

Appendix A Species and protein sequences used in this study.

| Protein name | Species | Accession number |
|--------------|--------------------------------|------------------|
| Drosha | <i>Pteromalus puparum</i> | PpDrosha* |
| | <i>Nasonia vitripennis</i> | XP_031778770.1 |
| | <i>Ceratosolen solmsi</i> | XP_011505854.1 |
| | <i>Copidosoma floridanum</i> | XP_014213264.1 |
| | <i>Apis mellifera</i> | XP_016766928.1 |
| | <i>Bombus terrestris</i> | XP_003394274.1 |
| | <i>Solenopsis invicta</i> | XP_025986293.1 |
| | <i>Tribolium castaneum</i> | XP_008199088.1 |
| | <i>Agrilus planipennis</i> | XP_018332033.1 |
| | <i>Drosophila melanogaster</i> | NP_477436.1 |
| | <i>Aedes aegypti</i> | XP_001653338.1 |
| | <i>Bombyx mori</i> | XP_012547603.1 |
| | <i>Papilio xuthus</i> | XP_013172751.1 |
| | <i>Tetranychus urticae</i> | XP_015787469.1 |
| Pasha | <i>Pteromalus puparum</i> | PpPasha* |
| | <i>Nasonia vitripennis</i> | XP_003424309.3 |
| | <i>Ceratosolen solmsi</i> | XP_011498331.1 |
| | <i>Copidosoma floridanum</i> | XP_014204775.1 |
| | <i>Apis mellifera</i> | XP_006559676.1 |
| | <i>Bombus terrestris</i> | XP_003397039.1 |
| | <i>Solenopsis invicta</i> | XP_011166887.1 |
| | <i>Tribolium castaneum</i> | XP_015836202.1 |
| | <i>Agrilus planipennis</i> | XP_018333701.1 |
| | <i>Drosophila melanogaster</i> | NP_001263150.1 |
| | <i>Aedes aegypti</i> | XP_021710923.1 |
| | <i>Bombyx mori</i> | XP_021208180.1 |
| | <i>Papilio xuthus</i> | XP_013167918.1 |
| | <i>Tetranychus urticae</i> | XP_015793480.1 |
| Exportin-5 | <i>Pteromalus puparum</i> | PpExp-5* |
| | <i>Nasonia vitripennis</i> | XP_008203379.1 |
| | <i>Ceratosolen solmsi</i> | XP_011499457.1 |
| | <i>Copidosoma floridanum</i> | XP_014204789.1 |
| | <i>Apis mellifera</i> | XP_396789.2 |
| | <i>Bombus terrestris</i> | XP_003398965.1 |
| | <i>Solenopsis invicta</i> | XP_011163732.1 |
| | <i>Tribolium castaneum</i> | XP_974696.2 |
| | <i>Agrilus planipennis</i> | XP_018329940.1 |
| | <i>Drosophila melanogaster</i> | NP_608339.2 |
| | <i>Aedes aegypti</i> | XP_001660023.2 |
| | <i>Bombyx mori</i> | XP_021204862.1 |
| | <i>Papilio xuthus</i> | XP_013174923.1 |
| | <i>Tetranychus urticae</i> | XP_015794798.1 |

| | | |
|----------------------------|--------------------------------|---------------------------|
| Loquacious | <i>Pteromalus puparum</i> | PpLoq* |
| | <i>Nasonia vitripennis</i> | XP_016839605.1 |
| | <i>Ceratosolen solmsi</i> | XP_011500654.1 |
| | <i>Copidosoma floridanum</i> | XP_014211571.1 |
| | <i>Apis mellifera</i> | AmLoq* |
| | <i>Bombus terrestris</i> | XP_012169164.1 |
| | <i>Solenopsis invicta</i> | XP_025989733.1 |
| | <i>Tribolium castaneum</i> | XP_008198430.1 |
| | <i>Agrilus planipennis</i> | XP_025832115.1 |
| | <i>Drosophila melanogaster</i> | NP_001285894.1 |
| | <i>Aedes aegypti</i> | XP_021701497.1 |
| | <i>Bombyx mori</i> | XP_012550850.1 |
| | <i>Papilio xuthus</i> | XP_013164367.1 |
| | <i>Tetranychus urticae</i> | XP_025017170.1 |
| Dcier-1 | <i>Pteromalus puparum</i> | PpDcier-1* |
| | <i>Nasonia vitripennis</i> | XP_001605287.1 |
| | <i>Ceratosolen solmsi</i> | XP_011502152.1 |
| | <i>Copidosoma floridanum</i> | XP_014210593.1 |
| | <i>Apis mellifera</i> | NP_001116485.2 |
| | <i>Bombus terrestris</i> | XP_020723074.1 |
| | <i>Solenopsis invicta</i> | XP_025991666.1 |
| | <i>Tribolium castaneum</i> | XP_008199045.1 |
| | <i>Agrilus planipennis</i> | XP_018328602.1 |
| | <i>Drosophila melanogaster</i> | NP_524453.1 |
| | <i>Aedes aegypti</i> | XP_001659747.2 |
| | <i>Bombyx mori</i> | KWMTBOMO06302† |
| | <i>Papilio xuthus</i> | XP_013163421.1 |
| | <i>Caenorhabditis elegans</i> | NP_498761.2 |
| Ago-1 | <i>Pteromalus puparum</i> | PpAgo-1* |
| | <i>Nasonia vitripennis</i> | XP_031783914.1 |
| | <i>Ceratosolen solmsi</i> | XP_011505420.1 |
| | <i>Copidosoma floridanum</i> | XP_014207835.1 |
| | <i>Apis mellifera</i> | XP_006571833.1 |
| | <i>Bombus terrestris</i> | XP_012170889.1 |
| | <i>Solenopsis invicta</i> | XP_025990461.1 |
| | <i>Tribolium castaneum</i> | XP_008196652.1 |
| | <i>Agrilus planipennis</i> | XP_018333020.1 |
| | <i>Drosophila melanogaster</i> | NP_001246314.1 |
| | <i>Aedes aegypti</i> | XP_021704825.1 |
| | <i>Bombyx mori</i> | NP_001095931.1 |
| | <i>Papilio xuthus</i> | XP_013176493.1 |
| | Dcier-2 | <i>Pteromalus puparum</i> |
| <i>Nasonia vitripennis</i> | | XP_031782799.1 |
| <i>Ceratosolen solmsi</i> | | XP_011494459.1 |

| | | |
|-------|--------------------------------|----------------|
| | <i>Copidosoma floridanum</i> | XP_023246881.1 |
| | <i>Apis mellifera</i> | XP_016773223.2 |
| | <i>Bombus terrestris</i> | XP_012163127.1 |
| | <i>Solenopsis invicta</i> | XP_025991526.1 |
| | <i>Tribolium castaneum</i> | XP_008201496.1 |
| | <i>Agrilus planipennis</i> | XP_018331371.1 |
| | <i>Drosophila melanogaster</i> | NP_523778.2 |
| | <i>Aedes aegypti</i> | XP_001652212.1 |
| | <i>Bombyx mori</i> | XP_021207858.1 |
| | <i>Papilio xuthus</i> | XP_013173437.1 |
| | <i>Caenorhabditis elegans</i> | NP_498761.2 |
| Ago-2 | <i>Pteromalus puparum</i> | PpAgo-2a* |
| | | PpAgo-2b* |
| | | PpAgo-2c* |
| | <i>Nasonia vitripennis</i> | XP_032455160.1 |
| | | XP_031786529.1 |
| | | XP_031786440.1 |
| | <i>Ceratosolen solmsi</i> | XP_011505420.1 |
| | <i>Copidosoma floridanum</i> | XP_014218615.1 |
| | | XP_023245295.1 |
| | <i>Apis mellifera</i> | XP_395048.4 |
| | <i>Bombus terrestris</i> | XP_012168271.1 |
| | <i>Solenopsis invicta</i> | XP_025997806.1 |
| | <i>Tribolium castaneum</i> | NP_001107842.1 |
| | | XP_008192985.1 |
| | <i>Agrilus planipennis</i> | XP_018319532.1 |
| | <i>Drosophila melanogaster</i> | AAO39550.1 |
| | <i>Aedes aegypti</i> | XP_011493002.2 |
| | <i>Bombyx mori</i> | XP_021205714.1 |
| | <i>Papilio xuthus</i> | XP_013179888.1 |
| R2D2 | <i>Pteromalus puparum</i> | PpR2D2-1* |
| | | PpR2D2-2* |
| | <i>Nasonia vitripennis</i> | XP_031785356.1 |
| | | XP_008214050.1 |
| | <i>Ceratosolen solmsi</i> | XP_011504752.1 |
| | <i>Copidosoma floridanum</i> | XP_014215309.1 |
| | | XP_014205151.1 |
| | <i>Apis mellifera</i> | XP_006560091.1 |
| | <i>Bombus terrestris</i> | XP_003395928.1 |
| | <i>Solenopsis invicta</i> | XP_011156338.1 |
| | | XP_011157520.1 |
| | <i>Tribolium castaneum</i> | XP_015835139.1 |
| | | NP_001128425.1 |
| | <i>Agrilus planipennis</i> | XP_018328507.1 |

| | | |
|-------|--------------------------------|----------------|
| | | XP_018330227.1 |
| | <i>Drosophila melanogaster</i> | NP_001162903.1 |
| | <i>Aedes aegypti</i> | XP_001655660.2 |
| | <i>Bombyx mori</i> | NP_001182007.1 |
| | <i>Papilio xuthus</i> | XP_013174949.1 |
| | <i>Tetranychus urticae</i> | XP_025017170.1 |
| Sid1 | <i>Pteromalus puparum</i> | PpSid1* |
| | <i>Nasonia vitripennis</i> | XP_031783168.1 |
| | <i>Ceratosolen solmsi</i> | XP_011495242.1 |
| | <i>Copidosoma floridanum</i> | XP_014207455.1 |
| | <i>Apis mellifera</i> | XP_006565236.1 |
| | <i>Bombus terrestris</i> | XP_012170554.1 |
| | <i>Solenopsis invicta</i> | XP_025995259.1 |
| | <i>Tribolium castaneum</i> | NP_001099012.1 |
| | | XP_015837738.1 |
| | | NP_001099128.1 |
| | <i>Agrilus planipennis</i> | XP_018319198.1 |
| | <i>Bombyx mori</i> | XP_004930736.1 |
| | | NP_001106735.1 |
| | <i>Papilio xuthus</i> | XP_013177205.1 |
| | | XP_013177321.1 |
| | <i>Penaeus vannamei</i> | ROT74464.1 |
| Ago-3 | <i>Pteromalus puparum</i> | PpAgo-3 |
| | <i>Drosophila melanogaster</i> | NP_001036627.2 |
| Piwi | <i>Pteromalus puparum</i> | Pp-Piwi-like1 |
| | | Pp-Piwi-like2 |
| | | Pp-Piwi-like3 |
| | | Pp-Piwi-like4 |
| | <i>Drosophila melanogaster</i> | AAD08705.1 |
| Aub | <i>Drosophila melanogaster</i> | NP_476734.1 |

*: Newly identified proteins in this study; †: Protein from silkbase;

Appendix B Sequences of the newly identified proteins in this study.

>PpDrosha

MDNVPLDTEYQYYWQADSSQAQNCOPYPNSSVSYPPSALPSQPPYNAPPPDYNGFNYSVPPPSYFVPPP
NSSIAYPAGTVTYSSYEYQSHTSVQENKDYTTSYAQAQSSTYNYNWDPKCPYQQDSKTDNTQWVAE
TSSWLSHTAVNSKSNSSADKGYKSVSNGLRRSKSPDSKFHSSKFTSRRKYDDYSNRYGDRNKERKSSR
DRFYKRRSRSLDRKSYASNNSYTNHSSARSERSERLERSESRSRRSRISRSRSKRSSRSQESHTSLQTSRST
NNIAKRKSLSEREMLLEKYRQNYCATSEDIKRKLNELSDGVLERKIWTRTAPADLYYSRDEKNNKLMK
ATGRLIQLCDDFKILIDRASSARELQPPYEPPPRKNRARLCRHKSEACSSSSSDSDSDSDEDDRTMEELM
AKKQHPQRLHPEMWFNDPGEEMNDGPLCRCSAKSRRSGIRHGIYAGEGAIEKCDPHTNADKLYHYRITIS
PPTNFLTKPTTIKHDEHEFIFEGFSMFHSHFPLVKLPTCKVIRFNIEYTYLMEEKMPENFTIQELDYFQDYLF
KEILELVDVFDLQAAQDKKGAQFHFMPRFVVDLPDNGQEILSMNEVLSYLIKSTLLIDPDELPLKVEMPO
FQWQNFADDEVKGMVVVTPGKKPCSVRVDQLDRNQADQPPGVIAYPEIVHFGIRPPQLSYAGNTDYQRA
WRDYVKFRHLLANMPKPSFEDKRKLEAKENKLQELRTQSKMKRDVTVDVTSEGFYRTGIMCDIVQHAM
LIPVLVCHLRFHKSLEDMLEQTLNYKFNRYLLQLALTHPSYRENFGTNPDHARNSTNCGIRQPEYGDRI
HYMNRKRGINTLINIMSRFGAKKETESSIAHNERLEFLGDAVVEFLTSIHLFHMFPDLEEGGLATYRAAIV
QNQHLLAVLAKKLNLEQYMLYAHGSDLCHDELRLHAMANCFEALMGALFLDGGIRIADTVFGETLFFKKE
LDLLNIWVNYPKHPLQEQEPTGDRQWIPSEFELLQKLTKEEGIGVEFSHIRLLARAFTDRSIGYTNLTLGSN
QRLEFLGDTVLQLIVSEYLYKFFPEHHEGHLSSLRSSLVNNKTQAVVCCDLGMTQYALYGNPKAELKTK
DRADLLEAFLGALYVDKGLLYCQVFCNVCFPRQLQDFIMNQDWNPKSKLQQCCLTLRTMDGGEPDIPV
YKVIIECKGPTNTRVYTVAVYFQKRLAKASGHISQEAEMNSAKEALEKSQDLFPQLDHQKRVIAKSMKM
QSWSQGGLKFRTIQSSKFDDRSSEDRNSIRYEDRRHDGRNEGRRRDYKYEDRQRDRNRHSSSDSDSHRKR
RSRSRERAKRSSRSRDRYDSVDRRLRDKPRSSSCSSSNSSNSNARSIRNKSAERHSSSDSTSNCTENIAAA
RGDNGEDKELKGANRKRSSVLSNASSNDASSNNISETDKKCVLPFSEKMKRLE

>PpPasha

MEVDVQKSDVVVEKKSEDETSEEMLTQESMDTCTTVLQEVQPSQSKETENPLISEDATDDDLRQFDVL
DDLERHTNQDGSDDGCSNDSMSDIPDDEIEAMLEEGLPDEFKGRSDKRSMDLYEEKEKLVLDEIGHN
HFDVLPQGWVQVTHNSGMPLYLHKQSRVCTLSKPYFLGPGSVRKHEVPVSAIPCLQYKRALKEEEEEKQ
RDKERMQAQAACSLPNAKIETIQENRATQSLDGENLRNYCQSLFRFKSIKVMRFKSWQRRKFTKNNKQ
RKQLERPTLPDGTKLITFPVGNNAVYSGNSTQEDESASRPPKHWIMNPSGKSYVCILHEYVQHALKKQPT
YKFKELENAATPYSAVVCINDMEYGSFGSSKKQAKANAARKTLEILIPQMRDKISGDTTDDNNSANSSQ
TIKASKVNSDADLSFFDEISINDPRVAEFCAKTTEPLPHAILITCLQRNFGLGDMEINYSVNTLKHQRNEFT
MRVGKHEATVICRNKKGKQRAAQAILQRLPHIHSWGSLLRLYGRSVKSFKEKKQLEQEITLLQGKA
AVNQPNHAILEKLRQEMRKLSDQRQAVKPIGKFIIPDMPTGSSSNLDNVDL

>PpExp-5

MEFGVGGDIAQISAELARLVELMSPNVSQQQRVEVLNACERFKESSPLCAQCGLFLAQKASNRSSIVRH
FGLQLMEHCVKYRWTQMSQTEKVFIKENAMKLLQEGTEPLLQEENHIKDALSRIVEMIKREWPPQWPQ
LLGELSQACTRGETQTELVLLVFLRLVEDVAILETLESNQRKDIYQALVTNMKEIFAFFLRLMELHFAEF
QKQSTLGRTEAAAHGRVQVVLTLVSGFVEWVPIHIMEDKGRLLQILCILLSHPTFRCPSECLQVQV
RKGKVEDRKQLMILFTEALGYMYAAATAAPPTGPSELQENHYLFLKLVQVLNGMATQLCSLWTKDD
ANGVRPVHFGFLDAVLTFTMHSSLTLVQLANTIWIMFFKHEYIKTDSLVLVSYIPKYVEYSGPKIVKVAFP
NKRHANGMASYFVDYDTEEEFRMFFHRLRMDLLEGFRNATMVAPLVTFAYVQWLVTKIRKGSNLAY
KSDPLDPEYLEWEALALALDSVMTRILLVSRPNVQTGLQLELCLGYSPQDPWILSALLSCISALFVMS
MSTGSMAMPGVAILPRVLEKIFAALVFNGPGETKENRSKASKNRRHAASLMVKIGLYPLLLPVFEQI
HSTVKNLVREPSQVSRMESLALYEALLISNHFCDYERQTRFVAEIIIGDTSTRLIAIGAQAFKSPVDLVHFL

GLDRPALENNRDERTGSNRSMLYCINTVLSVVKRAIPEDPDRAARGGFVAALSESGNPIYRNPATPHII
PVLPLFDLLRTMNALFTPAAALGLLSEGYRNVHGIIESEKANLLGLHTSNNNENSADSELQSAPLARMQGF
LTTIHNDNCYHMLGSGCHTIGRDLYQLPGLAAALLNSVFSNMEVIPDHRLRPIIRVFMKPFYSCPSAFYETV
LVPVLAHVSTHMYQRLSAKWQHMSLSYEAGSLDDNDTDAQEVIDDMLNRNLTRDFVDVLKVALVGGGA
ASDASPPDSMEQDSGGMAVDSPPARGSNSIVAIEVSELGSFILRHSSTFQSVVLCILGALSWNDSNASLRA
TMLTGPVVRALAAAEGILNPGMAAHIMMAILQGLQLHGQHEANQGSLITLGAQVYECLRPKFPNIEVMQ
QIPGVNADLQRFDEKMSAVTTTTKGNKVEKGKKDLFKKITNQLIGNIGQLFRKEVKIANLPRLEVSSKEI
VRVDEISKNSHDTGLTALFAGPT

>PpLoq

MEDLRQQVGSNMMDGGIDELRQGVGNMMVGGVVPVPHANNVHRRSKTRVTMHALMSEALPLDEAA
RLEMKALPNKTPVSVLQELLSRRGTTPKYELVQIEGAIHEPTFRYRVTVADVAMGTGRSKKEAKHAAA
KAVLDKLGIVNTENAEAPLNSIPDENFPLSSQNIQEIQSGYGEEKVVNNPIGSLQEMCMSRHWPPPKYSM
EGEEGLPHERQFTIVCSILKYRQVQGKSKKLAKRQAAHKMWQSLQDTNSNRTQGVDEDEIVQRNANV
NAHYADLKDSKIATLTPHSHKVSQFHKNLSSTGVKLFELQTHLFQNTCLNDGDVNLVQFLQEIASEQQ
FEVTVVDIEEKSITGKCQCLVQLSTLPVAVCYGSGMTSKDAQAAAAQNALEYLKIMTKK

>PpDicer-1

MAFFFNDNIHTKSFTPREHQVELLYSAKEKNIIICLGKITEQTFIVTKIIQELAINNRKTLSDNGKRTVYLLE
NEQACAIEKANHIEQLTDLKVLQCHNVNDFDSEEFKNTQVLILTVDMCSRLLMANKINPRQINLLIIDKCH
NLVTNHELSSILNIFRSCDNTTKIIGFAVPLYSLTKEPGALSIEIERIESLLQCQIETASDLLSILRYSPKPKY
LLAYKMEDEKCVQNDVKEFTYSTLNFLFDHRYDPTEIYSEEFLEDIKKIPNPVAKPCMIYEFLHILNTLGL
WAADRAALVLLMLIEKLIKIPYERHYLLFGVVALFLKIRAYCDDIFSGLTEAEKILKFSTPKIHRILIEVIK
TFSPPPKEEKEENVTVNNDVEKSDSANSKDQNHDKSVKNFHKKTDGGFKLRFRHSRGLTDPDLLCGVI
FVDNAVTAKVLFYLLNEMSKCDESLQFLSPLYTSEKTNDELFCGRDLELEYKKQEEVLKFKFRIHECNLLIS
TAILEEGIDIPKCNFVMRFEFPKNYQSYVQCKSRARATDALHVLMPVEIESEVYIQRLAHYHYIEKILLSKC
TSKEADECEETEADAYDSSLPEYKPLNEADAPKVTLNSAISLVNRYCAKLPSTFTFTKLTPEWKINNITVEN
KQMYICSLRLPINSPLKYSISSYPMPNKALAKRMAALQLCVLHKENEIDDYLLPVGKENFKAHPEDSEVP
TLPDENMNLTEARPGTTKRRQYKYKIADALIDCKPELEKPCYLHINMVLSCPLPEEQNTRGRKIYPPE
ESDIGFILTRKKIPNVCPPFIYTRSGEVRVDLKLKSKETVVLDESKIDKVVFLNYTFTNVLRQLKYLMLFD
PNVSENSYIIVPVKKQPDASDICVDWDFVECIYANRDSMPNCVSEKNRQNFVFEANKYHDAVIMPWYRN
QDQPQYFYVAEICNHLNPKSSFPGADYSTFEEYYSKKYNIQIQNLEQPLLDVDHTSARLNFLTPRYVNRKG
VALPTSSEETKRAKRENLEQKQILIGELCAIHFPASLWRQAVCLPCILYRINALLLADQIRRHVAQSISLGQ
ETLDEAFDWPPLDFGWSLADVLLKKSKEIEKTNSNKPEKEKSEKIKEETPAVKEKPERVKEEPLDDSGNCE
MEIGTWSNEMVNMAPMGGADADLAVQSNYEWGNIRYGSPTYDSEIDNSLENSCTDDDLGDDSDDDSDQD
DDEEGLTIA YTGDNV AEAFDKKGKAFEAKEKEKQSCVERQEKSSELMHWNFKGDLSDSEFYQLHRLAHIDK
AKMNEEILSQDMFIPCDKSVAFKRKTSKVLNNSSEVERRHESYVSVSNYFKHELIESGKKRCNNQGAEL
TKLVPKYNGSFSFDYQPELNGHSGPSPSLILQALTMSNANDGINLERLETIGDSFLKYAITTYLFCYNNI
HEGKLSHLRSKQVSNLNLRYLGRQKMLGESMIASKFEPHDNWLPPCYVVPKELEQALIESGVPSAMWNQ
ADIPPSRFSEINDLVRETEHKLEIISSELSQSDHSPLNGPNGLTGDPLRSFIPYNLITQHSIPDKSIADCVEALI
GAYLIACGPRGALLFMSWLGIHVLPSSEVTVISDSKPTERPPGSTPYVEREEEDGRTKWTQLRYKKLQEP
SPLFYNIPEPELELEIMLDGYDTLERSIGYTFRNRSYLLQAFTHASYQPNRLTDCYQRLEFLGDAVLDY
LITRHL YEDPRQHSPGALTDLRSALVNNTIFASLAVRCGFHXYFRHLSPLSTVIDRFVRIQEENGHSISEEY
YLLIGEKEFEEAEEAEDVEVPKALGDVFESLAGAIYLDNMSLDAVWAVYFEIMQSEIEQFSANVPKSPIRELL
ELEPETAKFGKPEKLADGRRVRVTVDVFGKGSFKGIGRNYRIAKCTAAKCALKKLKKSKQSCIKGRK

>PpDicer-2

MDQTPENEFKARPYQIYLYEKTIEQNSILYLPTGSGKTYIAVLLVKRLSGDVQRSYTEGGKRTIFIVNTVAL
VVQQTAFLSRHTGLVCKGYSGDMGVDFWSNTEWRNEIDTNQVLVMTAQIFLNLLIHGCMSLDKINLLIF
DECHRAVKDHPMRQIMQRFQDHPKNKLPRVLAMSASLLNANVPFGKIETTLRELEVTFQAKIITVESLAY
VTDYATNPKEYVEYYDTPQNISVLSEISSIVEYASSILKYVDLPNRMENPASSAIFKPVSKTIKLNRLSDVE
EHLTEMGLYGGSKSVLQHIIQLECLKRFSDEKQTIAIFDFIITQLVKIQKLLSDAMDKVSLPENIHRFSSSKV
LKLLEVLKTFYNNMENKTNFCCIIFVKRRFTAKVLYHLLLKLSSSDKNFEFLKPQYVMVGYNNDPYKNARE
TLCISKWNKEVLKFRNGTANCVVATDVVDEGVDIPSCTLVVRYAPMDFRAYIQSKGRARHSTSHFIILA
SSQDDYINRYRSFQHTEQFLRKALHGKSDNRTEPSKDDVDTLTYQYVIEPYIVTDANGTKSVITEQS AISLI
NEYCANLVKSKFIMLTPTWVKEDVNPSAYRVLLTLPISPLRDTIIGDVMQNVDAIKRSAALKACIELHKI
GELNDNLKPRKPEDIELHTKHLFPYYVADKDTTDGIPGTNTKKRHHELIYPEALCSAFPKNESLYLHIIDM
KPNYPRPNDNRHLVFYNLLSNSQTYGILSAKKIPEIPSPFIFMNIQDLQINMKSNNQLKSLTKAEIEQLKLFH
TLVFHHILKVIKDFMIFDTSNMDNNFLVVPINEKEQINWDIVNR YKTITSSRSENLVVKESEYDLALVTPYY
RASNMYIVTQVCEHLKAESSFPTSDYNSYVHYFRERHYIEIKNPSQPMLEVKPISSKINCIKPRSIKANLSKR
KRASLTEDFEEHLVPELCERIDFPSLYWLKATTLPSILHRISQLIAAEELRVKIAHEAKLDISSLETGKKWEP
MRIVDQFSQNSSESNDTTFDESLMDEEDPEAENVLVIDTSLEVDILSQERSLYSWSKEQEPTDLDRNAEQI
QLIDIEYYCQFMSGVTPEDNKNLKYQNAVKS VYVSPTNIPAPRLKMLTSSNPYGPSPVDILQSLTTKAGND
VFNLERVETLGDSFLKFAISLYLYQAYSTCGEGPLTYLKGKLVGNLNFYCSKQKNIAGR MHVEDFAPLS
NFVTPAYAAHRVLQQLRAEKVSANILYEIRVPAERFSGCISNSTTDMMQEKVLAWPSDNKVAHTGME
HFLGTQVVSDKSVSDCTEALIGTYLLHLGIK GALQIVKWFELPKSLDVNQYL YSEVQNPQLGDGDVNVH
MPWAQTIEERLGYKFRNRAFLQAFTHPSYMPNRQTASYQRLEFLGDAVLDFVLTVHIYETCGNLSPGEL
TDLRSALVNNITFACLA VRYGLHTALLALAPKLFDLVDRFVKFQEDRNYKIDDELLWVLEEEECNMAE
HVDVPKVLGDIYESVIGAVFLDSGKNFEVWVQVIYNLMKNEIDLFSKNVPKQPIRVIHETQGANPKFSSSQ
KVEKSSITMITLTVNVNGKKKFFHGFPGPTKKLAKCAASKQALKYLRYKN

>PpAgo-1

MYPVAPPPPPGPTTSTAVGATGNTGSNVVAPNSMGLVPTQQTHTPPQPPDLPMFACPRRPNIGREGRVIGL
RANHFQISMPRGFVHHYDINIQPKCPRKVNREIIE TMVHAYS KIFGTLKPVFDGRNNLYTRDPLPIGTEKL
ELEVILPGEGKDRVFRVVIKWWAQVSLYALEEALEGRTRQIPYDAILALDVVMRHLPSMTYTPVGRSFFSS
PDGYYHPLGGGREVWFGFHQSVRPSQWKMLNIDVSATAFYKAQPVIEFMCEVLDLRDINDQRKPLTDS
QRVKFTKEIKGLKIEITHCGTMRRKYRVCNVTRKPAQM QSFP LQLENGQTVECTVSKYFLDKYKMKLRY
PHLPCLQVGQE HKHTYLPLEVCNIVAGQRCIKKLTDMQTSTMIKATARSAPDREREINNLVRRADFNDS
YVQEFGLTISNNMMEVRGRILPPPKLQYGGRVSSLSGQTKQALPNQGVWDMRGKQFFTGV EIRVWAIA
CFAPQRTVREDALRAFTSQLQKISNDAGMPIIQPCFKYATGPDQVEPMFRYLKSTFQALQLVCVVLPG
KTPVYAEVKRVGD TLLGMATQCVQAKNVNKTSPQTL SNLCLKINVKLGGINSILVPSIRPKVFNEPVIFLG
ADVTHPPAGDNKKPSIAAVVGSMDAHP SRYAATVRVQQHRQEIIQELSSMVRELLIMFYKSTGGYKPHRI
ILYRDAVSEGGFLHVLQHELTAIREACIKLEADYRPGITFIVVQKRHHTRLFCADKKEQSGKSGNIPAGTT
VDVCITHPTEFD FYLCSHQGIQGTSRPSHYHVLWDDNHFESDELQCLTYQLCHTYVRCTRSVSIPAPAYY
AHLVAFRARYHLVEKEHDSGEGSHQSGCEDRTPGAMARAIVHADTKRVMYFA

>PpAgo-2a

MSKREKIHNWYKKLTSRFKKTKEAQDQAGPSQSQQEPERSQQSEPQSTDVAQETLLISASLEKGTCDQ
ADPSQSQISESQSTDVAPETT VIISASLEKDASDQAGPSQSQLEPGQFQQAQPSADVDQETSEFLATMSLE
EQSEDNSSKSKKKT PKLTVSADIANREVEQIPKRKDPK KAGTKGRPITVNTNMMAINVNQMNPNVVHYD
VGIVPNTPKYLMRPVFNEVKKTLFPERNPAFDGSRNAFSAGELPIEDPTTTEVVVYNEDGRVKKYKVTMK
VVNRIDLSWLKTFAPGSEDSRINNQVSIQALDVLK SAPALTSISVKN SFFTPPKGQVMSLGGGMDLWVGL
FQSAVLGWKPYFNV DV AHKPFKPKQSVL DLMKTICGCAQDQDQGGQQYGRQQGYGHRGGYGQQRGRH

GGYGQRGGYGQRGGYGQRGYGHQGGYGHQQAPANLSADLIYRYREDIRRYLKGLKVTIEIPGQPTTRRI
QRVNDLVICPRDNVFEHDGHRITVEQYYQLEKRYTIKHPDFPCLWVGGRDKNIHVPEICTVVGQAVQK
KLTDRTSSMIRSTASRPEDRKKKIMDAFNSIRHNEDPCMREFGISVSGEFETVPARVLNPPALTYQNQNV
RVAKGVWRASQFIQPSDLIADDNTWTVLNLNDRTRDDGLYRLVDTFKKIGKNVGMVGNPLSPFRSMQ
VRGQDTRDLMAYFNEMKTKEVKLVVVVVPEMKGYPYSKVKQMSSELRVGVLVLTQCLKSKTLFKLNDATAS
NILKINAKLNGTNHVFEHVSRPPCLKRPCMIVGADVTHPSPDATNIPSIAAVVASHDPNAFKYNVEMRL
QPPRQEIIGELAEIMKIQLEYFYTSTTQKPEIIFYRDGVSEGFQGMHSELLAIRKACQSLEDGYEPQITF
LVVQKRHHTRLFPTDPSNSDDKHGNVQAGTIVDTEITHPSHIDFYLVSHASIQGTARPTKYRCLWDDSDM
SEDEIENLTYFLCHMF SRCTRSVSYPTPTYAHLAAFRARALIQDVIDMNNLPQEQLRKLTIKDEV LKGC
PMFFV

>PpAgo-2b

MVKKKEERTKVEVKAEARTEVEVLEIQLRDHSRMFLIINNDHCRNKPLLGS DCHKLLNNKVF GINDSHNG
NSKNLLAADVVQETAKLSAVSLEKQSKDNSSKSEKTPKLTVSADTANKEVDQIPKRKDPK KAGTKGRPI
TVKTNMMAINFKQMNSDVVHYDVIDVPNTPKYLMRPVFLEAKKRLFPNKNPAFDGSKNAFSAGELPIKD
LSTTDVVVYNEDGREKKYTVTFKIVNRIDLSWLKTFEPGSQDSPRNQQISIQVLDVILRNAPSALPKSISVK
SSFFIPKQGVDP LGGGIDLWNGLFQSAVLGWKPYFNVDVAHKGFPPQSVLDMKTICGCDGQDQGG
PQYDRQRGYGRQGGYGQQGPTILSADLIHRHREEIAKFLKGLKVIIEIPGQPTSRR TQRVNDLVICPRDN
VFERDDGHRITVEQYYQLEKRYTIKHPDFPCLWVGGRDKNIHVPEICTIVGEQLIQK LDEKQTASMIKFA
AKGPEERKKKIMDGFNSMRYNQDPCMKEFGISISGEFETVQARVLEPPQLKYQRQNV RVANGFWTASEF
LKPSQSILDNKTW TILNLDNR TKDDR LYKLV DILKRMAISVSLPIGEPFTPF TKIQLKGHDIRGLMEYFNKM
KMNKIKLVVVV PDMKGPYSKVKQISELKVGVLTQCLKSKTLIYNLNDKTAGNILLKINAKLNGTNHVF
EQQKSLPACLKSQC MIVGADVTHPSPSK GIPSIAAVVASHDLNAFNYNVEIRLQPPNTEYIEELAEIMKIQ
IKFFYEHRTRQY PKKIIVFRDGVSESYFGQIMHIELQAIRKACQSIQADGLYKPQITFLVQKRHHTRLFPTDL
RNSDDKHGNVQAGTIVDTNITHPSHIDFYLVSHASIQGTARPTKYRCLWDDSHMSEDEIENLTYFLCHMF
SRCTRSVSYPTPTYAHLAAFRARALIQDVIDMNNLPQEQLK KMTIQEDILKRNPMFFV

>PpAgo-2c

MKGK KKKKQSGEDPQSASTSGGSQPNVPPQQQSAPQPQQSQQSAPPQQYVQDNAGS GRGKGQGRG
GGGGRVRS PETKAADV KQTTEQLSELSLKP KAGTPKSEKVPILTVPASIANIQAEKIPKRKDPK KAGTKGR
PISVKTNMLAINVQH IKSDCVHYDVIDVPNTPKYLMRPVFLEAKK LFPHRNPAFDGKKN AFSAGELPIK
DPATAEVVVYNEDGREKKYVTI KIANRIDLSWLKTVKPLD DNERNQISLQALDIIMRNAPALTSTP VGR
SFFTPPKGHVMTLGGGMDLWVGLFQSAVLGWKPF LNVDVAHKGFPPQSVLDMKTICRCDAQDQGG
RPQYGRQGGYGQQG PANLTADLIHR YREDIRKFLKGLKVTIEIPGQPTTRRTQRVNDLVKAPRDNQFE
YNGKTITVEQYYKHEKKYTIKHPDFPCLWVGKDKNVHVPEICTIVGGQATQKKLDETQTSSMIRYAAT
GTEDRKRKIMDAFN SMRHNQDPCMREFGVS VSGEFETVPARVLDPPQLTYQRQNV RVAKGVWRASQFI
KPSDLIEEDNTWTVLNLNDR TKDDGLYRLVDTLQKTGKTVGMFIGQPLSPFRSMQLRGPDTRELMGYFN
EMKTKNVKLVVVVVPEMKGYPYSKVKQMSSELRVGVLVLTQCLKSKTLFKLNDATAGNILLKVN AKLNGTN
HIFERNVSRPPCLGTPCMIVGADVTHPSPDATGIPSIAAVAASHDPNAFKYNVEIRLQPPKQEIIGELAEIMT
IQLKYFYNQTRRKPEIIFYRDGVSEGFQGMHAE LLAIRKACKTLEAGYEPKITFLVVQKRHHIRL FPTD
PRNSDDRNFN VQAGTIVDTEITHPSHIDFYLVSHASIQGTARPTKYRCLWDDSNMSEDDIENLTYFLCHMF
SRCTRSVSYPTPTYAHLAAFRARALTQDVIDMNNLPQEQLRKLTIKDEV LKGS PMFFV

>PpAgo3

MLARYHRQKTCSAGVAGQRLTLAPVVRR TGCLRRQARHRPFGCFPLHIGH RHKTKMGGKGAGRGNLV
EIIKKKMEEEKRLKEEQERARQLQELQQKQLEEKLQEEAAA VRQPSPTAGVGRGRGS LLEKLRSKMTPG
VDTSTPIPSTTASITSMQTTSTATSGRGALLS AMKKRSESAADSSSPVSTTASVTSMQTTSTTTGRGSVL

GAIKKRTGSDTLETSGEQTGVVGLSSLSLEETKSSLTEEEIVSRQGSSGKPISLTANYINLKVHKDKGLYQ
YEVKFSPPDVSRSRLRYKLLNQHLNDLGNVKNFDGGVLYLPIKLNQNRRTTFDSIHPIDNSTVCLTVIFQKKQ
AMKDNIFFNSLMNRVMKALTLVKIGRHDFNPTCAHQIPQHRLEVWPGYVTAINELEGLKLNLDATHR
VMRLDTRVRLMTDMYHKNPNNFKNAVANEIIGTSVLTRYNNRRTYKIDDIDWEKNPMLKFQRKDEEVSL
FDYYKHHYNITIQLDKQPLLVHRSKEKKTGETVEKISLLIPELCFLAGITDSIRSDFRITKDLDSQVTKVSPE
QRRQVIRKFIQQVNGNEVTKKLLSDWGLEIDNDTLDSLGRVLDPEELMFGNDYKTKSPNADWGKAATNS
KVLRTPNMPDWAVLCTQRNQRNCNDFVETFKGVCQKVNIRIGDPRVIFLRDDNTETYSSELRKLKLNRL
NMVVIIFPALRQDKYSVAVKRICCVDHPVPSQVIIAKTIAKPKLKSVEKIALQINCKLGGALWALKNPFT
NAMVCGIDVFHAGVGGGSKGSVAAFIASLDMKMLTSWHSRVCLQAPHQEIVDLLSQCLVSAINVYREKNG
QYPDRIFIFYRDGVGDGQLEAVKKEYEVRQFLNTCARLDPAYKPKLSVIIQKRVNTRIFEKRGQLGNPPPGT
VVDSTITRFLYDFYLVLPQLVRQGTVTPHYIVLHDGGDVKTDHMQRFTYKLCCHLYYNWPGTIRVPAPC
QYAHKLAYLVGQSIKSEPSHELNNLLYYL

>PpPiwi-like1

MNEKLLDANSVVAASSRLRQNPDNSMAAAVAALRKARAPPVSEVKTKEPHIISKSGTTGRKVVLNANYF
KLKSKTDWCLYQYRVDIAPDEERTMVRKSLLRTHKAALGAYLFDGTVLYTSNRLPNPMELNSIRQEDKQ
AMNIKIRLVGDVVKGDYHYFQFFNIMVRKCLENLNLQLVGRNYFDAAAKVEIAQYKLELWPGYVTSIRQ
HERDVLMAEISHKVMRNQTLVDILRDCHESQRNNYQKAFSATVIGSVVLTDYNNRRTYKVDVDYDYNKS
PSSTFKLRDGSISYATYYQQKYQVKIRDPRQPLLVSRSKARRRAGNDELVYLPELRCRSTGLTDDMRG
NFQLMKALAEHTRIAPEARMRLLAFNDRLRSKPEVIQDFKEWNFDLESNLVELPGRVLPLETTYFSGKK
VQLRQADWTVDVQDGPVSCRPLKDWVAIVPSKNAENAKSFIELMISVASKMKFAINRPLIAIDNDRSD
TYANTITTLTDKSVPLIFCVVSNNNQLRYTAIKKKLCVDRPIPSQVVMARTLVQKTPANTRSVATKVAV
QLNCKMGGIPWTTDIGQILPGLMVVGFVCHDKNGDFGALVASLDQNFGRFYSAVSTHAAGEELSNHFA
QLLGNAREFYKVNGLRNLPSRIIYRDGVGEGQLPFVYDYEITQVKDTLSKIYANPEHIKLGFIIVTKRLNS
RFFLNKNNPPAGTVIDDVITDPLKYDFFIVSQAVRQGTVAPTAYNVLEDGTGLKADQMQRPTYKLCCHMY
YNFSGTVRVPAPCQYAHKLAFSISQTIRQPADAKLASTLYFL

>PpPiwi-like2

MSDRGRDVRHGRGGTRPSYDEQQPGTSQGRPRLSSQRPEPYPRRPGPPAERGAEGYPPISPEYQRPSTSAG
PAYPSPAHDVAIRVTEHRSQPAYSDSMSDTASVAGSIDGAASVSSSIERGGMHGRRVLPANVITRPTTLES
KQGVTRPILLTANYFKLLSTTDWCLHKYRVDFIPEEDRKVVCKGLLRSHKELLGAYIFDGATMYSSKRL
PDPLELTAERAGDKQIMQIRIRHTGEMTRGDHDYLFQFFNIIMRKCLDYHLQLVGRNYFDAANKIEIREFR
LELWPGYITSIRQHEYDILMCSEITHKVMRNETLLHVLTHCLEENRQDYKNAFSKQVIGVTVLTDYNNNT
YRIDVDVFNVTPESTRKKTGETISYFEYKTRYNLRSQRLQPLLVSKTSKDRRAGKDELVYLVPELCR
STGMTESMRNNYRLMGALAQYTRVSPGDRIKLMFNTLHSPVNVVTELKNWNLTLDLDRQLVTLKGRV
LPADKIFYLHQDVSEVNTANWTNDLNRKLLKCGVLDNWVVVVPDRLMRDCQKFGVGCIIKVADAMG
FRINQPHYMEMRNDQAATYVNMLEQIKSTTNPQLIFCATTNNRMDRYSAIKKKCCVDRPVPTQVVIKAT
LQHKSSMSIATKVAIQMNCKIGGIPWSIPIPLKGLMVVGFVCHDKNDRNKDIGAMVASLEPNFGRYFSA
VSSHSSGEELSNMDSVNLCKALDQFKRYNQSLPKRIVYRDGVGDGQVPFVVSHEVQKIKSKLETYDDA
NDVKLAYIIVTKKINSRLFDGIRNPPPGTVVDDVITDPTKYDFMIIAQHVTQGTVTPTSYNVYDTCGMDA
DKLQRLTYKLCCHMYFNWSGTVRVPAPCQYAHKLAFLIAQCVHQPDDILKTLLYFL

>PpPiwi-like3

MSSRDVRRERGGTRPSYDEQQPGTSQERSFSSSQHEEFYPCRPGPSTEQISGIYPSSARGETVRAEYRSQPT
YPETASDTASDTASVAGSIDGAAAPSLTLGRGAMGGRRILPANIITRPSLETQKQVTVGQRIELTANYFKLL
STTDWCLHKYRVDFSPPEEDRKVVCKGLLRPHAEITIGPYIFDGSTLYTSTRLPEALELTSERLGDKQSILIKI
KHTGDMTRGDHDYIQQFFNIIMRKCLDYLELQLVGRNYFDARKKVELRDFRLELWPGYMTSIRQHEQDIL

MCAEITHKVMRNETLLDVLIQCHGENQQDYKNLFFKQVIGVVVLTDYNNNTYRIDDDVFNVTQSTFRK
KTGGEISYIEYYKSRYNLHIREMRQPLLVSXSKPKDRRAGKEELVYLVPELCRSTGLTDKMRNNFRLMSA
LAQYTRVLPGRKIDKLVFNKRLHNVENVGTELKRWNLQLDKKLVTLTGRVLPADNIVYYQNDQSGVN
NKANWTNDFRTRKLLKCGTLDNWWWVVPDCLKHECQNFVGMIRVASTMGFRIEQPSVKELRNDQITS
YTSMLNIKSTSNPMLIFCVATNNRADRYAAIKKKCCVDRPVPTQVVLAKSLKNKNAMSIATKIVIQMNC
KIGGIPWSIPIPLKSLMVVGFVCHDTITKNKDFGAMVASLDPHFGRYFSAVSSHTSGEELSNDLSVNLCK
ALAQFKHYNNALPKRIVYRDGVGDGVVFNHEVQQIKSKLETIYDDANDVKLAYIIVTKKINSRLFD
GIQNPPPGTVVDDVITDPTKYDFMIISQHVVTQGTPTAYNIVYDTCGMDADKLQRLTYKLCHMYFNWS
GTVRVPAPCQYAHKLAFLVAQAVHQSPDVVLETILHFL

>PpPiwi-like4

MPKYPGDKRRYYDDMSDRGRPMSRGRHSRRNPDEQQPSTSQETLRSSSQHREPYPRRPGPTEQISESY
PSSARGEAVRAEHRPQPTYSETASDTASVAGSVDGAAAPSSTLARGAMRGRVLPANIITRPTSLETQKG
VTGRPIELTANYFKLLSTTDWCLHKYRVDFSPPEEDRKKVCKGLLRPHRETLGAYIFDGSTLYTSTRLEAL
ELTSERLGDKQPMLIKIKHTGNMTRGDHDIYQFFNIIMRKCLDYLELQLVGRNYFDARNKVEVRDRLEL
WPGYITSIRQHEQDILMCAEITHKVMRNETLLDVLIQCHEANRQDYKNLFFKQVIGVVVLTDYNNNTYRI
DDVDFSVTPQSTFRKKNGEEISYVEYYQSRYNLRIKEMRQPLLVSXSKPKDRRAGKEELVYLVPELCRST
GLTDMIRNNFRLMSALAQYTRVSPGDRIKLLINFNKRLHSVQNVVTELNSWNLQLDKRLVTLTGRVLP
DNIVYYQNDQSGVNNKANWTNDFRNKLLKCGTLDNWWWVVPDRLKRDQNFVGMIRVASTMGFRI
EQPVVKELRNDQAASYADMLNIKSTSNPMLIFCVATNNRMDRYAAIKKKCCVDRPVPTQVVLAKSLQN
KNAMSIATKIVIQMNCIGGIPWSIPIPLKGLMVVGFVCHDTNTKNKDFGAMVASLDPHFGRYFSAVSS
HTSGEELSNDLSVNLCKALAQFKHYNNALPKRIVYRDGVGEGQVPFVFNHEVQQITNKLETVDNASD
VKLAYVIVTKKINSRLFAGYENPPPGTVIDDVITDPTKYDFMIISQHVVTQGTPTAYNIVYDTCGMDADK
LQRLTYKLCHMYFNWSGTVRVPAPCQYAHKLAFLVAQAVHQSPDVVLETLLYFL

>PpSid1

MHYARHLELLKIAFAVFIVILTVGSSKAETIPLTPIIPANYSYPYTYSLNKSIEYVFLYSENTANMLNAARIV
VRSEDAKPNYPLIVVVRQKKGILSWQIPLEVENKYLENPVLYNDTSRTLCPAKFYKTIRFDDSDDHVTVS
ISTASSKNITFNLNLTPITDFNMGSGETRQFSITPSQPVYYGYDFQKLTDSVSVLITVKSDEPACMTFSIQNTS
CPVFDLERSIQYAGYYQTVSYQGGITVPRKAYPKGFFIVLVNADDFECSGNTSIHYIREKRVNITISNSITI
QDYIVASFALSMLTFCFAYIIGAICCKIHRTRKLQEESIQDEIGRAQQTIQSPDAHTSGFQEIGEAVQIYLD
DDSSLEDDIDLMEDAYCDKDVVRTKLVLVCDLARKEPKILRKSRLYIYYLITVAIFYTAPVIQLVVTY
QNVLHVTVGNQDLCYYNFLCSHPFKMLSDFNHVFNSLGYVLLGLLFFIVCMRERFSEHSHNIDRFGFIPQH
YGLYYAMGFALMMEGVLSGSYHLCPNHSNFQFDTSFMYVIAVLCMVKIYQNRHPDINAQAPVTFGVLA
VTILLALVGVLDGELYFWIFFTIIHLSTCLWLSSQIYYMGRWKFNGGVFRMIMTMRHDARSIIYLFRL
YWRLLLLLVGNFCNLGLAIYGNMHRQDFATYLLMILMSNLILYTFYIVMKLCHGERILIQPLL YILLS
FVAWGGALYFFINKTISWALTPAQSRTHNRPCTLLNFYDYHDIWHFLSALAMFFSFMVLLTLDDDLDDA
HRSLISVF

>PpR2D2-1

MNKKTPVSILQEMMVKKGTSYELIHNGGGTHQNIFTYQVTCDGLSASGTGRCKKDAKHEAATAMLA
AIAKHIALPQLPATPCESPVRTPLPMTLPVAPKSPANVPFRNTIGELQELCSTNNLDDPQYKEISDVGPPHA
RIFTVQCLVSTFVEEGIAKTKKQAKHEAARKMLERISDIVPDDPNAIDSTDDENQKKEQMASKMAIARYP
FLSMVHSETGKSKINWGLKIVNFHQRLKNYFEDDKRNNVLEKLDVLSDLVKQYKDDMSEDNFSKLLKQEF
ESIQLNTAPDIHEIGYDGTETQAEVSALEKIIQTMKFFLQ

>PpR2D2-2

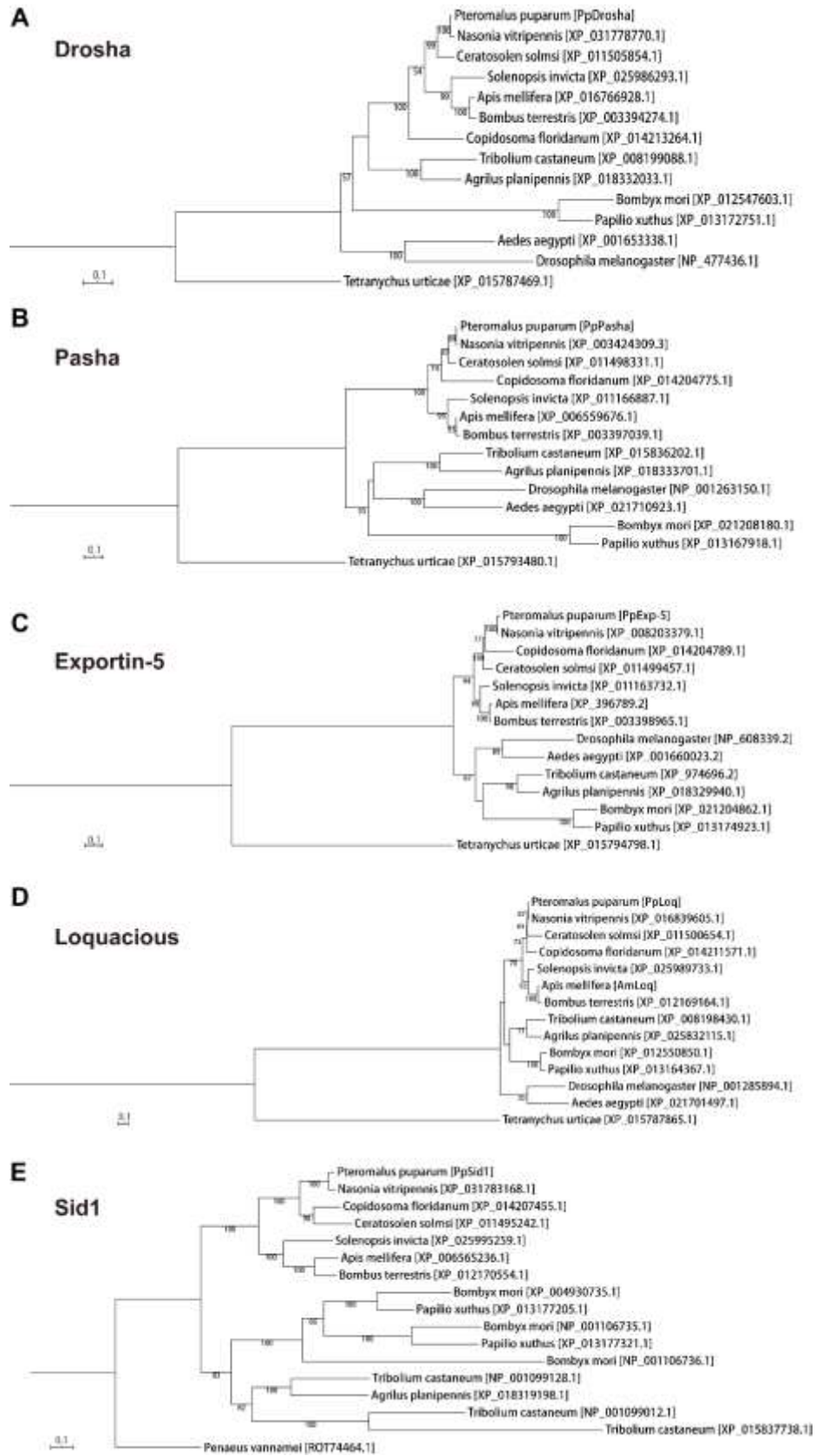
MSQSSGYFQMEAKIGSAPIYELCTTELKTIAPIFTFKVFCDGLTAYGSGKSKKEAKQNAAQNIERMSQRK
VVMDTSAATSKKERRAVQELMKLCYQLQLEPIYEEVSVRTGLPHAPIYSTKCTVSKFQQDGTARTKKQSK
QNAARKMIDQLKDTFNLVSLPHANGNNENKRIKKSMDNDVNTIEFSQIDAIDQALKLKINYQLKMESK
RKELLKNLSSTSVTLAILLSNLQKMSDLIQECQLNCSSQRITQLKQMFQTVMDTAKIDFHHMLLQTVEPAT
FMLIIQLNIVPDVIEMSIGKTREEAEWRTISKIATLKEILN

>AmLoq

MEEIHQPQQVGPNNMMPNVGGVPHNSNNVNRNRVRSSLHSLMIGKRPISEAAQLEMKSLSNKTPVSILQE
LLSRRGTPKYELIQVEGAIHEPIFRYRVTVADVAMGTGKSKKEAKHAAARAVLTKLRLNSESPPSPL
PNSIPDSENQELSGYGEEKIITNPICALQEMCMSRHWPPPKYTIENEEGLPHERQFTIVCSVLKYREVGQG
KSKKVAKRHAAHKMWQALHYMTTENPNIDEDEVLRNANVSARYADLKGSKISTLTIHSLKVSQFHKS
LKSSTGVKLFELQNTCLNDGDVNLVQFLQEIASEQQFEVTVVDIEEKSISGKCQCLVQLSTLPVAVCYGCG
VTSKDAQASAAQNALEYLKIMTKK

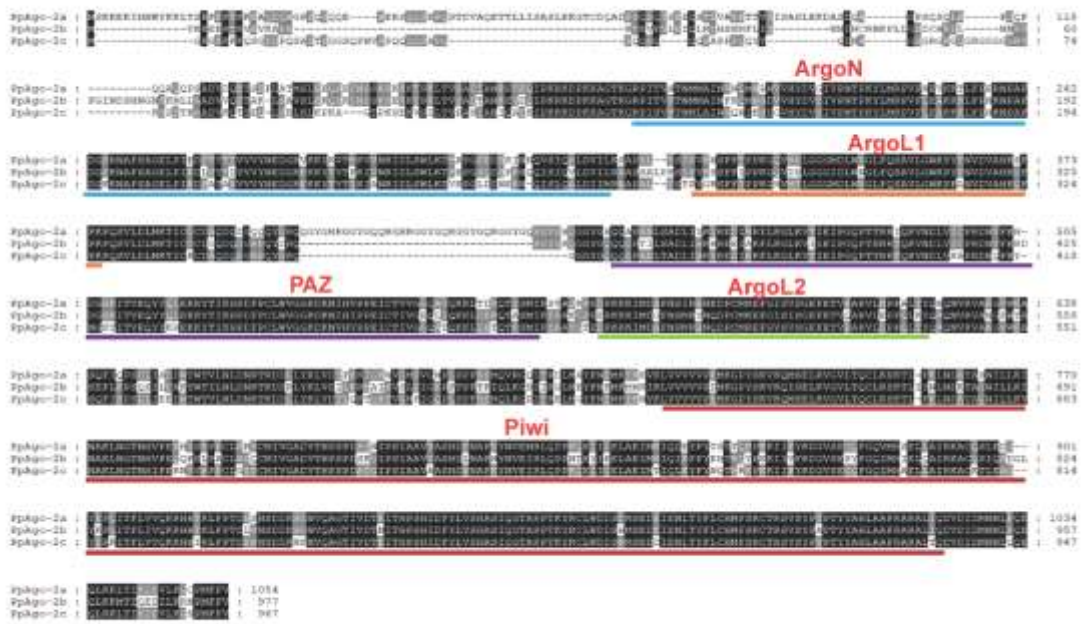
Appendix C Primers used in this study.

| Primer name | Forward sequence | Reverse sequence |
|--------------------|-------------------------|-------------------------|
| qDrosha | TCTCAAGCGGACTCAAGTT | TCGGAGGAATGTCGGTTATC |
| qPasha | AAACCACCGAGCCTTTACCT | CAGTCGCTTCATGCTTACCA |
| qExp-5 | GTCTGGATCGACCAGCTCTC | GGCGGTAGCACAGGTATGAT |
| qLoq | GTCGAGCGCAAAGGATTTAG | TGTGATAGCTTGCAGGATCG |
| qDicer-1 | GCTGGAGCTAGAGCCTGAGA | AGCTCTGCGACTTCTTGAGC |
| qAgo-1 | AGTTGGTAGTATGGACGCC | CAAGCTTCACGAATGGCAGT |
| qDicer-2 | CTGACAAGCAGCAATCCGTA | CCTCTCCGCAGGTTGAATAA |
| qR2D2-1 | AGTCACCAGCTAACGTTCCA | TGCTTTGCCTGTTTCTTGGT |
| qR2D2-2 | ACCTTGTCAGTTTGCCACAC | CGACAGAAGAATGGCAAGGG |
| qAgo-2a | GCTGACATTGCCAACAGAGA | CACCGGCCTCATCAAGTATT |
| qAgo-2b | GCTGACACTGCCAACAAGA | CCGTCAAATGCTGGATTTTT |
| qAgo-2c | TTCTCTGCCACATGTTCTCG | CATTGGGCTTCCCTTCAGTA |
| qSid1 | GCGAGGCATCTGGAACTTTT | CGCTTCGTACCACAATCCTG |



Appendix D Maximum likelihood tree of multiple sequence of alignment of Drosha (A), Pasha (B), Exportin-5 (C), Loquacious (D) and Sid1 (E) proteins. The tree is rooted at *Tetranychus urticae* or *Penaeus vannamei*. *Pteromalus puparum* proteins are pointed out by pentagram.

A



B



Appendix E Multiple sequence alignment of Ago-2 (A) and R2D2 (B) proteins. The lines below sequences indicate the ranges of predicted domains and the domain names are listed above the sequences. ArgonN: Protein argonaute, N-terminal; ArgoL1: Argonaute, linker 1 domain; ArgoL2: Argonaute, linker 2 domain; Piwi: Piwi domain; dsRBD: Double-stranded RNA-binding domain.

Appendix F Parameter estimation for hymenopteran Dicer and Argonaute genes under free-ratio model using codeml.

| Gene | Species | dN | dS | ω |
|----------------|------------------------------|----------|---------|----------|
| <i>Dicer-1</i> | <i>Pteromalus puparum</i> | 0.01059 | 0.11687 | 0.0906 |
| | <i>Nasonia vitripennis</i> | 0.00289 | 0.30766 | 0.0094 |
| | <i>Ceratosolen solmsi</i> | 0.03529 | 0.86671 | 0.0407 |
| | <i>Copidosoma floridanum</i> | 66.91498 | 0.07331 | 0.0011 |
| | <i>Apis mellifera</i> | 0.00922 | 0.13854 | 0.0666 |
| | <i>Bombus terrestris</i> | 0.00725 | 0.18267 | 0.0397 |
| | <i>Solenopsis invicta</i> | 0.01744 | 0.31744 | 0.0549 |
| <i>Dicer-2</i> | <i>Pteromalus puparum</i> | 0.01383 | 0.23354 | 0.0592 |
| | <i>Nasonia vitripennis</i> | 0.02244 | 0.05807 | 0.3865 |
| | <i>Ceratosolen solmsi</i> | 0.17033 | 1.134 | 0.1502 |
| | <i>Copidosoma floridanum</i> | 0.24268 | 1.63735 | 0.1482 |
| | <i>Apis mellifera</i> | 0.05915 | 0.12048 | 0.491 |
| | <i>Bombus terrestris</i> | 0.06887 | 0.35928 | 0.1917 |
| | <i>Solenopsis invicta</i> | 0.15945 | 1.46459 | 0.1089 |
| <i>Ago-1</i> | <i>Pteromalus puparum</i> | 0.00189 | 0.16145 | 0.0117 |
| | <i>Nasonia vitripennis</i> | 0.00001 | 0.10114 | 0.0001 |
| | <i>Ceratosolen solmsi</i> | 0.00188 | 0.96204 | 0.002 |
| | <i>Copidosoma floridanum</i> | 0.0041 | 0.74658 | 0.0055 |
| | <i>Apis mellifera</i> | 0.00124 | 0.11114 | 0.0112 |
| | <i>Bombus terrestris</i> | 0.00002 | 0.20237 | 0.0001 |
| | <i>Solenopsis invicta</i> | 0.01275 | 1.29289 | 0.0099 |
| <i>Ago-2</i> | <i>Pteromalus puparum</i> | 0.03996 | 0.26469 | 0.151 |
| | <i>Pteromalus puparum</i> | 0.05908 | 0.1959 | 0.3016 |
| | <i>Pteromalus puparum</i> | 0.03928 | 0.18797 | 0.2089 |
| | <i>Nasonia vitripennis</i> | 0.0425 | 0.22782 | 0.1865 |
| | <i>Nasonia vitripennis</i> | 0.08655 | 0.35726 | 0.2423 |
| | <i>Nasonia vitripennis</i> | 0.01077 | 0.18949 | 0.0569 |
| | <i>Ceratosolen solmsi</i> | 0.12709 | 1.02614 | 0.1239 |
| | <i>Copidosoma floridanum</i> | 0.05783 | 0.51774 | 0.1117 |
| | <i>Copidosoma floridanum</i> | 0.04508 | 0.53012 | 0.085 |
| | <i>Apis mellifera</i> | 0.07145 | 0.27662 | 0.2583 |
| | <i>Bombus terrestris</i> | 0.07728 | 0.25616 | 0.3017 |
| | <i>Solenopsis invicta</i> | 0.27479 | 1.28383 | 0.214 |

