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3	Supplementary Information:
4	The ubiquity and ancestry of insect <i>doublesex</i>
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6	Dana C. Price ^{1,*} Andrea Egizi ^{1,2} and Dina M. Fonseca ¹
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Order

Protura Protura Zygentoma Zygentoma Zygentoma Zygentoma Zygentoma Zygentoma Odonata Odonata Ephemeroptera Ephemeroptera Ephemeroptera Ephemeroptera Zoraptera Zoraptera Zoraptera Zoraptera Zoraptera Dermaptera Dermaptera Orthoptera Orthoptera Phasmatodea Phasmatodea Phasmatodea Phasmatodea Phasmatodea Phasmatodea Phasmatodea Phasmatodea Phasmatodea Mantodea Mantodea Mantodea Hemiptera Hemiptera Hemiptera Hemiptera Hemiptera Psocodea (Phthiraptera) Psocodea (Psocoptera) Hymenoptera Hymenoptera Hymenoptera Hymenoptera Hymenoptera Hymenoptera Hymenoptera Hymenoptera Raphidioptera Raphidioptera Megaloptera Neuroptera Neuroptera Strepsiptera Coleoptera Coleoptera Coleoptera Coleoptera Coleoptera Coleoptera Coleoptera Coleoptera Coleoptera Trichoptera Lepidoptera Lepidoptera Lepidoptera Lepidoptera Lepidoptera Siphonaptera Siphonaptera Siphonaptera Mecoptera Mecoptera Mecoptera Diptera Diptera

Diptera

Species Acerentomon sp. AD2013 GAXE01026579 4 Acerentomon sp. AD2013 GAXE01016537 1 Atelura formicaria GAYJ01004010 4 Atelura formicaria FCD0KP1ACXX:7:2202:17 Tricholepidion gertschi GAS001256612_2 Tricholepidion gertschi GAS001035827_5 Thermobia domestica GASN01031735 3 Thermobia domestica GASN01345779 5 Epiophlebia superstes GAVW01013444 6 Cordulegaster boltonii GAY001003812_6 Ephemera danica GAUK01006643 5 Isonychia bicolor GAXA01111429_5 Isonychia bicolor GAXA01058460 2 Baetis sp. AD-2013 GATU01002143 4 Zorotypus caudelli GAYA01001439 2 GAYA01396307 3 Zorotypus caudelli Zorotypus caudelli GAYA01394885 2 Zorotypus gurneyi GABA01003757 Zorotypus gurneyi GABA01001162 6 Apachyus charteceus GAUW01111996 3 Forficula auricularia GAY001017360 5 GAS001124537 1 Tetrix subulata Teleogryllus commodus GBHB01043004 5 Aretaon asperrimus GAZ001017692 2 Aretaon asperrimus GAWC01030481_5 Ramulus artemis GAWE01101273_2 Ramulus artemis GAWE01101272 2 Medauroidea extradentata GAWD01075183 3 GAWG01081702 1 Extatosoma tiaratum Sipyloidea sipylus GAWF01025783 1 Peruphasma schultevi GAWJ01240650 4 Timema cristinae GAVX01147206_3 Metallyticus splendidus GATB01322676 2 Mantis religiosa GASW01222710 6 GASW01220805 1 Mantis religiosa Cercopis vulnerata GAUN01115682 1 Okanagana villosa GAWQ01014169_1 Okanagana villosa GAWQ01014168 1 Notostira elongata GASV01006917 3 Xenophysella greensladeae GAYT01023103 2 Menopon gallinae GAWR01005396 5 Liposcelis bostrychophila GAYV01012131 4 Cotesia vestalis GAUP01001251 3 Chrysis viridula GATY01010708_4 Orussus abietinus GAUJ01013146 6 Tenthredo koehleri GAWW01003430 6 Microplitis demolitor JO916215 4 Sphaeropthalma orestes GAXP01022687 1 Sphaeropthalma orestes GAXP01017427 3 Osmia cornuta GAGH01066955 2 Xanthostigma xanthostigma GAUI01017467_1 Inocellia crassicornis GAZH01013937 4 Corydalus cornutus GATG01001294 4 GAYH01007785 3 Conwentzia psociformis Osmylus fulvicephalus GAYC01086997_6 Stylops melittae GAZM01003841 2 Meloe violaceus GATA01048201 6 Meloe violaceus GATA01002860 6 Gyrinus marinus GAUY01088671 2 Gyrinus marinus GAUY01088671 3 Meligethes aeneus GAPE01021632 6 Meligethes aeneus GAPE01021632 6 GATW01000814_5 Aleochara curtula JU430525_5 Pogonus chalceus GAFI01011207 3 Dendroctonus frontalis Annulipalpia sp. AD-2013 GATX01084595 6 GATC01014479 1 Nemophora degeerella Parides eurimedes GAXH01085310 6 Spodoptera litura GBBY01009273 Trichoplusia ni GBKU01001268 4 Helicoverpa armigera GBDM01028004 1 Ceratophyllus gallinae GAWK01054083 GAYP01008255 1 Ctenocephalides felis Oropsylla silantiewi GAWY01012876_1 GAYK01000638_3 Boreus hyemalis Boreus hyemalis GAYK01015512 3 Boreus hyemalis GAYK01017535 4 Bombylius major GATI01007470 4 Bombylius major GATI01007471 5 GAVA01013308 3 Triarthria setipennis Trichocera saltator GAXZ01015564 6 Liparia lucens GAZD01014428 3 Drosophila miranda GALP01005259 3 Sitodiplosis mosellana GAKJ01011849 2 Teleopsis dalmanni GBBP01063334 1 Bactrocera dorsalis GAKP01007048 1 GAMC01004171 3 Ceratitis capitata Anopheles quadrimaculatus GBTE01014262 1 Aedes albopictus GAPW01003320_1 Psorophora albipes GALA01000925_1 Anopheles funestus EZ974425 2 GAN001004847 1 Corethrella appendiculata GAWM01002290 1 Culicoides sonorensis Belgica antarctica GAAK01007185 1 Belgica antarctica GAAK01003177

GenBank accession OD1 sequence TTPLCARCKNHGLKIKLKGHKRYCRYIRCI-COECIATKAROIKMARO <OD2 singleton> PARLCAQCRNHGLRVPVKGHKRHCAYKQCT-CKECCLVKERQRVVALH 16:4233:6 <OD2 singleton> PSRLCAQCRNHGLRVPVKGHKRFCAYKLCH-CAECLLVKERQRVVALH <OD2 singleton> PSRLCAOCRNHGLRVPVKGHKRYCAYKLCT-CKECCLVKERORVVALH <OD2 singleton> TPPKCARCRNHRLKIPLKGHKRYCKFRYCK-CDKCRLTAERORVMAMO TPPKCARCRNHRLKIPLKGHKRYCKFRYCK-CDKCRLTAERQRVMAMQ SNRLCAQCRNHGLRVPVRGHKRYCTFRLCT-CRECRLVRERQRVVALH SNRLCAQCRNHGLRVPVRGHKRYCTYRLCT-CRECRLVRERQRIVALH <OD2 singleton> SNRMCAOCONHGLKIPVRGHKRFCKYRLCN-CONCLLVKERORIVALH RTPKCGRCRNHGAINELKGHKDACRWKDCS-CSRCALLIARQKINQDL AKPTCEKCKHHGVTVLLVGHKKRCPFTDNLKCAMCVKVDEKRLVMKEQ AKPTCEKCKHHGIIIFLEGHKNRCKYKDELDCEMCLAVDEKRRVMKDQ RIPMCGRCRNHGVINALKGHKDACRWKNCK-CSRCSLLIARQKINQDL AKPTCEKCKHHNIIVLLDGHKKRCRYRNNLDCEMCIKVDEKRKVMKDQ -----YRNCR-CNKCTLTAERORIMAAO KAPNCARCRNHYLKMGLKGHKRYCKFRYCE-CSKCRLTYERQRVMALQ TPPNCARCRNHGLKIPLKGHKRYCKYRYHD-CDKCLLTAERQRVMALQ RTPYCAKCRNHGLTVKLKGHKHKCRFRFHN-CPRCWVTTMRQHLTAIQ MNRLCALCRNHGLKIPVKGHKRFCGYRLCL-CKECCLVKERORVVAMH <OD2 singleton> TPPNCARCRNHRLKIGLKGHKRYCKYRYCT-CSKCRLTAERQRVMALQ TPPNCARCENHELKIGIKGHKEYCKYEYCT-CSKCELTAERORVMALO MNRLCALCRNHGLKIPVKGHKRFCGYRLCL-CKECCLVKERORVVAMH INRLCALCRNHGLKIPVKGHKRFCGYRLCL-CKECCLVKERQRVVAMH INRLCALCRNHGLKIPVKGHKRFCGYRLCL-CKECCLVKERQRVVAMH MNRLCALCRNHGLKMPVKGHKRFCGYRLCL-CKECCLVKERQRVVAMH LNRLCALCRNHGLKKPVKGHKRFCAYKLCV-CRECCLVKERQRVVALH TPPNCARCRNHRLKIGLKGHKRYCMFRHCT-CTKCRLTAERORVMAMO TPPNCARCRNHRLKIGLKGHKRYCKYRYCN-CDKCCLTAERORVMALO TPPNCARCRNHRLKIGLKGHKRYCMYRTCM-CTKCCLTAERQRVMAMQ TPPNCARCRNHGLINILKGHKRYCPFVDCT-CEKCSLTLERQKVMAKQ TPPNCARCRNHSLILPLKGHKRYCKYFDCD-CDKCELTAQRQKVMAKQ TPPNCARCRNHSLILPLKGHKRYCKYFDCD-CDKCELTAQRQKVMAKQ KORFCOFCRNHTKMIRTKGHKKECDYRFCK-CIKCRNTRERHKVMAKO -APNCARCANHCIREPLKGHKRFCGFRDCD-COKCOLTAERORVMA00 TPPNCARCRNHDIKIALKNHKRYCRYKHCK-CTKCILTLOROTVMAAO TPPNCARCRNHDIKVGLKNHKRYCKFKLCK-CAKCILTLQRQKVMAAQ TPPNCARCRNHRLKIALKGHKRYCKYRNCN-CEKCILTKDRQRVMALQ ALRSCARCSNHRLKIALKGHKHYCKYRLCN-CKKCRLTAERQKVMAKQ GTARCSRCLNHEVNMPLKGHKRYCFYKYCI-CTKCLSTAEROVIMAAE TPPNCARCENHELK TALKCHKEYCKYESCN-CEKCELTAERORVMALO TPPNCARCRNHRI.KTALKCHKRYCKYRNCN-CEKCILTKDRORVMALO SRPSCARCRNHDLHIRLKGHKRYCRYLACT-CPKCLLTADRORVMAKO RKPCCARCRNHRVKVALKGHKRICPYKSCK-CPKCEITADRQKITAMQ KPPNCARCRNHQLKIAVKGHKRYCRYRYCK-CDSCKLTADRQRVMARQ TPPNCARCRNHRVKIPLKGHKRYCKYRTCN-CQKCRLTAERQRVMAMQ TPPNCARCRNHRVKIPLKGHKRYCKYRTCN-CQKCRLTAERQRVMAMQ TRPNCARCRNHRVKVPLKGHKRYCKYRTCS-COKCCLTAERORVMAMO APPNCARCNNHKLKIPLKGHKRYCRYKRCC-CSKCLLTLSROODMAKO TPPNCARCRNHRMKVPLRGHKRYCNFRNCT-CQKCRLTADRQRVMAMQ TPPNCARCRNHLLKIPLKGHKRYCRYRTCA-CNKCCLTSERORVMAMO TPPNCARCRNHRMKVALKGHKRYCKFRTCK-CEKCRLTSERQRVMAMQ <OD2 singleton> TPPNCARCRNHRLKIALKGHKRYCKYRACK-CEKCRLTSERORVMAMO <OD2 singleton> TPPNCARCRNHRKKIALKGHKRYCMYRTCK-CEKCLLTTERQRVMAMQ <OD2 singleton> TPPNCARCRNHRLKVGLKGHKRYCTFRHCK-CEKCKLTAERQRVMAMQ TPPNCARCRNHRLKIALKGHKRYCKYRSCK-CEKCKLTSERQRVMALQ TPPNCARCRNHRKKITLKGHKRYCMYRNCK-CQKCQLTSERQRVMAMQ TPPNCARCRNHRIKIALKGHKRYCKYRYCN-CDKCRLTAERORVMALO APPNCARCRNHRVKVELKGHKRYCKYRLCN-CDKCRLTADRQRVMALQ APPNCARCRNHRLKVELKGHKRYCKYRYCN-CEKCRLTADRQRVMAMQ APPNCARCRNHRLKIELKGHKRYCKYRNCT-CEKCILTADRQRVMAQQ APPNCARCRNHRLKVELKGHKRYCRYRHCT-CEKCRLTADRQRVMALQ APPNCARCRNHRLKIELKGHKRYCKYRNCT-CEKCRLTADRQRVMALQ TPPNCARCRNHRLKIALKGHKRYCRFLYCK-CEKCKLTADRORVMAKO TPPNCARCRNHRLKIPLKGHKRYCRYLYCK-CEKCRLTADRORDMARO TPPNCARCRNHRLKIALKGHKRYCRFLYCK-CEKCRLTADRQRVMAKQ TPPNCARCRNHRLKIALKGHKRYCKFRNCN-CYKCILTAERQRIMAVQ <OD2 singleton> <OD2 singleton> TPPNCARCRNHGLKIALKGHKRYCKYRYCN-CEKCRLTAERQRVMALQ <OD2 singleton> TKPNCARCHNHGFKIOLKGHKRYCKFRNCN-CDKCRLTADRORVMALO TPPNCARCRNHSLKIALKGHKRYCRFRHCS-CEKCRLTAERQRVMALQ TPPNCARCRNHGLKITLKGHKRYCKYRYCR-CEKCRLTADRORVMALO TPPNCARCRNHGLKITLKGHKRYCKYRYCT-CEKCRLTADRORVMALO TPPNCARCRNHGDKVELKGHKRYCKYRYCN-CDKCNLTAERORVMALO TPPNCARCRNHGLKTTLKGHKRYCKFRFCT-CEKCRLTVEROINMAOK TPPNCARCRNHGLKITLKGHKRYCKFRFCT-CEKCRLTADRORVMALO TPPNCARCRNHGLKITLKGHKRYCKFRYCT-CEKCRLTADRQRVMALQ TPPNCARCRNHGLKIGLKGHKRYCKYRNCN-CEKCCLTAERQRVMALQ TPPNCARCRNHGLKIGLKGHKRYCRYRHCS-CEKCCLTAERQRVMALQ TPPNCARCRNHGLKIGLKGHKRYCKYRNCN-CEKCCLTAERQRVMALQ TPPNCARCRNHGLKIGLKGHKRYCKYRTCH-CEKCCLTAERORVMALO TAPNCARCRNHALKIPLRGHKRFCKYRYCG-CEKCRMTAERORDMAHK TPPNCARCRNHGLKITLKGHKRYCRYRYCN-CEKCRLTADROKIMALO TPPNCARCRNHGLKIGLKGHKRYCKHRYCN-CDKCRLTAERQRVMALQ KGSNCVRCRNHNLTVPLRGHKQYCPFAACT-CDKCCFTAEQQRQMRLL

OD2 Sequence DASLVFYOOMEHIMOKFKLPIESLPAIYALFKDANGSVEE-TCEKLMSAYEDLHELSVRLEAL AATT. IFOORLEHLINKFKLPVESLPATYAVIKDAKGCVEE-ASERIMAAYEELHEISVRLGII. ARERMLYESLVMMRQAFPVGEEAMPLLLCILKHSR-SVQE-ASLKIYQGHYDLNAKGFVGLDG -----KESLQTLLDMFRLPMETLPLIYVVLKDAR-SDVKEASNRIIEGGYWIMTQMAA EREQMLYESLMVMRQAFPVGEEAMPLLLCILKHSR-SVQE-ASWKICQGTRDLSARGIIGIEN EEGDVMMESLHALLEMFQLPLETLPLIYVVLKDARSDVKE-ASNRIIEAQSELRSIF----ARERMLYESLLVMRQAFPVGEEAMALLLCILKHSR-SVQE-ASWKIYQGTYDLTSRGFVGLDN ENVELLKDSLHALLDMFRLPLETLPLIYVVLKDARSDVKE-ASNRIMEGVYOCMNOGIIAPIG ENMDFIKDAIVALLOMFRLPVDTLPLIYAVLKGTRCDVKE-ASNOIODAODDLRAMALRMMYP ENMDFIKDAIVALLQMFRLPVDTLPLIYAVLKGNRCDVKE-ASNQIQDAQDDLRAMALRMMYP TRERMLYESLMQLRQLYPVPEASLPLLLCILKSSR-SIDE-ASAKIVQGTHELSTRGFPGIEA TRERMLYESLLELRQSFPVPEAGLPLLLCILKSSR-SVHE-ASAKIVQGTHDLTTSGLLGLDG DNIDSLRDSIOTLLDTFRLPMETLPLVYVVLKVSRSDVOE-AFNRISEGGYHMTSSDVPVAAA ORKNLMYSSLLELRSHHPVPDAALPLLLCILNISN-SVKE-ASEKTOKGLOELSLAAASSNEL DKTRVLQIINCLRSYHGLFSWESIPLVHIILEDCGMNYHA-ASSVIFNALRGQAVGRSSRGRT KKEKLSKEVQALMKSFHYKSWESVPLLEIILEDTNFDINR-ASDLILKAYESQYRINVNAEKV KKEKLSKEVQTLMKSFDYKSWESVPLLEIILEDTNFDINK-ASDLILKAYENQYQIKVEEEKV EKTRVLQIINCLRNFHGLSSWESIPLVHIILEDCGMNYHA-ASSVIFNALRRSCCGQEPPRKV KKEKLRKEVHALMKSFGLKSWESVPLLEIILEDANFDVNK-ASDLILKAYESQYRMKVSAEKV IKIQHIRESIHTIMTEFKLPSKSVVLIYTLLKLLRYSINN-VSHHINTSQTELEIIGNHEEQS ANGDVLRDSISVIINKFNLTLEAVPMIYALLKSAHFSVHO-VLENISNGKIPTSYCKINVVII ENFEVLRESIQALLEMFRYPVETLPLIYVILKYARSDIKE-ASNRILEAQEVVCAMGLRDAAR ----LPQSIQMLLDAFRLPMDALPLVYMVLKDSRSDVKE-ASKRILEGERHSSRSAPPSTD-VREOMLYESLLAMRRVFPVSDEAMPLLLCILKHSR-SLOE-ASWKIIOGTOSLISSGOKGLEL VSVEAMMDGVYTLLHMFHYHVEMLPLLLVVLSDAHCDVSE-AYNRILQVLGNHLDSVLPVTSS VSVEAMMDGVYTLLHMFHYHVEMLPLLLVVLSDAHCDVTE-AYNRILQEAYPRRVGNDEKCVT VSVEAMMDGVYTLI.HMFHYHVEMI.PLI.LVVI.SDAHCDVTE-AYNRILOGECAAA-----VRDQMLYESLLAMRRVFPVGDEAVPLLLCILKHSR-SLOE-ASWKIIQGTOSLISSGHKGYES VREQMLYESLLAMRRVFPVGDEAVPLLLCILKHSR-SLQE-ASWKIIQGTQSLISSGHKC*--VREQMLYESLLAMRRVFPVGDEAVPLLLCILKHSR-SLQE-ASWKIIQGTQTLISSGQKGTEA VREQMLYESLLAMRRVFPVGDEAVPLLLCILKHSRSLQEA-SWKIIQGTQSLISNGHKGLEAF VREQMLYESLLAMRRVFPVGEEAVPLLLCILKQSRSLHEA-SWKIIQGTQTLISRGHTGLDSY ESMEIPWDAALTLLEKFGLPAVSLPLIYVVLQYSRSDVSE-ASRHIVSGKYLLR------ESITVSRESILELLERFRLPWETLPLIYVVLDDARCDVSE-ASKRILKGYYVNTSHKVINTFK DTMDIPWEAVLALLOKFEFPAVSLPLIYVVLOYARSDVSE-ASRHILTGKYLIAMOFSP*---ENSKEPWDSILLLMHMLGLPVNTTPLLLIILKEVASDAYE-VCNRFFAAQEELRSTCPIKEEI RDSKELWESILHLLHWFSLPVNTTPLLHIILNEVATDEFD-AYNRILDAQEELRITCPIKEEI RDSKELWESILHLLHWFSLPVNTTPLLHIILNEVATDEFD-AYNRILDGKHSCTVSIIQNTKH TSLRIMWERILTVLIEKGYSLDTYPLLYIILKEVTSDVEE-VYOKIAOWTESSOFILVI*---GGEEVHWESITALLEMFOYRLGPCPLLHIILNKITPNIKE-AYSLILAGRPOLSFIN*----KSLENHRNCVMKLVKDFGLGIDAYPLAYVVFKIAGGDYEV-ATKMILEAOEEITKMSLLSASM KNLKNYRYYVMKLVTDFGLSIDSYPLVLVIFKILGGDYDL-AAKMITEGCDEVYPCLG----ENVEVLLECSTKLLERFWYSWDMLPLMYVILKDARADIEE-ATRRIAEANTEIRAVAFWKARR DSVEVLLECSSKLLELFQYPWDMVPLMYVILKDAGANLEE-AVRRIVEANNDIRALHFWKEVR ESMTDLLOLTLELLNKFEYTWDTIPLMYLILKSVMGDREK-ASRLIREVNDEIRKIYNYHCNR ENVEVILEYSTKLTERFRYPWEMT, PLMT, VILKDAOADLEE-ASBRTAEENTKPAKT, AVNPMTT ENVEVIJECSTKIJERFWYSWDMI.PLMYVIJKDARADIEE-ATERIAEGEENIJFKI.RIHISI. GDIEVLLDYSTKLLELFHYPWETLPLIYVIVKGLGPNVEE-AVRRIVEASNEIRMINFWKAFR DNVEIMLQYSMKLLELFQYSLKMLPLMYLIVKDSGANLEE-AVRRIVEANNEIRTMHFWKAMR DTVEIMLEYSTKLLELFQYPWEMLPLMYVILKDAGGNLEE-AMRRIVE-----HNVEVVLNYCHRLLDTFSYPWEMMPLMYVILKYASGDLTE-AVRSIDEGKRVVNEYSRIHNLN HNVEVVLNYCHRLLDTFSYPWEMMPLMYVILKYASGDLEE-AVRSIDEGKSVMSSIFYKMEMT NNIELILDCCHRLLDTFLYPWELMPLMYVILKYAGGNLDE-AIRSIREGKRVVNEYSRTHNLN VPSEAIIDKCHALLDKFKYPWEIMHLLITILRMCNYDYKL-AETQIQEGKEIVNEYSRTHNLN KQEEVVLEYCFRLLDRLKYPWEVMPLLYAILNVCDWNMDE-----INTEQIIDYSQKLMEIYKYPYEMMPLMYAILKSVNVDFIE-AQTRIDEGHQVVSDFLRKHNLN AQSTDLLEDCQKLLERFKYPWEMMPLMYAILKDARADLEE-ASRRIDEGKRVVNEYSRIHNLN RGAENLLEFCQRLKDKFQLSWKMISLVDVILKYAKEDQEE-AWRQIDEAFLEIRALAAVEAAR GOSTELLEDCOKLLDRFKYPWEMMPLMYVILKDAKADLEE-ASRRIDEGKRAVYEYSRIHNLN RSLCILHEICOKLHERYOWSWKMFPLVYLILKTHNEDYGD-ITRRIDEAYOEIRAIVAVETAK HQSTDLLEDCQKLLERFKYPWEMMPLMYAILKDARADLEE-ASRRIDEGRGFEILLDLCHRLR RGFEILLDLCHRLREKFNLSWRMIPLVDVILKYAKENQDE-AWRQIEEAFLYIKERTLAALEV VHSTDLMEDCLKLLERFRYPWEMLPLMYTILKDANGDLEE-AARRIDDGFIVLKTLLQTVVGY GQSTELLEDCQKLLDRFKYPWEMMPLMYVILKDAQADIEE-ASRRIDEDSNIKLLADLCSRLS DRSMDLLEDCHKLLEKFOYPYEMMPLMYAILKDARADLDE-ASRRIDEGKOVVNEYSRTHNLN PLFENMLDNCOKLIEKFOYSWEMMPLIYVILKYAKGNTEE-ASKSIEEGKWVITEYARONNLN HSLEALVENCHKLLEKFHYSWEMMPLVLVILNYAKSDIEE-ASRRIDEGKKIVNDYAWKHNLN VSLETLVENCHRLLEKFHYSWEMMPLVLVILNYAGSDLDE-ASRKIDEGKLIVNEYARKHNLN VSLENLVDNCNKLLEKFHYSWEMMPLVLVILNYAGSDVDE-ASRKIDEGKMIINEYARKNNLN VSLDALVENCHKLLEKFHYSWEMMPLVLVILNYAGSDLEE-ASRKIDEGKLMIDEYARKHNLN VSFETLVENCQKLLDKFHYSWEMMPLVLVILNYAGSDLDE-ATRKIDEGKMIINEYARKHNLN EEVKTLLEESHOILEOLKYPFEMLPFVFAILRKSDG-ISD-AIKNIHEGOLVVSEYSRMNNLN EKVKALLEESOOIVEOLKYPFEMLPFVYAILGLSGG-ISE-AIKNIHNGOLVVSEYSRMNNLN EEVKILLEESHQILEQLKYPFEMLPFVFAILRKSNG-ISD-AIKNIHEAVQAVQLSFFRGWFD SDARDYVQMCYRLLDHFNYPFEMMPLIYANLKYAGN-IDE-ASRRIEEGQWVVSEYSRLNNLN YSDKVVFDLSQKLLEKFNYPFEMMPLIYANLKYAGS-IDE-----YFDKVVFDLSFKLLKKFNCPIEMMPLIYANLRYADN-IDE-ASRRIEEDILRITYGRYHLVNL ONAELLLEYCOKLLEKFRYPWEMMPLMYVILKDARADIEE-ATRRIDEGOCVVNOYSRIHNLN OKAERLLEYCOKLLEKFRYSWKMMPVMYAILKDARADVEE-ASRRIDEAIRVVRPLTLHLRYP MSIDLILDYCOKLIEKFGYPWEMMPLMYVILKDAGVDIDE-ASKRIEEGIOVLKDSKMFDGID QNTELLLEYCQKLLDKFNYPWEMMPLMYAILKNENN-VEE-AERRIDEGRYVFKLAMRNHVMS TEDPOFLYNCEKLLEKLRYPWEMMPLIFAILKSARNDIEE-ALKLIEEGOHVVNEYYROHNLN LGQDVFLDYCQKLLEKFRYPWELMPLMYVILKDADANIEE-ASRRIEEGQYVVNEYSRQHNLN DNADMLLEKSOKMIERFNYPWEMMPLLYVILKYANGDLEE-ATNOLSEGODFIGKYSRMHNLN PPRDILIGNSEKLLOKFGYPYELMPLMYVIVRYAKADMEE-ASBSIDEGIRDTKDYKNLIDNI. LAODVFLEHCOKLLEKFRYPWEMMPLMYVILKDAGADIEE-ASRRIEEGOHVVNEYSROHNLN LAQDVFLEHCQKLLEKFRYPWEMMPLMYVILKDAGADIEE-ASRRIEEAKRIVNQTISLHWMD LPDDELVKRAQWLLEKLSYPWEMMPLMYVILKSADGDVQK-AHQRIDEGKRTIKAYELVKSSL RTDDELVRQSQQLLEKLHYPWEMMPLMYVILKGANGDVAK-AHQRIDEGQAVVNEYSRLHNLN RTDDELVKRSQYLLEKLHYPWEMMPLMYVILKGADGDVQK-AHQRIDEGMGVWRCWDFRWVVN LPDDELVKRAOWLLEKLGYPWEMMPLMYVILKSADGDVOT-AHORIDEGOAVVNEYSRLHNLN MONDILLEKSHOLLEKFNYPWELLHLMYTILKYSRANIEE-AARRIEEGOYVVNEYSKLHNLN RKTDRLIDNCDTLCSRYGYPWEMMPLVYTILALVNGDIEE-ACNRIDEGKMAILDYSNONNVS NNNDNFWDMADQMARDNNYPDEVTPLLHALIKSRKGKFHE-AQKLLDEGKMVVTEYSRRLNIN KDRKALFDCSSELLKKFRYPWEMMPLMYALSKYANGNIGE-VLQRIDEGQFAVNEQLRQQTLN

Order

Protura Collembola Archaeognatha Zygentoma Odonata Ephemeroptera Ephemeroptera Ephemeroptera Dermaptera Dermaptera Plecoptera Plecoptera Grylloblattodea Grylloblattodea Embioptera Mantodea Blattodea Blattodea Isoptera Hemiptera Hemiptera Hemiptera Hemiptera Psocodea (Psocoptera) Psocodea (Psocoptera) Neuroptera Neuroptera Neuroptera Coleoptera Trichoptera Trichoptera Trichoptera Lepidoptera Lepidoptera Lepidoptera Lepidoptera Siphonaptera Mecoptera Diptera

Species Acerentomon sp. AD-2013 Sminthurus viridis Lepismachilis y-signata Thermobia domestica Calopteryx splendens Isonychia bicolor Baetis sp. AD-2013 Eurylophella sp. AD-2013 Apachyus charteceus Forficula auricularia Leuctra sp. AD-2013 Cosmioperla kuna Grylloblatta bifratrilecta Galloisiana yuasai Haploembia palaui Empusa pennata Periplaneta americana Blatella germanica genome Prorhinotermes simplex Xenophysella greensladeae Planococcus citri Planococcis citri Acanthosoma haemorrhoidale Ectopsocus briggsi Ectopsocus briggsi Pseudomallada prasinus Conwentzia psociformis Euroleon nostras Lepicerus sp. AD-2013 Rhyacophila fasciata Platycentropus radiatus Hydroptila sp. AD-2013 Yponomeuta evonymellus Polyommatus icarus Dyseriocrania subpurpurella Trodia sylvina Ctenocephalides felis Panorpa vulgaris Bibio marci

[Accession] [frame]/[range] GAXE01105806.1 2/58-86 FCD0GUKACXX:7:1301:17137:108047:31 Contig 3838812 GASN01365969.1 4/81-127 GAYM01080693.1 4/36-82 GAXA01091320.1 6/79-119 GATU01010243.1 2/88-133 GAZG01080309.1 5/8-52 GAUW01001274.1 3/14-59 GAY001017359.1 5/107-152 GAUF01079838.1 5/141-187 GAYL01095698.1 6/28-74 GAWP01142412.1 3/58-104 GAWN01171825.1 4/47-93 GAZA01236824.1 5/84-130 GAWT01309511.1 4/111-157 GAWS01015027.1 4/1-44 JPZV01136765.1 2/621-668 GASE01075660.1 4/15-55 GAYI01136691.1 3/24-70 GAXF01116661.1 1/146-191 GAXF01103858.1 1/43-88 GAUV01009568.1 3/70-108 GAPT01011620.1 2/92-137 GAPT01002163.1 6/155-201 GAVV01019538.1 2/82-128 GAYH01007786.1 3/137-182 GAXW01087396.1 4/25-71 GAZB01138597.1 2/45-91 GAXX01046516.1 4/10-56 GASS01092876.1 3/184-204 GAVM01073298.1 1/112-158 GASG01093141.1 4/23-68 GAST01056365.1 4/55-86 GASY01015364.1 1/256-302 GAVB01150705.1 2/60-106 GAYP01006643.1 1/27-73 GAUH01062093.1 3/57-103 GATJ01056020.1 6/139-184

Sequence

sequence		new e-vai
RLPKCARCINHGL	KIPLKGHKRYCTNKYCDCP	8.5e-13
	KIPLRGHKRYCRFRTCKCQKCILTAERQRVMAMQ	2.0e-14*
APPNCARCRNHRL	KVELKGHKRYCKYRYCNCEKCRLTADRQRVMALQ	7.2e-28*
TPPKCARCRNHRL	KIPLKGHKRYCKYRFCNCDKCLLTAERQRVMALQ	1.3e-29
TPPKCARCRNHLY	KIPLKGHKRYCKFRYCKCDKCCLTAERQRVMAKQ	1.4e-29
TPPKCARCRNHRL	KIPLRGHKRYCNYRYCNCEKCKLTADRR	1.9e-24
	KIPVRGHKRFCKFRLCNCQNCLLVKERQRIVALH	1.2e-20
	KIPVKGHKRFCRNRTCNCAECKLVRERQRVVALH	1.9e-20
KTPNCARCRNHYV	KIKLKGHKRYCKYRNCRCNKCTLTAERQRIMAAQ	4.8e-28
	KMGLKGHKRYCKFRYCECSKCRLTYERQRVMALQ	2.3e-27
	KITLKGHKRFCAFRFCNCEKCCLTVERQRVMAAQ	3.0e-27
	KIALKGHKRFCRYRYCECDKCRLTAERQRVMAAQ	3.4e-29
	KIGLKGHKRYCKYRA <mark>S</mark> PCDKCCLTAERQRVMAMQ	5.4e-29
	KIGLKGHKRYCKYRHCTCDKCCLTAERQRVMAMQ	2.0e-30
	KIGLKGHKRYCRFRSCTCSKCCLTAERQRVMALQ	3.3e-28
	KIGLKGHKRYCKYRYCNCDKCCLTAERQRVMALQ	2.8e-30
	KIGLKGHKRYCKYRYCNCDKCCLTAERQRVMALQ	1.2e-28
	KIGLKGHKRYCTFRSCVCEKCVLTAERQRVMALQ	1.7e-15*
	KVQIKNHKNKCPFKDCLCKDCIPIIVRRDSTALA	7.6e-08
	KIPLKGHKRYCCFRDCDCEMCCLTAKRQKVMALQ	1.7e-27
	KIPLKGHKRYCRFWNCVCEKCVQTSTRQKVMARE	1.2e-22
	KNVLSGHKRYCLYLECKCEKCLATVGRQRSMAKN	4.9e-20
	KEEMKGHKRECERRHCSCDKCQATEKKRAKTRIQ	7.1e-12
	KEVLKGHKKKCIYRDCTCGSCMLSVNRREIAKKQ	8.3e-20
	KQSLKGHKRYCRFLTCDCSLCKLVVEKRVVMADQ	4.3e-15
	KIPLRGHKRYCRFRNCICHKCKLTAERQRVMAMQ	2.2e-28
	KIPLKGHKRYCRYKRCCCSKCLLTLSRQQDMAKQ	3.7e-22
	KIPLRGHKRYCRFRTCTCEKCRLTAERQRVMAMQ	7.1e-29
	KEPLKGHKRYCKFKKCQCEKCKLTAERQRVMAKQ	6.4e-28
	KIGLKGHKRYCKYRYCTCEKCRLTAERQRVMALQ	4.8e-31
	KIALKGHK	4.0e-08
	KIALKGHKRYCKFRHCNCDRCRLTAERQRVMALQ	3.6e-29
	KVELKGHKRYCKYRYCNCEKCRLTADRQRVMALQ	3.2e-30
	KIQLKGHKRYCKYRYCTCEK	1.1e-19
	KVALKGHKRYCRYRYCNCDKCRLTADRQRVMAMQ	2.5e-29 1.9e-30
	KEKLKGHKRYCKYRYCNCDKCKLTAERQRVMALQ	
	KIALKGHKRYCRYLYCKCEKCRLTADRQRVMAKQ	6.6e-29 1.5e-28
	KIALKGHKRYCTFKFCNCEKCRLTAERQRVMALQ	
TOPNCARCRNHCY	K <mark>VLLKGHKHYCRFRYCKCEKCVVTAKRQDIMANE</mark>	2.7e-25

HMM e-val

rder rchaeognatha lattodea lattodea	Species Meinertellus cundinamarcensis	[Accession]_[frame]/[range] GAUG01133643.1 3/44-81	HMM aligned sequence -TPKCARCRNHGV <mark>I</mark> SCLKGHKRLCRWRECRCPNCLLVVE	HMM e-va
lattodea lattodea		GAUG01133643.1 3/44-81	-TPKCABCRNHGVTSCLKGHKRLCBWRECRCPNCLLVVE	
lattodea				
	Periplaneta americana	GAWS01189376.1_5/2-47	-TPKCARCRNHGVISCLKGHKRLCRWRECQCPNCQLVVERQRVMAAQ	
	Cryptocercus wright	GAZN01162439.1_6/11-56	-RPKCARCRNHGMISWLKGHKRQCRFKSCVCAKCNLIAERQRVMAAQ	
alanoida	Calanus finmarchicus Calanus finmarchicus	GAXK01154012.1_1/25-70	-RPKCARCRNHGLISWLKGHKRSCQYRDCDCAKCNLIQERQRVMAAQ	
alanoida		GAXK01154011.1_1/25-70	-RPKCARCRNHGLISWLKGHKRSCQYRDCDCAKCNLIQERQRVMAAQ	
oleoptera	Aleochara curtula	GATW01017006.1_5/27-72	-VPKCARCRNHGMISTLRGHKKQCIYKNCSCAKCGLIKERQRIMAAQ	
oleoptera ollembola	Ips typographus	GACR01003199.1_2/53-98	-VPKCARCRNHGMISTLRGHKKQCIYKGCNCAKCGLIKERQRIMAAQ	
ollembola	Tetrodontophora bielanensis	GAXI01024987.1_5/8-53 GATD01092185.1_3/9-54	-TPKCARCRNHGVISCLRGHKKMCRWKECTCEMCELVAERQRIMAAQ	
ollembola	Pogonognathellus sp. AD-2013 Anurida maritime	GAUE01052744.1 3/128-173	-TPKCARCRNHGVISCLRGHKKMCRWKECSCEMCELVAERQRIMAAQ -APKCARCRNHGVISCLRGHKKMCRWKECDCELCELVAERQRIMAAQ	
iptera	Culicoides sonorensis	GAWM01017604.1 1/28-73	-TPKCARCRNHGVISCLKGHKRYCRWKECDCELCELVAERQKIMAAQ	
-	Trichocera saltator	GAXZ01130618.1 6/9-54	-TPKCARCRNHGVISCLKGHKKICRWKECCCPNCQLVVERQRVMAQQ	
iptera			-TPKCARCRNHGVISCLAGHKRLCRWRECCCPMCQLVVERQRVMAAQ	
phemeroptera phemeroptera	Isonychia bicolor Eurylophella sp. AD-2013	GAXA01106317.1_1/122-167 GAZG01012200.1 6/32-77	-SPKCARCRNHGVISGLAGHARLCRWRECRCQSCLLVVERQRVMAAQ	
rylloblattodea	Galloisiana yuasai	GAWN01063629.1_6/2-47	-RPKCARCRNHGVISSLAGHKRLCRWRECKCFSCLLVVERQRVHAAQ	
emiptera	Graminella nigrifrons	GAQX01025711.1 6/87-133	TPPNCARCRNHDIIVOLKGHKRYCGYISCRCEKCLLTAEROKVMAAQ	
emiptera	Cercopis vulnerata	GAUN01102337.1_4/98-144	TSPNCARCRNHGI INILKGHKRYCPYVDCSCEKCSLTMERQKVMAKQ	
emiptera	Macrosiphum euphorbiae	GAOM01011235.1 1/7-52	-RPKCARCRNHGTINILKGHKKICFIVDCSCERCSLIMERQKVMARQ	
emiptera	Cercopis vulnerata	GAUN01003127.1 4/8-53	-TPKCARCRNHGVISWLAGHKRRCSIKECVCPACHLIAERQKVMAAQ	
emiptera	Nilaparvata lugens	GAYF01062266.1 2/4-49	-HPKCARCRNHGVISCHKGHKKDCHKCSCSCA-CSLIAERQRVMAAQ	
emiptera	Acanthocasuarina muellerianae		TLPNCARCRYHKVIVGLKDHKHYCPNKTCKCEGCLNIEERQRVMAA	
-				
emiptera	Cercopis vulnerata Ganaspis sp. Gl	GAUN01092972.1_3/100-120	-PPNCARCRNHDIINILKGHKR	
ymenoptera ymenoptera	Ganaspis sp. Gl Ganaspis sp. Gl	GAIW01023601.1_3/89-135 GAIW01012036.1 5/215-260	TKPNCARCRNHKI IIKLKQHKNFCKYRYCMCEKCILLKERQRIMAEQ -STNCGYCRNHNI IIGLKGHKIFCKYRYCTCEKCILLKERQRIMAEQ	
ymenoptera ymenoptera	Leptopilina clavipes	GAXY01012036.1_5/215-260 GAXY01053312.1_1/30-75	-KPKCARCRNHGLISWLRGHKRECRYRDCFCVKCSLIAERQRVMAAQ	
-		GAXR01000833.1 2/51-96	-RPKCARCRNHGLISWLRGHKRECRIRDCFCVRCSLIAERQRVMAAQ	
/menoptera /menoptera	Stigmatomma oregonense Ganaspis sp. Gl	GAIW01012035.1 5/481-526	-STNCGYCRNHNI <mark>I</mark> IGLKGHKIFCKYRYCTCEKCILLKERQRIMAEQ	
-	Cotesia vestalis		-SPKCARCRNHGIISSLKGHKRSCRWKDCNCPCCLLVVERORVMAAQ	
/menoptera /menoptera	Cotesia vestalis Cotesia vestalis	GAUP01064990.1_1/26-71 GAUP01004774.1 1/35-80	-RPKCARCRNHGIISSLAGHKRSCRWKDCNCPCCLLVVERQRVMAAQ	
	Cotesia vestalis	GAKG01004932.1 5/23-68	-SPKCARCRNHGLISWLKGHKRECKIKECHCFKCSLIAERQKVMAAQ	
ymenoptera	Osmia cornuta	-	-RPKCARCRNHGLISSLAGHKRSCRWRDChCPCCLLVVERQRVHAAQ	
menoptera	Mischocyttarus flavitarsis	GAGH01047348.1_5/369-414 GAXM01016837.1 6/60-105	-SPKCARCRNHGLISWLKGHKRECKIRECLCPRCSLIAERQKVMAAQ	
ymenoptera	-	GAXP01010837.1_0/00-105	-RPKCARCRNHGVISGLAGHKRSCAWADCRCFCCLLVVERQKVMAAQ	
/menoptera /menoptera	Sphaeropthalma orestes Stigmatomma oregonense	GAXP01040437.1_2/70-113 GAXR01010442.1_4/19-64	-SPKCARCRNHGLISWLRGHKRECKIRECLCVRCSLIAERQKVMA	
-	Orussus abietinus	GAUJ01091006.1 5/57-102	-SPKCARCRNHGVISGLAGHKRSCAWKDCRCFCCLLVVERQRVHAAQ	
menoptera	Tenthredo koehleri		· · · · · · · · · · · · · · · · · · ·	
menoptera		GAWW01088411.1_2/215-260	-SPKCARCRNHGVISGLKGHKRSCAWKDCRCPCCLLVVERQRVMAAQ	
ymenoptera	Chyphotes mellipes	GAXL01043535.1_1/29-74	-SPKCARCRNHGVISGLKGHKKSCAWKDCRCPCCLLVVERQRVMAAQ	
ymenoptera	Pseudomasaris vespoides	GAXQ01011632.1_5/152-197	-SPKCARCRNHGVISGLKGHKRSCAWKDCRCPCCLLVVERQRVMAAQ	
ymenoptera ymenoptera	Pseudomasaris vespoides Pseudomasaris vespoides	GAXQ01011634.1_5/152-197	-SPKCARCRNHGVISGLKGHKRSCAWKDCRCPCCLLVVERQRVMAAQ -SPKCARCRNHGVISGLKGHKRSCAWKDCRCPCCLLVVERQRVMAAQ	
	Pseudomasaris vespoides Pseudomasaris vespoide	GAXQ01011633.1_6/152-197	-SPKCARCRNHGVISGLAGHKRSCAWKDCRCPCCLLVVERQRVMAAQ	
ymenoptera	Cotesia vestalis	GAXQ01011635.1_6/152-197		
ymenoptera	Mastotermes darwiniensis	GAKG01010867.1_5/96-130	-RPKCARCRNHGLISWLRGHKRECRYKECLCPKCSL	
soptera soptera	Prorhinotermes simplex	GAZE01022571.1_3/6-51 GASE01241946.1_5/28-73	-TPKCARCRNHGVISCLKGHKRLCRWRECQCPNCQLVVERQRVMAAQ -TPKCARCRNHGVISCLKGHKRLCRWRECQCPNCQLVVERQRVMAAQ	
-	-		TTPKCARCRIMECTPVKGHKRDCPHRECHCRDCYVVAKGGMYRALQ	
soptera	Nasutitermes takasagoensis	G5zW0JF02FLJ2Z_4/133-171		
epidoptera	Spodoptera exigua	GARL01010958.1_6/32-77	-VPKCARCRNHGLISSLRGHKKACAYRLCQCPKCGLIKERQRIMAAQ	
epidoptera	Polyommatus icarus	GAST01022246.1_6/28-73	-TPKCARCRNHGVISCLKGHKRLCRWRDCRCPSCLLVLERQRVMAAQ	
pidoptera	Athetis lepigone	GARB01040060.1_3/51-96	-VPKCARCRNHGLISSLRGHKKACAYRLCQCPKCGLIKERQRIMAAQ	
pidoptera	Yponomeuta evonymellus	GASG01107059.1_5/89-129	-TPKCARCRNHGVISCLKGHKRLCRWRDCRCPGCLLVLERQR	
intodea	Metallyticus splendidus	GATB01281539.1_1/31-76	-TPKCARCRNHGVISCLKGHKRLCRWRECRCPNCQLVVERQRVMAAQ	
ecoptera	Nannochorista philpotti	GADB01005720.1_1/40-85	-TPKCARCRNHNVIITDVKGHKKVCGWRNCRCPRCRLVDERQRVMAAQ	
coptera	Nannochorista philpotti	GADB01001122.1_6/166-211	-TPKCARCRNHGVISCLKGHKKLCRWRECQCPNCQLVVERQRVMAAQ	
coptera	Panorpa vulgaris Conventzia progiformis	GAUH01056862.1_5/41-86	-VPKCARCRNHGVISGLRGHKKECLYKGCLCAKCNLIKERQRIMAAQ -VPKCARCRNHGIISSLRGHKKICAYKSCRCAKCNLIHERORIMAAQ	
uroptera	Conwentzia psociformis	GAYH01081098.1_6/60-105		
uroptera	Osmylus fulvicephalus Tetrix subulata	GAYC01008034.1_6/33-78	-TPKCARCRNHGI SCLKGHKRLCPWKECKCNNCQLVLQRQKIMAAQ TPKCARCRNHGV SSLKGHKRLCRWRECSCASCQLVVERQRVMAAQ	
thoptera	Ceuthophilus sp. AD-2013	GASQ01096846.1_1/5-50		
thoptera		GAUX01271954.1_3/33-78 GAZT01143051.1_3/24-69	-TPKCARCRNHGVISCLKGHKRLCRWRECQCPNCQLVVERQRVMAAQ	
thoptera	Prosarthria teretrirostris Gryllotalpa sp. AD-2013	GAZT01143051.1_3/24-69 GAWZ01138841.1_4/2-47	-TPKCARCRHHGVISSLKGHKRLCRWRECVCASCQLVVERQRVMAAQ	
thoptera asmatodea			-TPKCARCRNHGVISCLKGHKRLCRWRE-QCPYCQLVVERQRVMAAQ	
	Timema cristinae	GAVX01115626.1_6/41-86	-TPKCARCRNHGVISCLKDHKRLCRWRECQCPNCQLVVERQRVMAAQ	
thiraptera	Menopon gallinae	GAWR01095172.1_5/146-191	-KPKCARCRNHGVISWLKGHKRECRYKDCTCARCILIAERQKVMAKQ -TPKCARCRNHGVISCLKGHKRLCRWRECQCPNCQLVVGRQKVMAAQ	
thiraptera ecoptera	Menopon gallinae Leuctra sp. AD-2013	GAWR01093074.1_6/87-132 GAUF01064618.1 6/8-53		
ecoptera socodea (Psocoptera)	Leuctra sp. AD-2013 Liposcelis bostrychophila	—	-TPKCARCRNHGVISCLKGHKRLCRWRECKCPNCQLVVERQRVMAAQ -TPKCARCRNHGVISCLKGHKKLCRWRECQCPNCQLVVERQRVMAAQ	
	Inocellia crassicornis	GAYV01083169.1_5/23-68		
aphidioptera		GAZH01032725.1_5/40-78	-IPKCARCRNHGMISSLRGHKKLCIYKNCRCAKCNLIKER	
iphonaptera	Ceratophyllus gallinae	GAWK01044190.1_6/53-98 GAZX01023775.1 4/32-77	-TPKCARCRNHGVISCLKGHKKLCRWRECQCPNCQLVVERQRVMAAQ	
iphonostomatoida	Caligus rogercresseyi	GAZX01023775.1_4/32-77 GAXC01041671.1 5/35-80	-RPKCARCRNHGVISWLKGHKRHCAYKDCSCCKCNLIAERQRVMAAQ -TPKCARCRNHGVISCLKGHKKLCRWRECRCAACLLVVERQRVMAAQ	
nysanoptera	Thrips palmi Physcophile fesciate			
richoptera	Rhyacophila fasciata	GAXX01069925.1_6/105-138	-EPKCASCKNHGVISDLKGHKRFCEWRDCECVKCR	
ygentoma	Occasjapyx japonicas Trichelenidien contachi	GAXJ01053929.1_3/25-69	PKCARCRNHGMISWLKGHKRHCRFKDCICAKCNLIAERQRVMAAQ	
	Tricholepidion gertschi	GAS001237223.1_5/12-57	-TPKCARCRNHGVISCLKGHKKLCRWRECQCPNCLLVVERQRVMAAQ	
ygentoma ygentoma	Occasjapyx japonicas	GAXJ01108043.1 3/66-111	-TPKCARCRNHGVISCLKGHKKLCRWRECQCPNCLLVVERQRVMAAQ	

Order	Species	NCBI accession	Sequence	HMM e-val
Protura	Acerentomon sp. AD2013		AATLIFQQRLEHLLNKFKLPVESLPAIYAVLKDAKGCVEEASERLMAAYEELHEISVRLGLL	9.8e-14
Collembola	Anurida maritima		MSVDGMKESLHALLEMFRFPLESLPLVYRILKDYRADFRLASSKIMEAQTELRSLALLEAAR	2.7e-17
Collembola	Pogonognathellus sp. AD-2013	_	GAEPLVDGMKESLQTLMDMFPNPLPLVYLILKDSRADFKLASTRILEAQAELRSIALREAAR	7.8e-11
Collembola	Orchesella cincta	_	TAIDEMKESVKALMEMFRLPSETDFLIYRILKDYRGDFKQASAKIFEAQCELRSWELLREAA	8.6e-12
Diplura	Occasjapyx japonicus	-	DNVEEVNQALKVILKLFHLPDESCHLLWVILKDAKGDLMLASSRIMEAQSELRSMAFQDVTR	1.7e-15
Archaeognatha			ENVENVKESVYGLMDRFRLPSECLPLLYVILKDARFDIKEAANRIVDAQADLHAVTLREAQY	1.6e-20
Archaeognatha	Lepismachilis y-signata		VSLETLVENCNKLLEKFHYSWEMMPLVLVILNYAGSDLDEASRKIDEGQWRSQTFCCAAN*-	5.6e-27*
Zygentoma	Tricholepidion gertschi		EEGDVMMESLHALLEMFOLPLETLPLIYVVLKDARSDVKEASNRIIEAQSELRSIF	3.3e-23
Zygentoma	Thermobia domestica	_	ENVELLKDSLHALLDMFRLPLETLPLIYVVLKDARSDVKEASNRIMEGVYQCMNQGIIAPIG	2.4e-22
Odonata	Calopteryx splendens	_	DNMNFIKDAILALLOMFRLPVETLPLIYAILOGARCDVKEATNOIONAOEHLRAMALRMMYP	1.1e-19
Ephemeroptera	Isonychia bicolor	_	DNIDSLRDSIQTLLDTFRLPMETLPLVYVVLKVSRSDVQEAFNRISEGGYHMTSSDVPVAAA	1.2e-20
	-	_		2.0e-14
Plecoptera	Leuctra sp. AD2013 Perla marginata		DGHEVPFESIRKLLERFRLPSEAQPLLLAILKNSDYDSNEASKQIEIGGYNSSHPQGKAISV PYPDVHLGSIAKLIDMCHLSPETQPLILAILKLSKFDVKEAYKQIMAGGYFYNKNCTSN	2.0e-14 3.6e-10
Plecoptera				6.6e-20
Mantophasmatodea Grylloblattodea	Tanzaniophasma sp. AD2013 Galloisiana yuasai	_	ENMOGLREAIHTLLKIFQLPLETLPLLYVVLKDARSDFTEAYNRIVEARNELETLARREAAC	2.3e-15
	-		DHGKLLWESIQALREMFHLSQDSVPLIFLVLNFSHFDVMEASIRIKEGMFVIIEYLVGTCIV	
Phasmatodea	Aretaon asperrimus	_	VSVEAMMDGVYTLLHMFHYHVEMLPLLLVVLSDAHCDVSEAYNRILQVLGNHLDSVLPVTSS	4.3e-18
Phasmatodea	Ramulus artemis	_	VSVEAMMDGVYTLLHMFHYHVEMLPLLLVVLSDAHCDVTEAYNRILQGECAAA	1.5e-18
Blattodea	Blaberus atropos	_	ESLEVTLQSMHYLMNTFRMPLEALPLVYVVLQLSHSDVSEATARIIKGSYST	6.0e-16
Blattodea	Periplaneta americana	_	QMLLEMFRFPPVALPLIYVVLQVSQSDVNVAYNRIIQAQEQLRSMALREAAR	1.6e-15
Blattodea	Blatella germanica genome	_	ENQEVPLQSMQYLMDFFRMPMEALPLIYVVLQLSHSDVKEAATRIIKGSYSNVIFHI*	6.3e-06*
Thysanoptera	Frankliniella occidentalis	_	ENGRVHRESIQRLVEKFNFPPVALPLIYVVLKDHGSDYEEVKN	1.1e-12
Thysanoptera	Frankliniella cephalica	_	ENGRVHRESIQRLVEKFNFPPVALPLIYVVLKDHGSDYEEVKNLLLEAASELDETSYNQERR	6.3e-16
Thysanoptera	Thrips palmi	_	ENGRVHRESIQRLVEKFNFPPVALPLIYVVLKDHGSDYEEVKNLLLEAASELDQSGFNQERG	1.4e-14
Thysanoptera	Gynaikothrips ficorum	-	-NERVPWQSTMRLLQMFNFPENALPLIYVVLADHKADEDKVMALLEDGEF*	3.1e-07
Megaloptera	Sialis lutaria	_	HSLDVVLERCTRLLDTFMYPWEMMPLMYVILKYAGGNIDEAIRSIREGKRVVNEYSRIHNLN	8.4e-25
Megaloptera	Corydalinae sp. KMRSPBM-2012	_	-NIELILECCHRLLDTFLYPWELMPLMYVILKYAGGNLDEAIRSIREGKRVVNEYSRTHNLN	2.1e-24
Strepsiptera	Mengenilla moldrzyki	JP085896_4	NETERILDNSKKLLETFKFHSEMMPLMYVILKDARTDIRKATRRIDEGWQVVNEYSIKNNLN	5.7e-22*
Strepsiptera	Stylops melittae		FXQKLMEIYKYPYEMMPLMYAILKSVNVDFIEAQTRIDEGHQVVSDFLRKHNLN	3.0e-16
Coleoptera	Meloe violaceus		RGAENLLEFCQRLKDKFQLSWKMISLVDVILKYAKEDQEEAWRQIDEAFLEIRALAAVEAAR	2.3e-08
Coleoptera	Meloe violaceus		RGAENLLEFCQRLKDKFQLSWKMISLVDVILKYAKEDQEEAWRQIDEAFLEIRALAAVEAAR	2.4e-08
Coleoptera	Dendroctonus ponderosae	_	DRSMDLLEDCHKLLEKFHYPYEMMPLMYAILKDARADLDEASRRIDEGKQVVNEYSRTHNLN	1.9e-25
Coleoptera	Onthophagus nigriventris	-	IGQKDLIQESLQLLEKFRYSWEMMPLIYAIVKDTPD-LEEASKRIDEGKDAEQLVDFLIK*-	3.1e-14
Coleoptera	Onthophagus nigriventris	_	*RR*TTSGFFNKIKDRFHLSWKMISLIHVILKNAKDDQEKAFRQIDEAFLEVQALAKYYPTP	5.9e-05
Trichoptera	Hydroptila sp. AD2013	_	ALMESILENCYILLERFNYSFEMMPLIYAILKATHIDIDEASRRIDEGQTFYLYVKDVMTAA	1.3e-19
Trichoptera	Platycentropus radiatus	_	ASLETLLENCHKLLEKFHYSWEMMPLMYAILKDAHSDLEEASRRIDEGKWVINEYARQNNLN	9.9e-29
Lepidoptera	Dyseriocrania subpurpurella	_	TSLETLVENCQKLLKKFHFSWEMMPLVYAIVKHAQSNLDEASQRIEEALSTSIGHYERPLTT	2.2e-16
Lepidoptera	Dyseriocrania subpurpurella	_	TSLETLVENCQKLLKKFHFSWEMMPLVYAIVKHAQSNLDEASQRIEEGYHFSHYYHYTLLRE	7.1e-18
Lepidoptera	Polyommatus icarus		VSLESLVENCHKLLEKFHYSWEMMPLVLVILNYAGSNIDEASRKIDEDVKHVVCHES	3.4e-19
Lepidoptera	Yponomeuta evonymellus		LVENCNKLLEKFHYSWEMMPLVLVILNYAGSDLDEASRKIDEGKMRVDEYARKHNLN	1.4e-22
Lepidoptera	Empusa pennata	_	ENIAVSRESILELLDRFRLPWETLPLIYVVLDDARCDVNEASKRILKGCYAKANDKVIVNAI	2.1e-17
Siphonaptera	Ctenocephalides felis	_	EKVKALLEESQQIVEQLKYPFEMLPFVYAILGLSGG-ISEAIKNIHNGQLVVSEYSRMNNLN	1.6e-15
Mecoptera	Bittacus pilicornis	_	SEVNLLLDHCQKILEKFNYPFEMMPLIYAILKDAGD-IEEASRRIDEGFLVYERLREEIRIM	1.1e-20
Mecoptera	Boreus hyemalis		YSDKVVFDLSQKLLEKFNYPFEMMPLIYANLKYAGS-IDE	3.6e-10
Mecoptera	Boreus hyemalis	-	YFDKVVFDLSFKLLKKFNCPIEMMPLIYANLRYADN-IDEASRRIEEDILRITYGRYHLVNL	7.4e-10
Diptera	Bombylius major	· · · · · · · · · · · · · · · · · · ·	QKAERLLEYCQKLLEKFRYSWKMMPVMYAILKDARADVEEASRRIDEAIRVVRPLTLHLRYP	2.0e-20
Diptera	Bibio marci	_	VDVLLECSTKLLELFQYPWEMLPLMYVIVKDAGLNLEEAVRRIVEANNDIRALYFWKAVR	1.8e-19
Diptera	Bibio marci	GATJ01051986_1	LDDTAVVDHCQTILKRYDYPGEMLPLIYAIVQYCPN-LEEAIRRIDEGYRFYLLYTKLNSQV	3.7e-11

Atelura formicaria GAYJ01004010* Thermobia domestica GASN01031735* Tricholepidion gertschi GAS001256612* Ephemera danica GAUK01006643* Isonychia bicolor GAXA01111429* Baetis sp. AD-2013 GATU01002143* Atelura formicaria FCD0KP1ACXX 7 2202 17946 4233 61 Thermobia domestica GASN01345779 Tricholepidion gertschi GASO01035827 Isonychia bicolor GAXA01058460

ARERMLYESLVMMRQAFPVGEEAMPLLLCILKHSRS-VQEASLKIYQGHYDLNAKGFVGLDG Zygentoma ARERMLYESLLVMRQAFPVGEEAMALLLCILKHSRS-VQEASWKIYQGTYDLTSRGFVGLDN Zygentoma EREQMLYESLMVMRQAFPVGEEAMPLLLCILKHSRS-VQEASWKICQGTRDLSARGIIGIEN Zygentoma TRERMLYESLMQLRQLYPVPEASLPLLLCILKSSRS-IDEASAKIVQGTHELSTRGFPGIEA Ephemeroptera TRERMLYESLLELRQSFPVPEAGLPLLLCILKSSRS-VHEASAKIVQGTHDLTTSGLLGLDG Ephemeroptera QRKNLMYSSLLELRSHHPVPDAALPLLLCILNISNS-VKEASEKIQKGLQELSLAAASSNEL Ephemeroptera -----KESLQTLLDMFRLPMETLPLIYVVLKDARSDVKEASNRIIEGGYWIMTQMAA----Zygentoma ENVELLKDSLHALLDMFRLPLETLPLIYVVLKDARSDVKEASNRIMEGVYQCMNQGIIAPIG Zygentoma EEGDVMMESLHALLEMFQLPLETLPLIYVVLKDARSDVKEASNRIIEAQSELRSIF-----Zygentoma DNIDSLRDSIQTLLDTFRLPMETLPLVYVVLKVSRSDVQEAFNRISEGGYHMTSSDVPVAAA Ephemeroptera

* - high-confidence EST contig

¹ - transcriptome sequencing read obtained from SRA accession SRR921568

>Sipyloidea_sipylus_contig22122

>Lepismachilis_y-signata_contig_383881

Supplementary Figure Captions

Supplementary Figure S1. High-confidence *doublesex* transcripts reported in this study. Highly conserved amino acid positions are in red text. The conserved Lysine residue characteristic of the *dsx* DM domain (OD1, see Results and Discussion) is highlighted in yellow. Deviations from the canonical Cys-His zinc finger motif are highlighted in blue. Also included are OD2-encoding singleton domains recovered for taxa that possessed a high-confidence EST, to illustrate evidence for putative *dsx* paralogy.

Supplementary Figure S2. All type-A OD1-encoding singletons recovered in our analyses. Highly conserved amino acid positions are in red text. The conserved Lysine residue characteristic of the Type-A *dsx* DM domain (OD1, see Results and Discussion) is highlighted in yellow. E-values were calculated against our profile HMM using HMMer v3.1; asterisks indicate e-value was calculated via NCBI Conserved Domain Database search.

Supplementary Figure S3. All type-B OD1-encoding singletons recovered in our analyses. Highly conserved amino acid positions are in red text. The conserved Isoleucine residue characteristic of the Type-B *dsx*-lik*e* DM domain (OD1, see Results and Discussion) is highlighted in yellow. E-values were calculated against our profile HMM using HMMer v3.1; asterisks indicate e-value was calculated via NCBI Conserved Domain Database search.

Supplementary Figure S4. All dimerization (OD2) domain-encoding singletons recovered in our analyses. Highly conserved amino acid positions are in red text. E-values were calculated against our profile HMM using HMMer v3.1; asterisks indicate e-value was calculated via NCBI Conserved Domain Database search.

Supplementary Figure S5. Alignment of both putative *doublesex* transcripts from each of the Zygentoma and Ephemeroptera species reported in this study. The high-confidence ESTs (black text) share similarity with each other, while the alternate motif (red text) segregates.

Supplementary Figure S6. De-novo assembled contigs referenced in this study.