

Supplementary Material S1 Immunomodulatory (IM) peptides applicable as immunoadjuvants with a length of 10 and 15 amino acid residues encrypted in various arthropod AMPs. IM peptides with a score ≥ 0.7 were retrieved using the VaxinPAD program.

SPECIE	Name of parent peptide	Antimicrobial peptide	Peptide ID	Proposed IM sequence from Vaxinpad
<i>Hyas araneus</i>	Arasin 2	SRWPSPGRPRPFGRPNPIFRPRPCICVRQPCCDTY	HAara2-163	PCICVRQPCP
<i>Portunus trituberculatus</i>	PtesALF1	YEALVTSILGKLTGLWHNDSVDFMGHICYFRRRPKIRRFKLYHEGKFWCPGWAPFEGRCKYCVVF	PTesalf1-457	ICYFRRRPKIRRFKL
<i>Macrobrachium rosenbergii</i>	MrH4	MTGRGKGGKGLGKGGAKRHRKVLRDNIQGITKPAIRRLARRGGVKRISGLIYEETRGLKVFLENVIRDAVITYTEHAKR KTVTAMDVVYALKRQGRPLYGFGG	MRh4-679	KPAIRRLARR
<i>Scylla paramamosain</i>	Arasin-likeSp	SPRVSRRYGRPFGRPFVGGQFGRPGVCIRSPCPCANYG	SPAara-291	CVCIRSPCPC
<i>Scylla paramamosain</i>	Sp-ALF1	YETLIASVLGKLTGLWHNNSVDFMGHTCHFRRRPKVRKFYHEGKFWCPGWAPFEGRSRTKSRSRSGSSREAIKDFVRKA LQNGELITQQDATVWVN	SPalf1-452	RRRPKV RKFK
<i>Calliphora vicina</i>	Alloferon	HGVSGHGHGVHG	CVa11-91	SGHGHGVHG
<i>Limulus</i>	LALF	DGIWTQLIFTLVKNLATLWQSGDFQLDHECHYRIKPTFRRLKWKYKGFWCPSWTSITGRATKSSRSGAVEHSVRNFV GQAKSSGLITQRQAEQFISQYN	LPalf1-	YRIKPTFRRLKWK

<i>polyphemus</i>			455	YK
<i>Pachycondyla goeldii</i>	Ponericin W3	GIWGTLAKIGIKAVPRVISMLKKKKQ	PGponW3-227	VPRVISMLKK
<i>Pteromalus puparum</i>	PP113	GKWGWIIYITILFADVGGFKSSRHPEERRVQERRFKRITRGPD	PPp113-266	RVQERRFKRI
<i>Ixodes ricinus</i>	DefMT5	GFFCPYNGYCDRCRKKLRRRGGYCGGRWKLTCICIMN	IRdefmt5-704	RCRKKLRRRG
<i>Lachesana tarabae</i> vi	Latarcin-1	SMWSGMWRRKLKKLRNALKKKLGKE	LTlat1-381	GMWRRKLKKL
<i>Mesobuthus marten</i> sii	BmKbpp	FRFGSFLKKVWWSKLAKKLRSKGKQLLDYANKVLNGPEEEAAAPAE	MMbmK-503	SFLKKVWWSKLAKKL
<i>Gastrophysa atrocyanea</i>	Diapause-specific peptide	AVRIGPCDQVCPRIVPERHECCRAHGRSGYAYCSGGGMYCN	GAdia-145	VRIGPCDQVCPRIVP
<i>Lachesana tarabae</i> vi	Latarcin-5	GFFGKMKEYFKKFGASFKRRFANLKKRL	LTlat5-387	FKRRFANLKK
<i>Oiketicus kirbyi</i>	Cecropin	WKPFKKIEKAVRRVRDGVAKAGPAVAVVGQAT	OKcec-603	WKPFKKIEKAVRRVR
<i>Lucilia sericata</i>	LSerStomox1	AGFRKRFNKLVKVKHTIKETANVSKDVAIVAGSGVAVGAAMG	LSerstox1-548	RKRFNKLVKVKHTI
<i>Galleria</i> a	AGm1	ENFFKEKERKGQRIRDAIISRRPRVETLAQAQKIIKGGD	GMagm	RIRDAIIS

<i>mellonella</i>				1-684	RR
<i>Callinectes sapidus</i>	Callinectin	WNSNRRFRVGRPPVVGRPGVCFRAPCPCSNY		CScal-89	CVCFRA PCPC
<i>Rhipicephalus haema physaloides</i>	Rhamp	ERILDLRKTCKKCKNGEVLGCVSGHGPPGCSENECGMGRPKACFFDCHYGCWCTGKLYRRKRDRKCVPKHECLL		RHrha-419	RRKRDR KCVP
<i>Parabuthus schlechteri</i>	Parabutopon	FKLGSFLKKAWSKLAKKLRAKGKEMLDYAKGLLEGGSEEVPGQ		PSpa-351	SFLKKA WKSCLA KKL
<i>Scylla paramamosain</i>	Sp-ALF2	YEALVASILGKLSGLWHS DTVDFMGHTCHIRRRPKFRKFKLYHEGKFWCPGWTHLEGNSRTKSRS GSARDAIKDFVYKALQNKLITENNA AAWLK		SPalf2-453	HIRRRPK FRK
<i>Scylla paramamosain</i>	Sp-ALF3	YEALVASILGKLSGLWHS DTVDFMGHTCHIRRRPKFRKFKLYHEGKFWCPGWTHLEGNSRTKSRS GSARDAIKDFVYKALQNKLITENNA AAWLK		SPalf2-453-	TCHIRRR PKFRKF KL
<i>Ixodes scapularis</i>	Scapularisin-3	AFGCPFDQGTCHSHCRSIRRRGRRCASF AKRTCTCYQK		ISsca3-531	IRRRGRR CASFAK RT
<i>Macropis fulvipes</i>	Macropin 1	GFGMALKLLKKVL		MFmac1-176	GMALKL LKKV
<i>Vaejovis punctatus</i>	VpAmp1.0	LPFFLLSLIPSAISAIKKI		VPamp1-709	IPSAISAI KK
<i>Drosophila melanogaster</i>	Cecropin-C	GWLKKLGKRIERIGQHTRDATIQGLGIAQQAANVAATARG		DMcccC-120	LKKLGK RIERIGQ HT
<i>Androctonus</i>	Androctoni	RSVCRQIKICRRRGGCYYKCTNRPY		AAaand-	RQIKICR

<i>australis</i>	n			324	RRG
<i>Cryptotympana dubia</i>	Cryptonin	GLNGLALRLGKRALKKIIKRLCR		CDcry106	RALKKIIKRL
<i>Bombyx mori</i>	Moricin 2	AKIPIKAIKTVGKAVGKGLRAINIASTANDVFNLKPKKRKH		BMmor2-64	NFLKPKKRKH
<i>Scylla serrata</i>	SsALF	QYEALVASILGKLSGLWHSDTVDFMGHTCHIRRKPKFRKFKLYHEGKFWCPGWITHLEGNSRTKSRSGSTREATKDFVHK ALQNKLITKNSADAWLKG		Ssal f-495	TCHIRRKPKFRKFKL
<i>Litopenaeus vannamei</i>	Histone H2A	SGRGKGGKVKGKSKSRSSRAGLQFPVGRIHRLLRKGNYAERVGAGAPVYLAAVMEYLAAEVLELAGNAARDNKKTRIV PRHLQLAIRNDEELNKLLSGVTIAQGGVLPNIQAVLLPKKTEKK		LVhis-638	KKTRIVPRHLQLAIR
<i>Derma-centor silvarum</i>	Ds-defensin	VPAESEAAHLRVRRGFGCPLNQGACHNHCSIRRRGGYCSGIIKQTCTCYRN		DSdef-706	LRVRRGFGCP
<i>Portunus trituberculatus</i>	PtALF7	QYEALTAAILTKLSKMWHSDDLNFLGHTCHVSRTPTVKRFKLYWKGKFWCPGWAPFSGTSRTKSRSGSAREATKSFVDQ ALQRRLLITQQEADLWLKG		PTalf7-456	VKRFKLYWKG
<i>Scolopendra mutilans</i>	Scolopin-1	FLPKMSTKL RVPYRRGTKDYH		SMscol-594	TKLRVPYRRG
<i>Periplaneta americana</i>	ISGCock Contig13_4 610?1	YPCKLNLKLGKVPFHF		PAisg13-726	YPCKLNLKLGKVPFH
<i>Galleria mellonella</i>	Gm1	ENFFKEIERAGQRIRDAIISAAPAVETLAQAQKIIKGGD		GMgm1-136	QRIRDAIISA
<i>Acalolepta</i>	Luxoriosin	SVRTQDNAVNRQIFGSNGPYRDFQLSDCYLPLETNPYCNEWQFAYHWNALMDCERAIYHGCNRTRNNFITLTACKNQ		ALI	RAIYHG

<i>luxurio sa</i>		AGPICNRRRH	ux-2	CNRT
<i>Bomby x mori</i>	Moricin	AKIPIKAIKTVGKAVGKGLRAINIASTANDVFNFLKPKKRKA	BM mor- 72	FNFLKPK KRK
<i>Vespa tropica</i>	Mastoparan -VT3	INLKAITALAKLL	VT mas 3- 505	LKAITAL AKK
<i>Bomby x mori</i>	CECD_BO MMO Cecropin-D precursor	GNFFKDLEKMGQVRDAVISAAPAVDTLAKAKALGQ	BM cec- 70	QVRDA VISA
<i>Acanth oscurri a gomesi ana</i>	Gomesin	QCRRLCYKQRCVTYCRGR	AGg om- 328	RRLCYK QRCV
<i>Mischo cyttaru s phthisi cus</i>	Mp_mastop aran MP	INWLKLGKKMMSAL	MP mas- 180	LKLGKK MMSA
<i>Rhipice phalus micropl us</i>	Microplusi n	HHQELCTKGDDALVTELECIRLRISPETNAAFDNAVQQLNCLNRACAYRKMCAATNNLEQAMSVYFTNEQIKEIHDAATA CDPEAHHEHDH	RM mic- 375	ECIRLRIS PE
<i>Mesob uthus eupeus</i>	Meucin-25	VKLIQIRIQYVTVLQMFSMKTQ	ME meu 25- 416	VKLIQIRI WI
<i>Drosop hila melano gaster</i>	Drosomyei n-2	DCLSGKYKGPCAVWDNEMCRRICKEEGHISGHCSPLKWCCEGC	DM dro2 -110	PCAVWD NEMCRR ICK
<i>Oxyope s</i>	Oxyopinin- 2	GKFSVFGKILRSIAKVFKGVGKVRKQFKTASDLDKNQ	OKo xy2-	AKVFKG VGKVRK

<i>kitabensis</i>				371	QFK
<i>Pacificastacus leniusculus</i>	Astacidin 1	FKVQNQHGGVVKIFHH		Plast-417	FKVQNHGQVVKIFH
<i>Oxyopes takobius</i>	Oxt 4a	GIRCPKSWKCKAFKQRVLKRLLAMLRQHAF		OTOxt4a-429	RVLKRL LAML
<i>Lachesana tarabaeivi</i>	Latarcin-2a	GLFGKLIKKGFRKAISYAVKKARGKH		LTlat2a-382	IKKFGRK AISYAV KK
<i>Scolopendra subspinipes mutilans</i>	Scolopendin 1	MDSFQKIEKIGEGTYGVVYKADKVSGRVLVKKIRLENESEGVSTA		SSMsc o1-529	RLVALK KIRL
<i>Macrospis fulvipes</i>	Macropin 2	GTGLPMSERRKIMLMMR		MFmac 2-177	ERRKIM LMMR
<i>Ceratitiscapitata</i>	Ceratotoxin C	SLGGVISGAKKVAKVAIPIGKAVLPVVAKLVG		CCcerC-94	VISGAK KVAKVA IPI
<i>Podisus maculiventris</i>	Thanatin	GSKKPVPPIYCNRRTGKCQRM		PMtha-315	KKPVPPI YCNRRTGK
<i>Lachesana tarabaeivi</i>	Cytoinsecto toxin 1-13	GFFGNTWKKIKGKADKIMLKKAVKIMVKKEGISKEEAQAKVDAMSKKQIRLYLLKHYGKKLFFKRPKNCDQ		LTCyt1-13-628	LKHYGK KLFK
<i>Bactro</i>	Bactrocerin	VGKTWIKVIRGIGKSIKWQ		BDb	IKVIRGI

<i>cera dorsalis</i>	-1			ac1-56	GKS
<i>Bactrocera dorsalis</i>	Bactrocerin-1	VGKTWIKVIRGIGKSKIKWQ		BDb ac1-56-	WIKVIRGIGKSKIKW
<i>Vespa magnifica</i>	Vespid chemotactic peptide 5g	FLIIRRPVLGLL		VM ves-579	FLIIRRPVL
<i>Simulium bannanense</i>	SibaCec	GKLTGDKLKRGAKKALNVASKVAPIVAAGASIAR		SBsi b-711	KLKRGAKKAL
<i>Ixodes sinensis</i>	Ixosin-B	QLKVLDLWGTRSGIQPEQHSSGKSDVRRWRSRY		ISix oB-169	KSDVRRWRSR
<i>Hyalophora cecropia</i>	Cecropin-A	KWKLFKKIEKVGQNIRDGIIKAGPAVAVVGQATQIAK		HCC ecA-322	WKLFFKIEKVGQNIR
<i>Haploelma hainanum</i>	Oh-defensin	MLCKLSMFGAVLGVPACAI DCLPMGKTGGSCGGVCGCRKLT F KILWDKKFG		5HH oh-d-431	GCRKLT F KIL
<i>Periplaneta americana</i>	ISGCock_Contig12_4 176	VGRKHSILNCIPYLKKKKIMRL		PAis g12-732	LNCIPYLKKKKIMRL
<i>Bombyx mori</i>	Attacin	QAGSFTVNSDGTSGAALKVPLTGNDKNVLSAIGSADFNDRHKL SAASAGLALDNVNGHGLSLTGTRIPGFGEQLGVAGK VNLFHNNNHDL SAKAF AIRNSPSAIPNAPNFNTLGGGV DYMFKQKVGASLSAAHSDVINRNDYSAGGKLN LFRSPSSSLD FNAGFKKFDT P FYRSSWEPNVGFSFSKFF		BM att-73	SAKAF AIRNSPSAIP
<i>Litopenaeus stylirosus</i>	Ls-Stylicin1	SSFSPPRGPPGWGPPCVQQPCPKCPYDDYKCPTCDKFPECEECPHISIGCECGYFSCCEPKPVCEPCESPIAELIKGGYKG		LSst yl-174	PCVQQP CPKC

<i>Hogna carolinensis</i>	Lycotoxin-2	KIKWFKTMKSIAKFIAKEQMKKHLGGE	HCl yc2-363	KIKWFK TMKS
<i>Galleria mellonella</i>	G. mellonella moricin- like peptide A	KVNANAIKGGKAIGKGFKVISAASTAHDVYEHKNNRH	GM g.me A-137	AIGKGF KVIS
<i>Cicada flammata</i>	Cicadin	NEYHGFVDKANNENKRKKQQGRDDFVVKPNNFANRRRKDDYNENYYDDVDAADV	CFci c-619	NFANRR RKDD
<i>Antheraea pernyi</i>	pro-ApCec	APEPKWKFFKKIERVGGQNIRDGIIKAGPAVAVVGQATNIAKG	APp ro-763	WKFFKK IERVGQ NIR
<i>Drosophila melanogaster</i>	CEC3_DR OVI Cecropin-3 precursor	GWLKKIGKKIERIGQHTRDATIQGVGIAQQAANVAATAR	DM cec3-121	LKKIGK KIERIGQ HT
<i>Heliothis virescens</i>	Cecropin-B	KWKVFKKIEKVGRNIRDGIVKAGPAIAVLGQAN	HVc ecB-155	KWKVFK KIEK
<i>Lachesana tarabae</i>	Cytoinsecto toxin-1a	GFFGNTWKKIKGKADKIMLKKAVKIMVKKEGISKEEAQAKVDAMSKKQIRLYLLKYYGKKALQKASEKL	LTc yt1a-467	MLKKAV KIMV
<i>Locusta migratoria</i>	Locustin	ATTGCSCPQCII FDPICASSYKNGRRGFSSGCHMRCYNRCHGTDYFQISKGSKCI	LMI oc-179	HMRCYN RCHG
<i>Manduca sexta</i>	Bactericidin B-3	WNPFKELERAGQVRDAIISAGPAVATVGQAAAIARG	MSb acB3-312	QVRDA IISA
<i>Hyphtria cunea</i>	Hyphancin IIIIE	RWKFFKKIERVGGQNVRDGLIKAGPAIQVLGAAKAL	HCh ypE-166	WKFFKK IERVGQ NVR

<i>Pandin us imperat or</i>	Pandinin-1	GKVWDWIKSAAKKIWSSEPVSQKGGQVLNAAKNYVAEKIGATPT	PIpa n1- 365	KVWDWI KSAAKK IWS
<i>Oxyope s kitaben sis</i>	Oxyopinin- 2d	GKFSVFSKILRSIAKVFKGVGKVRKGFKTASDLDKNQ	OKo xy2 b- 374	AKVFKG VGKVRK GFK