

Table 4. Proteins locating to nucleocapsids and unknown localization

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
Identified by using multiple techniques						
9	p78/83			MTNRRYESVQSYLFNNRNNKID AHQFFERVDATAEAQIIKNNIYDN TVVLRNDVLLNILKLANDVFDN KAYMYVDDSEVSRHYNAVVK MKRLVIGVRDPSLRQSLYNTIAY IERLLNIGTVNDSEITMLIADFYD LYSNYNIELPPPQALPRSRRPSV VQPAAPAPVPTIVREQTKPEQIIP AAPPPPPSPVPNIPAPPPPPPSMS ELPPAPPMPTEPQPAAPLDDRQQ LLEAIRNEKNRTRLRPVK <u>PKTAP</u> <u>ETSTIVEVPTVLPKETFEPKPPSA</u> SPPPPPPPPPPPAPPAPPPMVDLSS APPPPPLVDPSEM <u>LPPPAPSLSN</u> <u>VLSELKSGTVRLKPAQKRQSEII</u> <u>PKSSTTNLIADVLADEINRRVA</u> MAKSSEATSNDEGWDDDDNR PNKANTPDVKYVQALFNVTSS QLYTNDSDERNTKAHNILNDVE PLLQNKTQTNIDKARLLQDLAS FVALSENPLDSPAIGSEKQPLFET NRNLFYKSIEDLIFKFRYKDAEN HLIFALTYHPKDYKFNELLKYV QQLSVNQQRTTESSA		1
49	PCNA					2
54	vp1054					3
61	FP25K		<u>MDOFEQLINVSLLKSLIKTQ</u> IDENVSDNIKSMSEKLKRL EYDNLTDSVEIYGIHDSRL NNKKIRNYYLKKICALLDL NFKHVISSFDKNHVAKL CDATRAKEWQTKSERRRL KNFNLNINYDGPVKIFVAA TAEQKLLLKKTRDALLPFY KYISICKNGVMVR <u>DEKSR</u> VFIVKNEQNIEYLKANKYY AFHSDSVDNFESENDSEKM LQNLI		4	
65	DNA polymerase		MKIYPYNELKTRFAEYAKP GEFNITSADTFRIIRL <u>HYDE</u>	MKIYPYNELKTRFAEYAKPGEF NITSADTFRIIRLHYDEKQGCLFA		

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			<u>KQGCLFAFCNTNIKERVVLQ</u> FYFKVKLNLYSYKQCYDK <u>HIFPSCRNKCISYTTFVAPG</u> VEGNYLNKINVIKYERNKA <u>APSDNAACLDKFLHNVNR</u> VHMQTPEVGAYMRFKKTQRCQN QRCQNNYVGGSTTRMFNL QHFNEDFELVDEMTLTSGI MPVLCYDIETHSDGHNMS KASVDCIMSIGFVVYKNDE YAKFCFMYHKLPTQIPETY DDDTYVVMFQNEIDMITAF FDMIKITNPDVILDNGDVF DLPIYLGLRNKTKMLLKRY <u>DLPAAAPTTKLFINKLGNK</u> VDTYYFNYIYHIDLKYKFFFSS DSNQHKVENFQLNTISSYY LGENKIDLPWTEMVKMYN TRRLDVIAKYNVQDCMLPI <u>KLFVKLKMADSVYSQCILH</u> <u>RLCTDDVICNISHLISVACF</u> YAAITNTRINESTGKEEPDP YFFNKNLDSIISGQFKADK AAAGISNLKRKLPLKNIPK DAINLG PANQTVKYKGKVLKP VLKPRAGIYKNAFSLDFNS LYLTIMIAICA CLSNLILCED GNVYLNHNSRAIVVKLLLK LLSERCKFKKNRDNQSESA FLYDLYDQKQNSVKRTAN <u>SIYGGYGFYKVLANYITRV</u> GRNQLRLAISLIEGLSNDPE ILEKFNLGSITFKVYYGDTD STFVLPTFNYNEISNETDTL KQICTHVETRVNNSF DGY KMAFENLMKV LILLKKKK YC YLNSEN KIVYKGWLVK <u>KDMPVFMRIA FRTAVEQIL</u> RHLDMDKCLQSLQTSFYE YYDEFAKSKSLTDYFSM YNDNPGKKRKSTDNEGP SPKRRVITVARHCREILVN KG TDVF PGNGDRIPYLLIDI EGKVTEKAYPLRLFPVK	FCNTNIKERVVLQFYFKVKLNLYS YKQCYDKHIFPSCRNKCISYTTF VAPGVEGNYLNKINVIKYERNK AAPSDNAACLDKFLHNVNRVH MQTPFVEGAYMRFKKTQRCQN NYVGGSTTRMFNLQHFNEDFEL VDEMTLTSGIMPVLS CYDIETHS DGHNMSKASVDCIMSIGFVVYK NDEYAKFCFMYHKLPTQIPETY DDDTYVVMFQNEIDMITAFFDM IKITNPDVILDNGDVF DLPYI LG RLNKT KMLLKRYDLPAAAPTTK LFI NKLGNK VDTYYFNYIYHIDL YKFFSSDSNQHKVENFQLNTISS Y YLGENKIDLPWTEMVKMYNT RRLDVIAKYNVQDCMLPIKLFV KLKMADSVYSQCILHRLCTDDV ICNISHLISVACFYAAITNTRINES TGKEEPDPYFFNKNLDSIISGQFK ADKAAAGISNLKRKLPLKNIPK DAINLG PANQTVKYKGKVLKP RAGIYKNAFSLDFNSLYLTIMIAI CACLSNLILCEDGNVYLNHNSR AIVVKLLLKLLSERCKFKKNRD NQSESAFLYDLYDQKQNSVKRT ANSIYGGYGFYKVLANYITRVG RNQLRLAISLIEGLSNDPEILEKF NLGSITFKV VYGD TDSTFVLPTF NYNEISNETDTLQKIC THVETRV NN SFTDGYKMAFENLMKV LILL KKKKYCYLNSEN KIVYKGWL KKDMPVFMRIA FRTAVEQILRH LDMDKCLQSLQTSFYE YYDEFA KS KSLTDYFSM TYNDNPGKKR KSTDDNEGPSPKRRVITVARHCR EILVNKG TDVF PGNGDRIPYLLI DIEGVTEKAYPLRLFPVKMRI SWIKHMGILCTFMNELL EIFGDE QKD KIAK CFTA IMQKYM QNQL YDRKEPV LVKINQKKCSVKRKR DDDDDDDDDDDDCDSSDSEN <u>DTQCANNTYKFCLYKMKK</u>		

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			MRISWIHKHMGILCTFMNEL LEIFGDEQKDKIAKCFTA IQYMQNQLYDRKEPV INQKKCSVKRKRDDDDDN DDDDDDDCDSSDSE CANNTYKFCLYKMKK			
66	93 kDa		<u>MQRWPKYGGTDVNTRTV</u> <u>HDLLNTINTMSARIKT</u> <u>LERYEHALREIHKVVVILKPSA</u> <u>NTHSFEPDALPALIMQFLSD</u> <u>FAGRDINTLTHNINYKYDY</u> <u>NYPPAPVPAMQPPPPPQP</u> <u>APPQPPYYNNYPYPPYPF</u> <u>STPPPTQPPESNVAGVGGS</u> <u>QSLNQITLTNEESELALF</u> <u>KNMQTNMTWELVQNFVE</u> <u>VLIRIVRVHVNNVTMINV</u> <u>ISSITSVRTLIDYNFTEFIRC</u> <u>VYQKTNIRFAIDQYLCTNIV</u> <u>TFIDFFTRVFYLVMTNFQF</u> <u>TTFDQLTQYSNELYTRIQT</u> <u>ILQSAAPLSPPPTVETVNSDI</u> <u>VISNLQEQLRKERALMQQI</u> <u>SEQHRIANERVETLOSOYD</u> <u>ELDLKYKEIFEDKSEFAQQ</u> <u>KSENRKIKQLERSNKELN</u> <u>DTVQKLRDENAEIRLSEIQL</u> <u>OKGDLDEYKNMNRLNED</u> <u>IYKLKRRIESTFDKDYVETL</u> <u>NDKIESLEKOLDDKQNLNR</u> <u>ELRSSISKIDETTORYKLDA</u> <u>KDIMELKQSVSIKDQEIAM</u> <u>KNAQYLELSAIYQQTVNEL</u> <u>TATKNELSqvATTNQSLFA</u> <u>ENEESKVLLLEGTLAFIDSFY</u> <u>QIIMQIEKPDYVPISKPQLT</u> <u>AQESIYQTDYIKDWLQKLR</u> <u>SKLSNADVANLQSVSELSD</u> <u>LKSQIISIVPRNIVNRILKEN</u> <u>YKVKVENVNAELLESVAV</u> <u>TSASVALVQQYERSEKQN</u> <u>VKLRQEEFEIKLNDLQRLLE</u> <u>QNQTDFESISEFISRDPAFN</u> <u>RNLNDERFQNLRQQYDEM</u>			

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			<u>SSKYSALETTKIKEMESIAD</u> <u>QAVKSEMSKLNTQLDELN</u> <u>SLFVKYNRKAQDIFEWKTS</u> <u>MLKRYETLARTTAASVQP</u> NVE ‡ SKLNTQLDELNSLFVK sequence obtained by MS/MS			
67	lef3				X	
70	hcf-1	1		MDSLANLCLKTLPYKFEPKFLR TKYCDACRYRFLPKFSDEKFCG QCICNICNNPKNIDCPSSYISKIP <u>KKENKEIYITSNKFNKTCNECN</u> QQSNRRCLISYFTNESCKELNCC WFNKNCYMCLEYKKNLYNVNL YTIDGHCPSPFKAVCFSCIKRIKTC QVCNQPLLKMYKEKQEERLKM QSLYATLNDVDLKILDIVDVN YSRKMLCAQCHIFARCFCNTNT MQCFCPRQGYKCECICRRSKYF KNNVLCVKSKAACFNKMKIKR VPWKHHSVDTFKSIYKLINV		
77	vlf-1	1	MNGFNRNENNFWNKIKI QSAPRFESVFDLATDRQRC TPDEVKNNSLWSKYMFPK PFAPTTLKSYSRSFIKIVYC SVDDVHLEDMSYSLDKEF DSIENQTLLIDPQELCRRML ELRSVTKETLQLTINFYTN MMNLPEYKIPRMVMLPRD KELKNIREKEKNMLKNVI DTILNFINDKIKMLNSDYV HDRGLIRGAIVFCIMLGTG MRINEARQLSVDDLNVLIK RGKLHSDTINLKRKRSRNN TLNNIKMKPLELAREIYSR <u>NPTILOQISKNTSTPFKDFRR</u> LLEESGVEMERPRSNMIRH YLSSNLYNSGVPLQKVAKL MNHESSASTKHYLNKYNIG LDETSSEEENNNDDDAQ HNRNSSGSSGESLLYYRNE ‡ sequenced by MS/MS			5
89	p39		<u>MALVPVGMAPRQM</u> RVNR CIFASIVSFDACTYKSPCSP	MALVPVGMAPRQM R VNR <i>C</i> IFA SIVSFDACTYKSPCSPDAYHDD		6

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
			DAYHDDGWFICNNHLIKRF <u>KMSKMVLPIFDEDDNQFK</u> MTIARHLVGNKERGIKRILI PSATNYQDVFNLSMMQA EQLIFHLIYNNENAVENTICD <u>NLKYTEGFTSNTQRVIHSV</u> <u>YATTKSILDTTNPNTFCSRV</u> <u>SRDELRRFDVTNARALRGG</u> <u>AGDQLFNNSYSGFLQNLIRR</u> <u>AVAPEYLQIDTEELRFRNC</u> <u>ATCIIDETGLVASVPDGPEL</u> <u>YNPIRSSDIMRSQPNRQLQIR</u> <u>NVLKFEGDTRELDRTLSGY</u> <u>EYPTTYVPLFLGYQIINSEN</u> <u>NFLRNDFIGPRANPNATLGG</u> GAVAGPAPGVAGEAGGGI AV	<u>GWIFICNNHLIKRFKMSKMVLPIF</u> <u>DEDDNQFKMTIARHLVGNKERG</u> <u>IKRILIPSATNYQDVFNLSMMQ</u> AEQLIFHLIYNNENAVENTICDNL <u>KYTEGFTSNTQRVIHSVYATTKS</u> <u>ILDTTNPNTFCSRVSRLDELRF</u> <u>VTNARALRGAGDQLFNNSYGF</u> <u>LQNLIRRVAPEYLOIDTEELRF</u> RNCATCIIDETGLVASVPDGPEL YNPIRSSDIMRSQPNRQLQIRNL <u>KFEGDTRELDRTLSGYEEYPTY</u> VPLFLGYQIINSENNFLRNDFIPR ANPNATLGGGAVAGPAPGVAG EAGGGIAV		
95	helicase			MIDNILQFFLKNVPQDKTYEINN LQDANHLIIRNRTGTRRLFEYV NNFQQFLNTIRNNFNGPCAKHD MGASCEDTEEPAEKHAQTLGD HDWVLESNDFCIFVKPFILKKHY DIIKYINFEDFFKSTDPGYINKC VQAGDYYYYWPNWPKKQAFSFN GWQLFLNIKFGIVIEPTIPIHNKK LGPVDLFVFDPKCFLNVELSLRT NHDPPQTLFVNNGTKFDDSHED LFILKMADGTVATCKVNGELVN SDKNFFNYIRDDINLEECITVPKY KHIVNVNLKSLRVFENNNFDKN DVDLSDTRSRKPRIVPIISASSEN ADYIQTQINLGLIAIHENMVKVL ATHERANDPNLLQQYFEKSKFK NFDFLIYVLWKILTKNENFSYRE TDIKLFLELLCESLFAACDKEALN EALKRCEPYKKQEKIVFNRACN HWFDFFDTKLCSVSLGYYFGIHY MIYLTQSAKNEILDHDELWAYT YENVMALNLPPDIVCKGFFRK <u>L</u> <u>ENVVTGVNLVFNGKHYQIVKKE</u> DDLFKLTKSNCYKLSNIKFNNW KYLYLTTHGVYNVFTNSFHSSC PFLLGTTLPQTFKKPTDEKYLPE DAFNYMLSTSADELSIYRTYHIA	X	

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				KMCRDVKMLKTNTAIVNYMGN CNTCQADMRLVALNNLFRDLWN LDDENLITLALYVNKNRVSMDL HNLKCKPCRSTVSGSRPKCKCY KKIKINRKALKVCLMADMFGN AELSELIWMLIFTNKTYVSTTLIR TNSEFVNQHGEFFSKEHNKIIQY LYRTIHIEYVDMILMDKFNDKR LFLTELRDDVAREPDVQFEESDN ISKFYTHADALMILKKYNVW WDKIILARSTDDLPTWLTRFY RIIMSKVDLKEYSYNLLKKIVEG YLYFKRFTNFNHNANAIMLMHFA ASLAIPVDYGGKAIYMPGEPGSG KSSFFELLDYLVLMHKFDDNH SGESNKETSDKEVSKLNSQLYTI NELKQCSESYFKKHADSSKSDS KSRKYQGLLKYEANYKMLIVN NKPLYVDDYDDGVQDRFLIVYT NHKFVDSVKFAGSVYEHIKSKQ FPIESMYYESLVTVPVRLFLSHVL MYRRDPKTGFVVYKTLLSNDP MHKHNLMCLSTNNSPLYALIYIL NIKTVRSATITIGEDKMEEMIGIA VQHFKNFLHPSFVQYNYKKNIN ASSSKSFVFNEQVLLQQIKNFK NNYNKTTNVFYNMTMALNRND LNTSVPNFVC		
100	p6.9		MVYRRRRRS <u>STGTTYGSTR</u> RRSSSGYRRRP <u>GRPRTYRR</u> SRS <u>SSTGRRSYRTRYY</u>	MVYRRRRRS <u>STGTTYGSTR</u> RRR SSGYRRRP <u>GRPRTYRR</u> RSRSST GRR <u>SYRTRYY</u>		7
101	BV/ODV-C42			MSAIALYLEINKLRLKIDEPMQL AIWPQLFPLLCDEHQSVQLNTD VLINFMMHVARKSQNTILNNNA AIAQYAAGNADVVAAPASAQP TPRPVINLFARANAAAPAQPSEE LINMRRYRNAARKLIHHYSLNS TSSTEYKISDVVMTMIFLLRSEK YHSLFKLLETTFDDYTCRPQMT QVQTDTLLDAVRSLEMPSTTID LTTVDIMRSSFARCFNSPIMRYA KIVLLQNVALQRDKRT <u>TLLELLI</u> <u>ERGEKIQMLQPQQYINSGETEIPF</u> CDDAEFLNRLKHIDPYPLSRM		8

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				YYNAANTMFYTTMENYAVSNC KFNIEDYNNIFKVMENIRKHSNK NSNDQDELNIYLGVQSSNAKRKY		
102	13.1 kDa		MIASINDTDMTDNNMSQARRN ARRNRRNRPPARPSAQTO <u>MAAVDMLQTINTAAASOTA</u> <u>ASLLINDITPNKTESLKILST</u> QSVGARSLLPEMQANASTI KLNRIETVNVLDFLGSVYD NTIQVIVTE ‡ sequence obtained by MS/MS	MIASINDTDMTDNNMSQARRN RRNRPPARPSAQTO <u>MAAVDMLQTINTAAASOTA</u> <u>ASLLINDITPNKTESLKILST</u> ESLKILSTOSVGARSLLPEMQAN ASTIKLNRIETVNVLDFLGSVYD NTIQVIVTE		
104	p87		MNDSNSLLITRLAAQILSRN <u>MQTVDVIVDDKTLSEEKI</u> DTLTSMSVLA VNSPPQSPPR VTSSDLAASIKNNS <u>KMVG</u> <u>NDFEMRYNVLRMAVVFK</u> HYPKYYNETTAGLVAEIES NLLQQYQNYVNQGNYQNIE GYDSLLNKAEECYVK <u>IDRL</u> <u>EKESIKKIMDDTAFEREQE</u> <u>AERLRAEQTAANALLERR</u> <u>AOTSADDVNRADANIPT</u> <u>AFSDPLPGPAPRYMYESSE</u> <u>SDTYMETARRTAEHYTDQ</u> <u>DKDYNAAYTADEYNSLVK</u> <u>TVLLRLIEKALATLKNRLHI</u> <u>TTIDOLKKFRDYLNSDADA</u> GEFQIFLNQEDCVILKNLSN LASKFFNVRCVADTLEVM LEALRN <i>NIELVQ</i> PESDAVR <u>RIVIKMTQEJKDSSTPLYNI</u> <u>AMYKSDYDAIKNKNIKTLF</u> <u>DLYNDRLPINFLDTSATSPV</u> RKTS <u>GKRS</u> AEDDLLPTRSS KRA NRPEINVISSEDEQEDD DVEDVDYEKES <u>RRKLED</u> EDFLKLKALEFSKD IVNEK <u>LQKIIIVVTDGMKRLYEYCN</u> <u>CKNSLETLP</u> SAANYGSLLK <u>RLNLYNLDHIEMNVNFYEL</u> LFPLTLYNDNDNSDKTLSH QLVNYIFLASNYFQNCAKN	MNDSNSLLITRLAAQILSRNMQT VDVIVDDKTLSEEKI <u>DTLTSMSVLA VNSPPQSPPR</u> VTSSDLAASIKNNS <u>KMVG</u> <u>NDFEMRYNVLRMAVVFK</u> HYPKYYNETTAGLVAEIES NLLQQYQNYVNQGNYQNIE GYDSLLNKAEECYVK <u>IDRL</u> <u>EKESIKKIMDDTAFEREQE</u> <u>AERLRAEQTAANALLERR</u> <u>AOTSADDVNRADANIPT</u> <u>AFSDPLPGPAPRYMYESSE</u> <u>SDTYMETARRTAEHYTDQ</u> <u>DKDYNAAYTADEYNSLVK</u> <u>TVLLRLIEKALATLKNRLHI</u> <u>TTIDOLKKFRDYLNSDADA</u> GEFQIFLNQEDCVILKNLSN LASKFFNVRCVADTLEVM LEALRN <i>NIELVQ</i> PESDAVR <u>RIVIKMTQEJKDSSTPLYNI</u> <u>AMYKSDYDAIKNKNIKTLF</u> <u>DLYNDRLPINFLDTSATSPV</u> RKTS <u>GKRS</u> AEDDLLPTRSS KRA NRPEINVISSEDEQEDD DVEDVDYEKES <u>RRKLED</u> EDFLKLKALEFSKD IVNEK <u>LQKIIIVVTDGMKRLYEYCN</u> <u>CKNSLETLP</u> SAANYGSLLK <u>RLNLYNLDHIEMNVNFYEL</u> LFPLTLYNDNDNSDKTLSH QLVNYIFLASNYFQNCAKN	9	

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			<u>ENYMRETFNVFGPFKQIDF</u> <u>MVMFVIKFNFLCDMRNFA</u> <u>KLIDELVPNKQPNMRIHSV</u> <u>LVMRDKIVKLAFSNLQFQT</u> <u>FSKKDKSRTNTKHIQLRIML</u> MNANYNVI				
109	44.8 kDa		MECPFQIQVCISDRFFAFPH <u>NLVEPQSDVGNKLIENLIV</u> YVPTDDDRLYIDKKQFPKF NSVLVYR HEHDVNIDSRS P <u>KKTASATIVYWNPLVPITEI</u> <u>GAGETR</u> VFSVLLTNLFYC NTMIVHHENPKCPIEFTYPE TDMQSACSALLKNRNGQS VPPP I KSNLRIACEIPLSHF KELVESNDFLLCFNLETST <u>MVKIILSLKRIFCIFQYRKQP</u> <u>ARYVINLPHEEDNLYNKL</u> NWERTRR L MKGDVPSNCA <u>TVNRSSLKYIKQAQSLLGIP</u> DYSQTVVDFVKMFQKIIFP YQLVPNVIIKLN N FDQMVS SAPNKAEPYKKIRLFCKND SIAISSGIVPINMPDFSPPN <u>TFDYSDYANRTNINFVTQR</u> <u>VLTDGGFSSGITVTPVKYN</u> YYL ‡ PDFSPPNTFD Y SDYANR; TNINFVTQR sequences obtained by MS/MS	MECPFQIQVCISDRFFAFPHNLV <u>EPQSDVGNKLIENLIVYVPTDDD</u> RLYIDKKQFPKFNSVLVYRHEH DVNI D SRS P KKTASATIVYWNP L NNLFY VPITEIGAGETRVFSVLLTNLFY CNTMIVHHENPKCPIEFTYPE D MQSAC S ALLKNRNGQSVP P PIKS NLRPIACEIPLSHFKELVESNDFL LCFNLETSTMVKILSLKRIFCIFQ YRKQPARYVINLP H EEDNLYN K LNWERTRRLMKGDVPSNCATV NRSSLKYIK <u>QAO</u> QSL L GIPDYSQ T <u>VVDFVKMFQKIIFPYQLVPNVII</u> <u>KLNNFDQMVSSAPNKAEPYKKI</u> RLFCKND S IAISSGIVPINMPDFSP N <u>PPNTFDYSDYANRTNINFVTQRV</u> LTDGGFSSGITVTPVKYN Y YL			
114	49 kDa			MSKYALLQKMIINELLFLNDNV <u>NYATNKI</u> LSKDQANGELHKLLA MLLNYKKS N EAPNIKFDL K NL SFMLENKDKIDIIQFDDIK <u>NYVQ</u> <u>PAIVNLFESHNRS</u> LN N YSTE L ST LLEDGNENLVPNITDIDNIKLSH MQARLLCYTAVVESRNSKPW KAIFNNDTNVLTDSFFNHIMNIL NM I KTNQGS L AHN V V V HIE N I QM N LENKLKPRSI T IE T VD K DK K F ENKHSDIEVCYVMNNSLHPQDV NSQQTLMCSSFVELNVLPFCLY NDTLPPDQS M SVFNLYKFE Q TK N K SV S KPSRLGNVMFVN S LD K P			

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				ITRETIVNIINSYHNACQNLKRSG HRVVGDYRAYERDYKLAALDFI <u>I</u> LMLVTSITHRTLKYNMNLNIHEK MFQELKTVVCKHNAAKLYDILI NYDINKEPLNNFRYNYEVL		
129	p24					10
132	25.1 kDa		MSDKPTKKGGSHAMTLR ERGVTKPPKKSEKLQQYK <u>K</u> AIAAEQTLRTTADVSSLQ NPGESAVFQELERLENNAV <u>V</u> LENEQKRLYPILDTPLDN FIVAFVNPTYPMAVFVNTD YKLKLECAR <u>I</u> RS DLLYKKN <u>N</u> EVAINRPKISSFKLQLNNV ILD TIE YDLQNK <u>V</u> L TITA <u>P</u> VQDQELRKSIYFNILNSD SWEVPKYMKKLFDEMQUE PPVILPLGL	MSDKPTKKGGSHAMTLRERG VTKPPKKSEKLQQYKK <u>A</u> IAAEQ <u>T</u> LR TTADVSSLQNPGESAVFQEL ERLENNAV VVLENEQKRLYPILDT PLDNFIVAFVNPTYPMAYFVNT DYKLKLECARIRSDLLYKKN VAINRPKISSFKLQLNNVILD TIEYDLQNKVLTITAPVQDQELR KSIYFNILNSDSWEVPKYM FDEMQUEPPVILPLGL		
142	49 kDa		MSGGGNLLTLERDHFKYL LTSYFDLK <u>D</u> NEHVPSEPMA <u>F</u> IRNYLNCTFDLDDAVLM NYFNYLQLSQLK <u>H</u> LGST <u>S</u> TNIFKFVKPQFRFVCDRT <u>V</u> DILEFDTRMYIKPGTPVY <u>A</u> TNLETNSPRKMMFLY <u>E</u> FGKVFKNKIFVNINNYGC VLAGSAGFLFDDAYVDWN GVRMCAAPR <u>L</u> NNMHPF LYLLGEDMAK <u>H</u> FVDNNILP <u>P</u> PHPSNAKTR <u>K</u> INNSM FMLKNFY <u>K</u> GLPLFKSKYTV <u>S</u> TKIVTRKPNDIFNEIDKEL NGNCPFIKFIQRDYIFDAQFPP DLLDLLNEYMTK <u>S</u> SIMKIIT <u>K</u> VFIEENPAMS GEMSREII DRYSVDNYR <u>K</u> LYIKMEITN <u>Q</u> FPVMDHES SYIFVSKDF <u>L</u> OLKGTMNAFYAPK QRLS <u>I</u> LA VNRLFGATE TIDFH PNL <u>L</u> VYRQSSPPV R <u>T</u> GDVYVV <u>D</u> KNEKVFLV KHVF SNTVP <u>A</u> YLLIRGDYESS SDLKSLR <u>D</u> LNPWV QNTLLK LIPDSVQ Q	MSGGGNLLTLERDHFKYL LTSYFDLK <u>D</u> NEHVPSEPMA <u>F</u> IRNYLNCTFDLDDAVLM NYFNYLQLSQLK <u>H</u> LGST <u>S</u> TNIFKFVKPQFRFVCDRT <u>V</u> DILEFDTRMYIKPGTPVY <u>A</u> TNLETNSPRKMMFLY <u>E</u> FGKVFKNKIFVNINNYGC VLAGSAGFLFDDAYVDWN GVRMCAAPR <u>L</u> NNMHPF LYLLGEDMAK <u>H</u> FVDNNILP <u>P</u> PHPSNAKTR <u>K</u> INNSM FMLKNFY <u>K</u> GLPLFKSKYTV <u>S</u> TKIVTRKPNDIFNEIDKEL NGNCPFIKFIQRDYIFDAQFPP DLLDLLNEYMTK <u>S</u> SIMKIIT <u>K</u> VFIEENPAMS GEMSREII DRYSVDNYR <u>K</u> LYIKMEITN <u>Q</u> FPVMDHES SYIFVSKDF <u>L</u> OLKGTMNAFYAPK QRLS <u>I</u> LA VNRLFGATE TIDFH PNL <u>L</u> VYRQSSPPV R <u>T</u> GDVYVV <u>D</u> KNEKVFLV KHVF SNTVP <u>A</u> YLLIRGDYESS SDLKSLR <u>D</u> LNPWV QNTLLK LIPDSVQ Q		

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
			‡ ILSILAVNR; TTVDILEFDTR; LYLLGEDMAK; SKYTVVNSTK sequences obtained by MS/MS			
144	ODV-EC27	2		MKRIKCNKV <u>RVTVEIVNSDEKIQ</u> <u>KTYELAEFDI</u> KNLSSLESYETLK IKLALKYMA <u>M</u> LSTLEMTQPLL EIFRNKA <u>DTRQIAAVVF</u> STLAFIH NRFHPLVTNFTNKMEFVVETN DTSIPGEPI <u>LFTENEGV</u> LLCSVDR PSIVKMLSREFDTEALVN <u>FENDN</u> CNVRIAKT <u>FGASKR</u> KNTTRSSD YESNKQPNYDM <u>DLSDFSITE</u> VE ATQYL <u>TLLLTVEHAYLHYYIFK</u> NYGVFEYCKSLTDHSLFTNKL STMSTKTSNLLSKFKFTIEDFD KINSNSVTSGFNIYNFNK		11
147	IE1				X	
Identified by using one technique						
5	12.4 kDa			MYRTSR <u>INNAPVVASOHDYDRD</u> <u>QIKRE</u> NSLRRNVHDLCTRSGTS FDCNKFLRSDDMTPVVTITPKR TADYK <u>ITEYVGDVKT</u> IKPSNRPL VESGPLVREA <u>AKYGECIV</u>		
14	lef 1	2				
22	44 kDa		MYRVLIVFFLFVFLYIVYQP FYQAYLHIGHAQ QDYN DT <u>LDDRM</u> DYIESVMRR <u>HYV</u> <u>PIEALPAIRFDTNLGTLAGD</u> <u>TIKCM</u> SVPLFVS <u>DI</u> LPMD CSQICDNP <u>SAAYFFVNETD</u> VFVVNGHRLTVGGYCSTN SLPRNCNRET <u>S</u> VILMSLNQ WTCIAEDP <u>RYAGTDNMT</u> <u>OLAGRQHFDRIMP</u> GQSDR NVLFDR <u>LLGREVN</u> TTNTF <u>RRSWDELLEDGTRRF</u> EMR CNARDNNNNLMFVNPLNP LECLPNVCTNVS <u>NVHTSVR</u> PVFETGE <u>CDCGDEAVTR</u> VT <u>HIVPGDRTSMCASI</u> IDGLDK STASYRYR <u>VECVNLYTSIL</u> <u>NYSNNKLLCP</u> SDTFDSNTD			

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
			AAFAFEVPGSYPLSRNGINE PTYRFYLDTRS <u>RVNYNDVR</u> GQLS			
30	54.6 kDa	1				
39	p43		MDKRANSRKPFLYNEDY YCEKP KRYFHTNKVIFEKL DSYATNINRCRELLTDFFD YCLPKYYRRKNKFTLLFRL LEPVIKQAGASALTTVSND QCRWLEINQFSGWEQRDN QYAHKWLIKVVGADMGQ <u>QILFIIKQVTKK<u>KTCNLGF</u></u> <u>HNYYKLFRRCLSILLFKHK</u> EVFVQCLQVILKAAMPVK KKGVVKSNYAFAVTNALH YYIVDNPHLLCK <u>DINVAIK</u> VRRLLIK <u>HEMPLTEKQIKLS</u> FEKCSKGIEVPLYEKLLL HMMRINDDNLQWPSLNN KKIMEWNANRCFDESNKIL HVYIGQYTSSCRRVKKSF FKYNGWNRQGRFCRTEKF CHLVNLQLNKGSKKLKR VQRKLINCK			
58	6.8 kDa			MFSSRKRRVAKRAFN <u>AKSKKF</u> <u>IGEVVSTRKRLKQNTNTPHYDT</u> SEDEDENYYNY		
59	8.2 kDa			MYQIPDMLYNEKMP <u>PRAKKLFV</u> <u>ETFTKYHKMNGGDEDIAMHKA</u> RKALEEKYV <u>KIDTLQNLGFRAK</u> PPTR		
74	30.5 kDa	1				
79	12.1 kDa	2				
86	PNK/PNL		MLHVSRLLANGGVNLCD KFKVKIKNYTEHDLMVLN YESFERDRDHVVVECRGL ILNSRTYAVVSRSFDRFFNF QELLQNIGGEDAHHK <u>LFQS</u> <u>KENFKFYEKIDGSLIKYKY</u> NGEWHASTRGSAFAENLC VSDVTFKRLVLQALQLDEA HNQFQALCNEYLDCASTH MFELTSKHNRIVTVYDEQP TLWYLASRNNETGDFYFC			

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
			SNLPFCKYPKCYEFTSVQE CVEHAAQLKNLEEGFVYY DKNNAPLCKIKSDVYLN HKNQSRAENPTKLAQLVIN GEHDDFLALFPHLKSVIKP YVDARNFTNESTINIMVS GLTLNQQRNFNELVQTLPW <u>KCLAYRCRKAQTIDVESEF</u> LKLTEPEKIKMKNIIKFVST KQALNNKLA PTIKLPSKQ LLVLIGISGSGKSTYAKSLK GYTEINRDDVR <u>VKLFLNGD</u> <u>YTKLN AFYNQSRKCQTK</u> <u>EEOITKMCIEQFLKA AKCG</u> ANVVVSDTNLNTQSVDM WQKMAATHNYHFLTRLM DVSLETALERNYKRSDKFP LNPETIKKQYKKFLKVNNF EYYVPVGDKFPRAVLCDL DGTVALPTNRSFYDFDNRV AQDEARLDVITCVKYLAN CHDAIIVFMSGRSERVICEQPT RNWIEKYFDIKSYKLFMRP SDDTCKDYLLKLKFNNYI RGKYNVIAVFDDRPCVVR MWQDLKIPTVFNVCRDYL EF			
88	cg30			MEFVKLQCNCFSVAEIKNYFLQ PIDRLTIIPVLELDTCKHQQLCSMC IRKIRKRKKVPCPLCR <u>VE SLHFN</u> <u>VYSVNRNVVDVIKCSASSVAQW</u> NKINANFDAASLASVLFEKSLLD DAEDNNANADDTMLSEAQAILK KLQVDIAEQTQLNIKQQLDL DK LQQTSVSMQEKL DKIKSDYNNM HKSFKELQLKRITTEALKSLND DYAKLASKNAKLSSENKVLSNK NIELIKHNLLQNEYTTLQSYKC ITNATITTNVTINVD		
92	p33			<u>MIPLTPLFSRYKD SYLLYSFRLID</u> LLRASKSTH LTKLLSSQATYLYH FACLMKYKDIQKYEV VQLIEWA INASPDMDLQQFRIEFMDKTTEL NLRSCQPKSFTYTFTTIWDTMHF		

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
				LSLIIDDMVYTRDKSSLDFVMQ QLKTMKVLFYNVFFILQCAMCR DHYMNVKGFIYHIELIEALDKE KYGTDTFVDSYQQETAGADVA VVSNNMLMKNLMAYVSMTFH NHINDYKWIQRNKKPPAHYERM TWGEYKKLLNLQ		
114	49 kDa			MSKYALLQKMI <u>NELLFLNDNV</u> <u>NYATNKLF</u> SKDQANGELHKLLA MLLNYKKSNEAPNIKFDLKNL SFMLENKDKDIIQFDDIK <u>NYVQ</u> <u>PAIVNLFESHNRSLNNY</u> TELST LLEDGNENLVPNITDIDNIKLSH MQLARLLCYTAVVESRNSKPW KAIFNNDTNVLTDSFFNHIMNIL NMIKTNQGSLAHNVSVVYHIENI QMNLLENKLKPRSITIETVDKDKF ENKHS DIEVCYVMNNSLHPQDV NSQQTLMCSSFVELNVLPCLY NDTLPDDQSMSVFNLYKFEQT NKS VSKPSRLGNVMFVN SLIDKP IT <u>RETIVNIINSYHNACQNLKRSG</u> HRVVGDYRAYERDY <u>KLAALDFI</u> <u>ILMLVTSI</u> HRTLKYNMLNIHEK MFQELKTVVCKHNAAKLYDILI NYDINKEPLNNFRYNYEVL		
133	alk-exo			MFASLTSEQ <u>KLLLKYKFN</u> NYVKTIELSQAQLAHWRS NKDIQP <u>KPLDRAEILRVEK</u> ATRGQS <u>KNELWTLLRDR</u> NTASASSNSSGNMLQRPAL LFGNAQESHV <u>KETNGIMLD</u> HMREJIESKIMSAVVETVLD CGMFFSPLGLHAASP <u>DAYF</u> SLADGTWIPVEIKCP <u>NYR</u> DTTVEQMRVELGNGNRKY RVKHTALLVNKKGT <u>PQFE</u> MVKTDAHYKQMQRQMYV MNAPMGFYVVKF <u>KQNLV</u> VVSVP <u>RDETFCNKELSTEN</u> NAYVAFAVENSNCARYQC ADKRR <u>LSFKTHSCNHYSG</u> QEIDAMVDRGIYLDY <u>GHL</u> KCAYCDFSSDSRETCDSVL		

ORF	Identity	Library screen	SDS-PAGE - MALDI	MUDPIT MS/MS	Western blot	Ref.
			KREHTNCKSFNLKHKNFD NPTYFDYVKRLQSLLKSHH FRNDAKTLAYFGYYLTHT GTLKTFCCGSQNSSPTKHD HLNDCVYYLEIK ‡ sequence obtained by MS/MS			

MALDI, matrix-assisted laser ionization desorption; MUDPIT, multidimensional protein identification technology; MS/MS, tandem MS.

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