---	---	---	---	DADEDEDGAWLN	---	YPVVRPHSSIE	---	NS	---	---	---	---	---	VYNTRNL
Anoplophora	QOE	---	SNNYROPWNRYEPNFDVDINQDD	---	---	---	S-ODQDEENWLDNPV	---	YPHATGY	---	DKEL	---	---	---	---	---	GPXYMIDQS
Leptinotarsa	QOPSTI	---	NYRYPWNRYEPNLDLDINQDD	---	---	---	S-ODQDGENWLDNPV	---	YPHATAF	---	EKDL	---	---	---	---	---	GPXYMYEPS
Aethina	D	---	NYRPWQNRLEN	---	---	---	E-LNDNDESWLETPV	---	YPHVS	---	NDI	---	---	---	---	---	GPQYMIDE
Hycleus	---	---	RPWRLDDE	---	---	---	S-NEQDEIWLDPV	---	YPQING	---	NDI	---	---	---	---	---	GPXYMFD
Tenebrio	N	---	GYRPWSGEP	---	---	---	N-NIEEEENWLDAPV	---	YPHAGH	---	INDI	---	---	---	---	---	GPSYLLDE
Tribolium	N	---	GYRPWGLEP	---	---	---	D-NLEEEENWLDAPV	---	YPHATGH	---	NDL	---	---	---	---	---	APSYLMDE

**Figure S2.** Sequence comparison of Coleoptera baratin precursors.

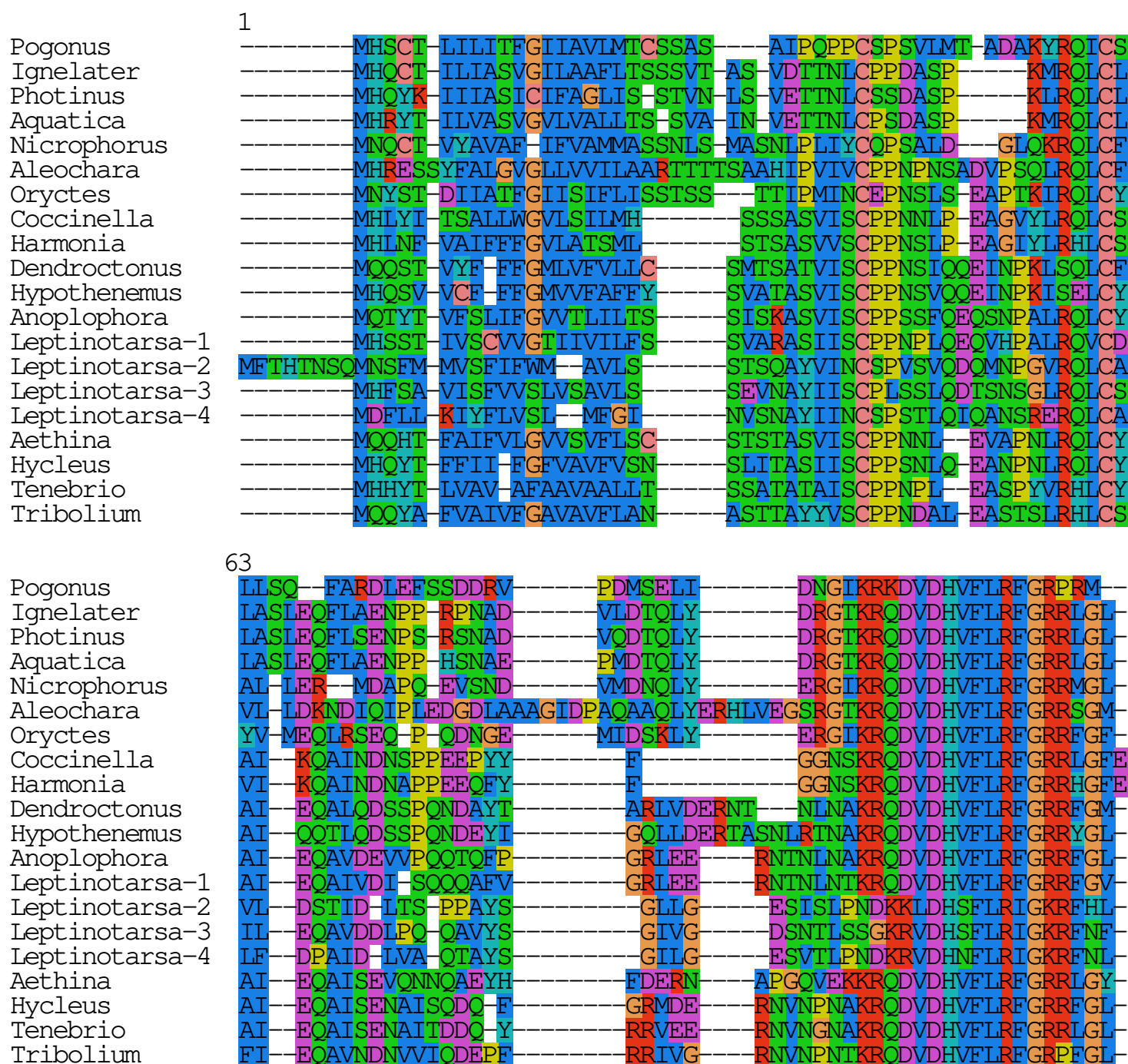
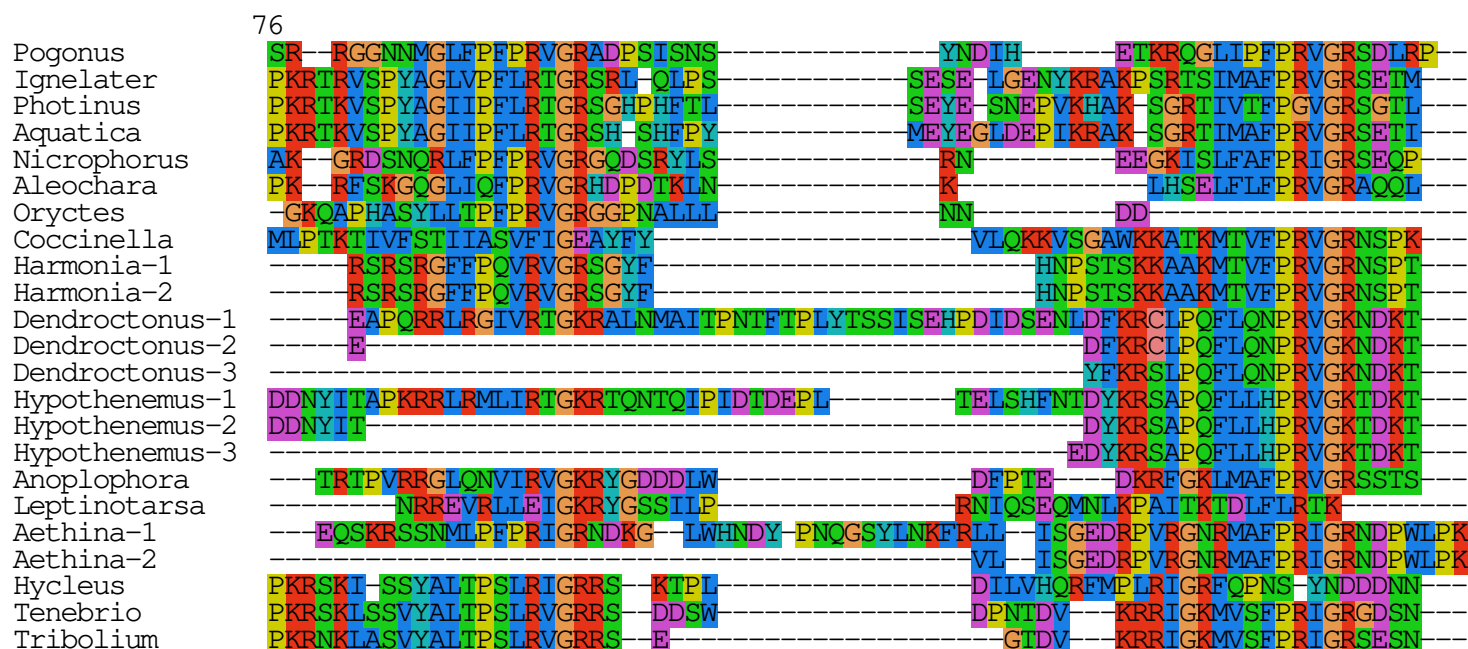
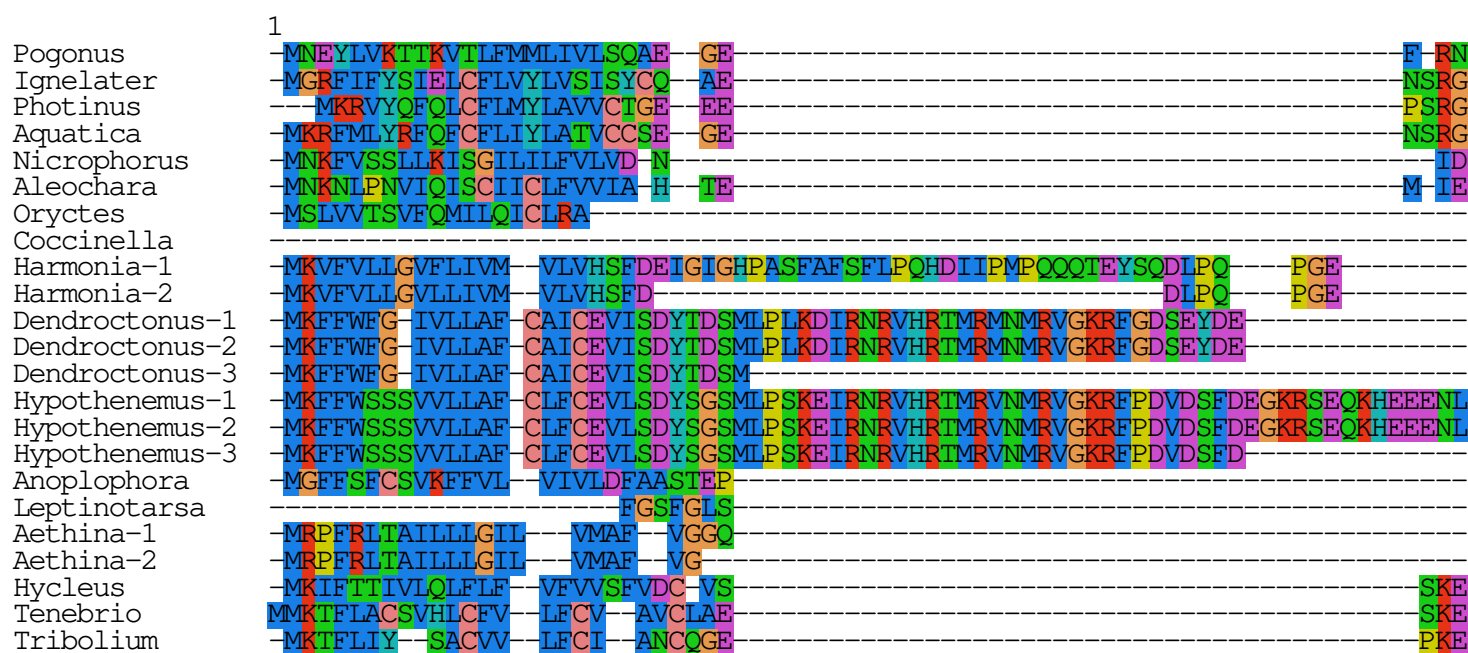


Figure S3 Sequence comparison of Coleoptera myosuppressin precursors.



Figure S4. Sequence alignment of parts of the orcokinin B precursors from the species indicated. Note that the putative convertase cleavage sites consists of a single Arg residue in most species, but that in the two Coccinellids, *Harmonia* and *Coccinella*, all have been replaced by a Lys-Arg pair, while in *Aethina* several single Arg convertase cleavage sites have similarly been mutated into Lys-Arg doublets.



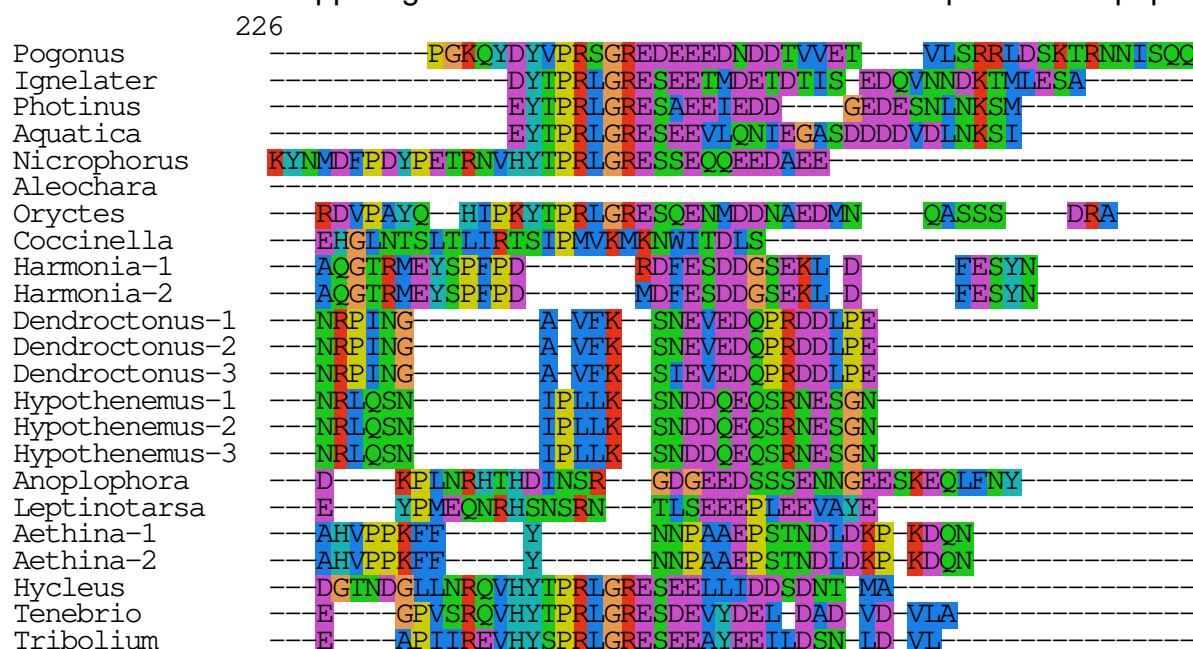


Figure S5. Alignment of periviscerokinin precursors.

ARCHOSTEMATA

Micromalthus debilis (GDOQ01020265.1)

MALLWALVLFPAAILASGTEQEPGIYPQYYSRYPDREPPFAFPEEFQSDQRKTVQKSYEEIAEDLSKLSNNLQ
QKLRFASALLKQENPYETDPADRIEISNLQNSLCILNAEIDNYFRQKAGVFSKGNTQAKRSVWDSKKSTFDSAG
RTSPIELANKSPNIYKSNNSLANAYAQPLFTMKFDATPTDGRTPVQRKRREITKAQLENLLQDEIEKQEEEEWE
RIEGEDDFKKRSQNPPTVVLKSGEVDIPKRVFPNPWGKRLAQFPVNRDLLGVWESKRDMPFDPWGGKRARVP
FDSWGGKRDRTAFDSWGGKRDRLPFDSWGGKRSSRVKLSPWAEKRPKQRKSQFYSWGGKRSKYDATLV*

Priacma serrata (GACO01006445.1)

-----GGDDTDQKRTKFFSWAGKRTNDGTEPSFEFLPSFKRSHSSPNIVVKPREMALPKWTFPHSWGKRAVQV
ESERLPISRVPFNAWGGKRSSDHEDNEFLPLHQEMTSLPHNPEGEFFKSVQTIETEVEVSQLGWKEKSVLTLRN
IKMLPL*

ADEPHAGA

Pogonus chalceus (JU438197.1)

MYKNSVIQTSKILLQIILIYILIQFANTNCMTIPNDNSNMLQTKSIRNRLNLLEDDNIFGTETPTDTKTYNSGN
DMENNNNGDIFTKETDNENIFTEDSDIDDKKLVNKIVSLLDPNEVTPKMIEKLSQFKTFYLSNSNDKTLINL
CTLSRNYINDVCLLIQSSNRSNGLYNPYIYQRCVKLLPLLDYFHCVNEKQNYDETAKENEFKRVPFHAWGGKRF
RQVATEKVQSSKVPFNAWGGKRMNYDDGDEYCKRSKSKFHSWGGKRTVRETANVL*

Gyrinus marinus (GAUY02012896.1)

-----QMLMDASKQVDDAKHVKFLGWIKRFNSNANGNNIPKRIPFHAWGGKRSNQVGTEKIQASRVPFNAWGGKRTYNMNSDESTPDNMYEINTPSSVLPYLVYKRSRTKFHSWGGKRYNY*

Figure S6. Predicted partial or complete Coleoptera leucokinin precursors as obtained from conceptual translation of TSA transcripts found at NCBI. The nucleotide identifiers are indicated in parentheses. Lys-Arg and Gly-Lys-Arg processing sites are indicated in purple and the amino acid sequences of the mature peptides are in green.

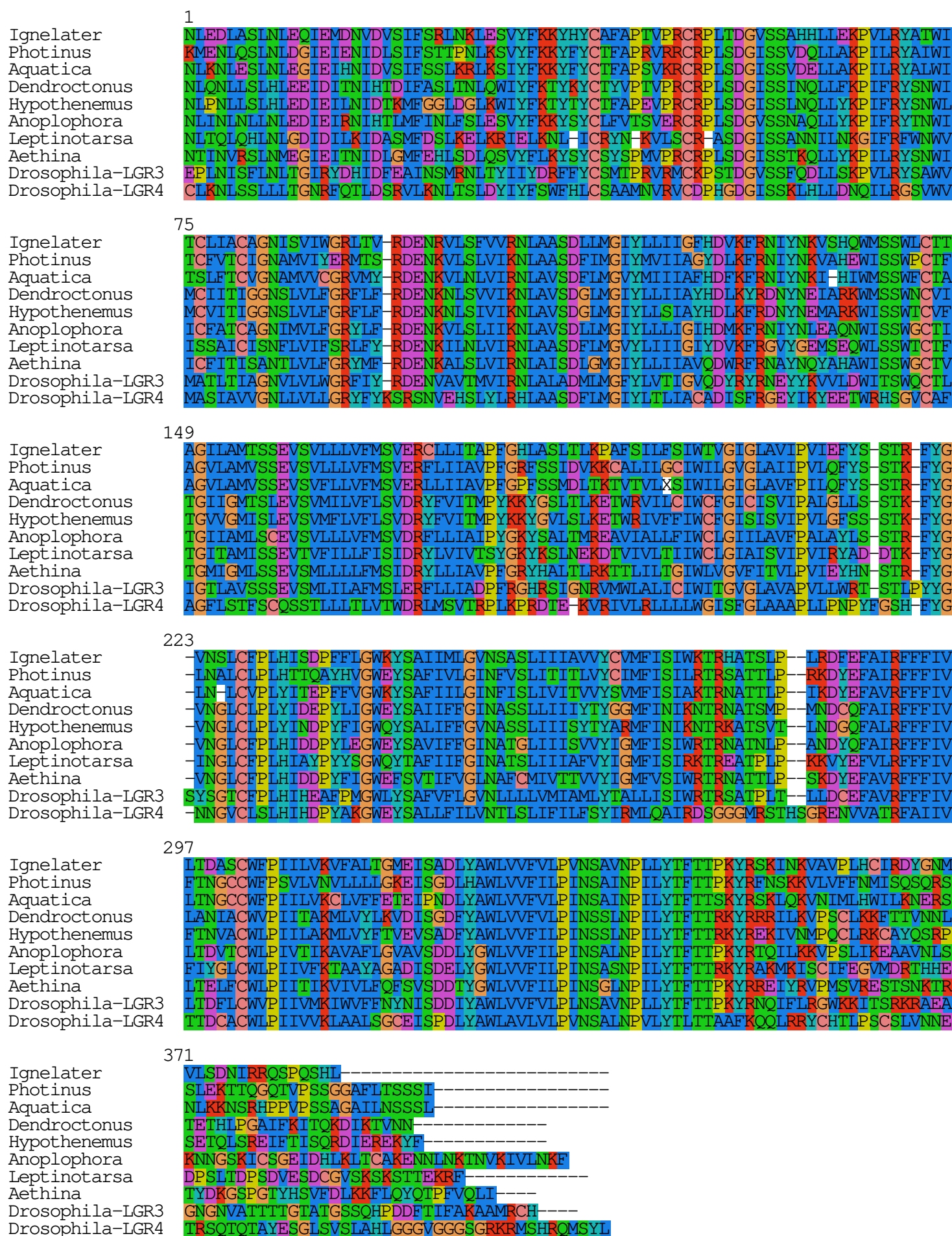


Figure S7. Alignment of N-terminal parts of Coleoptera LGR3's; for comparison the LGR3 and LGR4 from *Drosophila* have been added.

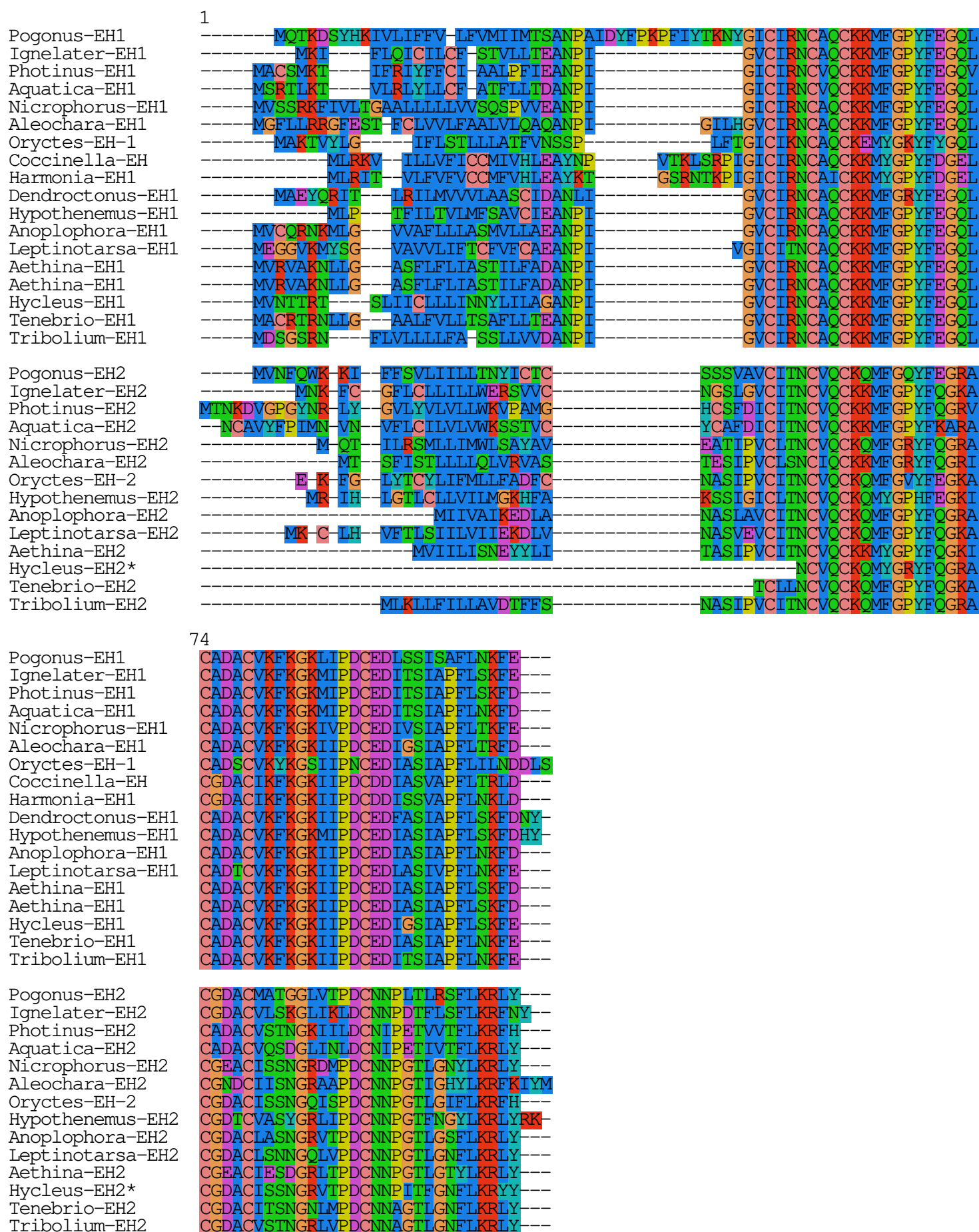


Figure S8. Alignment of Coleoptera eclosion hormone sequences. The asterisk for the second *Hycleus* eclosion hormone indicates the conceptual translation of a pseudogene.

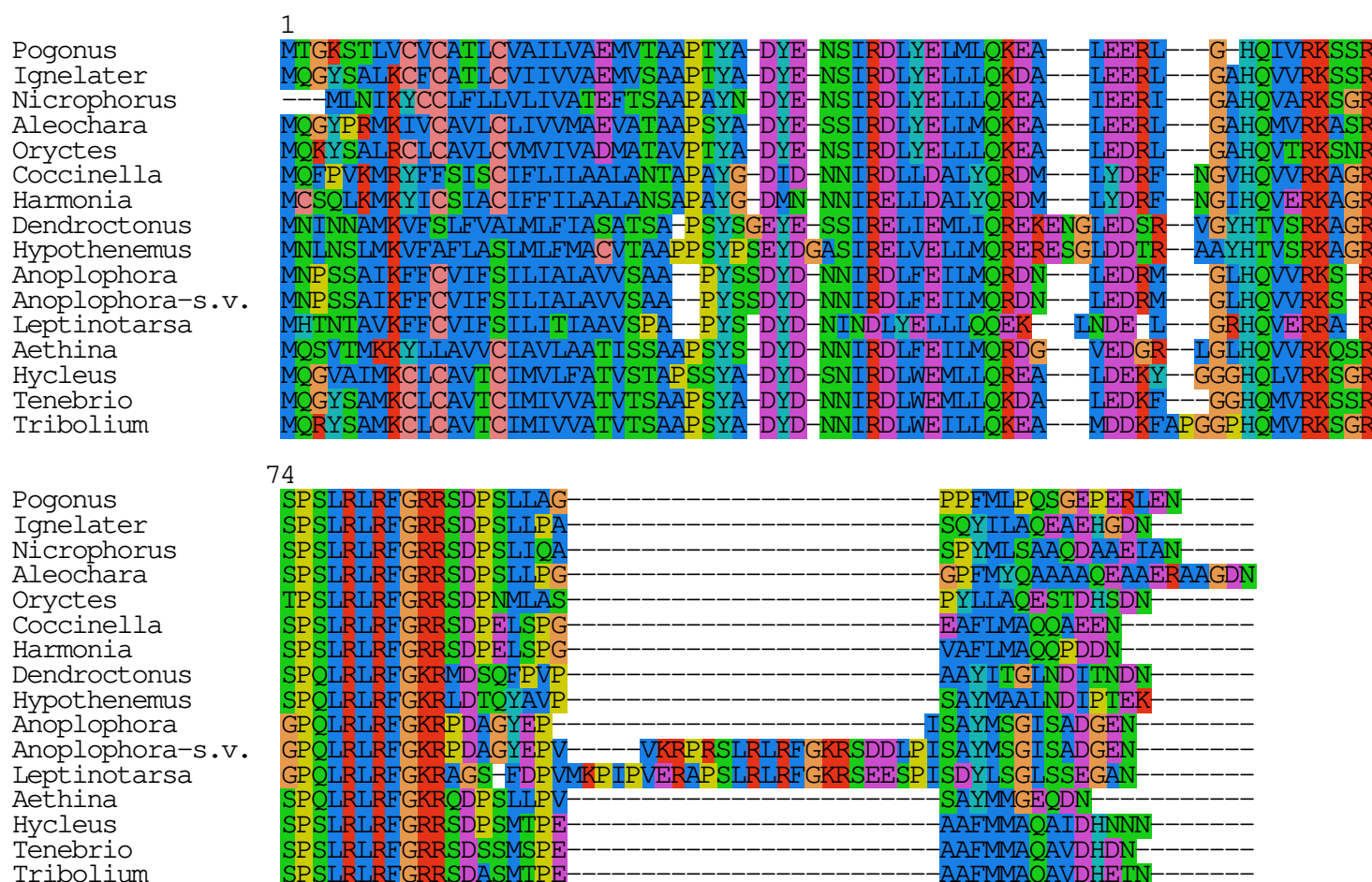


Figure S9. Alignment of Coleoptera sNPF precursors. Note that these sequences are well conserved. The apparent exceptions of *Leptinotarsa* and *Anoplophora* are due to the appearance of an extra exon. *Anoplophora* s.v., *Anoplophora* splice variant.

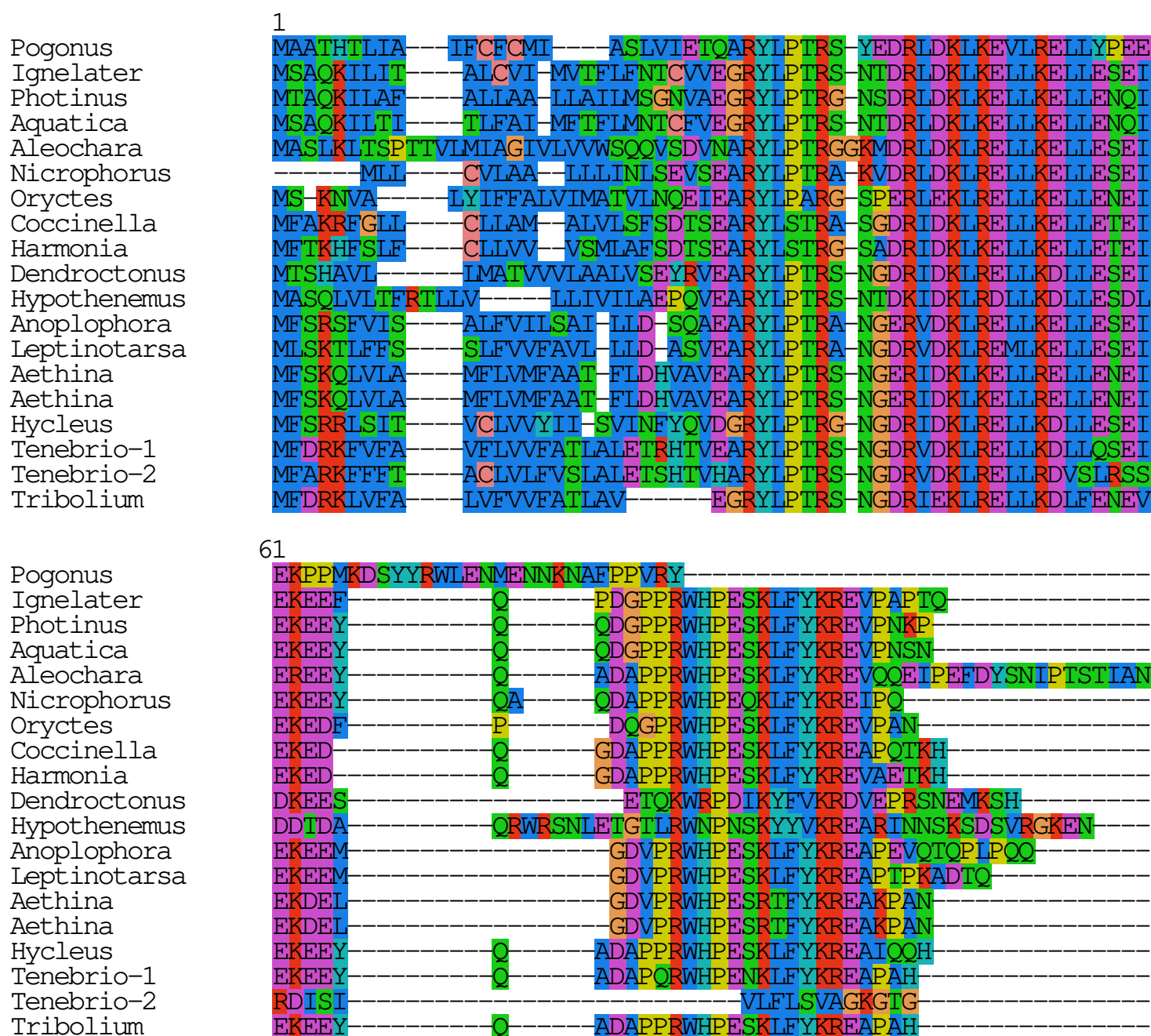


Figure S10. Alignment of proctolin precursors.



Figure S11. Alignment of RFLamide precursors.

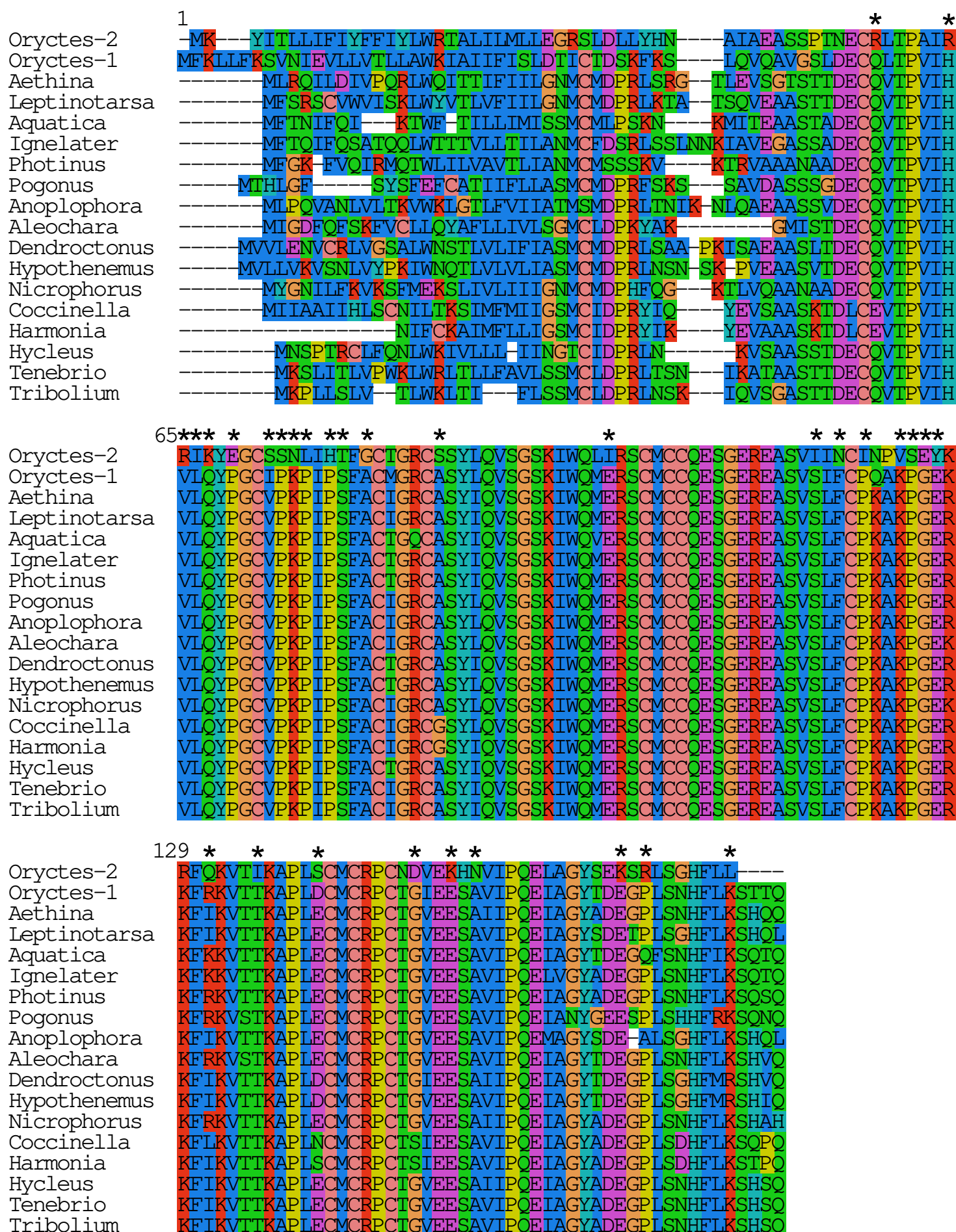


Figure S12. Sequence comparison of Coleoptera bursicon A sequences. Note that the first sequence, that of the second *Oryctes* bursicon A has a large number of amino acid residues that are different from the consensus sequence; those are indicated by the asterisks on top of the sequences.

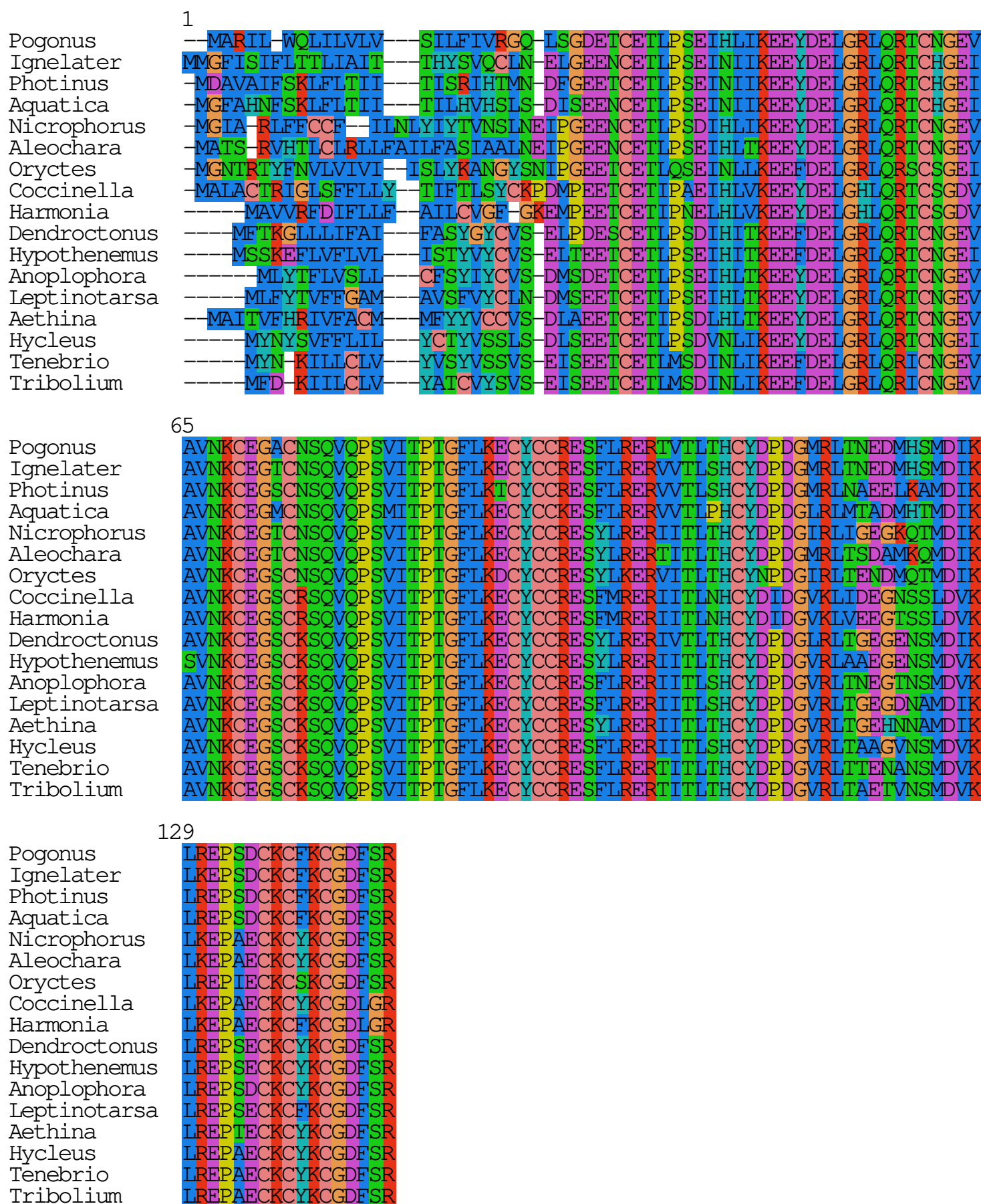
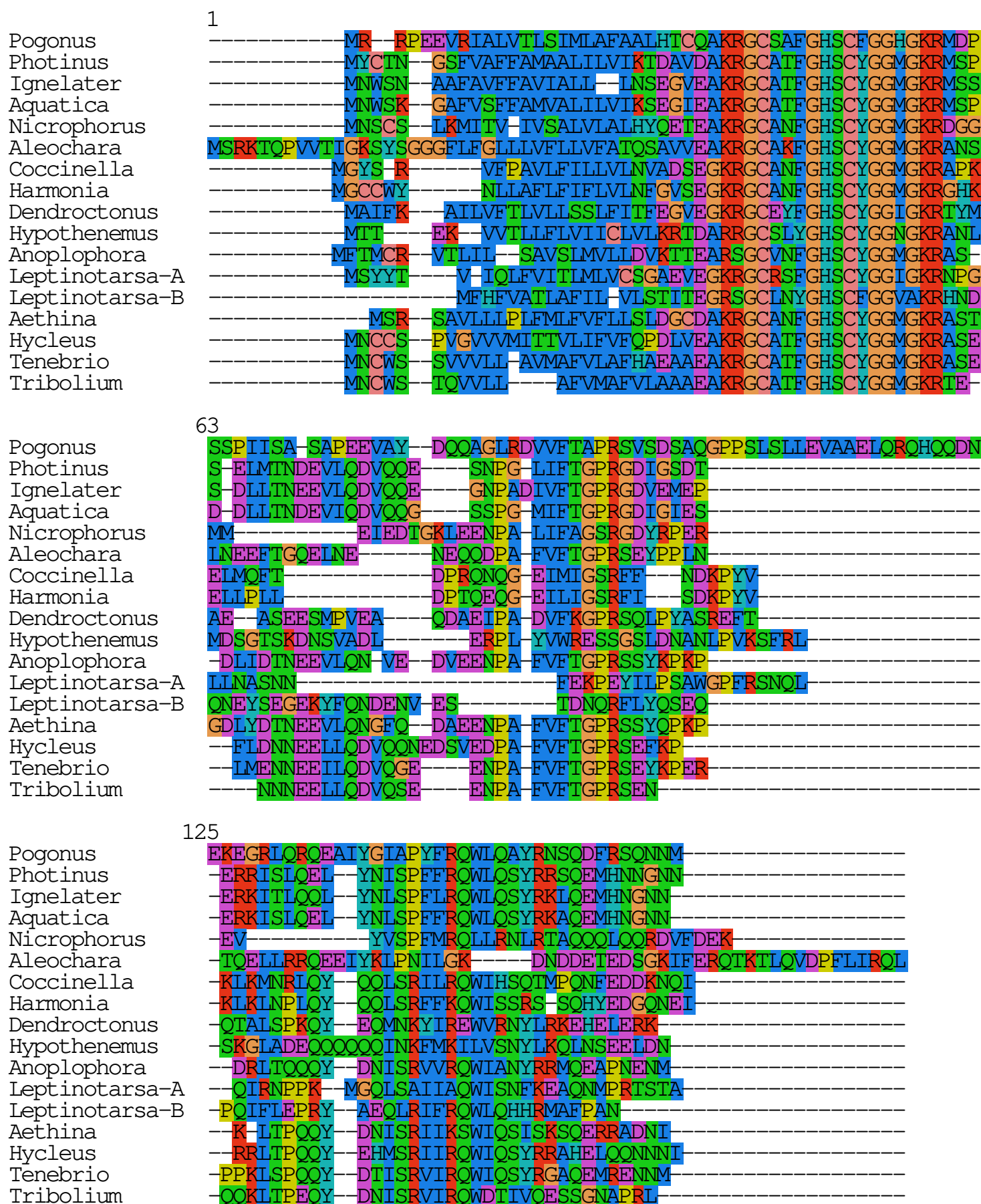


Figure S13. Sequence comparison of Coleoptera bursicon B sequences.



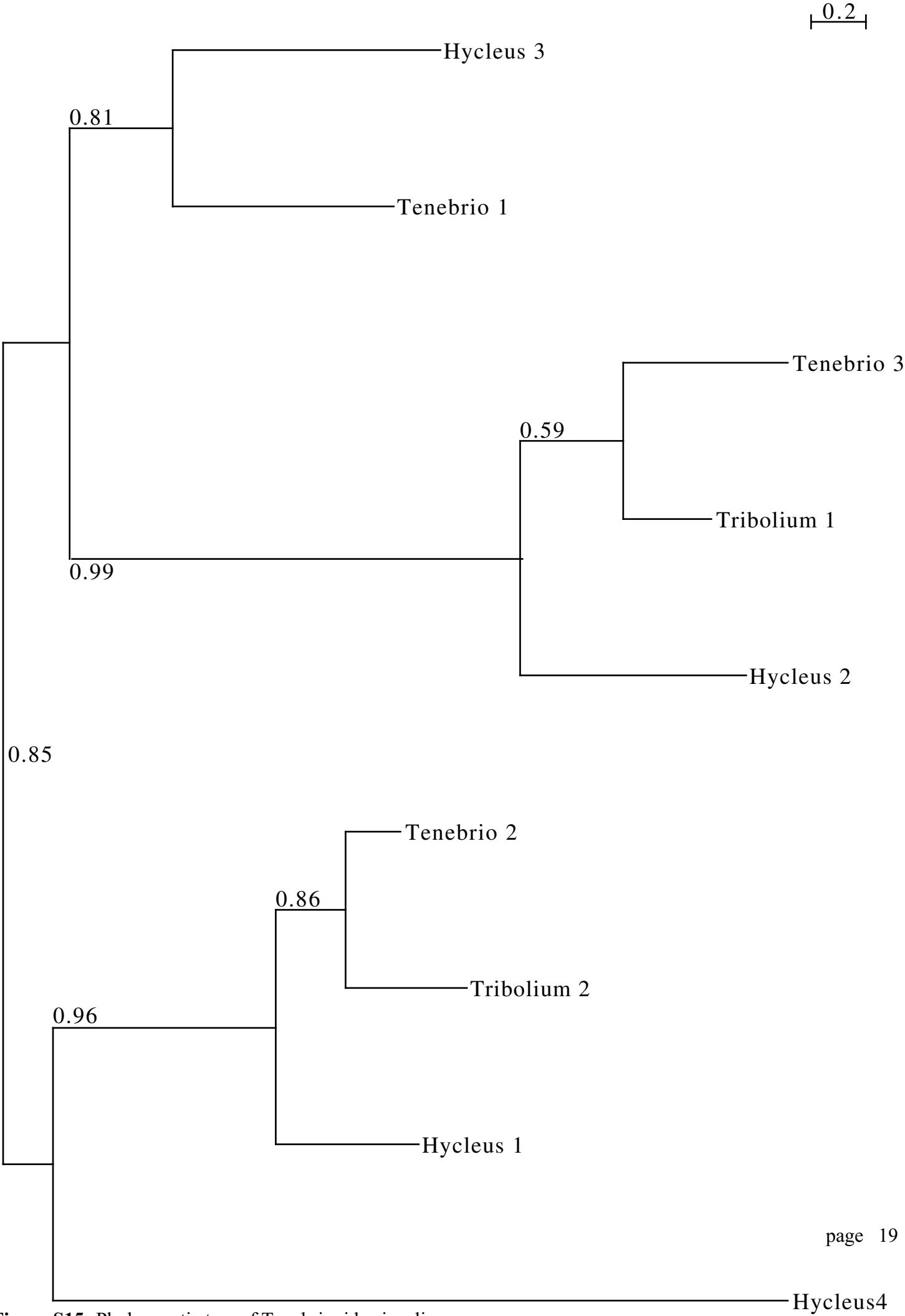
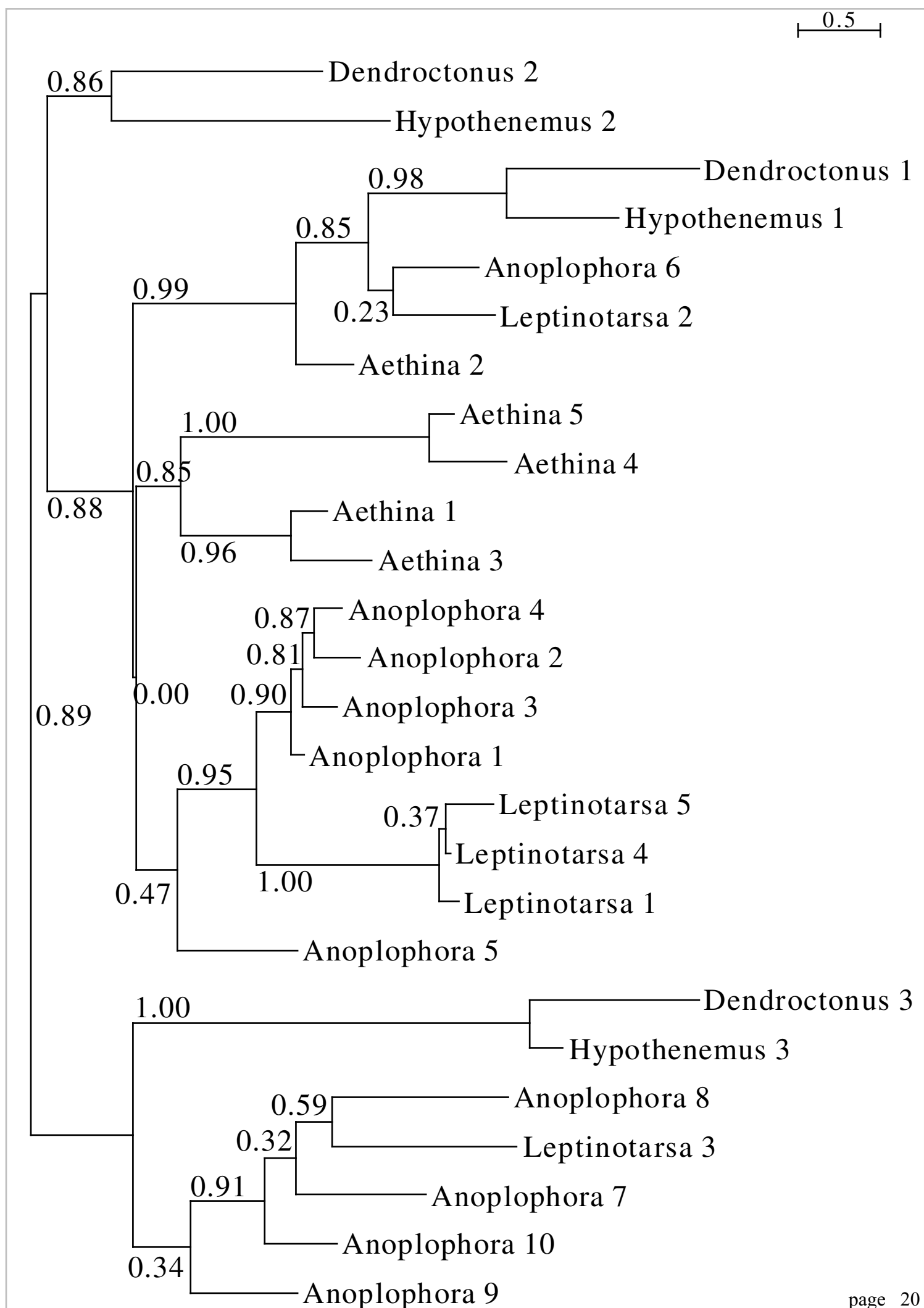


Figure S15. Phylogenetic tree of Tenebrionidea insulins.

**Figure S16.** Phylogenetic tree of insulins from *Dendroctonus*, *Hypothenemus*, *Anoplophora*, *Leptinotarsa*, *Aethina*

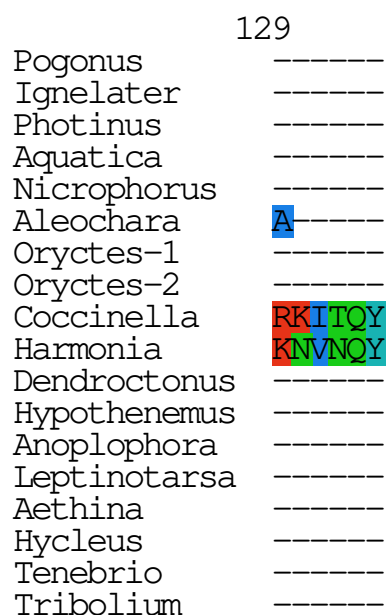
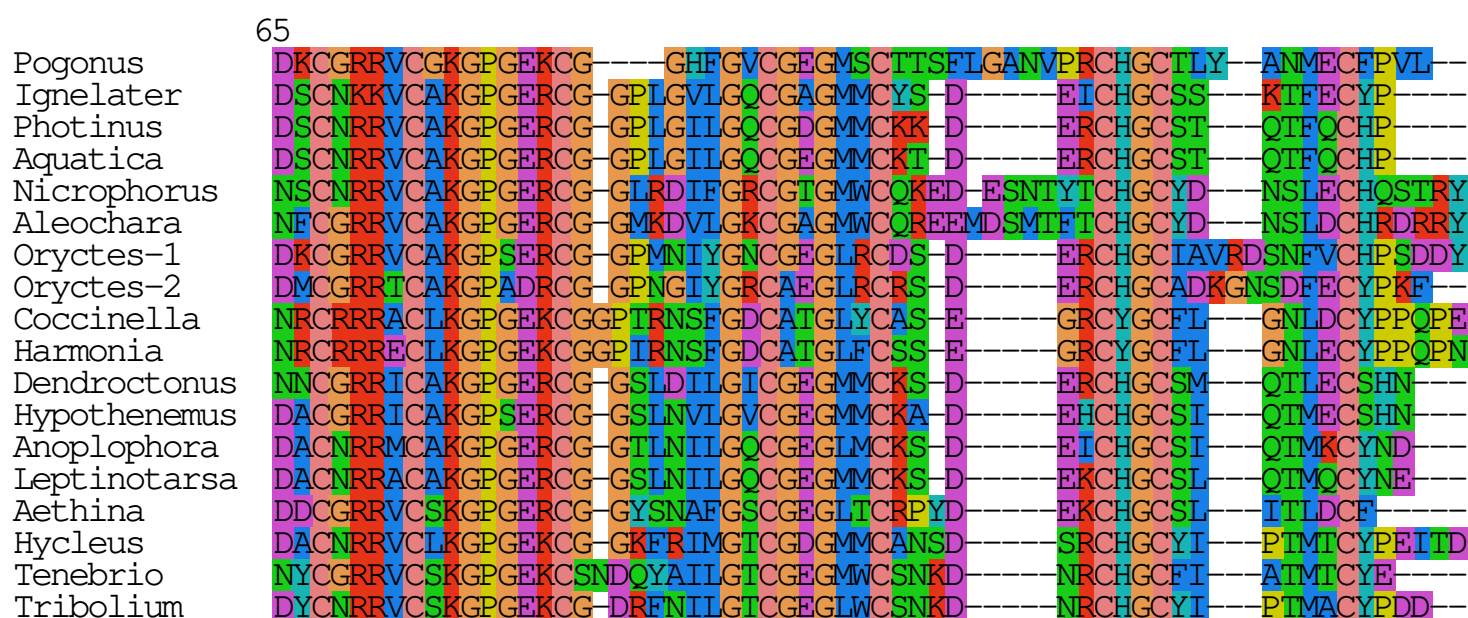
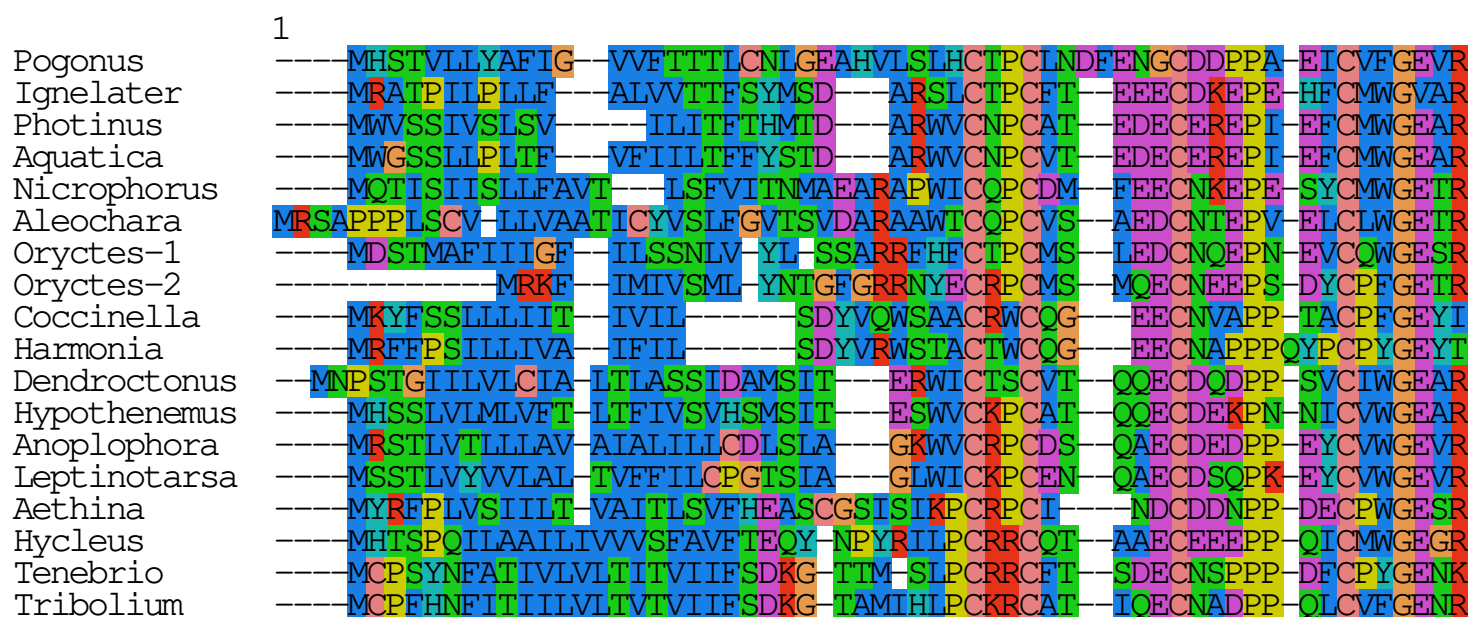


Figure S17. Alignment of Neuroparsin precursors.

Pogonus	---MNNQQ---VISSLVFVVLIVGINAGLEKLD---SNTKISYDM---VGHRL---KR---ST
Ignelater-1	---MPSYRLECCAAFIIVISCVITAAIAEDHNENDKLA---NTATPQKYQWYV---SPRLGRKKKHVSPF
Ignelater-2	---MGRSGLECCAVFLIVSCVFALIIAEEHNGDQLT---NTAVSRDLQWYVRRKPGVWLTPRVGRKKRSASPY
Ignelater-3	---MSTSSRLYKNVLTFRSVHKNHDIIVAIAEDHNENDKLA---NTATPQKYQWYV---SPRLGRKKKNVSPL
Photinus-1	---MGQFCLKCCVLLIT---FLVVVTIANQYNGNHHI---LMAVQLYSNKKPTVLLSSNAQRQKRN---SQY
Photinus-2	---MKQPCATCTVIVAIVT---MAVSQVHTEVS---
Photinus-3	---MKQFYATCAILLTVVS---MAIA---
Aquatica	---MGQSSLKCCAVIFVILNLLFAITTTADQSGDHRK---VIGLOWYVNKKPNTVLSSHILRKKRN---SNY
Nicrophorus-1	---MVRQAIFVLAVLILV---AICHSA---D---NDERSIDSYSKRKMALWF---GPRLGRKKRNPSS
Nicrophorus-2	---MHRQVIVLSVLLVSALLISA---A---
Nicrophorus-3	---MHRQAIVLSVLLVSALLITSA---A---
Oryctes-1	---MEHLPVWSCVLLLLCSICVFTEQA---NQADTRLKLHTASPFWF---GPRLGRNKRNPFS
Oryctes-2	---MDRSLYTFCALLLIFTVIA---D---SSNEKTVDEYAKRKQISPLWF---GPRLGRSKRNOHF
Aleochara	---MDRLVLFICVLLITITQTISS---S---DQOEATGVDGQTNKKYARIWFEP---PRMGRRKRHPT
Harmonia	---MNRFIWFNCILLFNTFDGSAVSL---NDYE---LVEFRKPHHKKHYPAFYSWEGSTMK---EKREPID
Coccinella	---MERFVWFNCVLLLLYTIDGSAVSL---NDYE---LVEFHKPYHKRYPSFYSWEGANIK---DKAEPID
Dendroctonus-1	---MIRISVINSEFLLLIIVLYYAIDSVDSS---FDPAGDHERKESSPMWF---GPRIGRKKRNPKE
Dendroctonus-2	---MGFSVTVICVAIFFLLILRANPSASIEHGNNINGDPSETEINSRGSLSQENYSPAWMDRKF---GKQKINDLL
Hypothenemus-1	---MSNISVFNCVLLAILLYTTEIAES---LDAPN---DRKTSQAMWF---GPRIGRKKRNPID
Hypothenemus-2	---MLRVTCIILVETIVTISMS---STFVIL---SNNIESDATINTAVEKFPWINSRKS---KT
Anoplophora-1	---MAKVTLISFMVFLTLALYFNGVLTAN---SDELIDAKRRDGEKKYSPLWF---GPRIGRKKRNPTE
Anoplophora-2	---MDRIIVACASLIILNIYSHMVLATHDYDS---SLYSSKYREIDSKAAPLWF---SSNLGRKKRNLKN
Leptinotarsa-1	---MSKLSVTFAAFIIMLYFKSVASA---EVDEKRRDEAAKYSPLWF---GPRIGRKKRNLID
Leptinotarsa-2	---MDRIVLVSCVWILIGNFCFELILASQRFDG---SVETKMKEEEESSKMPAWF---SPRLGRKKRNSKE
Aethina-1	---MKQSTKINFALIFMSLIILEICCAESHGNN---LSADKPEHEKHA---SELWHGPKLGRKKRNPSE
Aethina-2	---MTRFTLVNCAVLLIYLEIAL---CSQPT---NVVAALDKQRDDEVKLDNLWF---GPRIGRKKRNDSE
Aethina-3	---MRQSTRINCALVFISLVILEVYCTGSHNYE---NLPDKSGKKTGYEKYISHLWSGARLGRKKRNLDD
Aethina-4	---MKQLTRINVALIFMSLVILEVCCIESQHGN---LSADKSEKPTYH---DKHASQSLPKLGRKKRNPSE
Aethina-5	---MSQLAL---TCTIIMA---MCLHIVLAAPSN---D---VTIDKRNQKTANQMWFGPRLGRKKRNPTE
Hycleus	---MDRINLVNCAVLCIIVLYFGIVLAVPHYE---SAQD---EGGDKENFWFGPRLGRKKRNPQSN
Tenebrio	---MERIILVNLAVLCVAILLSEVLSVPHYG---SHQSVRRERNDDKQSYMFWFGPRLGRKKRNPSSN
Tribolium	---MERFILLNIVLVCVAVLFEVTLSTPHE---SSVPNERNDSSKETYFWFGPRLGRKKRNPSSN

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Pogonus	---NEDITPGLYYNKDDLHATAELIQENP---WILANIPD---MOK---ROMYFTPRIGRDLSEENSNS
Ignelater-1	---EE---TYELSSLDKEQLESLLGNIQEPPEWTAYTISD---GKR---N---SNEFVPRLGREITIEIAEPD
Ignelater-2	---EENNYRTPSFDREQLESIMDTIQEPP---WILVTINE---GKR---HTVNFTPRLGRESSEENEPD
Ignelater-3	---EE---THELSSLNKEQLESLLDNIQEPPGWTAYTISD---GKP---N---SNEFVPRLGREITIEETDPD
Photinus-1	---EDNYRIPAPLDKEQLEALIEAVEEPP---WTVLTFNE---GKR---HTVNFTPRLGRESSEEEAE
Photinus-2	---KLMNFPADNEQVEITMIEATQEHF---WAFITFGG---KKR---SVSFIPRLGRELGDENME
Photinus-3	---DE---RNEFTIGG---KKR---GVSFIPRLGRELGDENAE
Aquatica	---DDAFITISSPLEKEQMEAFILKALQEP---WT---LTFNE---GKR---HTVNFTPRLGRESSEEEENE
Nicrophorus-1	---DEL---LKNINLDREQLVALLEMLQESP---WAVVAINE---GKRH---NKMNFTPRLGRESGEDLAGS
Nicrophorus-2	---FDGQND---YR---SMFN---K---GKP---STYDFVPRLGRESGEDMNAY
Nicrophorus-3	---IDGRND---YQ---SMFT---K---GKP---STYDFVPRLGRDSGEDVNAY
Oryctes-1	---PDT---YSLSILITCEDVEE---YVKVIP---WDLVTKMISE---GKRDI---KSRMNAYIPRLGRESSEEFISS
Oryctes-2	---DDI---EILLASECDEIDEYIRSIPOSM---FERCSINEDKR---DKP---KSKQFSPRLGRDSYEDYFGM
Aleochara	---MEY---YKTFPMDHQOLETFLEFMOQSP---WADFTLPPNEN---AKR---YDVNFVPRLGRESSEEMGNA
Harmonia	---DEY---LKEFPDLDRK---SFSS---WLLR---MND---GKR---AGRIFPRLINRESNDNELSF
Coccinella	---DEY---VKYSDLDK---SFSS---WLLR---MND---GKR---AGRIFPRLINRESNDNELSF
Dendroctonus-1	---NEI---FIN---SPDQPTVNLDDMLRETP---MVVLAFFSEA---F---KQHNFFVPRLGRESGESLNGY
Dendroctonus-2	---RDKNKMVHILETILDEYL---CSIFTGNDG---MOKRDGHLYTTPRLGRNVEVDNAR
Hypothenemus-1	---PKT---YIN---DKNQPMDLYEALRESP---LVVAVNEA---N---KQHNFFTPRLGRESGETILPNW
Hypothenemus-2	---DVVWF---TGKIKVSKLLEYLQEYP---WTVIALDSRSG---IEKRSNSVNFTPRLGRNLDEEGYSA
Anoplophora-1	---NIF---KT---LDREDLDA---YALKNP---WPLMTFNN---GKR---QMDFTPRLGRDSMEDQEER
Anoplophora-2	---SV---TQHELIDLWEALAKSA---TALVTIDGNDEFVCRSMPPNNRFYLVEGVDKSEFNPRLGKRIE---EIPGN
Leptinotarsa-1	---ESY---EN---ADNENVEELDFIRKSP---WTIMLFNH---GRG---HIVNYPKFSKESPDGINEA
Leptinotarsa-2	---ETIY---R---SSEQELADLLEAFEDSP---LAFIAVNGNEKFICKNIPGKR---SNFIPRLGRNVEDDFSGS
Aethina-1	---DLF---REE---LEQKEQANLFDMLQDTP---WTVVAVGD---GKR---HVSSFTPRLGRELQDDFGSN
Aethina-2	---DH---YNY---LNAEQIONILEAIEQSP---YTIVILNP---TKR---HTVNFAPRLGRDSGEDD
Aethina-3	---EVE---REE---QKE---OLDNLLQILQDNP---WAVVAMTD---GKR---HVSSFTPRLGREVDDEFSGN
Aethina-4	---DVE---REE---MEQKEQANLFDMLQDTP---WAVVAVSD---GKR---HVSSFTPRLGRELEEDFGSN
Aethina-5	---AEGYDY---LEADEIQSILEAIEGSP---YSLVMNP---NRH---KSLSFAPRLGRESSEES
Hycleus	---DET---YKN---SETEQLATIMDAVQDSP---WALLAF---K---GKR---HTLSFTTPRLGRESSEELTN
Tenebrio	---IDF---YRN---REREQLATLLDVIQDSP---WAVVAV---N---GKR---HVVNFTPRLGRESSEEFVSS
Tribolium	---DDI---YQD---MOKEELVSLTDAIQDVP---WAVVAVNDLLE---GKR---HVVNFTPRLGRESSEEFVNN

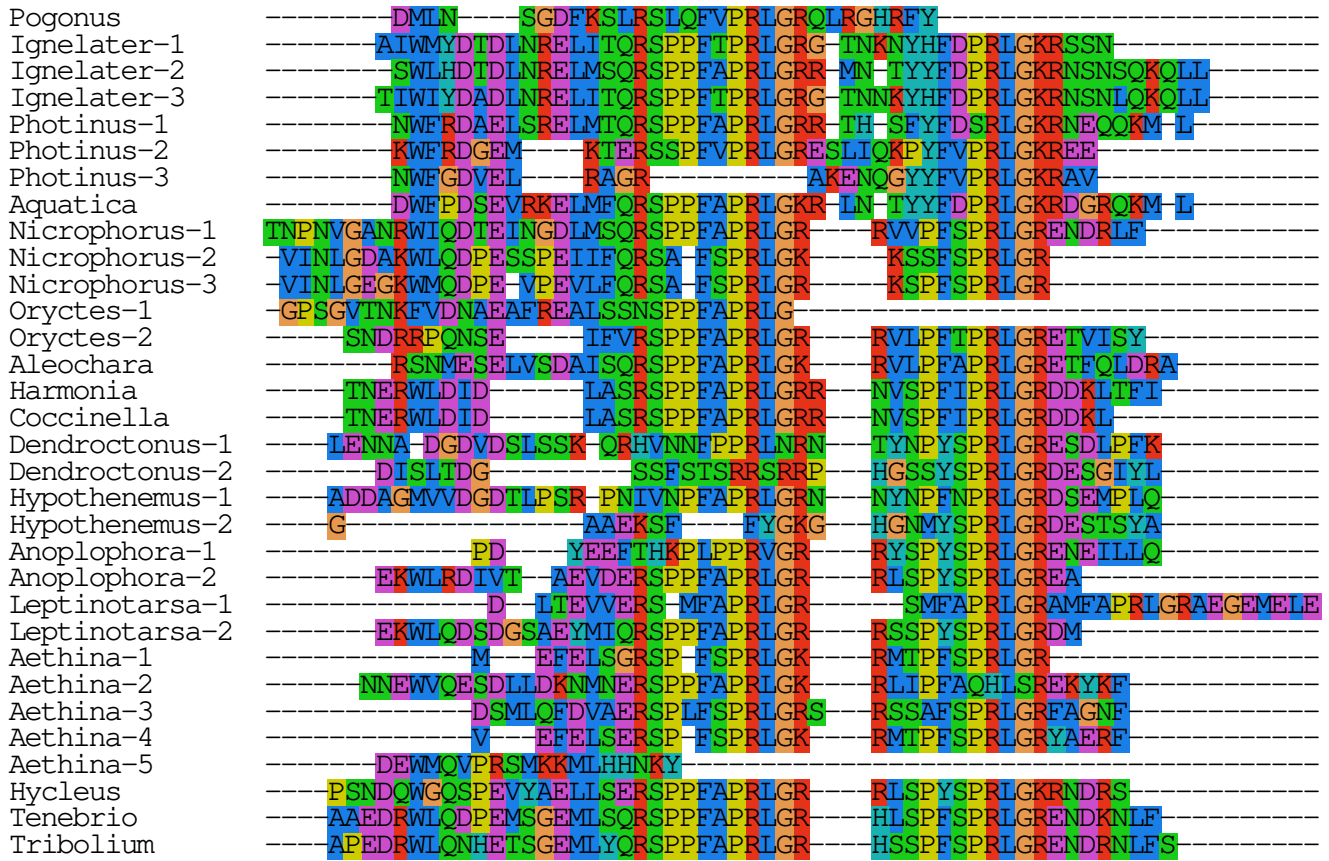


Figure S18. Alignment of pyrokinin precursors.



Figure S19. Alignment of relaxin precursors.

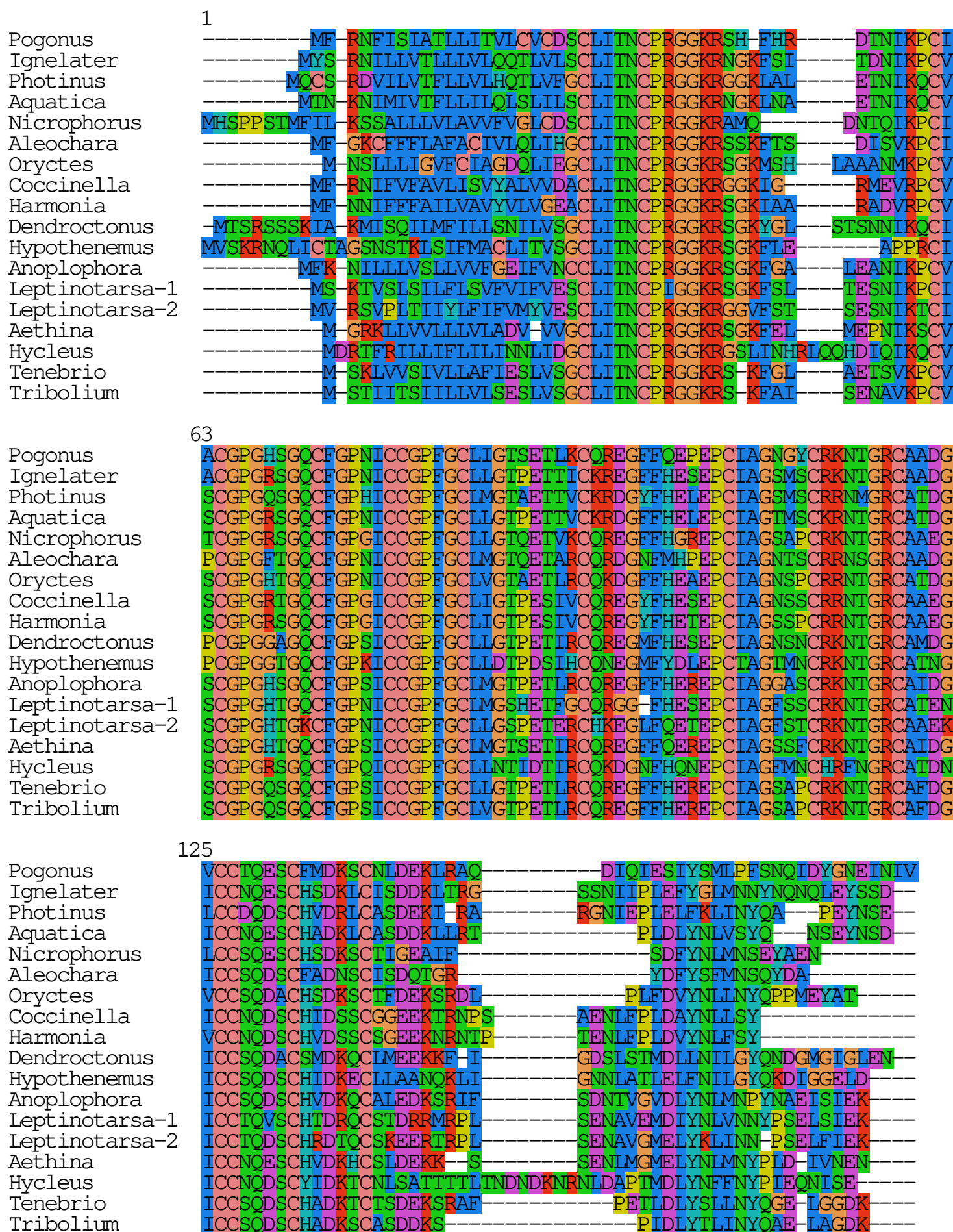


Figure S20. Alignment of vasopressin precursors.

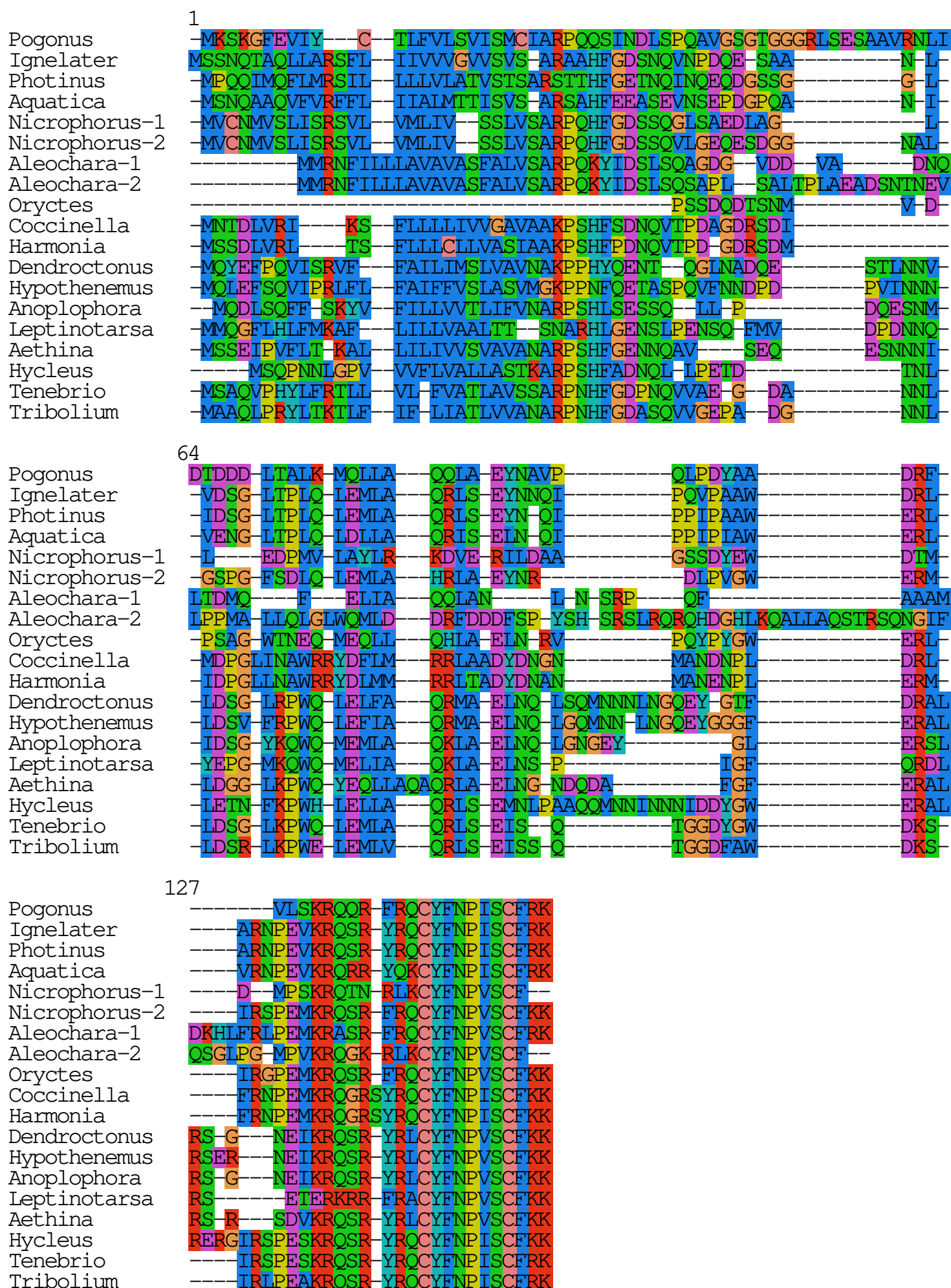


Figure S21. Alignment of allatostatin CCC precursors.

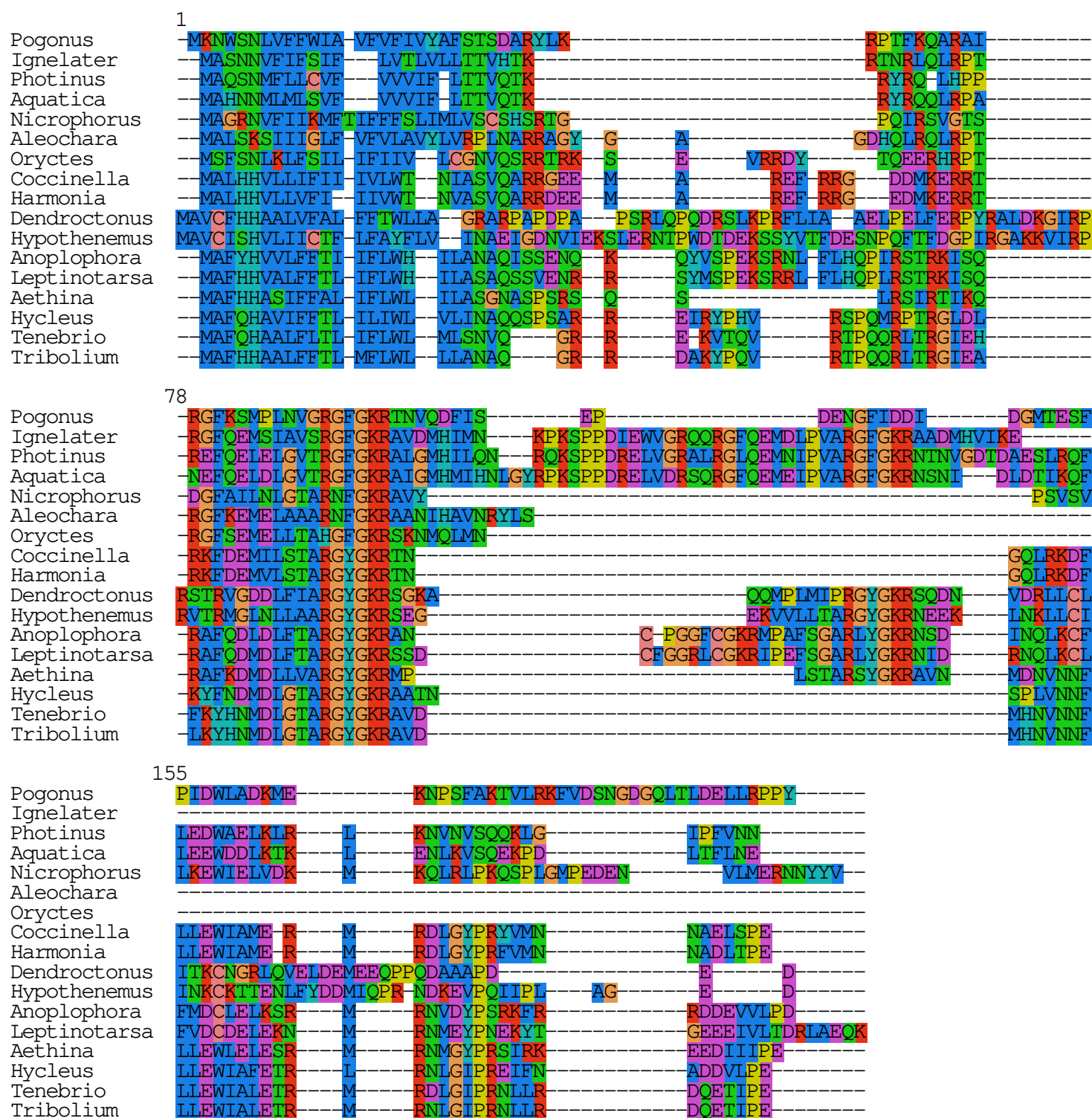


Figure S22. Alignment of allatotropin precursors.

Pogonus

MKTLIILMVISFAFATPYQYIRNRQYQIKHFKNSNQIARTHELPSLQTLFKSFLLKRC SNLGNDDCSNGYLPE
IGNDDDYLGSGNTPGKRC SNLGNDDC

Ignelater

MNGIILLFGIVLSLDYTWGSPTKRCANMFDEGCINGGVGGAGRDEEWIKDPGNTPGKRCANMFDEGCINGGIGG
SGRDEEWIKDPGNTPGKRCANMFDEGCINEGIGGAGSDEEWIKNPGNTPGKRNLFFKNIPRLYSRSVHKNHQGS
TRRFG

Photinus, pseudogene: has no signal peptide and is not expressed

QIRPKCANLFDESCLNKNLIDSALDEQWLSNGHVPGKRNLNDEGKYTFL

Aquatica

MVKYLVSFLLIISISLGSKYVNSARIKRCANLFGEDCINGGTIGGGPDQLWITGNGKQKFLHNLPKVNPKCVNA
FDTNCINNRRLLQEASNQAWSTYPESTYRKENIQDKKVLNMLYLILKGHSIPTKRCANLMDESCINGGIVGSGDD
EELTGDGS PGR

Nicrophorus

MIRFMVLFILVLGTIDAKPKPCHHRDQACLENQVAIHRPPGIFKSLSDFFNRVDIVTSKRASC SNLYDEGCING
GLPGAGDDEDWLNGLGKRASC SNLYDEGCINGGLPGAGDDEDWLNGLGKRAGHKDWMNKSIGKRASC SN
LYDEGCINGGLPGAGDDEDWLNGLGKRASC SNLYDEGCINGGLPGAGNDEDWLNGLGKRASC SNLYDEGC
INGGLPGAGNDEDWLNGLGKRASC SNLYDEGCINGGLPGAGNDEDWLNGLGKRASC SNLYDEGCINGGLP
GASEDQAWLTGNGLGKRVNIPQPGYNKNFKNKNSKTPIMY

Aleochara

MIRIITLPVVFSAVFLISDGAPCANLFDEGCTYRLLKESAMDQGWLRKNHLGKR TFQEHTQIGIDFFHRLHNI
LNDRCNNFFDINCIENPIENHQRVDVNPDAIRAFALQLHEDILNRCAQNAMDKNCFIQNFTPKDDDDKWIA

Oryctes

MKLLTLILLSVGTINGGYIGKKYCPGLFGDSCASGGLDGSADDEDWLNKPNNPGKRCIANVFDGSCSYGATLPG
AAADEYWKQHGS PGKRC SNVFDSCSNGAVVNGGSDDEFLNGNGNPGKRC SNLFDSCSNAGLIGAGDDESWLN
GGGS PGKRC SNLFDSCSNGGLIGAGDDENWLNNGGS PGKRYLPKQWYKAWGPGGVKTAKRHVSDKVVHESLLRK
MAKIRHM

Coccinella

MRKQQFFLMLVLTIHLSQLIECRSYSAAILEEVLRRNSHQNLSPEDVHLFNELYNRLAMPVRKRCANMFDEGCS
NIGIHGSGSDEDWIKVNS PGKR

Harmonia

MKFLFALIMTTIYFLLVESRSFSSLSFEEAVGRNSHPKLSREDIELFNELYRRLSLPVRKRCANMFDEGCSNI
GIHSGSDEDWLKDNS PGKR

Dendroctonus

MNRIITLLAISVALSAAMSVPNVYHKTSNSLKQEQLAALINQLLREKQVYQEVKRECLRMGPLGCLNDPRKHLL
TMYNPGKRCANFGDDGCLSGGVQAGSDSDWLASGFTPGKRCANFGDEGCVTGGVIGAASDSDWIGGGFS PGKR
CANLGDEGCANGGVPGAAADTDWIDGGYS PGKRCANLGDEGCVTGGVSGAGSDSDWISGGFS PGKRCANLGDEG
CANAGVGGAGSDADWLSGDFN PGR

Hypothenemus

MKCIFIVLAVSVCLSFVRSFPNNPDYRQSTQLKQKKLSEIIEDLMRDKERYRGSIREYDGC GSAGVPAGSDS
DWIGGGFT PGKRCANFGDDGCANGGIPGANADQDWLDGGFS PGKRCANLGDEGCANGGVEGAGSDADWLNNGFN
PGRKR

Anoplophora-1

MNFLLAITAVLLGTSQGMYLQRNPHHQAVARQSRNEPTVLKSLGDFFRMHAAATKRCAYLLEDSCNNGGIPG
AGSDNDWLNQGFN PGKRGLNLFEEGAAYNGLSGSGADSDWLNNGFN PGKRCVNTMDESCSNGGIPGSGSDSDWL

DGGFN**PGKR**SLNLFEEGIVNKGVS~~GAAADNDWL~~NGGFN**PGKR**CANTMDES~~C~~NGGGIPGSGEDRDWLDDGSAN**PGKR**RLNGIEGVPGSGNDNDWLHGDNT**PGKR**

Anoplophora-2

MKCLLAITVLLGSSQGFYVERNPHQQLAIARQSRNQPTIFQSLGDLFHRMHATAT**KRC**AYLLDEA~~C~~NNGGIPGAGSDSDWLTQGFN**PGKR**SSKVLEEEPDNDLNPEPDDDRPNGDFNLGKH~~C~~VNAMDES~~C~~DNEEIPGAGPERDWLH**R**GFHPGKPGLHPFEEDNDVPVAAADGGWLSSFSIV~~C~~PNFMDV~~C~~GNVGGKRVPGSGADRDWLEEGGN**PGKR**RLNGIPGSGKDNDWLHGDNT**PGKR**

Leptinotarsa

MNTSTLFLIVLLGFTTAVPLQQWHANKYSSEYHKPNQRGLFSSLGKLFNPSSN**KR**ILNSRGGFSSGERSLNLFD DSVANSKISGSGSDSDWINGGFS**PGKR**CANLMGES~~C~~NNGGVPGSGSDDDWIHGGFS**PGKR**SLNLFDDGAANSKISGSGSDTEWIDGGFS**PGKR**CANLMDES~~C~~SNGGVPGSGSDDDWIHGGAT**PGKR**SLNLFDDGAANSKISGSGSDSEWIHGGFS**PGKR**SLNLLNYASNSKIPGSGSDSDWLNLDGFNS**GRR**GKSSSEKIGHVFGG

Aethina

MRGIIIVVLAMVLA~~VT~~LARYVRRHPADVAMTRAKIHL**KR**NINDPELFKQLGDFHRMHASIR~~C~~SPGLLDEG~~C~~SN GGIPGSGTDADWINNYG**PGKR**CVNILSES~~C~~DNGGIPGAGSDDDWIHDNS**PGKR**CANILDEG~~C~~NNGGVPGSGSDADWINNNS**PGK**

Hycleus-1

MKAIFIIVLVFLSLANCYYLAPHANRPVKVPLRNLNDLLYRLNGLAL**RR**CANLFD~~ES~~CANGMSGAGADDDWL NGNTN**PGKKR**CANLFD~~ES~~C~~S~~NGDINGAGSDDDWLNGDTN**PGRR**

Hycleus-2

MKFVKILLILLTSTY~~A~~IDIHQSNRKSLLKLLINSENYGFLGRL**KRRH**CANLYDE~~SC~~INVHLPLAKIDEDWL RKGFN**PGK**

Tribolium-1

MKTAFLLLFVLM~~PA~~VL~~C~~YYVHPNLHYFPARVNYRSAPAKSWGALFHRLQLAS**KRC**ANTFDE~~SC~~INDVINGAGSD EAF~~LN~~GGDN**PGKR**C~~VNT~~FDE~~SC~~SNGDINGAGSDDDWLHGDDT**PGRR**

Tribolium-2

MRPVLLVLMIFS~~V~~SYARYLEPYNYGYPRPLVD~~FL~~NRLSLSDKI**KRC**GNTFDE~~SC~~CANLPIIGASSDES~~W~~LAHSS**P**
GKRCANVWGESCINGGIIGGGSDQSWLQGDDN**PGRR**

Tenebrio-1

MKA~~AF~~VLLAIALPAAY~~C~~FYLQPNYHVPARLGTRNAPVYKSLGDLFHRHLAS**KRC**VNTVDE~~SC~~INGGGNDAGND ED~~FL~~NGGDT**PGKR**CANLYDE~~SC~~SNGGINGAGADDDWLHGGNN**PGRR**

Tenebrio-2

MKLSLLIFAVALSTACGLYMPKHRPRDEEVFQSLASFFNQLGRRANM**KRC**ANTFDE~~SC~~LN~~GP~~IGGATSDENWL SNGS**PGKR**C~~SN~~IFGSS~~C~~VDGGTAGAGADEDFLGGGG**PGRR**

Figure S23. Sequences of the various Calcitonin B genes from Coleoptera. Mature peptide sequences are in light blue the conserved C-terminal proline in dark blue, Gly residues that expected to be transformed into C-terminal amide in green and convertase cleavage sites and dibasic amino acid residues anticipated to be removed by carboxypeptidases in red. Signal peptides are highlighted in yellow.

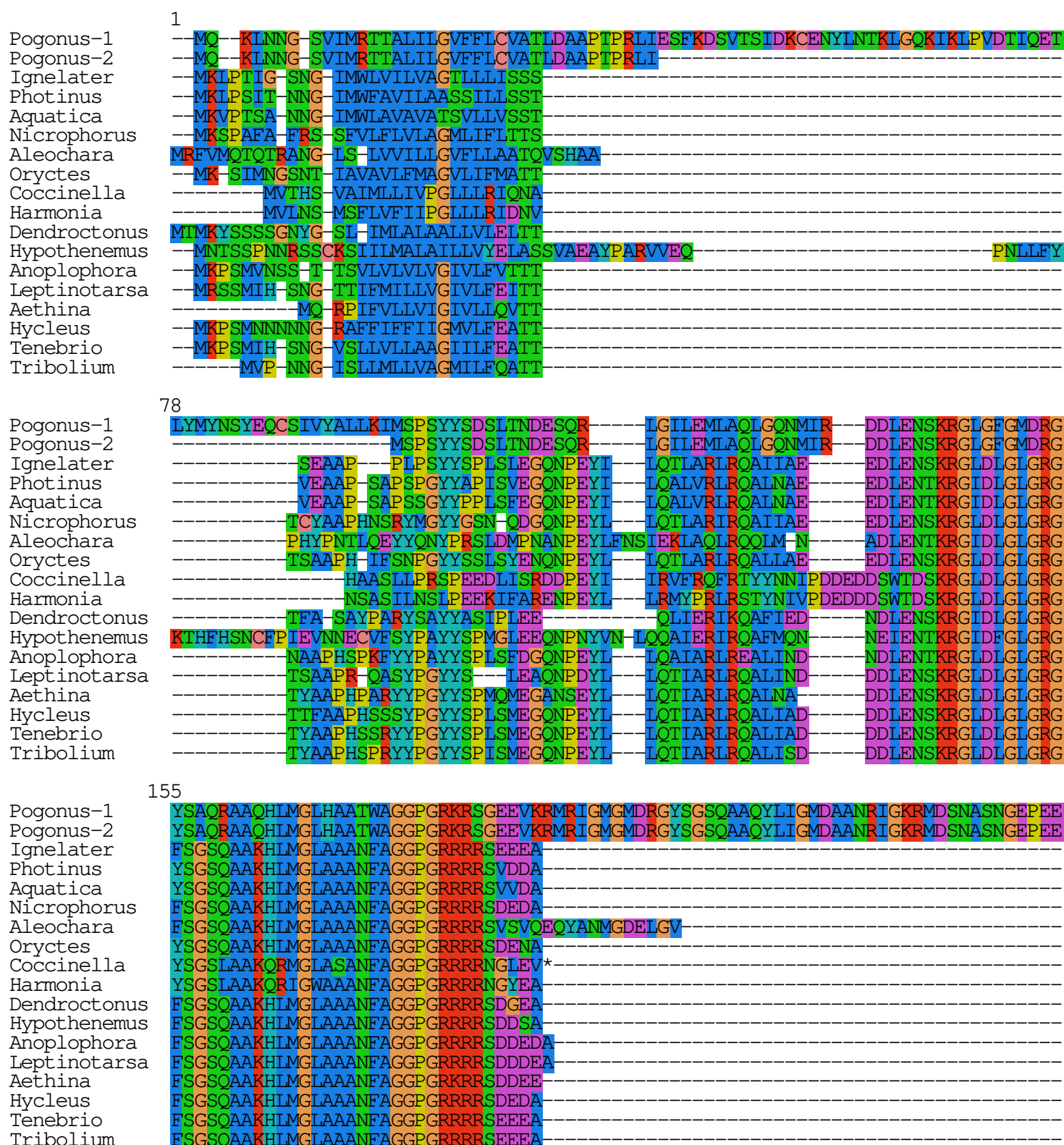


Figure S24. Alignment of DH31 precursors.

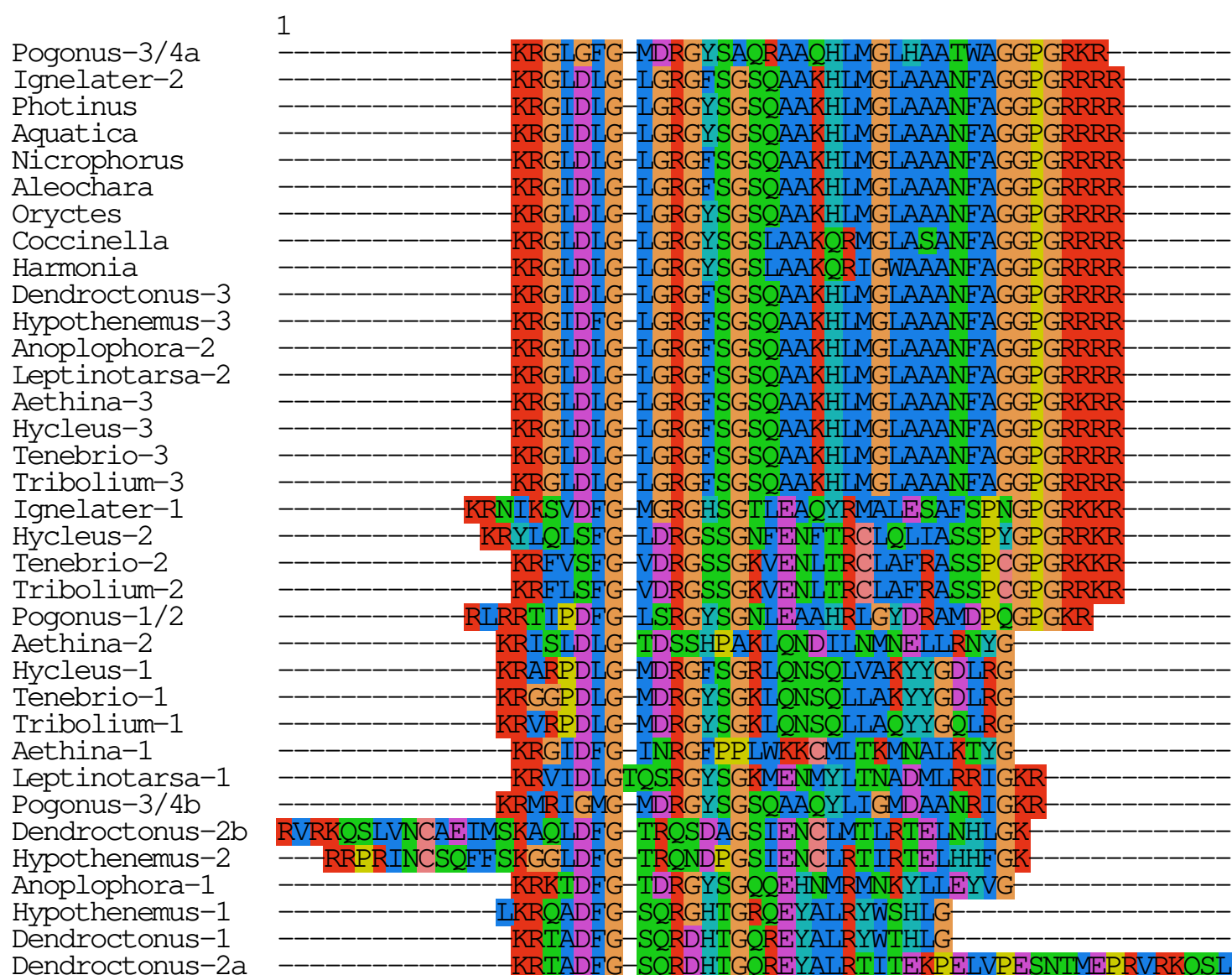


Figure S25. Alignment of DH31 and other putative neuropeptides encoded by transcripts from Coleoptera DH31 genes. Numbers behind the names indicates the transcript number (some species seem to have only one transcript). When this number is followed by a letter it indicates the putative peptide on that particular transcript. For example Pogonus-3/4a indicates DH31 itself that is coded by two transcripts in Pogonus and a refers to the first putative neuropeptide on this transcript. Pogonus-3/4b refers to the second putative neuropeptide on this transcript. Note that sequence conservation is very strong for DH31, but much less for the other putative neuropeptides encoded by these genes.

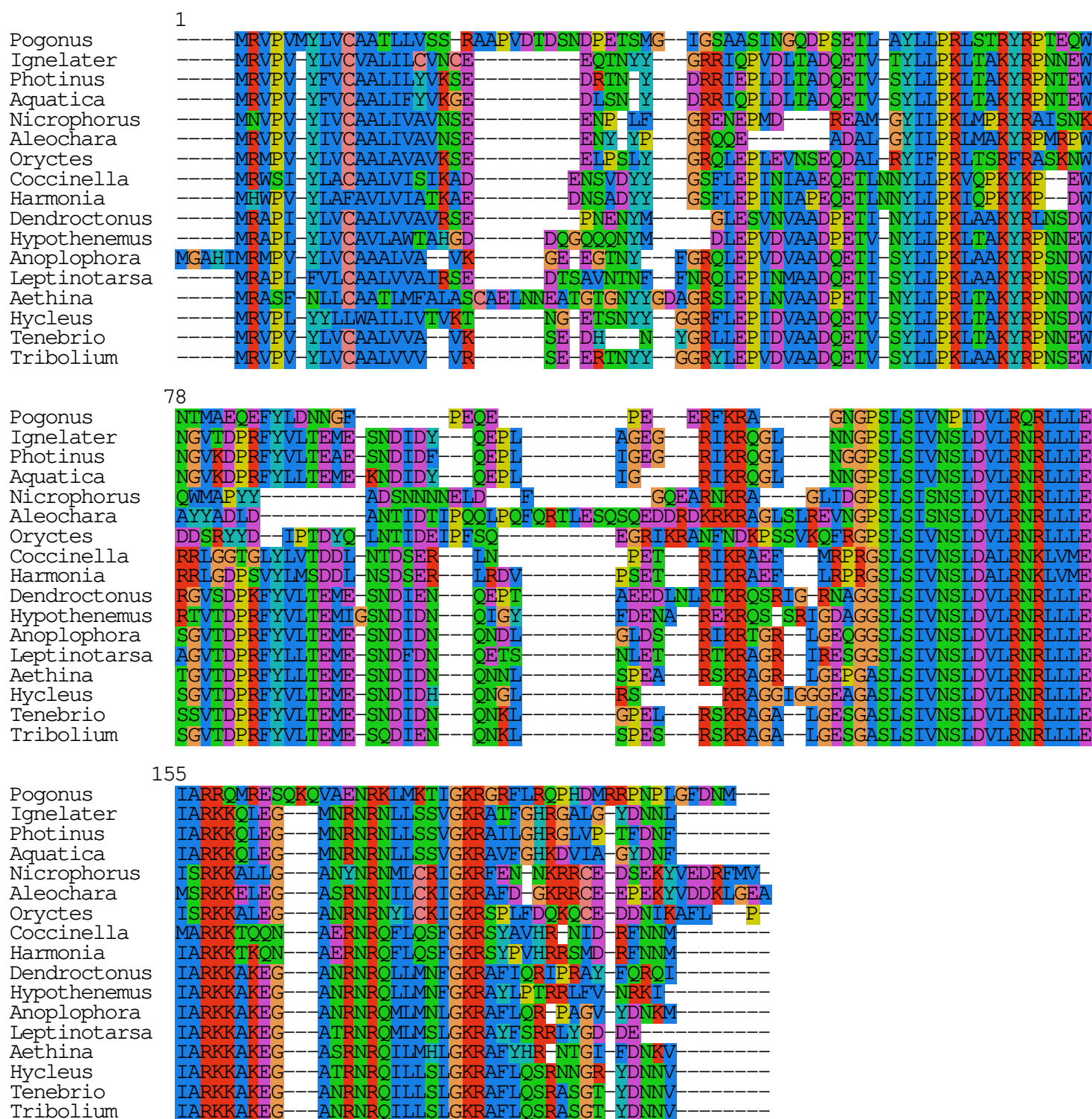
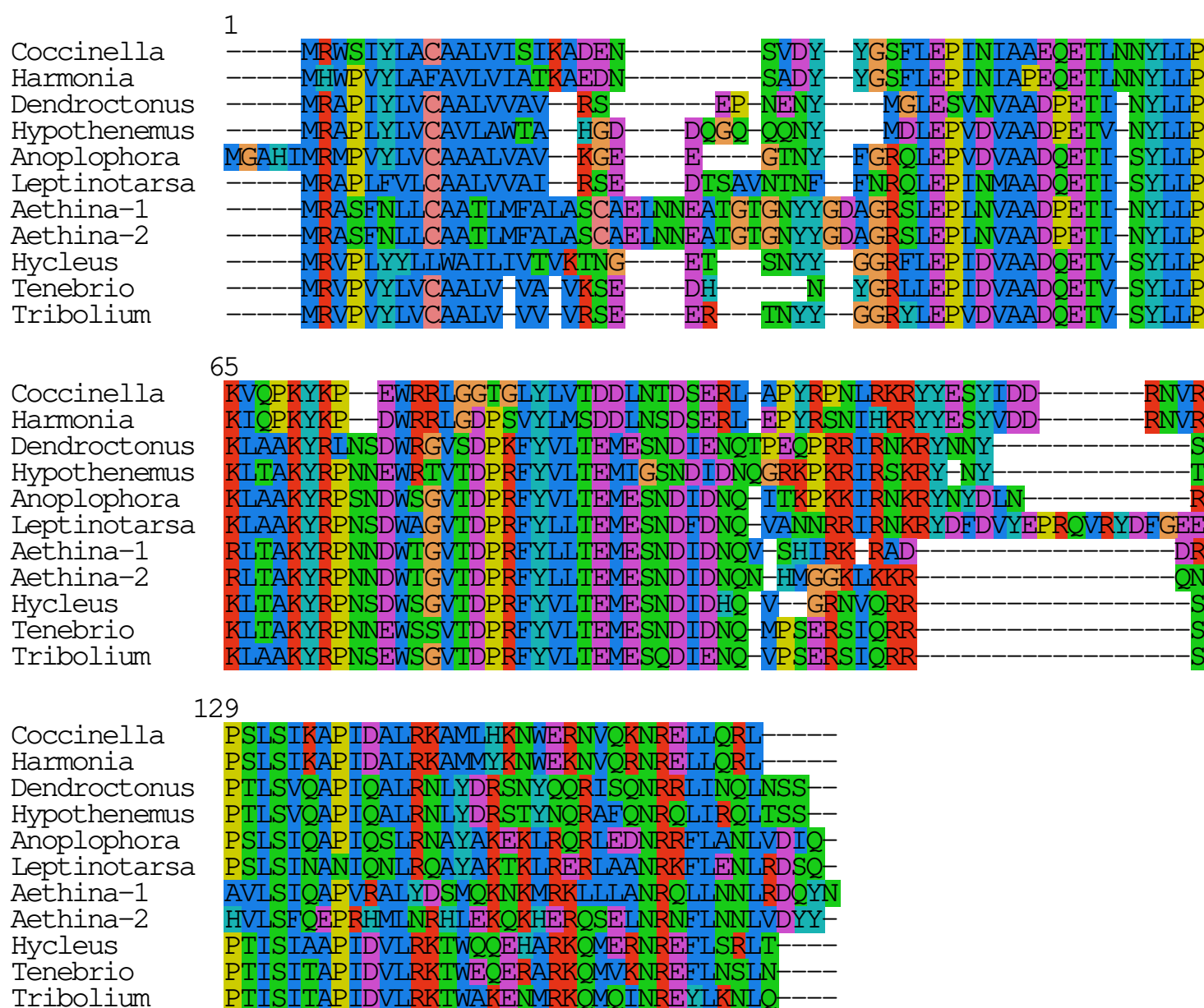


Figure S26. Alignment of DH47 precursors.

**Figure S27.** Alignment of DH37 precursors.

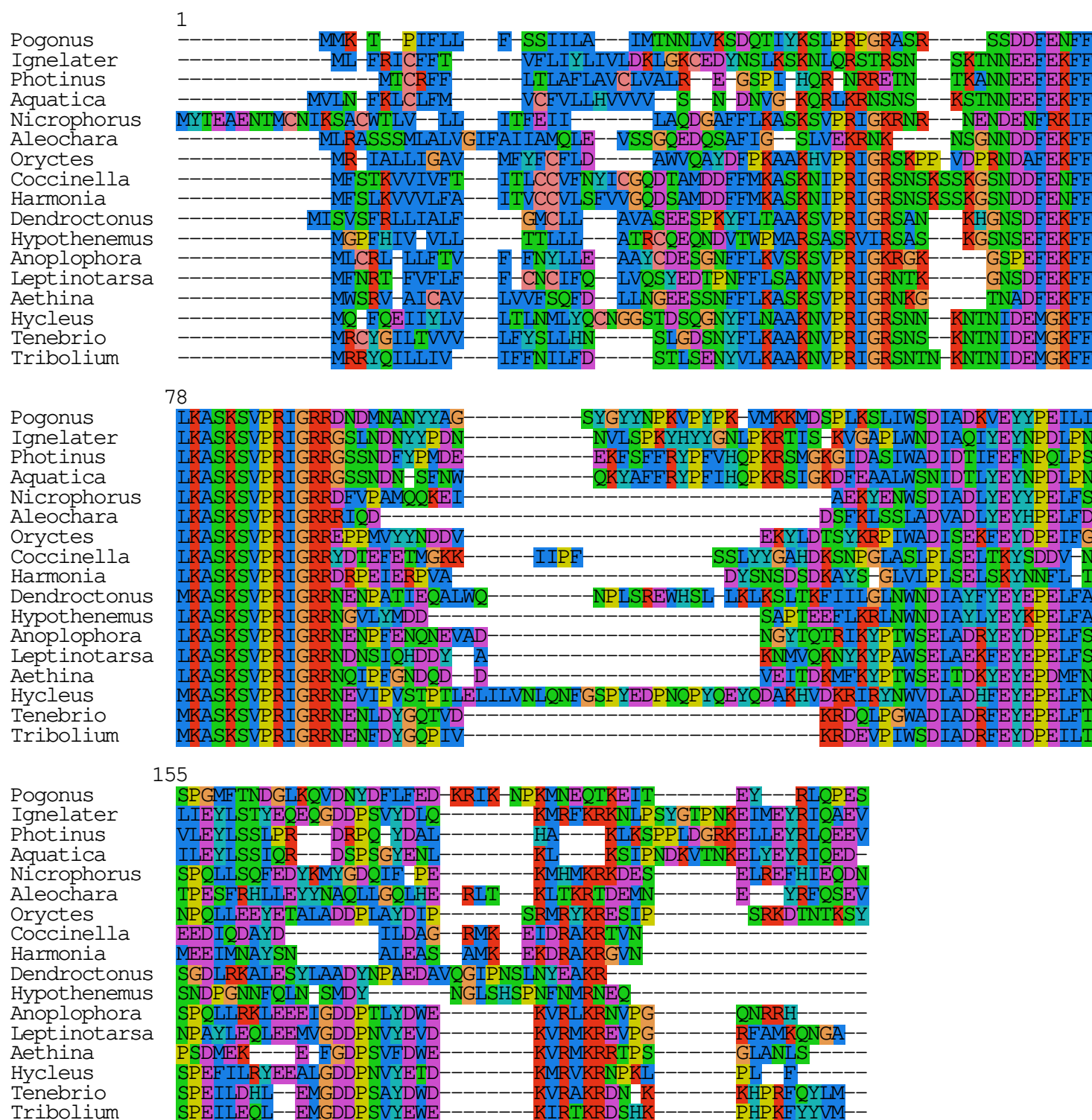


Figure S28. Alignment of ETH precursors.

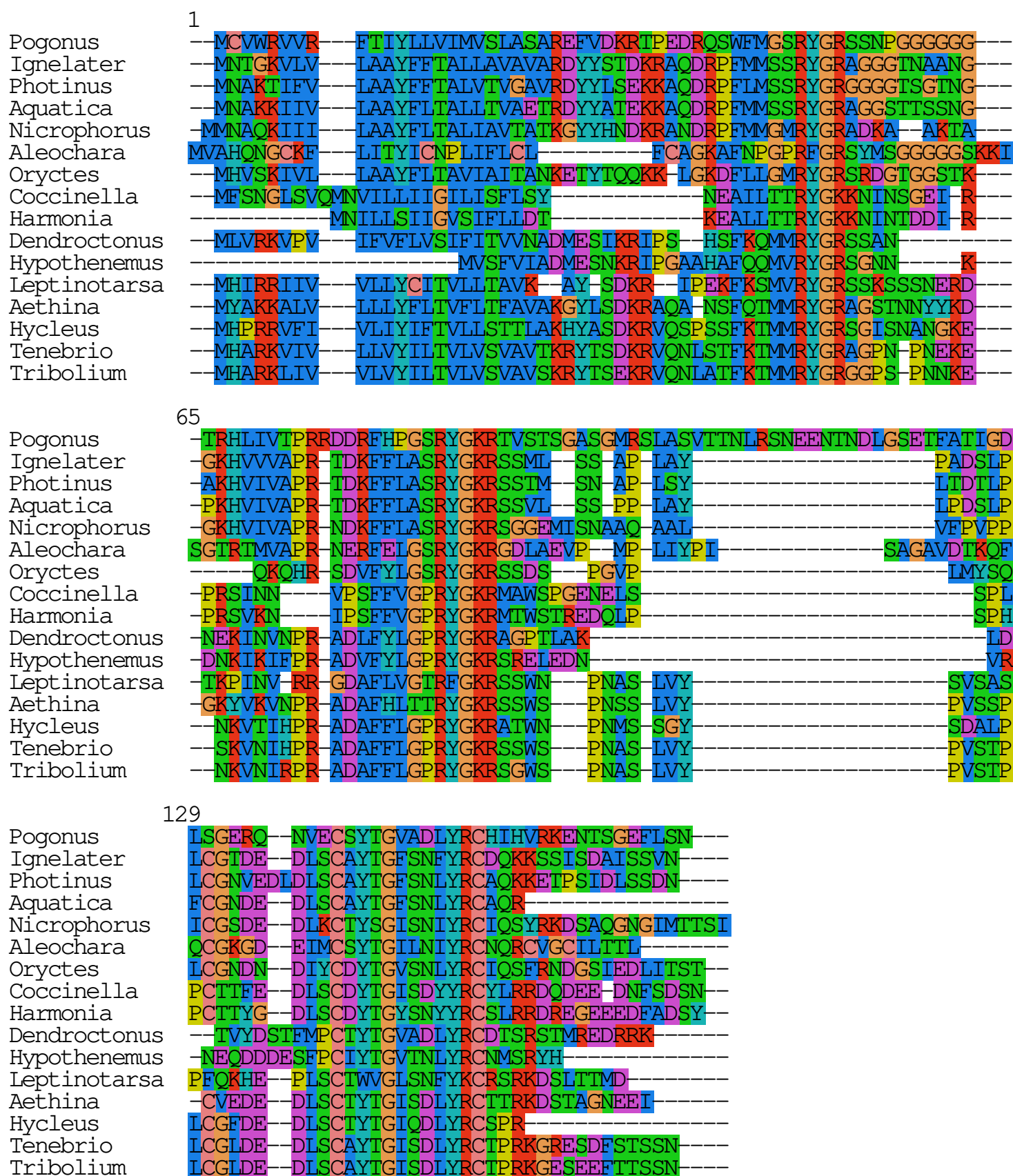
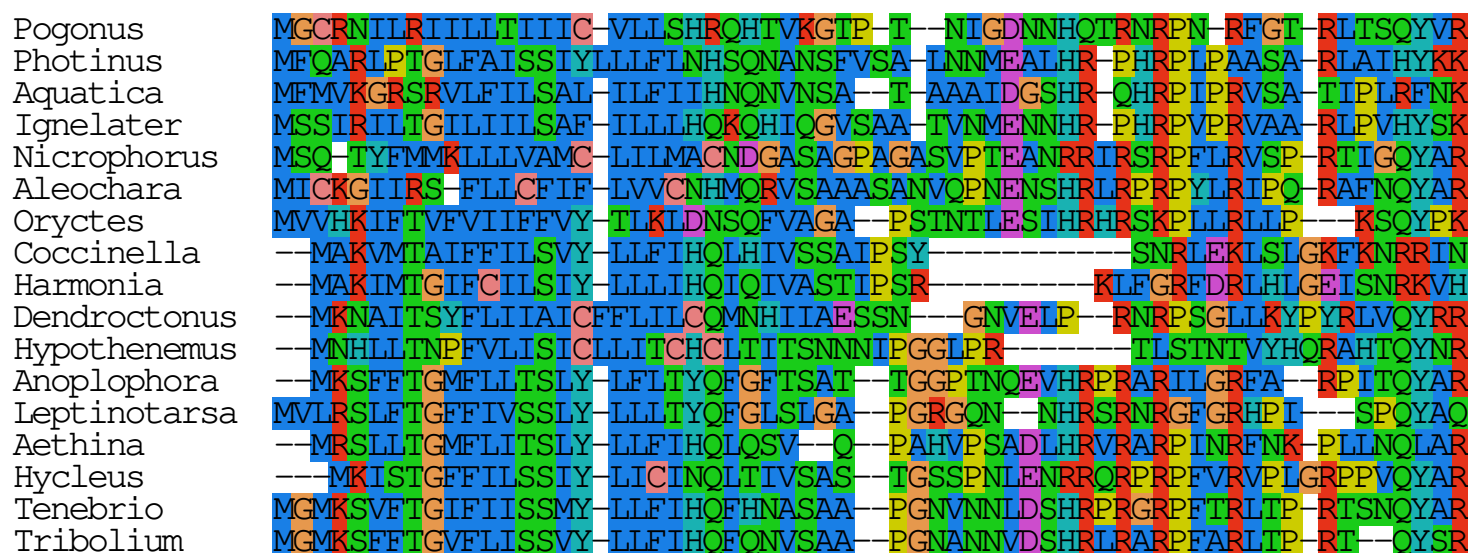
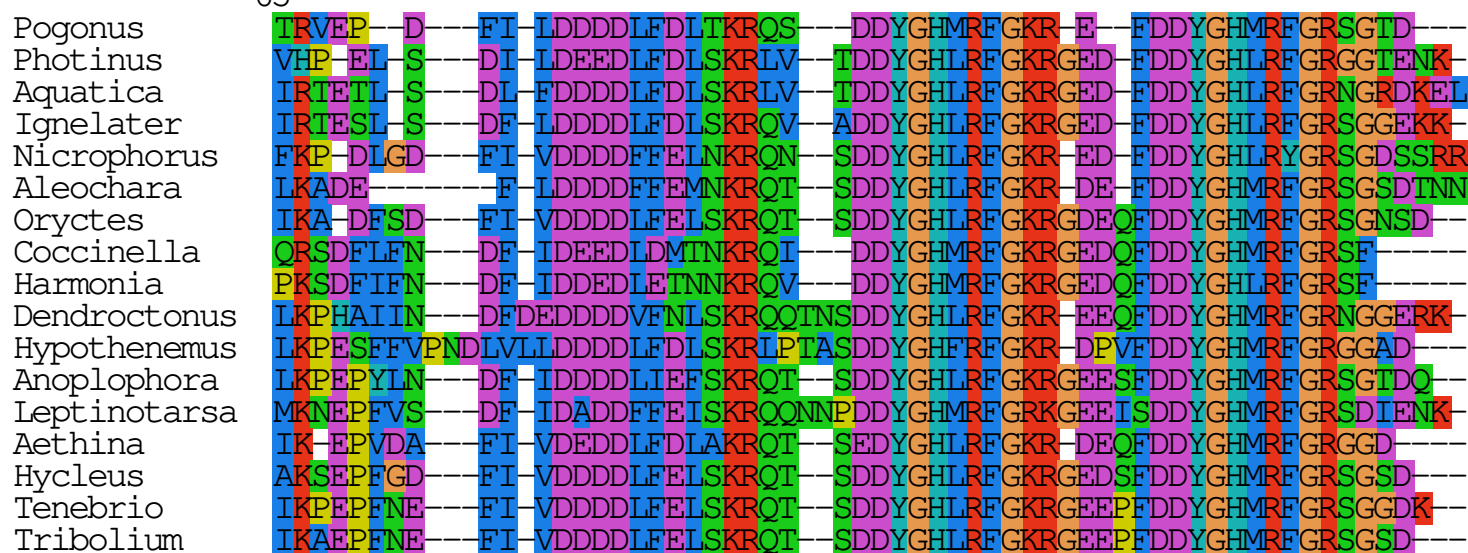


Figure S29. Alignment of RYamide precursors.



65

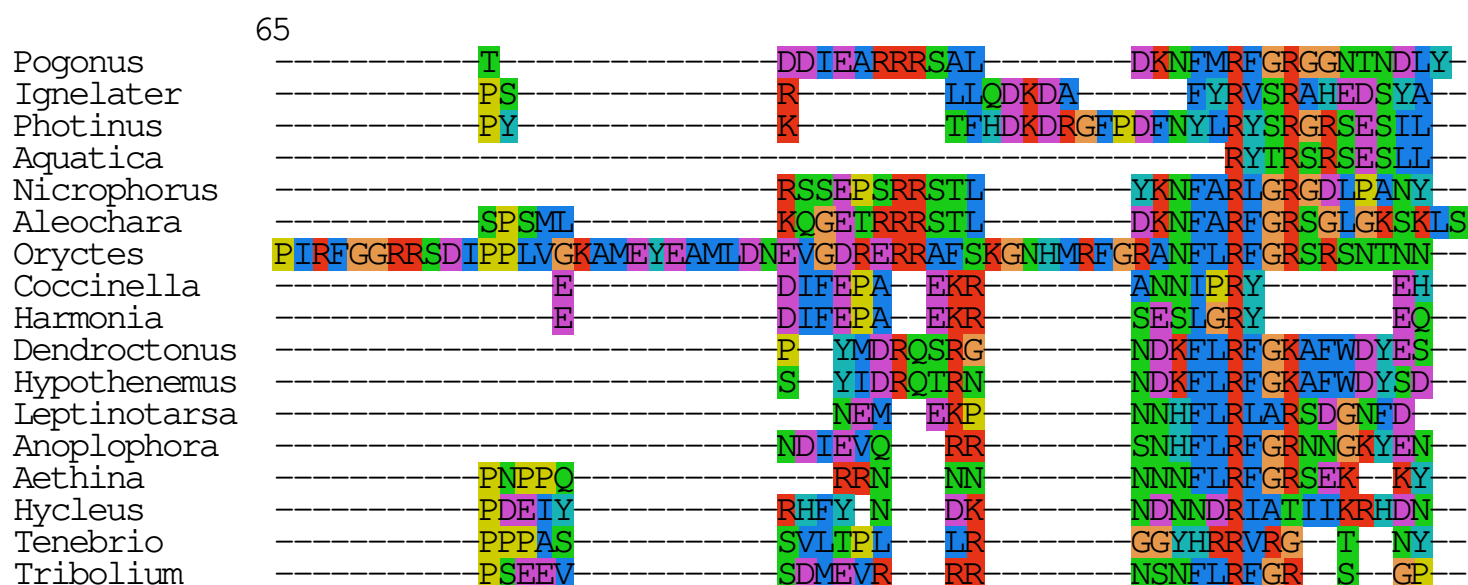


129

Pogonus
Photinus
Aquatica
Ignelater
Nicrophorus
Aleochara
Oryctes
Coccinella
Harmonia
Dendroctonus
Hypothenemus
Anoplophora
Leptinotarsa
Aethina
Hycleus
Tenebrio
Tribolium

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Figure S30. Alignment of Sulfakinin precursors.



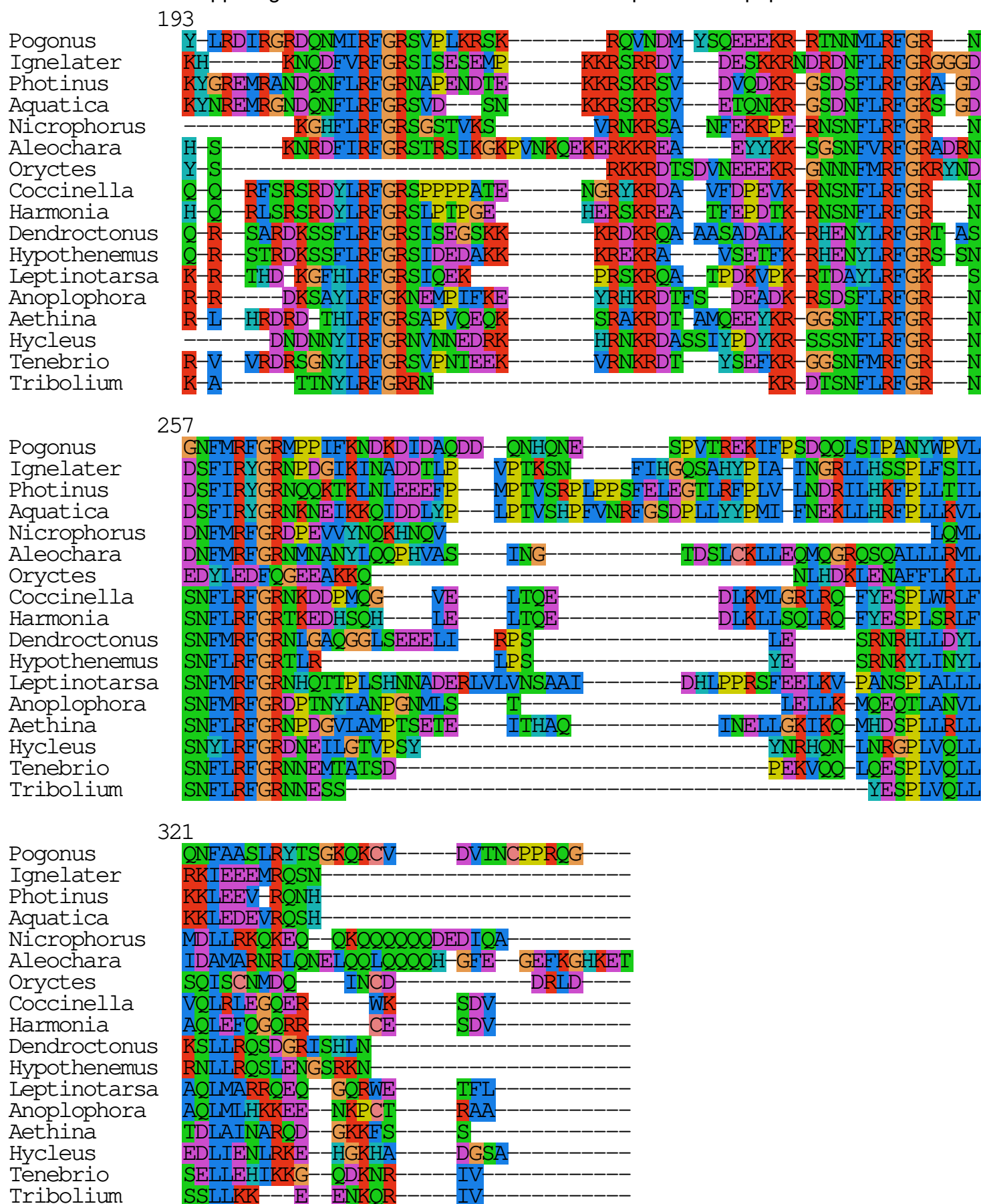


Figure S30. Alignment of FMRFamide precursors.

1

Pogonus	MACSFFNVLRFFWLVI	CIFA	VVVKPEDPD	SKISMCELSLAC	LLLLPP	DSMQVQALKRELF	RYLEEK
Ignelater	MAYFFSPKILISCGI	LWFT	LLIGIKSEERENSL	SACEIEKLLKP	ILSPDPERTI	QAQALRQRLIN	NLLRQG
Ignelater-s.v.	MAYFFSPKILISCGI	LWFT	LLIGIKSEERENSL	SACEIEKLLKP	ILSPDPERTI	QAQALRQRLIN	NLLRQG
Photinus	MRCPPKFLFCCGI	LWFT	LFGEIKSED	TPISSCEIERLLKP	ILSPDPERTI	QAQALRMLRM	NLLIRQG
Aquatica	MLCFFPSKILFCCGI	LWLL	FFDAIKSED	PESPLSTCEIERLLKP	ILSPDPDRTI	QAQALRNRLIN	NLLIRQG
Nicrophorus	MAK	VPI	VLLI	AVFIIRAHCC	EYMDLDEIFARIF	CPEKATTMQA	QALRHDIT
Aleochara	MG	TLIVGVVV	LA	LSCA LAPOLASA	QSDDEEINSMLEK	FAPESSSMOSRA	LRREVLRRL
Oryctes				INAQEE	CDTNLENMTSLVSP	DEPOTIQLQSL	RNELIRRIKEA
Coccinella	MVSGFRKPFVAV	VFLSVFIIMVQSD	E		CNMDFERTLEILFAP	QEQSPQIRSLRRYLI	SIFHNM
Harmonia	MVSREPLKLLI	VFLSVVYCMVQSD	E		CNLDERTLEILFAP	QEQSPQIRSLRRYFI	SILHSM
Dendroctonus	MGNLRLINLSV	WCVLVGCALIDNKV	SDNS		CDLDIPESLRILLDP	QETDSIQVQALRRQL	LLRKLQ
Dendroctonus-s.v.	MGNLRLINLSV	WCVLVGCALIDNKV	SDNS		CDLDIPESLRILLDP	QETDSIQVQALRRQL	LLRKLQ
Hypothenemus	MGSLGLIQLSM	WCILVGFALIDNKV	SDNS		YDFDTAEALRILLPE	ENDSIQVQALRRQL	LLKQLSNL
Anoplophora	MSFLSQ	TWVSLAA	LVLV	GSYGVNSDNT	CDIDVENTLRTLSP	QEN	SMQVQALRRDVL
Anoplophora-s.v.	MSFLSQ	TWVSLAA	LVLV	GSYGVNSDNT	CDIDVENTLRTLSP	QEN	SMQVQALRRDVL
Leptinotarsa	MAAF	EP	IFISVVL	FISA	MLYEVL	PKS	CDIDIEKTLGTLN
Aethina	MRKP	IVGF	WCI	LIVD	AVLLQVNS	EN	CDGEIENALRKLL
Aethina-s.v.	MRKP	IVGF	WCI	LIVD	AVLLQVNS	EN	CDGEIENALRKLL
Hycleus	MG	SFIVRIGV	LMVA	LIVTVKSD	D		YDFDIENALRTLTP
Hycleus-s.v.	MG	SFIVRIGV	LMVA	LIVTVKSD	D		YDFDIENALRTLTP
Tenebrio	MAVF	GAPKFLIGTGI	IMS	A	LFFMVKSDE	T	CDIEIENTLRTLTP
Tribolium	MLF	GAPKFFFGTGV	LVFA	LIFVVKSD	E		CDIEIENTLRTLTP
Tribolium-s.v.	MLF	GAPKFFFGTGV	LVFA	LIFVVKSD	E		CDIEIENTLRTLTP

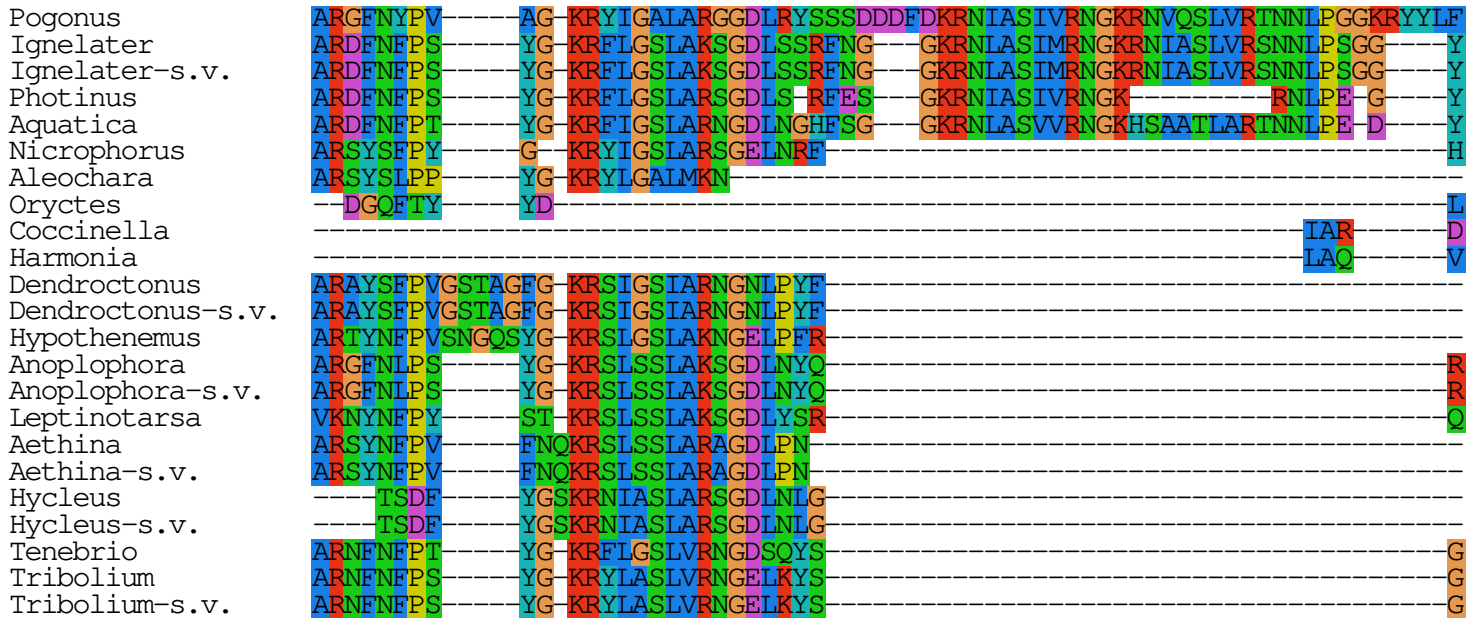
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Pogonus	LEEDEL		SKRSFASLVRDGNLP	YPKRSIEALARS	GYLKSATDDF		
Ignelater	FEEQVDNG		IEEKRSISSLARWNDLPM	KRNLOALARAGYLRSL	PDDD		EIDD
Ignelater-s.v.	FEEQVDNG		IEEKRSISSLARWNDLPM	KRNLOALARAGYLRSL	PDDD		EIDD
Photinus	LDEQVDNG		LEEK	GWDDMAT	KRSLSLAREGYLRPS	EE	SADD
Aquatica	LDEQIGNE		LDDKRSISALARWNDMP	KRNLOALAREGYLRSL	ED		NEDD
Nicrophorus	LAREGINEDDELVR	RVDEM	NKKESYDDDYRMAAF	GKRNLAALARAGYIRTL	IPDED		N
Aleochara	IEREQDDDI		ANYKRSINSLARLGTIPG	KRNLAALAKGGYIRTL	IPDSSTEQ		EQ
Oryctes	KOREELENEI		LQYKRSLSLARWNNMPG	KRNLESALARAGYIRTL	PPTN		E
Coccinella	VARAEE		IEGWNTYVY	PKRSLEALARAGYLHTL	PDDE		DNQ
Harmonia	VARAEE		AEGWNTYVY	PKRSLEALARAGYLHTL	PDEE		DNQ
Dendroctonus	DCHNL	ELAELEDKNY	KSLASLAGWGNIPAEH	KRNLEALARAGITWYKNP	PEI		DQS
Dendroctonus-s.v.	DCHNL	ELAELEDKNY	KSLASLAGWGNIPAEH	KRNLEALARAGITWYKNP	PEI		DQS
Hypothenemus	LQPNV	EEFAEQPDDGNYRRTIAS	LDLWDLPEEH	KRNIEALARSGMACRMF	PEL		EP
Anoplophora	LDRLDQEEDE		RNLKRSLSLAAWNLP	E	HKRNLEALARAGYLKTL	PDDE	D
Anoplophora-s.v.	LDRLDQEEDE		RNLKRSLSLAAWNLP	E	HKRNLEALARAGYLKTL	PDDE	D
Leptinotarsa	AEKQEE		DENKRALVALAKYNDLPE	HKRNLEALARAGYIKTL	PDEE		D
Aethina	LEKFEE	EED	PNLKSSLSALARWNNLP	E	KRNLESALARAGYIKTL	PDVE	D
Aethina-s.v.	LEKFEE	EED	PNLKSSLSALARWNNLP	E	KRNLESALARAGYIKTL	PDVE	D
Hycleus	FDNGNIDV		DIGNDVGVIDRW	N	KRNLEALARAGLVRTL	PITQDNNDDNNEHDNDE	
Hycleus-s.v.	FDNGNIDV		DIGNDVGVIDRW	N	KRNLEALARAGLVRTL	PITQDNNDDNNEHDNDE	
Tenebrio	LEKANL	DDD	MNYKRSLSLAQWNL	PD	KRNLEALARAGYVRTL	PSQD	D
Tribolium	LDRADD	EDE	MNYKRSISSLAQWNL	PG	KRNLEALARAGYIRTL	PNPD	E
Tribolium-s.v.	LDRADD	EDE	MNYKRSISSLAQWNL	PG	KRNLEALARAGYIRTL	PNPD	E

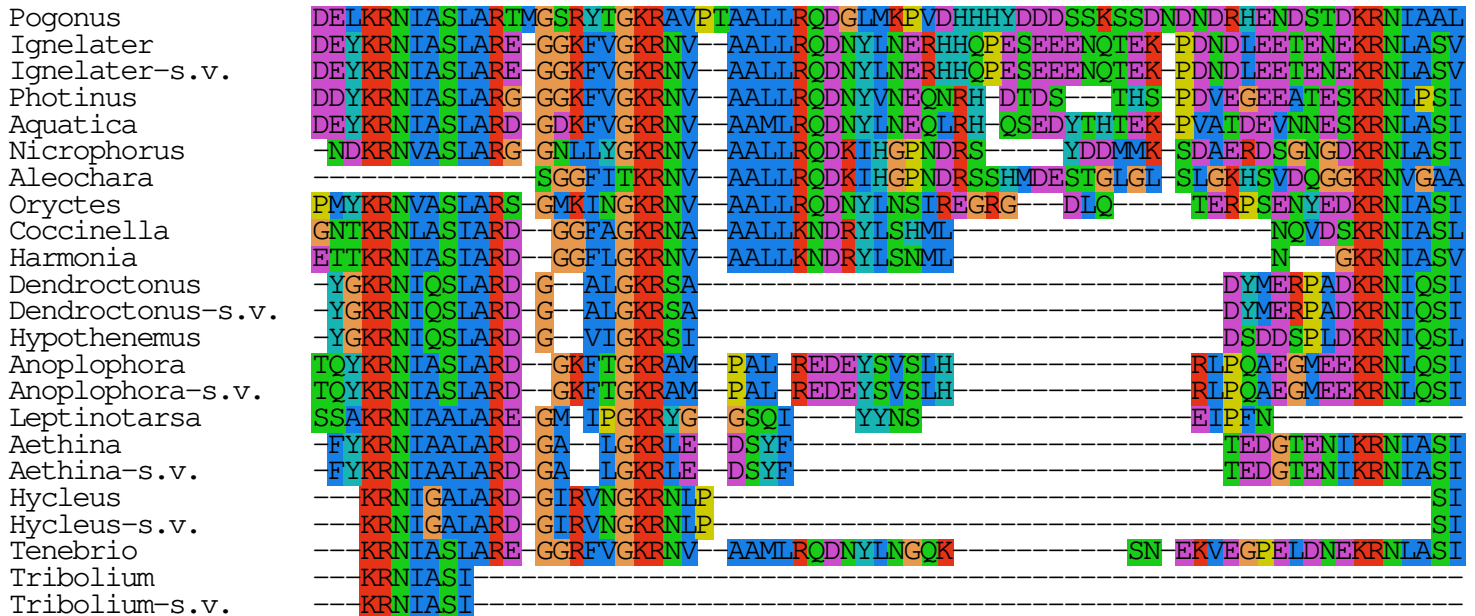
145

Pogonus	TDDDVKRSIANL	AKNGQLP	TFORPEDE	QOTQKRGIESL	ARNGLHGX	RK	LENLFENLYGKRNIASL
Ignelater	NDPNLKRSIQYL	VKNGQLP	YRS	EESKRGIQAL	ARNGELHNT	RD	LEELIDELYEKRNI
Ignelater-s.v.	NDPNLKRSIQYL	VKNGQLP	YRS	EESKRGIQAL	ARNGELHNT	RD	LEELIDELYEKRNI
Photinus	GD	LKRSIQSL	ARNQFPV	C	S	DD	LKRSISSLARNGELINN
Aquatica	TDLSLKRSIQSL	ARLNQFP	YQT	EPPKRSISSL	ARNGEFDG	RN	LEIDELYEKRNI
Nicrophorus		GKRSIANL	AKNGQLPN	YQN	DAEKRGIESL	ARNGELHNT	RE
Aleochara	EEEQDKRSIESL	VRAGQMPN	VAYD	AE	SKRGIESL	ARNGELHNT	RNLGSL
Oryctes		PDYKRNEMITRS	VRNTQSPV				
Coccinella	TDSNDKRSLSAL	AKNGQFPLHQR	R	DEESYKRS	SGPPNSGDISKEI	QKMLDDLYNN	
Harmonia	TDSNDKRSLSAL	AKNGQLPVHRL	R	DEESFKRS	SGTPVSTNDINKEM	QKMLDDLYNN	
Dendroctonus	EDANYKRSMEVL	LKNGPAPLMPES		LENQKRGIESL	ARNGDFLRR	QP	FQSFLNSQDYKRNIASL
Dendroctonus-s.v.	EDANYKRSMEVL	LKNGPAPLMPES		LENQKRGIESL	ARNGDFLRR	QP	FQSFLNSQDYKRNIASL
Hypothenemus	QDSTYKRSIEIL	AKNGQHP	PLVROP	LENQKRGIESL	ARNGLHRP	PN	YOIQOQGNMDYKRNL
Anoplophora	GDSNYKRSIANL	AKNGQLP	M	S	DGQKRGIESL	ARNGLRPPK	KD
Anoplophora-s.v.	GDSNYKRSIANL	AKNGQLP	M	S	DGQKRGIESL	ARNGLRPPK	KD
Leptinotarsa	ADANYKRSIANL	AKNGQLP	TR	K	EDQKRGIESL	ARNGLHQN	RE
Aethina	ANDSFKRSIANL	AKNGQLPHH			DDKRGIQSL	ARNGEIHTK	PTS
Aethina-s.v.	ANDSFKRSIANL	AKNGQLPHH			DDKRGIQSL	ARNGEIHTK	PTS
Hycleus	DDGNFKRSLSL	VRNGQLPH			QNKRGIQSL	ARNGELIQH	QRNNFDD
Hycleus-s.v.	DDGNFKRSLSL	VRNGQLPH			QNKRGIQSL	ARNGELIQH	QRNNFDD
Tenebrio	EDPNYKRSLATL	AKNGQLP	T	YQN	ND	SKRGIESL	ARNGELTTR
Tribolium	EDPNNKRSLSL	AKNDQLP	TTFQ	N	ESKRGIESL	ARNGELHNR	RD
Tribolium-s.v.	EDPNNKRSLSL	AKNDQLP	TTFQ	N	ESKRGIESL	ARNGELHNR	RD

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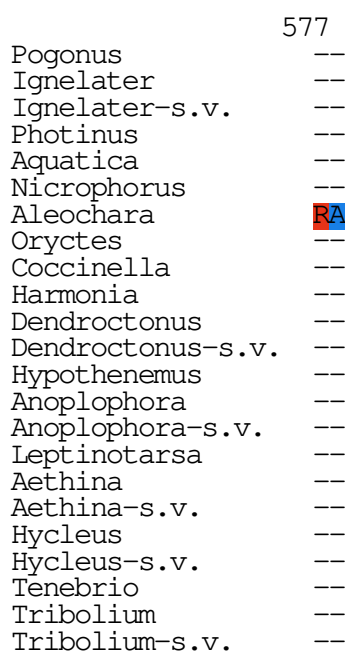
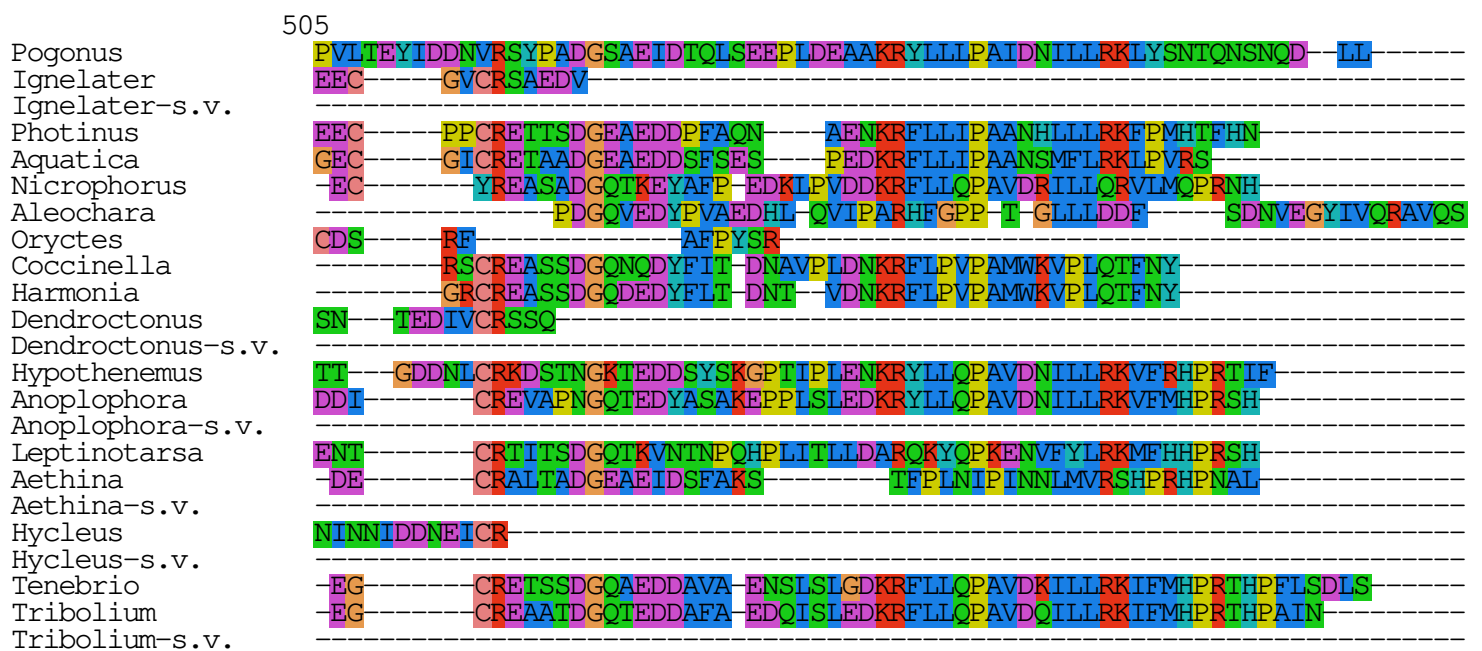
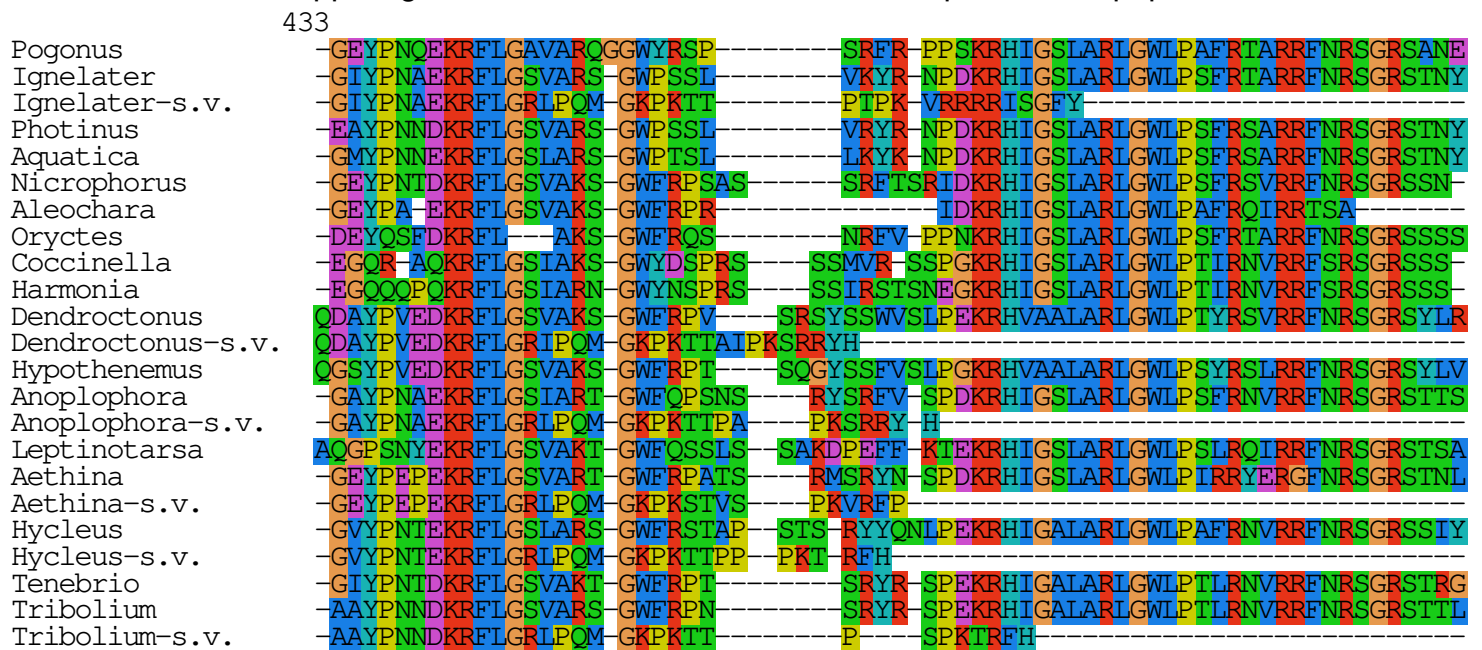
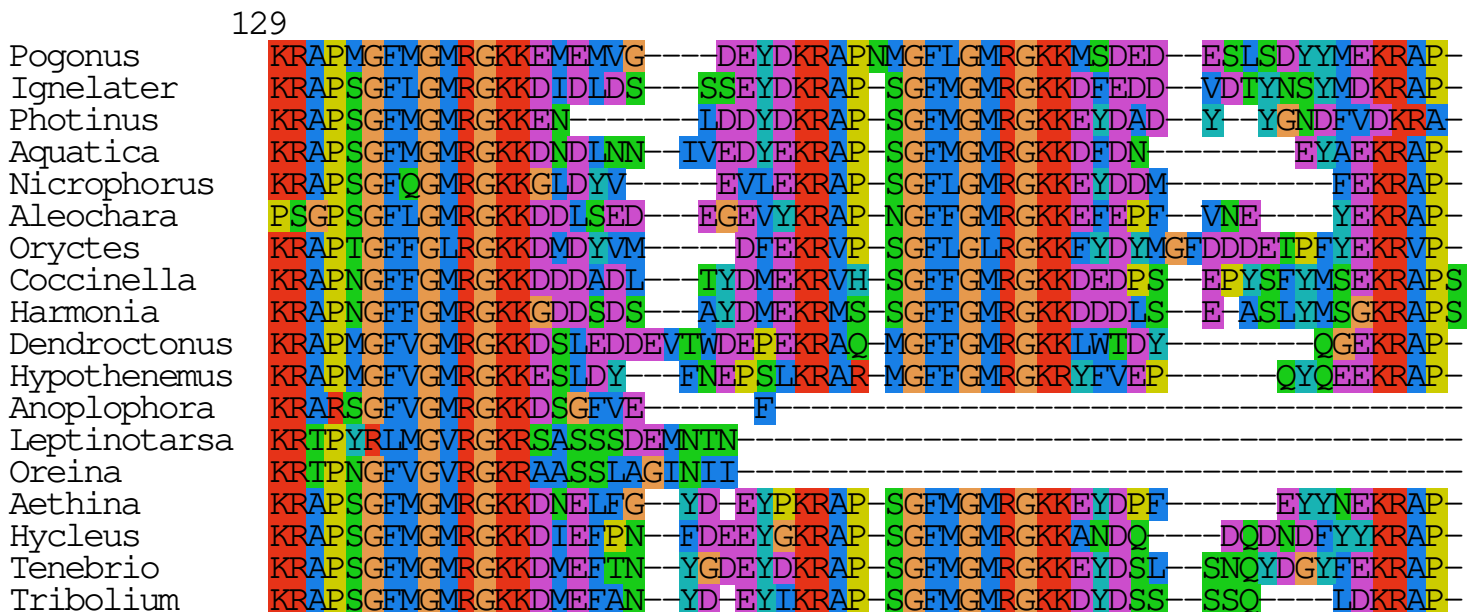
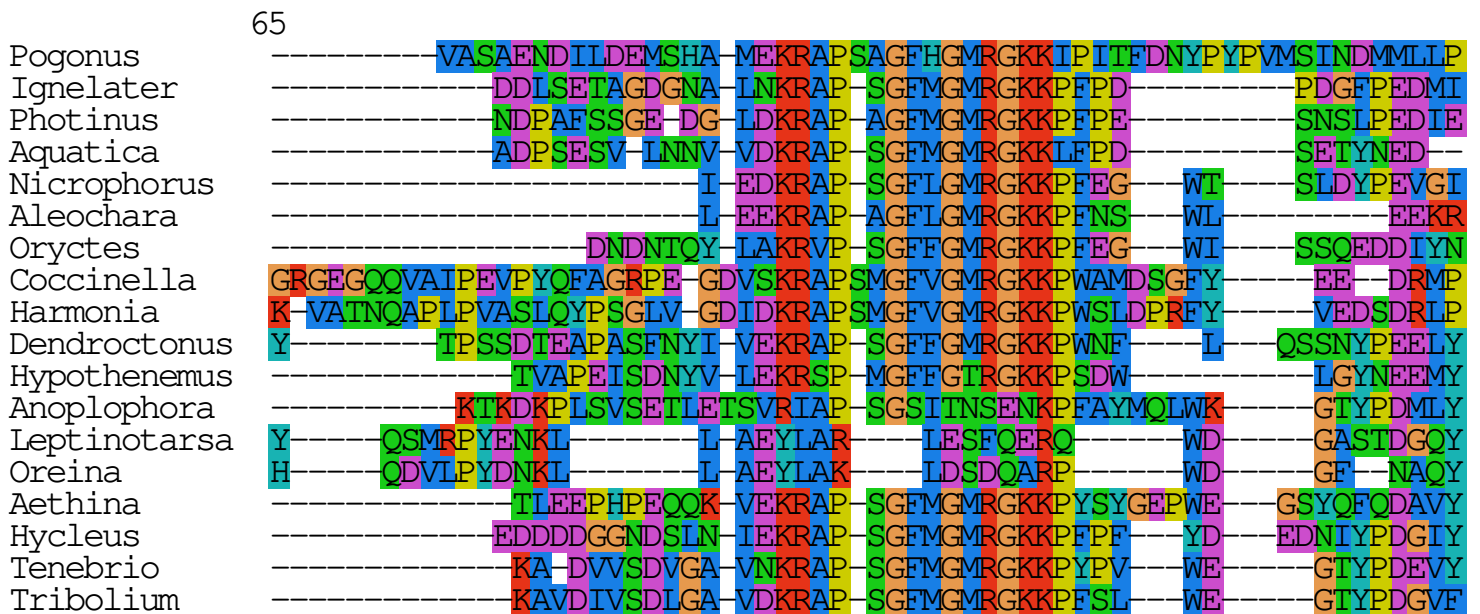


Figure S32. Alignment of NPLP1 precursors; s.v., splice variant.



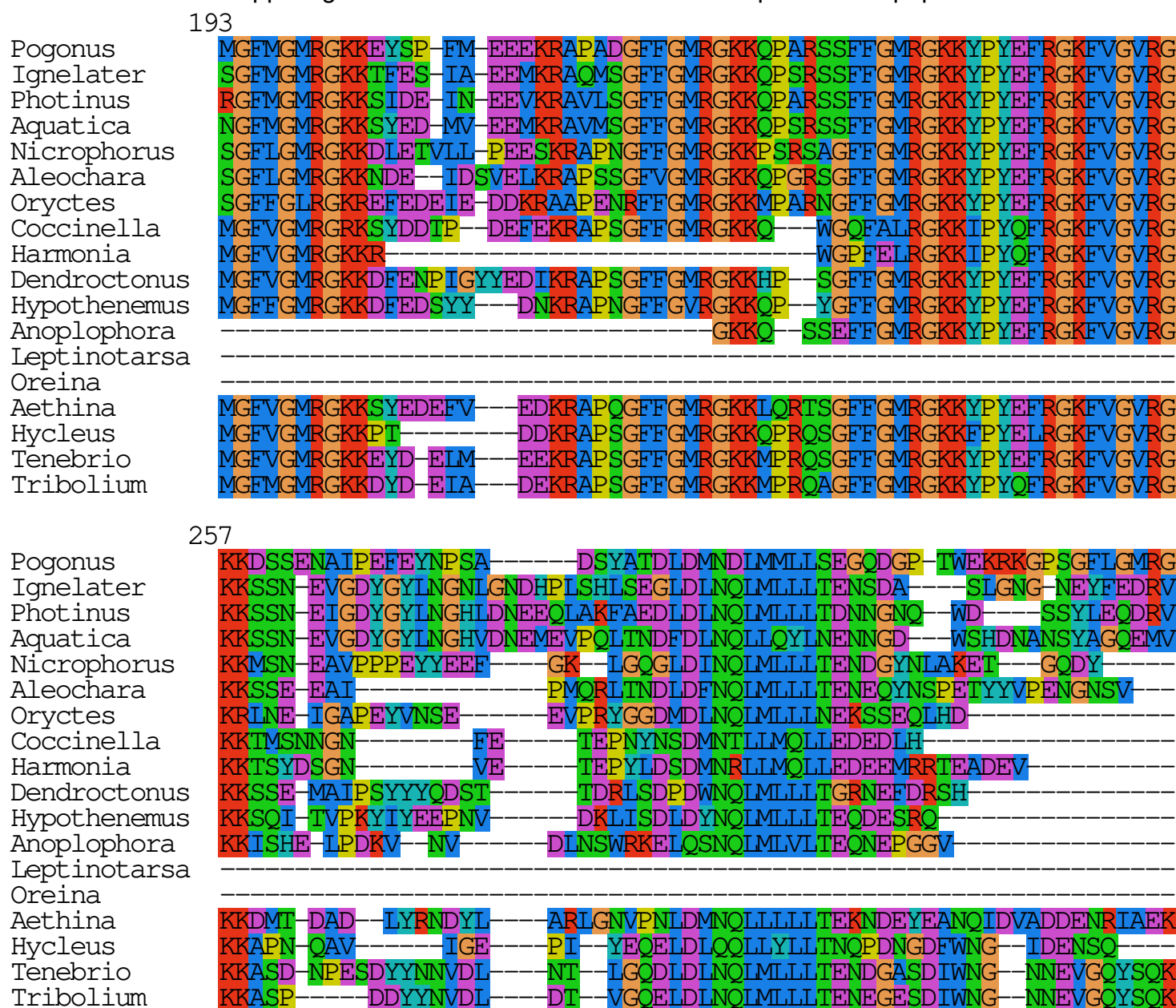
**Figure S34.** Alignment of Tachykinin precursors.



Figure S35. Alignment of CCAP precursors.

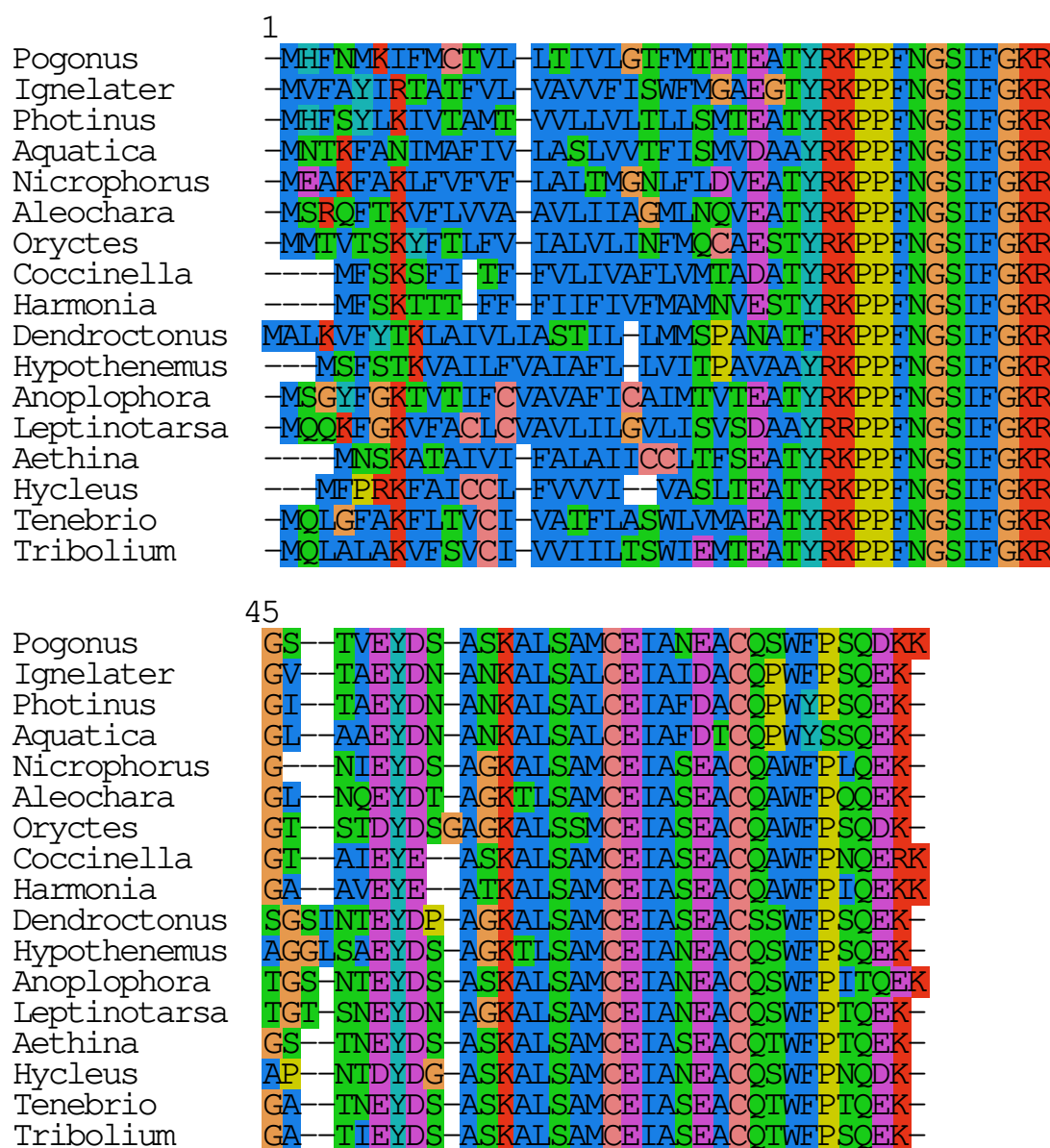
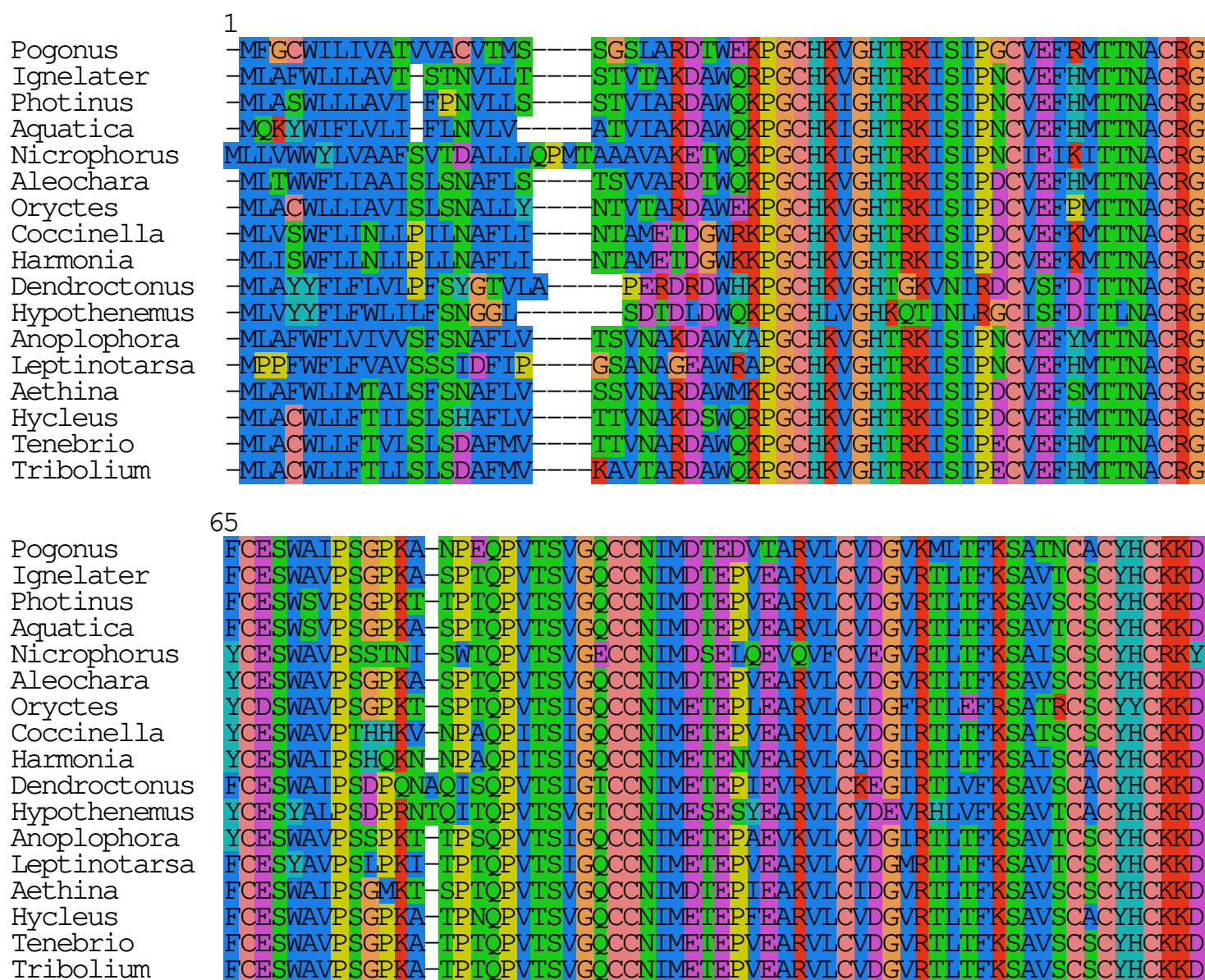


Figure S36. Alignment of SIFamide precursors.

**Figure S37.** Alignment of GPA2 precursors.

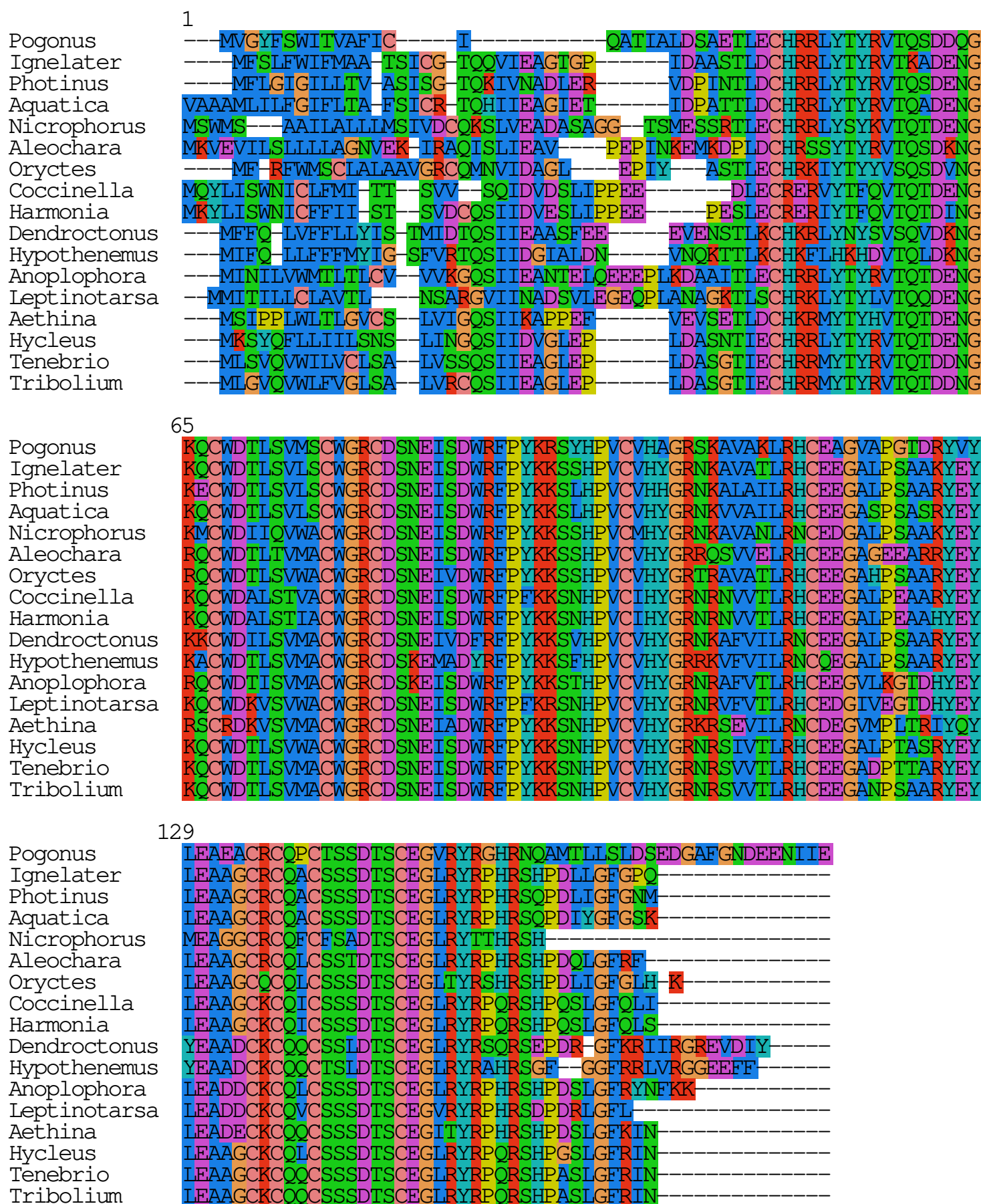


Figure S38. Alignment of GPB5 precursors.

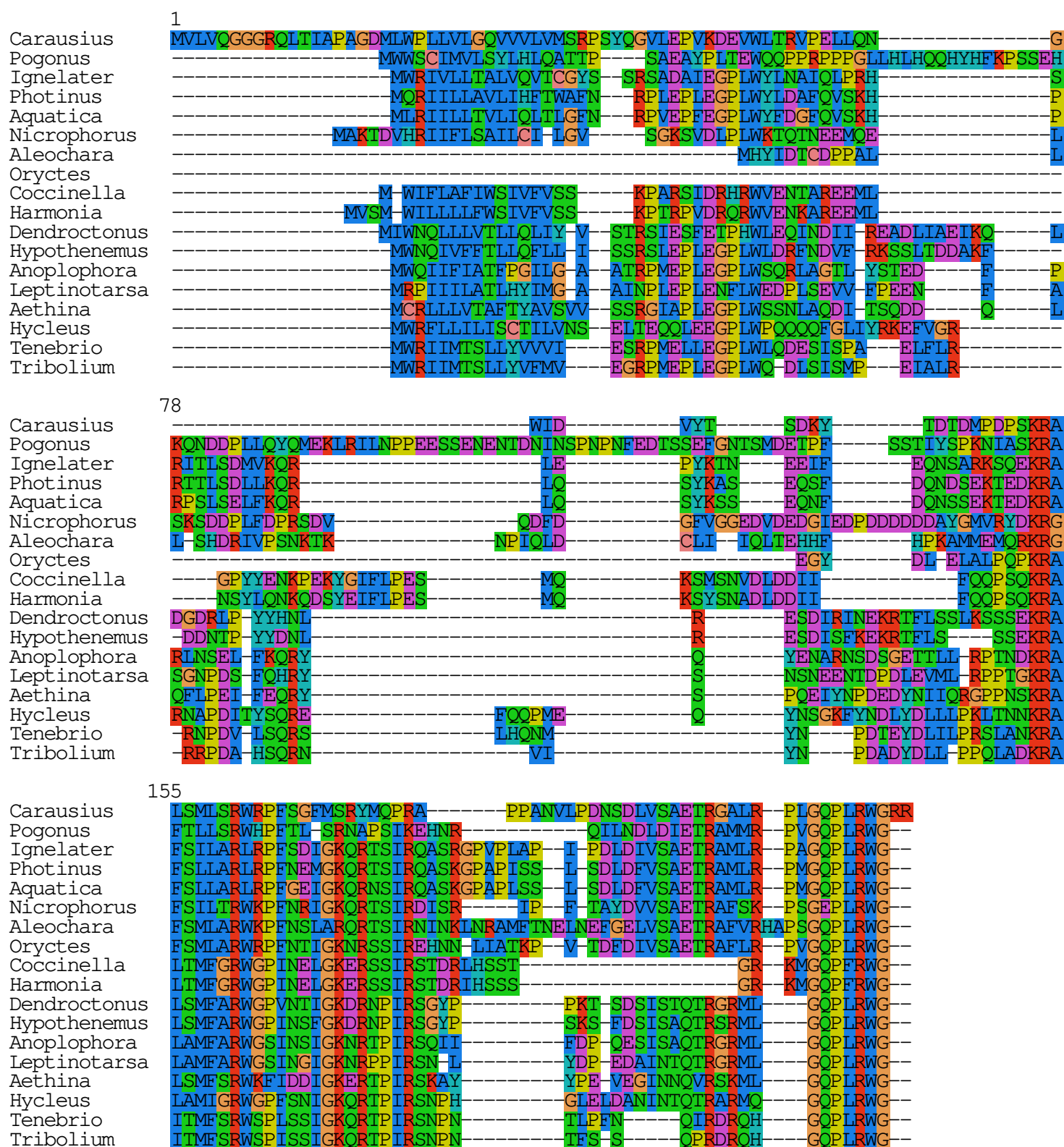
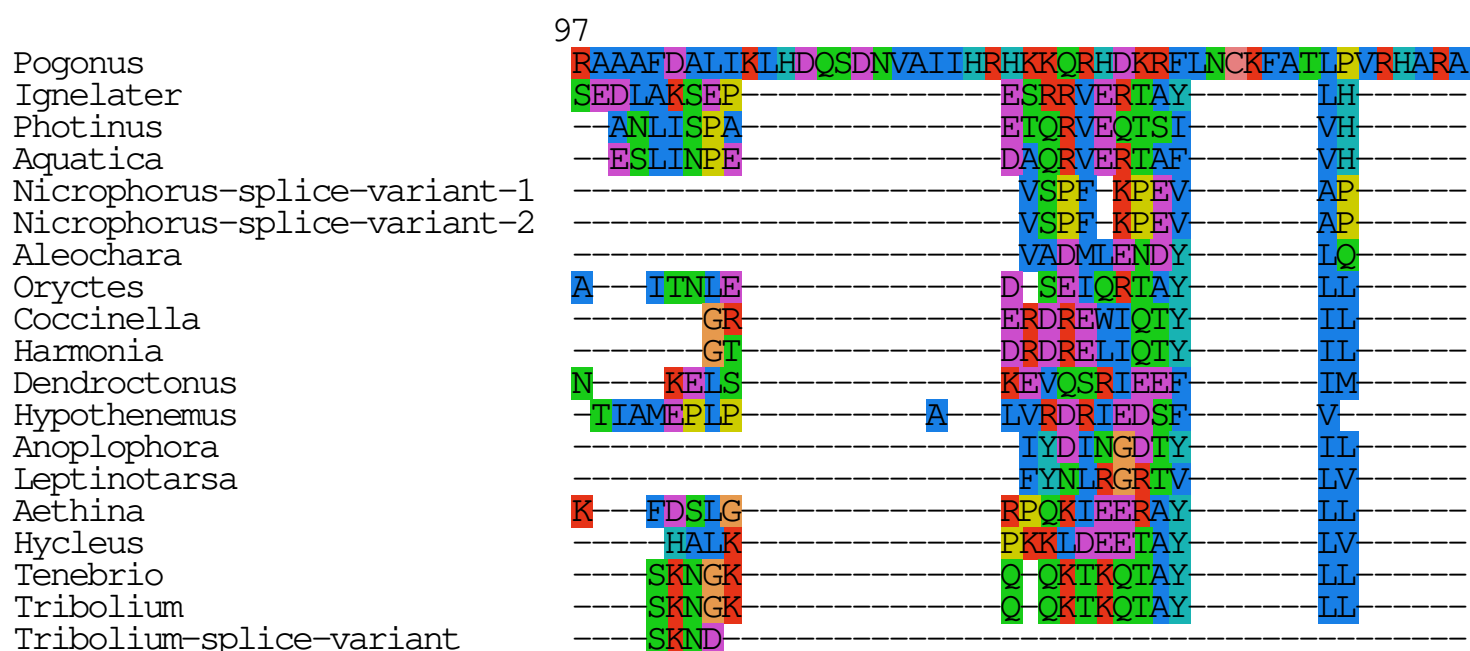
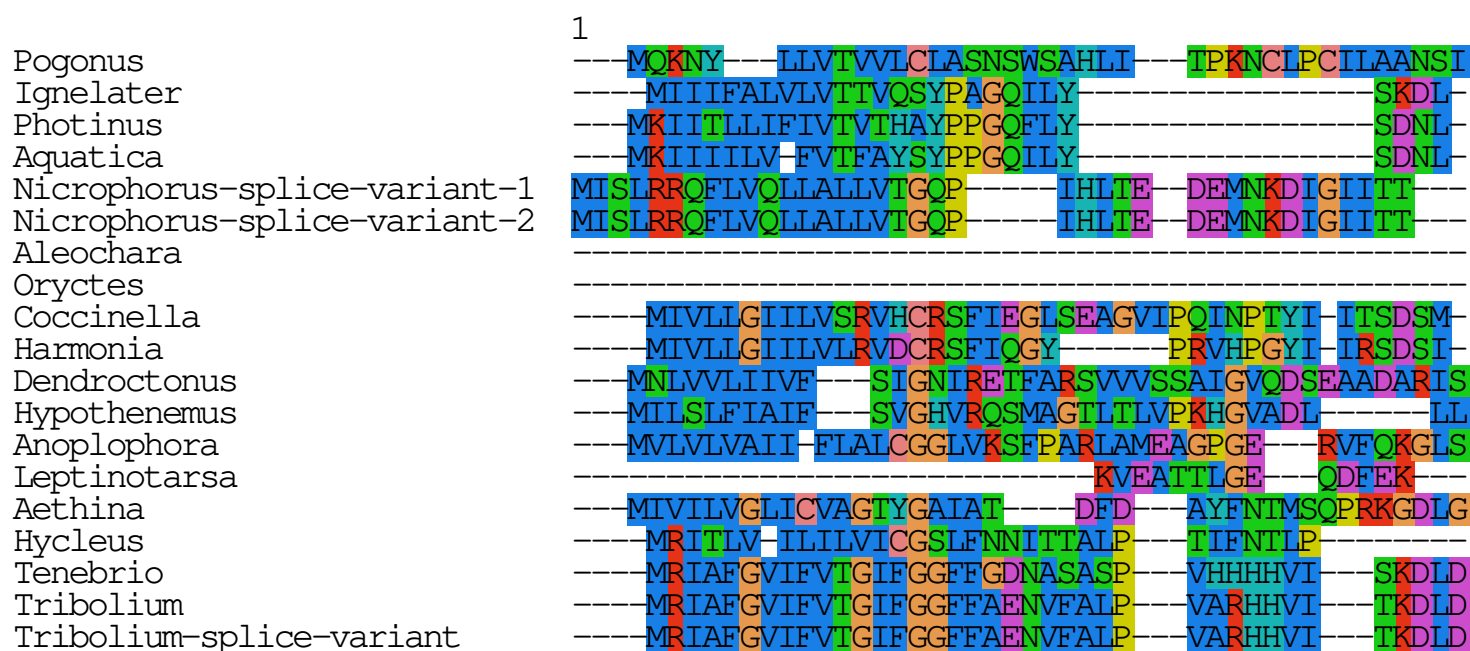
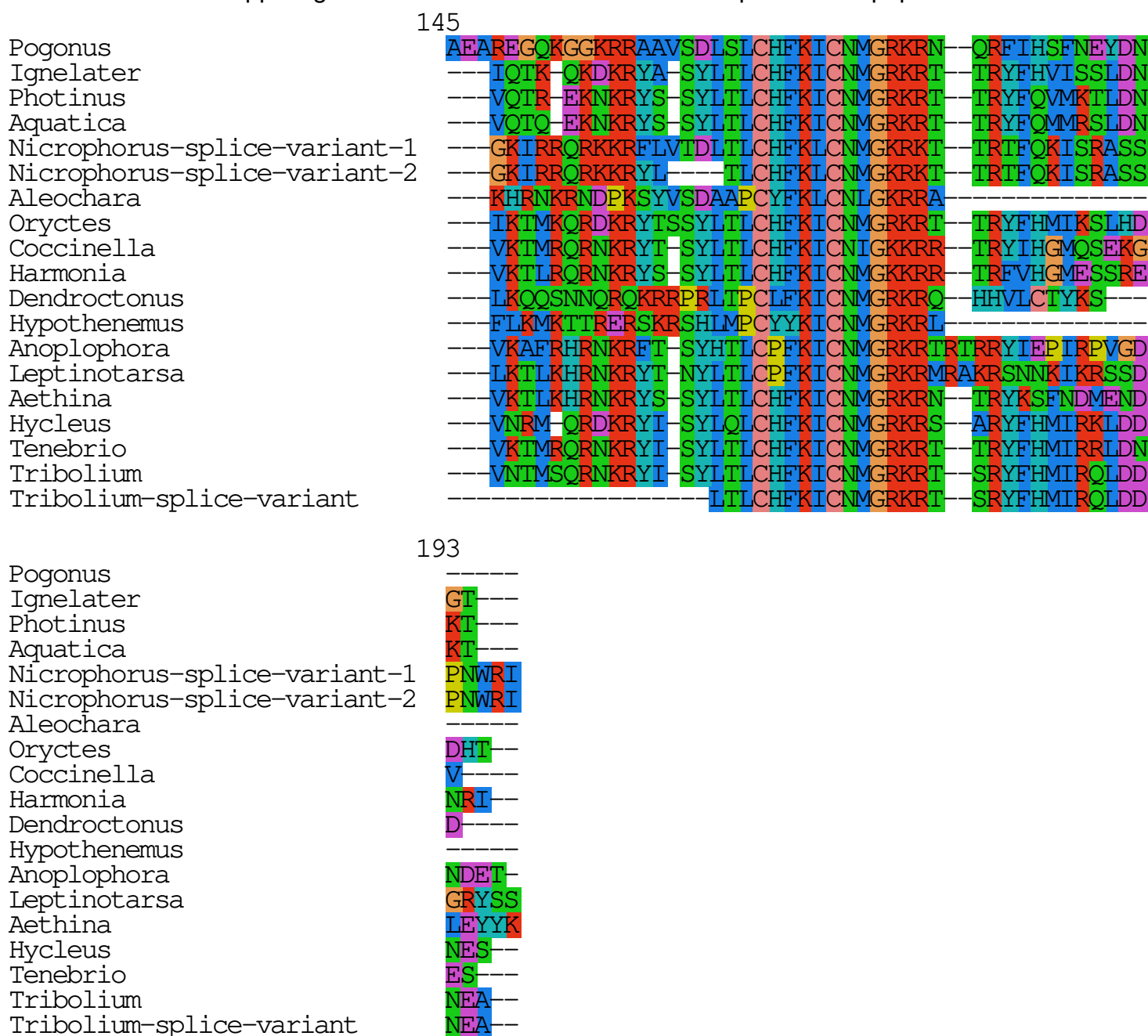


Figure S39. Alignment of Hansolin precursors.



**Figure S40.** Alignment of CNMamide precursors.

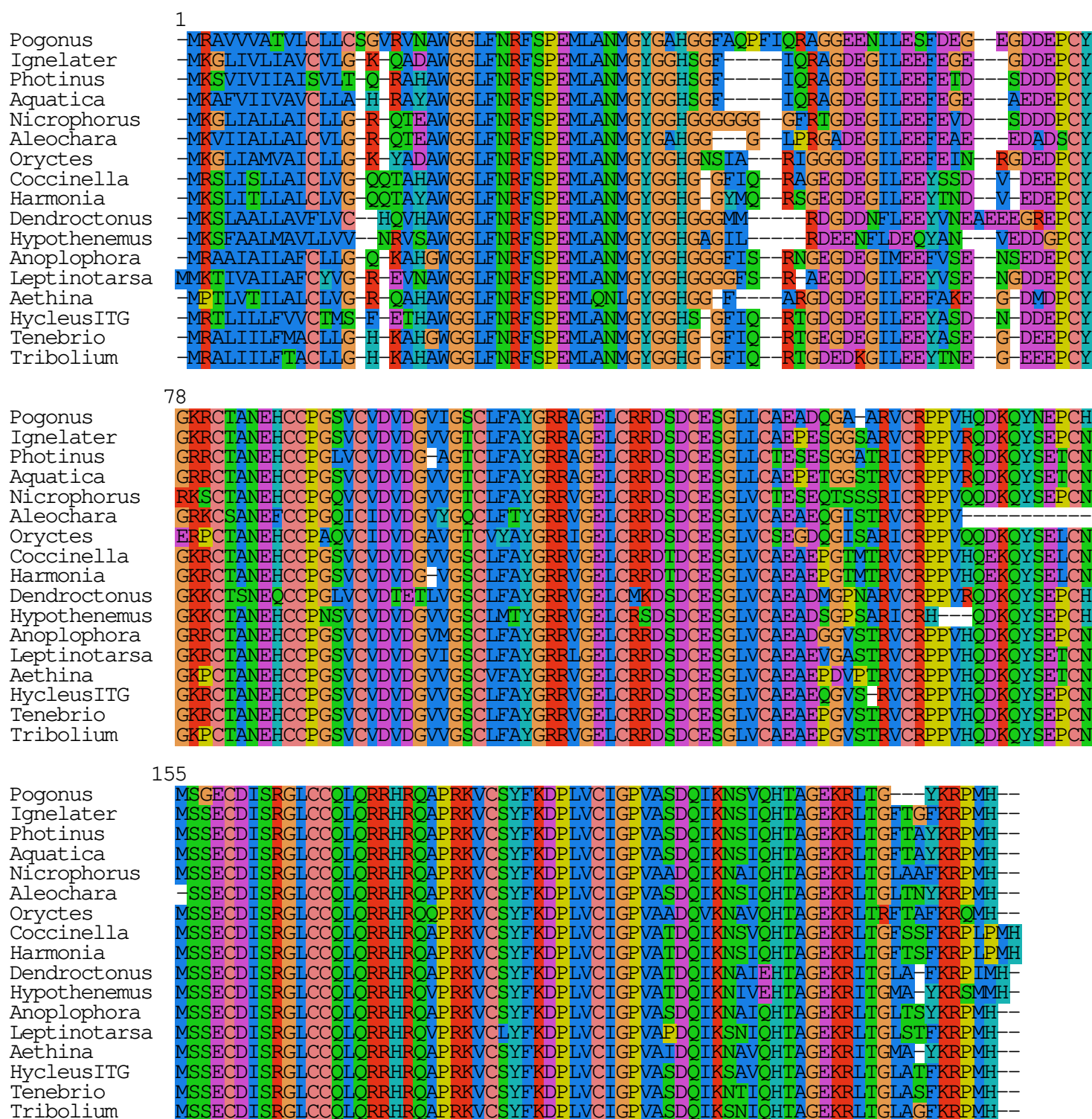


Figure S41. Alignment of ITG-like precursors.

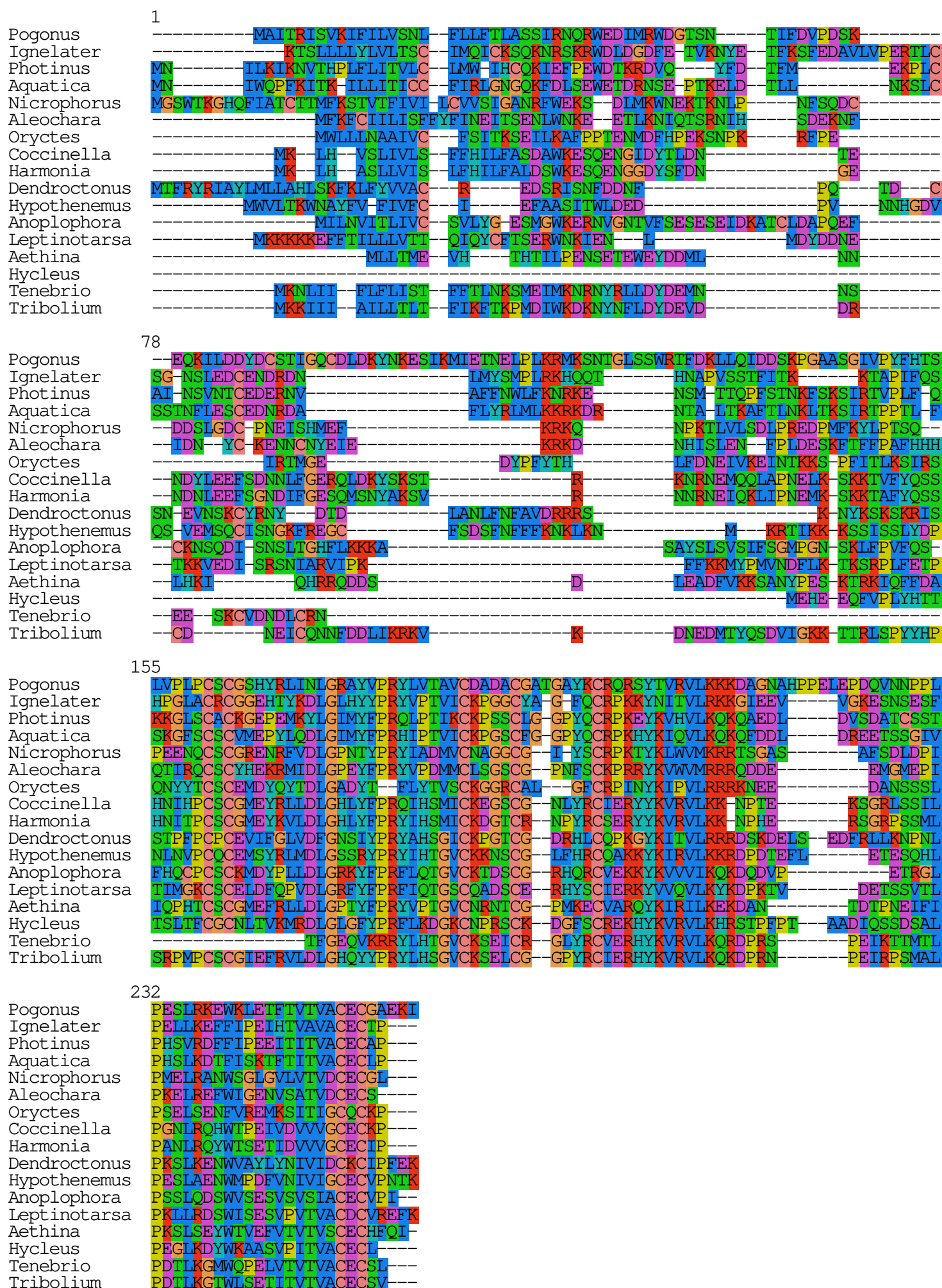


Figure S42. Alignment of PTH precursors.

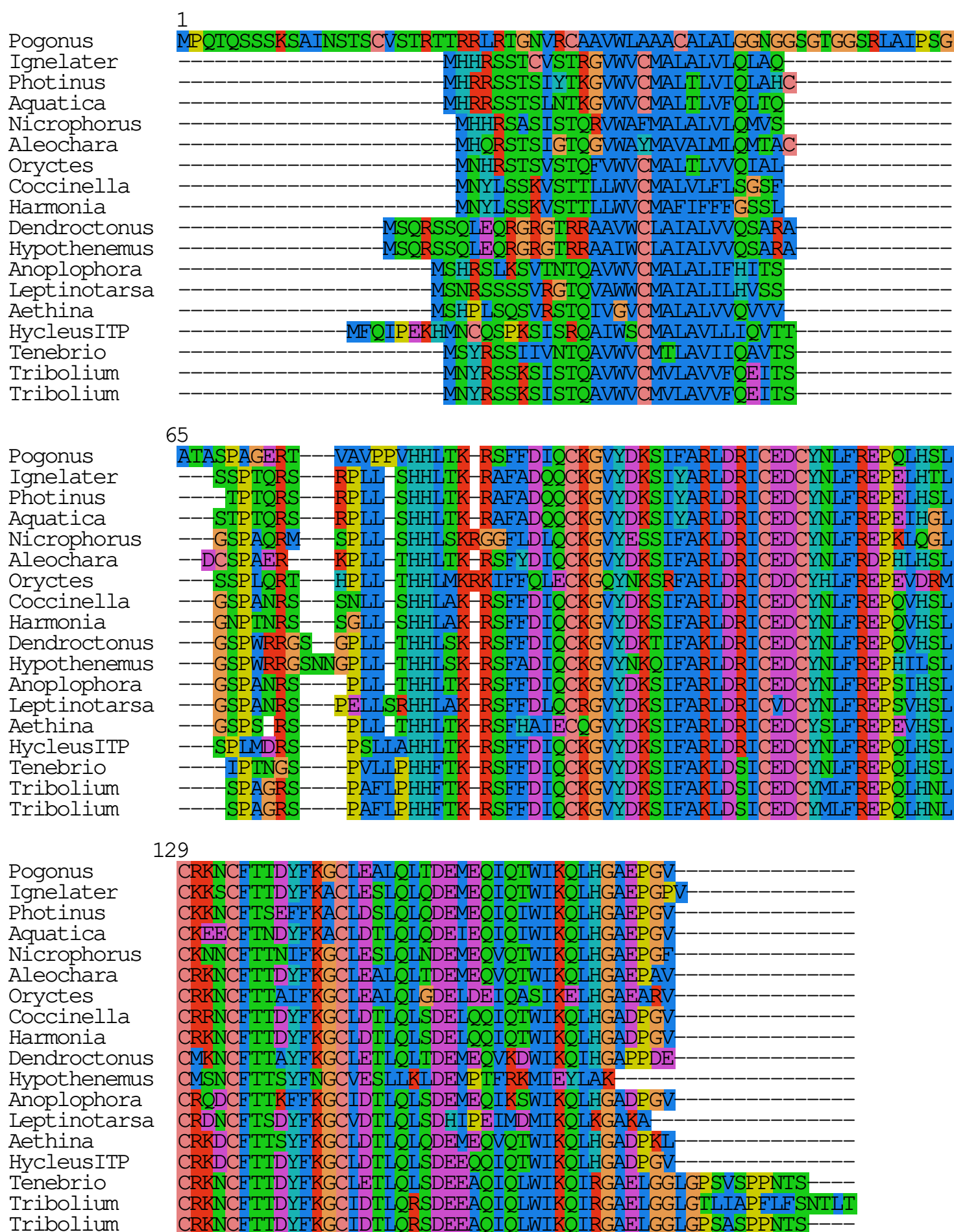
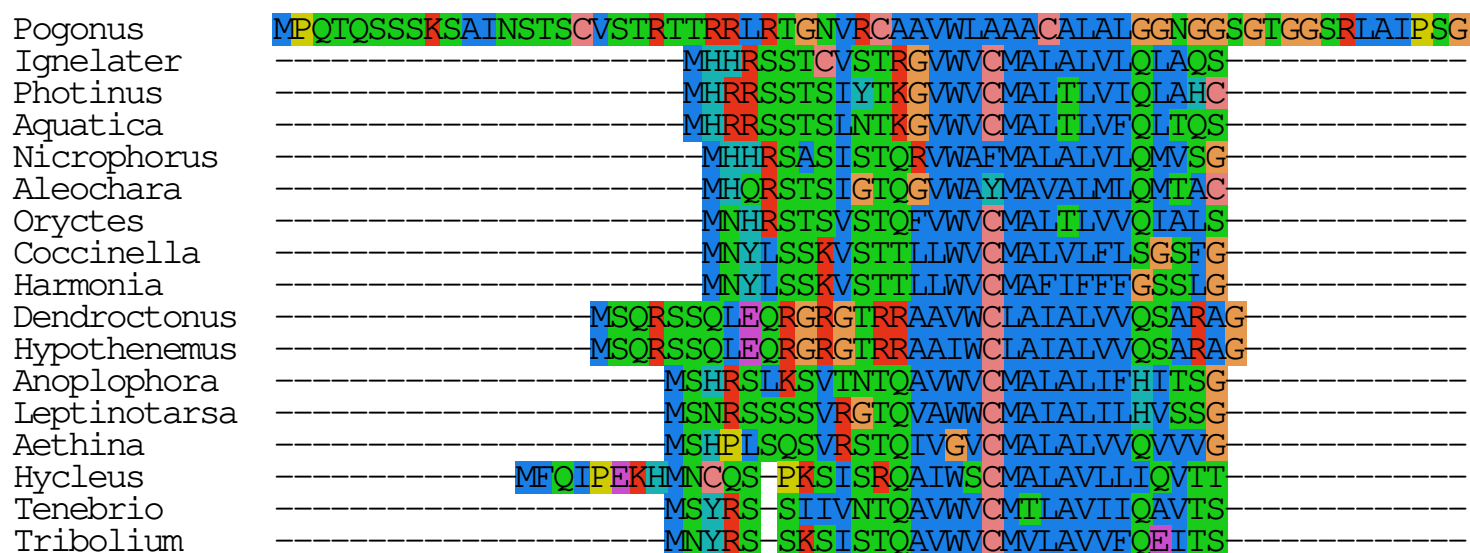


Figure S43. Alignment of ITP-A precursors.



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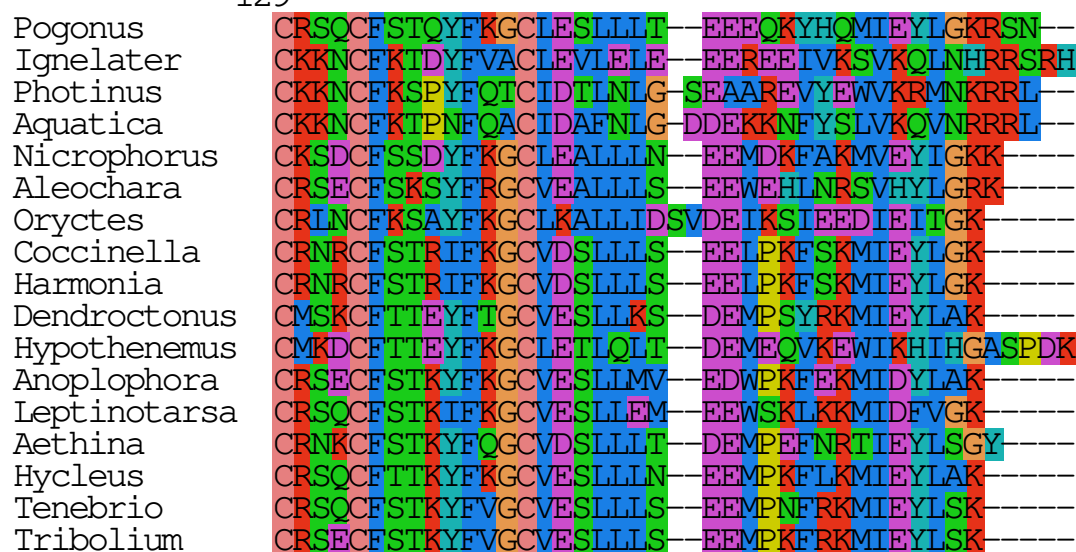


Figure S44. Alignment of ITP-B precursors.