Supplemental Figure 1.

Sequence alignment of IK regions of UNC-89, obscurin and SPEG using ClustalW.

UNC-89	NDEKLKTEPLSADTLREFKYQHKWLERRVFVQQTP-SEQILEAILGPATAQAQQN	54		
OBSC	LKSMPAEEAHFINTKQLKFLLARSRWQRSLMSYKSILVMRSIPELLRGPPDSPSLGVARH			
SPEG	KTQAKGAEVSTDHLKLFLSRRRWQRSQISYKCHLVLRPIPELLRAPPERVWVTMPRR	57		
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UNC-89	APVAPEGRRPAEIYDYLRIQPKKPPPTVEYVPQPRKEHPPFIDEFGQLIDG	105		
OBSC	LCRDTGGSSSSSSSSDNELAPFARAKSLPPSPVT	94		
SPEG	PPP-SGGLSSSSDSEEEELEELPSVPRPLQPEFSGSRVSLTDIPTE	102		
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UNC-89	-DAFDRPEGTGFEGPHRQPPQIPPQPQRPNQAAHDSRRHEQQPQHQ	150		
OBSC	HSPLLHPRGFLRPSASLPEEAEASERSTEAPAPPASP-EGAGPPAAQGCVPRHSVIR	150		
SPEG	DEALGTPETGAATPMDWQEQGRAPSQDQEAPSPEALP-SPGQE-PAAGASPRRGELRRGS	160		
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UNC-89	GQPQRIPVDQYGRPLVDPRYLNDPSHRPSSLDDAPFYVDKYGNPVHFDKYGRPMAPQ	207		
OBSC	HGALAPGSRRHP	174		
SPEG	SAESALPRAGPRELGRGLHKAASVELPQRRSPSPGATRLARGGLGEGEYAQRLQ	214		
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UNC-89	NLEKRKLIPQDKGETPSHSKKEKTQHPVATPILASPGGDQQQQKIP	253		
OBSC	-ARRRHLLKGGYIAGALPGLREPLMEHRVLEEEAAREEQATLLAKAPSFETA	225		
SPEG	-ALRQRLLRGGPEDGKVSGLRGPLLESLGGRARDPRMARAASSEAAPHHQPP	265		
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UNC-89	MRMIRGERREIEE-EIANRILSDISEEGSIAG	284		
OBSC	LRLPASGTHLAPGHSHSLEHDSPSTPRPSSEACGEAQRLPSAPSGGAPIRDMGHPQ-	281		
SPEG	LENRGLQKSSSFSQGEAEPRGRHRRAGAPLEIPVARLGARRLQESPSLS-	314		
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UNC-89	SLASLEDFEIPKDFQVEASEPSTPTLTPEVTIRETIPKPTPSPTSPQKSPVP	336		
OBSC	GSKQLPSTGGHPGTAQPERPSPDSPWGQPAP-FCHPKQG	319		
SPEG	ALSEAQPSSPARPSAPKPSTPKSAEPSATTPSDAPQPPAPQPAQD	359		

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UNC-89	VTYSDSILAGLPAADKKVLEDA	368
OBSC	SAPQEGCSPHPAVAPCPPGSFPPGSCKEAPLVPSSPFLGQPQA	362
SPEG	KAPEPRPEPVRASKPAPPPQALQTLALPLTPYAQIIQSLQLSGHAQGP	40
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UNC-89	ENDPSIPVGAPLFLEGLHGSDLTIDTTSASGLIKVTSPAINLSPNPKSPRRSTPGTKSPV	428
OBSC	PPAPAKASPPLDSKMGPGDISLPGRPKPGPCSSPGSASQA	402
SPEG	SQGPAAPPSEPKPHAAVFARVASPPPGAPEKRVPSAGGPP	44
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UNC-89	VLSPRQEHSMEVLIATKRGKPGFLPPGELAEDIDDEDAFMDDRKKQVKPKDHDGENDFKD	488
OBSC	SSSQVSSLRVGSSQVGTEPGPSLDAE-GWTQEAEDLSDS-TPTLQR	446
SPEG	VLAEKARVPTVPPRPGSSLSSS-IENLESEAVFEAKFKR	485
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UNC-89	EKERLEKDKNRRTVNLDDLDKYRPSAFYKDDSDFGHPGY	52
OBSC	PQEQATMRKFSLGGRGGYAGVAGYGTFAFGGDAGGMLGQGPMWARIAW	494
SPEG	SRESEEDGIYRPSPAGTPLEL	530
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UNC-89	DIDATPWD	535
OBSC	AVSQSEEEEQEEARAESQSEEQQEARAESPL-PQVSARPVPEVGRAPTRSSPEPTPWE	551
SPEG	VRRPERSRSVQDLRAVGEPGLVRRLSLSLSQRLRRTPPAQRHPAWE	57
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UNC-89	SHYQIGPDTYLMAARGAAFNSRVRNYREELFGMGAPTVKQGFLGVRNRDITVRERRRYTD	595
OBSC	DIGQVSLVQIRDLSG	566
SPEG	ARGGDGESSEGGSSASA	593
UNC-89	ILRETTQGLEPKSHEQSTALLQKAPSATAIERIKADIEKVTPCATKKNDDGT	64
OBSC	DAEAADTISLDISEVDPAYLNLSDLYDIK-YLPFE	600
SPEG	RGSPVLAMRRRLSFTLERLSSRLQRSGSSEDSGGASGRS	630

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UNC-89			647
OBSC	FMIFRKVPKSAQPEPPSPMAEEELAE	FPEPTWPWPGELGPHAGLEITEESEDVDALL	657
SPEG	TPLFGRLRRATSEGESLRRLGLPHNQLAA	QAGATTPSAESLGSEASATSGSS	682
UNC-89			647
OBSC	AEAAVGRKRKWSSPSRSLFHFPGRHLPLDEPAELGLRERVKASVEHISRILKGRPEGLEK		717
SPEG	APGESRSRLRWGFSRPRKDKGLS	PPNLSASVQEELGHQYVRSES	726
UNC-89		647	
OBSC	EGPPRKKPGLASFRLSGLKSWDRA	741	
SPEG	DFP	729	