

Supplemental Figure S3. Alignment of ICL centrin sub-families of *Paramecium*.

The 35 centrin isoforms were aligned by CLUSTALW and are visualized by BOXSHADE.

The different accession numbers of *Paramecium* centrins (named Ptcent_icl or Ptcent) are listed in Table 1.

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Ptcent_icl1a 1 -----MARRGQQPPPQ---APPAQKNQPGKFNPAEFVK
Ptcent_iclld 1 -----MARRGQQPPPQ---APPAQKNQGTGFNPAEFVK
Ptcent_icl1b 1 -----MSRRGQQPPPQ---QOAPP-QKNQAGKFNPAEFVK
Ptcent_icl1c 1 -----MARRGQQPPPQ---QOAPPQTKNQAGKFNPAEFVK
Ptcent_icl1f 1 -----MARRGQQPPPQ---QOAOQPAQKNAQGKFNPAEFVK
Ptcent8 1 -----MRSKAQA-AFVTQKQDQPKQGNVRKPQER
Ptcent15 1 -----MRGKQQA-AFVTQKQDQPKQGNVRKPQER
Ptcent10 1 -----MRAKP-QOVAQKQDQPKQGNVRKPQER
Ptcent12 1 -----MRGKP-QOVAQKQDQPKQGNARKPQER
Ptcent_icl1e 1 -----MSKKO-APVAQKPVGRQQQVNRPQDR
Ptcent_icl1g 1 -----MSKKO-APVAQKPVGRQQQVNRPQDR
Ptcent18 1 -----MNRKO-APVAQKPVGRQQQVSRKQDR
Ptcent_icl5a 1 -----MSRTNKQVEQAPANKKVQAF--AKPQKPKE
Ptcent_icl5b 1 -----MSRTNKQVEQSPVNKKVQAF--AKPQKPKE
Ptcent_icl6a 1 -----MSRANKQAE-APAQNKKQKGAPAAKPQKPKE
Ptcent_icl6b 1 -----MSRANKQAE-APAQNKKQKGAPAAKPQKPKE
Ptcent_icl7a 1 -----MNQQQLLDKDETALQKPVWKNH
Ptcent_icl7b 1 -----MNQQQLLDKDETALLQKPVWKNH
Ptcent_icl8a 1 -----MSRREQIEKDQKALLAKPWKNH
Ptcent_icl8b 1 -----MSRRDQIEKDQKALLAKPWKNH
Ptcent_icl3a 1 -----MSRQAPSQAQKVDPKRAAPPPPQ-KGPKPNAEDYAKP
Ptcent_icl3d 1 -----MSRQAPSQAQKVDPKRAAPPPPQAKGPKAFAEDYAKP
Ptcent_icl3e 1 -----MSRQAPQSQKNDKK--APPPQQPKGPKPNAEDYAKP
Ptcent_icl3f 1 -----MSRQONSKQHPQPQAOQNQQQRNPQPPQPKSKVNAEDYAKP
Ptcent_icl3b 1 -----MRPNPRQSKESKNVQQPQPPPPQSKPVQEDKQFNPDNYVR-
Ptcent_icl3g 1 -----MRSNPRQSKESKTVQPPPPPPQ-PKVQEKKQFPNDTYYVR-
Ptcent_icl10a 1 -----MNRKKNTTSVPTLQSKINQSLVPPQIIMSP-QSYSTYIPPEKFDPNDYIT-
Ptcent_icl10b 1 -----MNRKKNTTSVPTLQSKVNPOYAPPQQLSP-QSYSTFTIPPEKFDPNDYIT-
Ptcent_icl11a 1 MRNQRKSQSPPRTTOSIMRDSPNTQLEDESISTTRIPQNRESTRARMKINQSHNNALAQQGIRKQIA-ERASANGGNPNEFPKKFVT-
Ptcent_icl11b 1 MRNQRKSQSPPRTTNQSVMDSPNTQLEDESISTTRVPQNRESTRARMKINQSHNNALAQQGIRKQIA-ERTSAKGGNPNFPKKFVT-
Ptcent_icl11c 1 MRNQITSHSPFRTTOSIIRDSPHTLQDESRISNRRVPQNRESPARMLINQSHNNIQCQVVKQIA-ERTSKGGNPNFPKKFVT-
Ptcent_icl19b 1 -----MSRTQSRQQAQOPTQQAQKN-QPPAPLPATKQKAPAVPQPVQQQPAQGDASKYVK-
Ptcent_icl19a 1 -----MNRNSRQQAQOPTQQAQSKT-QPPPPAQAQAKGKGPAPAQQAVQQQPGGFASKYVK-
Ptcent_icl19c 1 -----MSRSQQRQQAQOPTQQAQSKT-QPPPPAQAQAKGKGPAPAQQAVQQQPGGFASKYVK-
Ptcent_icl19d 1 -----MSRSQQQKAQQAQTTQQANKS-QVQPAATRGKSPATAASQQPVQPAASGFDPSKYVK-
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Ptcent_icl1a 31 PGLTEEVIEIKEAFDLFDIDGQSIDPPELKAAT-TSLGFEAKNOTIYOM-SIDPDTDGSCOIDFEEFLKLMTRARIERDSKADIOKVFN
Ptcent_iclld 31 PGLTEEVIEIKEAFDLFDIDGQSIDPPELKAAT-TSLGFEAKNOTIYOM-SIDPDTDGSCOIDFEEFLKLMTRARIERDSKADIOKVFN
Ptcent_icl1b 31 PGLTEEVIEIKEAFDLFDIDGQSIDPPELKAAT-TSLGFEAKNOTIYOM-SIDPDTDGSCOIDFEEFLKLMTRARIERDSKADIOKVFN
Ptcent_icl1c 31 PGLTEEVIEIKEAFDLFDIDGQSIDPPELKAAT-TSLGFEAKNOTIYOM-SIDPDTDGSCOIDFEEFLKLMTRARIERDSKADIOKVFN
Ptcent_icl1f 31 PGLTEEVIEIKEAFDLFDIDGQSIDPPELKAAT-TSLGFEAKNOTIYOM-SIDPDTDGSCOIDFEEFLKLMTRARIERDSKADIOKVFN
Ptcent8 29 PGLTEEVIEIKEAFNLFDTEGIGRUDPELKAAT-QSLGFDQDNPIETNM-AELENEDGRIDIDFDCEFLDAITSKIGNRHRGINKLFD
Ptcent15 29 PGLTDDEIDEIREAFNLFDTEGIGRUDPELKAAT-QSLGFDQDNPIETNM-AELENEDGRIDIDFDCEFLDAITSKIGNRHRGINKLFD
Ptcent10 27 PGLTDDEIDEIREAFNLFDTEGIGRUDPELKAAT-QSLGFDQDNPIETNM-AELENEG-TDIDFDCEFLDAITSKIGNRESKDCINKLFD
Ptcent12 27 PGLTDDEIDEIREAFNLFDTEGIGRUDPELKAAT-QSLGFDQDNPIETNM-AELENEG-TDIDFDCEFLDAITSKIGNRESKDCINKLFD
Ptcent_icl1e 27 PGLTEEVIEIKEAFNLFDTEGIGRUDPELKAAT-QSLGFDQDNPIETNM-AELENEG-TDIDFDCEFLDAITSKIGNRESRDCINKLFD
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Ptcent18 27 PGLTEEVIEIKEAFNLFDTEGIGRUDPELKAAT-QSLGFDQDNPIETNM-AELENEG-NEIDFDCEFLDAITSKIGNRESRDCINKLFD
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Ptcent_icl5b 29 NYLTPPEINLRETFDFDDK-GTIDA-EIKVY-EDLGVDARNEKEFYOM-QDENFGC-CSIDFTVNIIDRIGNNK-KDGAFLFQ
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Ptcent_icl7a 22 TTELIN-HINELYRLFLNLYCNPRP-RRIDLDKDVMTAQBLGLVKSPIVANT-SQSDQRGDGCVDFEEVRE-TTKI-GNTFDQKGRCQ-LFT
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Ptcent_icl8b 23 PIPTY-HINEFYEMFLVLAEPB-KKADVDRDLVTAK-MGLLTD-KFPII-AHA-DPPIASYY-ADFETEITSDITAKI-GHPFDQKGRVQ-LFK
Ptcent_icl10a 38 PHLTKSEVIEVKCAFDFDNGSGSIDPPELLE-EAEFASCIKYHNNKFIYO-LGELEDTDNSCGIDFEEFLH-AATAKVSDKDTRE-CIOKVFN
Ptcent_icl10b 39 PHLTKSEVIEVKCAFDFDNGSGSIDPPELLE-EAEFASCIKYHNNKFIYO-LGELEDTDNSCGIDFEEFLH-AATAKVSDKDTRE-CIOKVFN
Ptcent_icl3e 37 PHLTKSEVIEVKCAFDFDNGSGSIDPPELLE-EAEFASCIKYHNNKFIYO-LGELEDTDNSCGIDFEEFLH-AATAKVSDKDTRE-CIOKVFN
Ptcent_icl3f 44 PHLTKSEVIEVKCAFDFDNGSGSIDPPELLE-EAEFASCIKYHNNKFIYO-LGELEDTDNSCGIDFEEFLH-AATAKVSDKDTRE-CIOKVFN
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Ptcent_icl3g 41 QNLSRDEVIELKKAQFLFDODDGGCTIDPAPLKCAFEDLGLRAQN-KM1YO-LGELEDTDNSCGDQGFSFDNPIK-AATAKQNLKETRGSIM-TEN
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Ptcent_icl11b 51 GSTKRDILYKEIFDLSNNNGVHQPMDL-KAFASACKYQPKQI-IYQ-IADEDQDOSIIIBEREVVR-IMSMNPGEKODIDDFENIYF
Ptcent_icl11c 87 KDIKES-IILIKOVFDYDSEQAGI-SPNDLQELI-SSCC-YHTPKETIYE-FSELDDEDELGGIT-BYFLGILNQDKSKS-RKTIRRVYR
Ptcent_icl11d 87 KDIKES-IILIKOVFDYDSEQAGI-SPNDLQELI-SSCC-YHTPKETIYE-FSELDDEDELGGIT-BYFLGILNQDKSKS-RKTIRRVYR
Ptcent_icl11e 87 KDIKES-IILIKOVFDYDSEQAGI-SPNDLQELI-SSCC-YHTPKETIYE-FSELDDEDELGGIT-BYFLGILNQDKSKS-RKTIRRVYR
Ptcent_icl11f 87 KDIKES-IILIKOVFDYDSEQAGI-SPNDLQELI-SSCC-YHTPKETIYE-FSELDDEDELGGIT-BYFLGILNQDKSKS-RKTIRRVYR
Ptcent_icl11g 56 PGLSKDEVIEKIKECFDIFDDDK-GSISPNMKNIAIALC-MEQSAEEIVNMQDLDQDGSSLIDFREFIN-FGFSGTIED-EQV-EKLYQE
Ptcent_icl19a 56 PGLSKDEVIEKIKECFDIFDDDK-GSISPNMKNIAIALC-MEQSAEEIVNMQDLDQDGSSLIDFREFIN-FGFSGTIED-EQV-EKLYQE
Ptcent_icl19c 53 PGLSKDEVIEKIKECFDIFDDDK-GSISPNMKNIAIALC-MEQSAEEIVNMQDLDQDGSSLIDFREFIN-FGFSGTIED-EQV-EKLYQE
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Ptce <i>_ic11a</i>	121	LFD S RAGVILKDRLRVAK-ELGET DDSELCEMIDRAD---S G D S QVTFEDFYNIMTKKTTA-----
Ptce <i>_ic11d</i>	121	LFDS S RACVIIKLDRLRVAK-ELGET DDSELCEMIDRAD---S G D S QVTFEDFYNIMTKKTTA-----
Ptce <i>_ic11b</i>	122	LFDS S RACVITLKDRLRVAK-ELGET DDSELCEMIDRAD---S G D S QVTFEDFYNIMTKKTTA-----
Ptce <i>_ic11c</i>	123	LFDS S RACVIIKLDRLRVAK-ELGET DDSELCEMIDRAD---S G D S QVTFEDFYNIMTKKTTA-----
Ptce <i>_ic11f</i>	123	LFDS S RACVITLKDRLRVAK-ELGET DDSELCEMIDRAD---S G D S QVTFEDFYNIMTKKTTA-----
Ptce <i>_8</i>	120	LFDD D GSNTINLNNLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_15</i>	120	LFDD D GSNSINLNLLNKLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_10</i>	116	LFDD D GSNSINLNLLNKLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_12</i>	116	LFDD D GSNSINLNLLNKLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_ic11e</i>	116	LFDD D GSNSINLNLLNKLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_ic11g</i>	116	LFDD D GSNAINLNLLNKLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_18</i>	116	LFDD D GSNAINLNLLNKLKRVS K -ELGET TAEELCEMERA--SNG-REI E EDFYNVKLT M -----
Ptce <i>_ic15a</i>	118	IYDP D TGFIDFTNLKRVS K -ELGET ND E LEM M HHIHILRK T S PEC I SE F EYEL T A P RRY-----
Ptce <i>_ic15b</i>	118	IYDP D TGFIDFTNLKRVS K -ELGET ND E LEM M HHIHILRK T S PEC I SE F EYEL T A P RRY-----
Ptce <i>_ic16a</i>	120	IYDP D TGFIDFTNLKRVS K -ELGET ND E LEM M HHIHILRK T S PEC I SE F EYEL T A P RRY-----
Ptce <i>_ic16b</i>	120	IYDP D TGFIDFTNLKRVS K -ELGET ND E LEM M HHIHILRK T S PEC I SE F EYEL T A P RRY-----
Ptce <i>_ic17a</i>	113	IVD D GK C TDKD L KK I SD-E H LNF S Q D -DE V I H NVAG---YDAED T b O E K YAK L TNR R K I TE V L R T K -----
Ptce <i>_ic17b</i>	113	IVD D GK C TDKD L KK I SD-E H LNF S Q D -DE V I H NVAG---YDA D T b O E K YAK L TNR R K I TE V L R T K -----
Ptce <i>_ic1a</i>	113	IID D GK C TDKGDL H K I SE-E R FN T ED D DE I I H NVAG---YEA D V b E K E K Y G K V Q R Q V E Q E I Y R N K I -----
Ptce <i>_ic18b</i>	113	IID D GK C TDKGDL H K I SE-E R FN T ED D DE I I H NVAG---YEA D V b E K E K Y G K V Q R Q V E Q E I Y R N K I -----
Ptce <i>_ic13a</i>	129	LYDNK E CR I WD L LRV A Q-D L GE E T D EE I Q H M E KK A D L -----D D E V T F DD F Y N M T O E G K Q -----
Ptce <i>_ic13d</i>	130	LYDNK E CR I WD L LRV A Q-D L GE E T D EE I Q H M E KK A D L -----D D E V T F DD F Y N M T O E G K Q -----
Ptce <i>_ic13e</i>	128	LYDNK E CR I WD L LRV A Q-D L GE E T D EE I Q H M E KK A D L -----D D E V T F DD F Y N M T O E G K Q -----
Ptce <i>_ic13f</i>	135	LYDNK E CR I WD L LRV A Q-D L GE E T D EE I Q H M E KK A D L -----D D E V T F DD F Y N M T O E G K Q -----
Ptce <i>_ic13b</i>	132	LFDLNRCR I WD L LRV S V-D L GE E ND E E V KK I ER K A D L -----D D E V T F DD F Y N M T G V I Y -----
Ptce <i>_ic13g</i>	131	LFDLNRCR I WD L LRV S V-D L GE E ND E E V KK I ER K A D L -----D D E V T F DD F Y N M T G V I Y -----
Ptce <i>_ic110a</i>	144	QFD D Y K GY I IID D RE A S-E C EN E K D E L LEN I IK A CD P --- G N T R K Q G I R M K S L Q K N -----
Ptce <i>_ic110b</i>	144	QFD D Y K GY I IID D RE A S-E C EN E K D E L LEN I ON I Q S C D P --- G N T R K Q G I R M K S A Q K R -----
Ptce <i>_ic111a</i>	179	KMDKN N K G F I I LODL R Q V Y K L R E E D E V V L A I F K T D S ---N Q D K T F E D F Y N M T K V I Y -----
Ptce <i>_ic111b</i>	179	KMDKN N K G F V I L Q D L R Q V Y K L R E E D E V V L A I F K T D S ---N Q D K T F E D F Y N M T K V I Y -----
Ptce <i>_ic111c</i>	179	K D KS N K G F V I L Q D L R Q V Y K L R E E D E V V L A I F K T D S ---N Q D K T F E D F Y N I T K N V I Y -----
Ptce <i>_ic19b</i>	146	EFDSGGQS K T Y E D K R I ND-LSERVTDELRMEYADK---KDESLSWDEBKAVVQEYPNQA-----
Ptce <i>_ic19a</i>	146	EFDSGGQS K T Y E D K R I ND-LSERVTDELRMEYADK---KDESLSWDEBKAVVQEYPNQA-----
Ptce <i>_ic19c</i>	144	EFDSGGQS K T Y E D K R I ND-LSERVTDELRMEYADK---KDESLSWDEBKAVVQEYPNQA-----