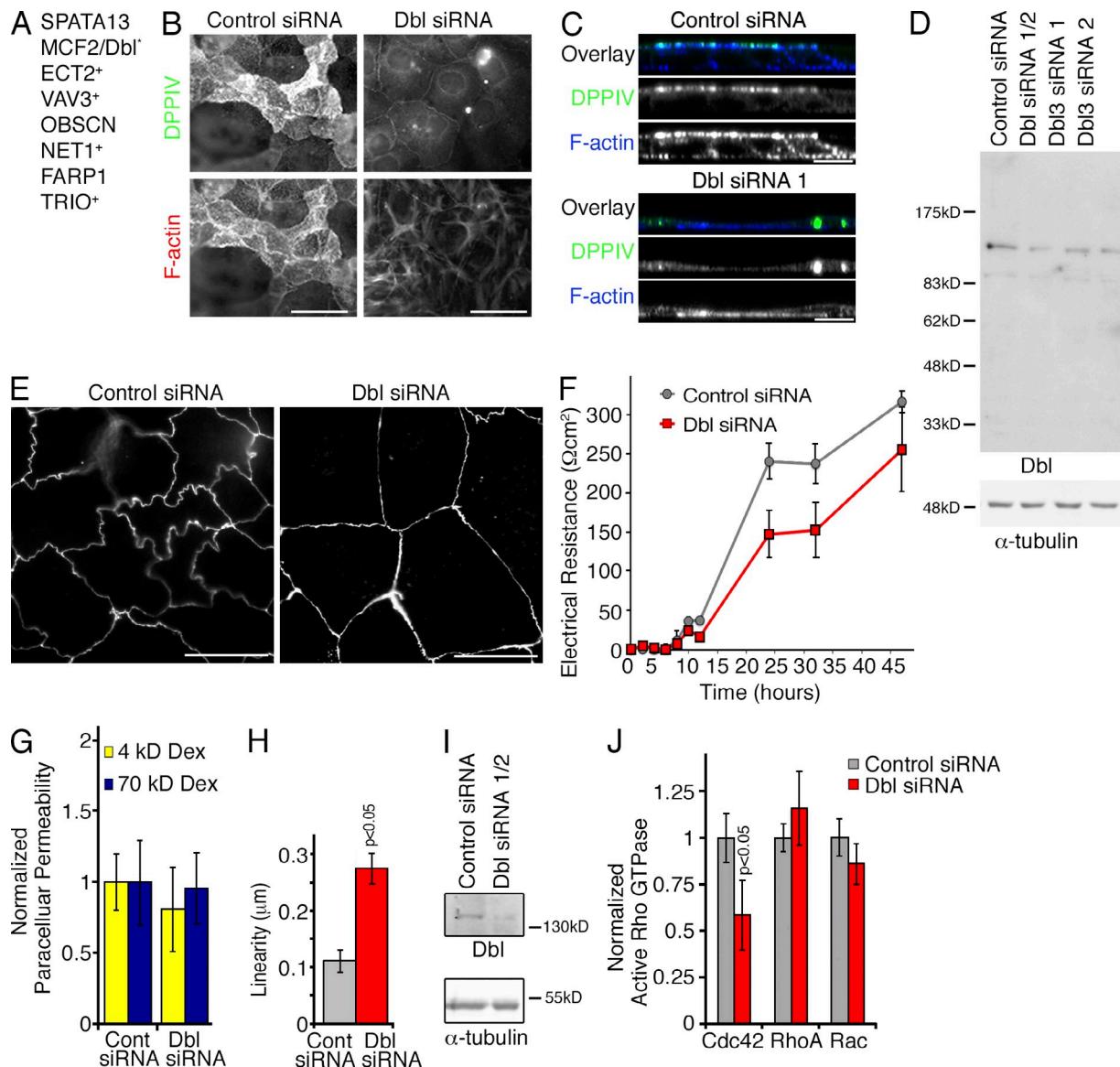
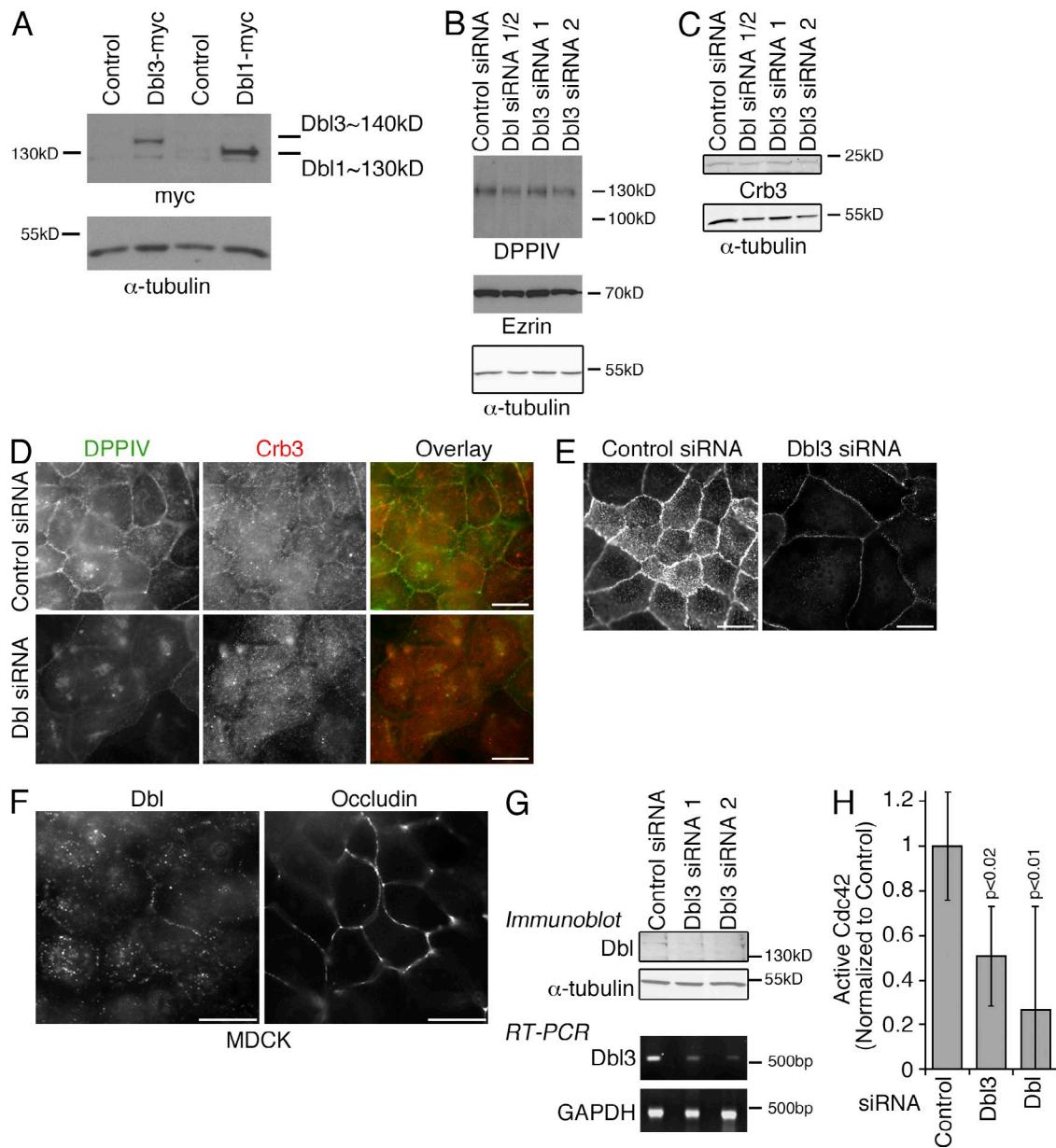
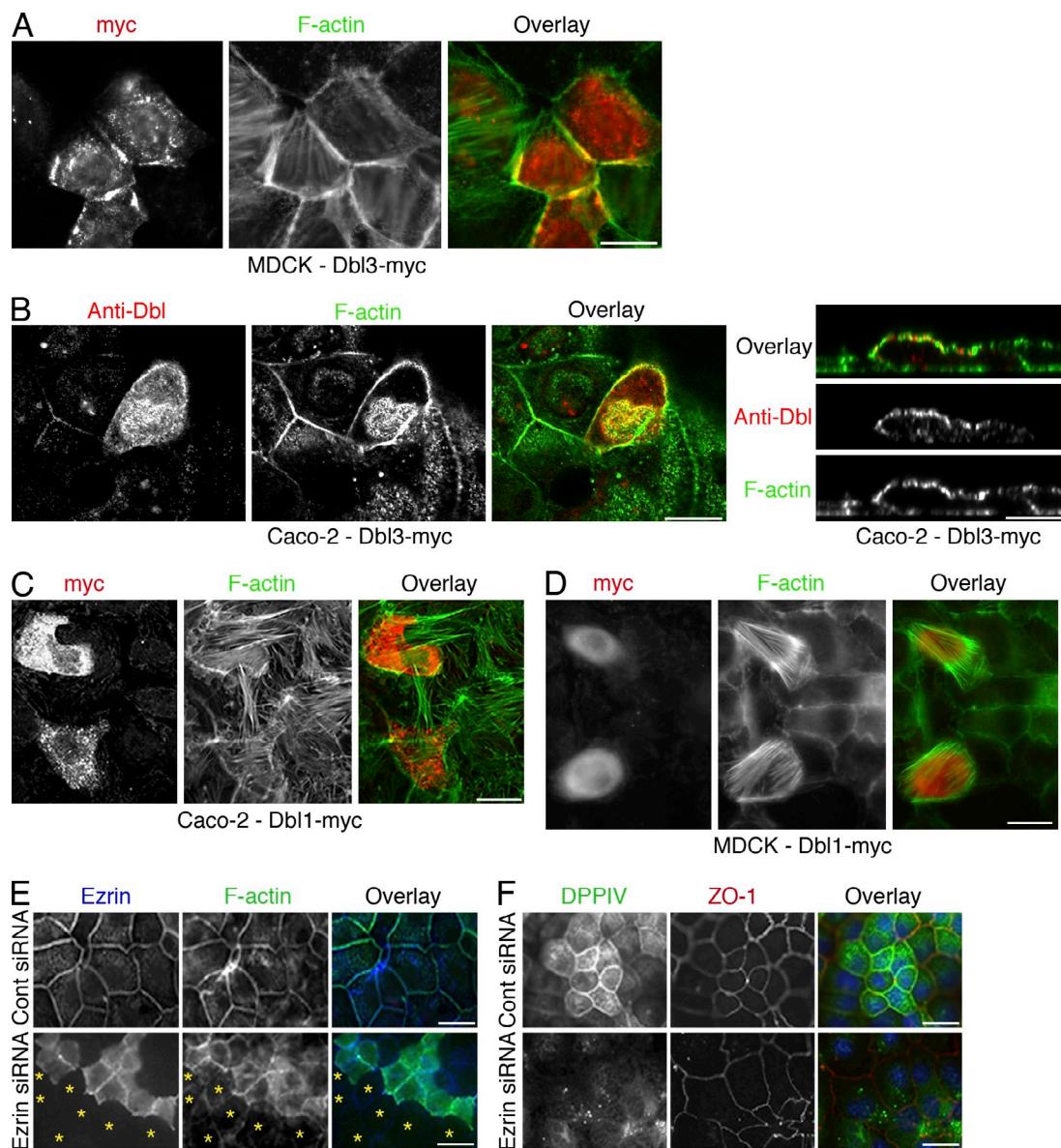


Zihni et al., <http://www.jcb.org/cgi/content/full/jcb.201304064/DC1>

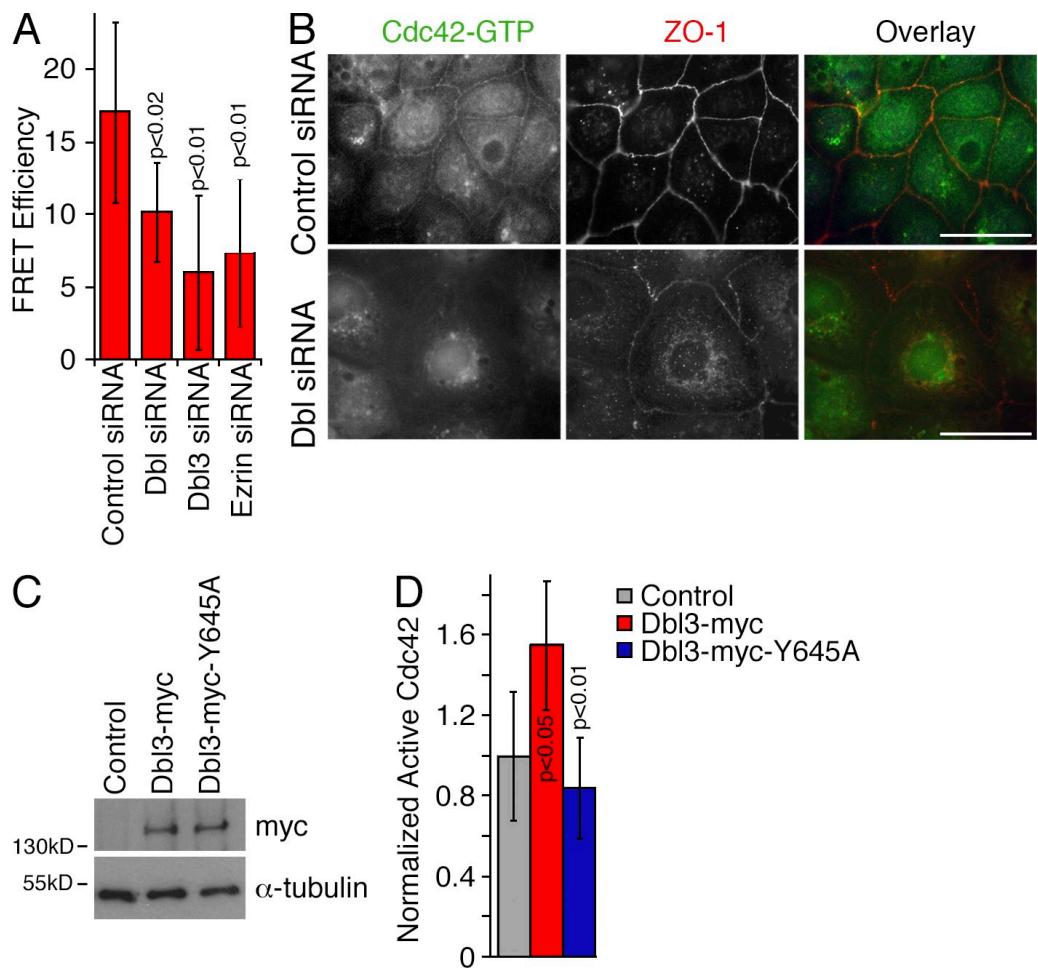
**Figure S1. Dbl regulates apical differentiation but not epithelial barrier formation.** (A) Caco-2 cells were screened for GEFs regulating junction formation and apical differentiation using a functional siRNA screen. The table shows positive hits of the screen that were confirmed in two independent rounds of screening. (B and C) Caco-2 cells transfected with control or Dbl-specific (targeting all isoforms) siRNAs were processed for confocal microscopy as indicated. Shown are confocal zy sections from the apical membrane (A) and z sections to illustrate the observed cell flattening and reduced accumulation of apical F-actin and DPPIV. (D) Caco-2 cells were transfected with siRNAs as indicated and analyzed by immunoblotting total cell extracts with anti-Dbl antibodies. (E) Caco-2 cells transfected with control and Dbl-specific siRNAs were fixed and stained to the tight junction protein ZO-1. Shown is an epifluorescence image. (F and G) Caco-2 cells transfected with siRNAs were subjected to a calcium switch protocol. Tight junction formation was then assayed by measuring transepithelial electrical resistance and fluorescent dextran permeability. Note that the effect of Dbl on junction formation was measured at the same time as the effect of SH3BP1; hence, the control values are the same as those previously published in Elbediwy et al. (2012). (H) The linearity of tight junctions was determined by measuring the length of linear junctional fragments in images such as those in E. Shown are means  $\pm$  SD ( $n = 10$ ). (I) Levels of Dbl depletion and active Rho GTPases in Caco-2 cells were measured after transfection of siRNAs (J shows means  $\pm$  SD,  $n = 3$ ). Bars: (B) 30  $\mu\text{m}$ ; (C) 10  $\mu\text{m}$ ; (E) 20  $\mu\text{m}$ .



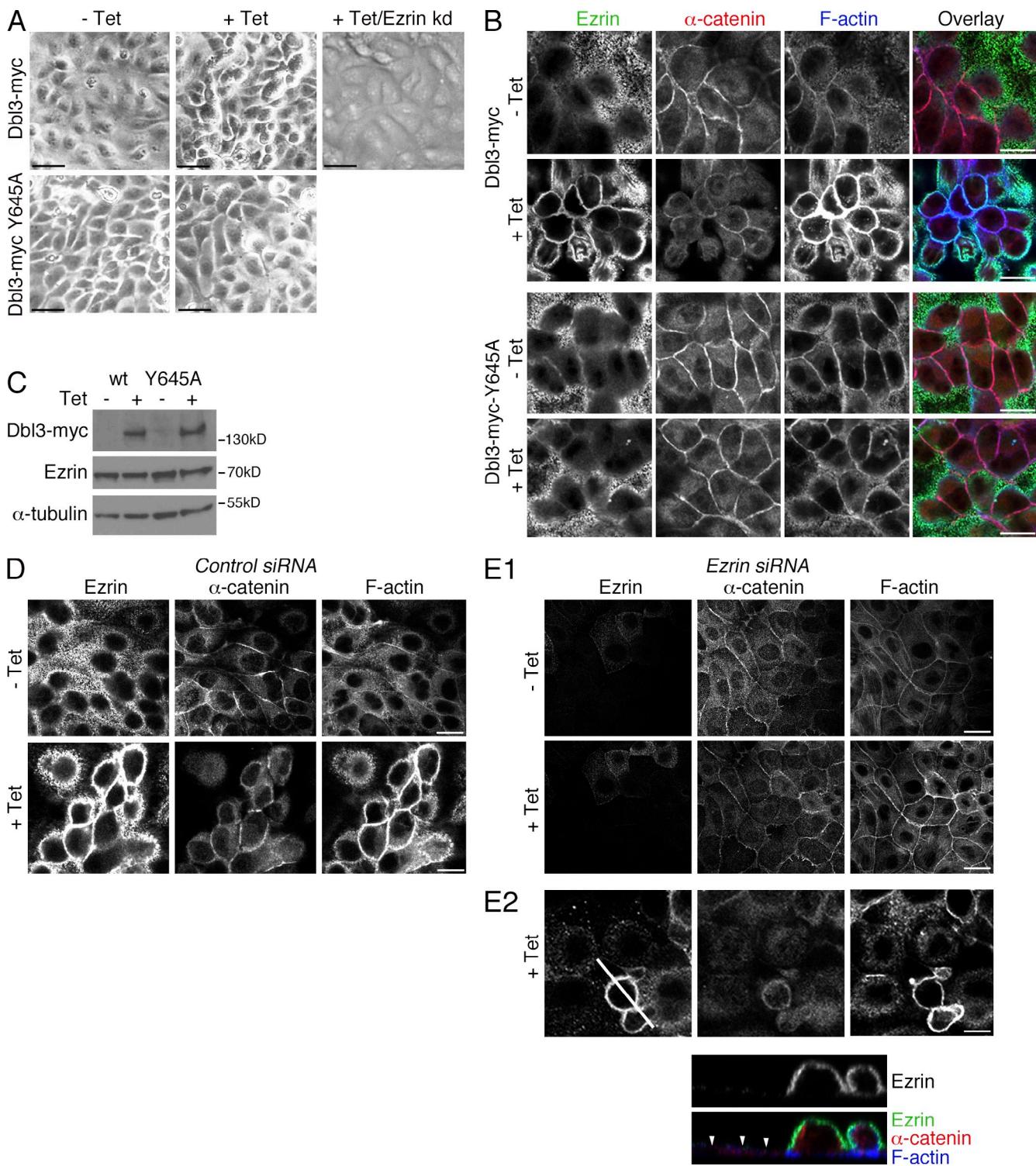
**Figure S2. Analysis of Dbl expression and depletion.** (A) MDCK cell lines stably expressing Dbl-Myc isoforms were processed for immunoblotting using anti-myc antibodies. (B and C) Total cell extracts of Caco-2 cells that had been transfected with control siRNAs and siRNAs targeting all Dbl isoforms or Dbl3 only were immunoblotted with antibodies against DPPIV, ezrin, Crb3, and  $\alpha$ -tubulin. (D and E) Caco-2 cells transfected with siRNAs were processed for immunofluorescence using antibodies against the apical marker DPPIV and the apical polarity regulator Crb3 (D) or ezrin (E). Shown are epifluorescence images. (F) Expression of endogenous Dbl in MDCK cells was analyzed by epifluorescence using antibodies against Dbl and occludin. (G and H) MDCK cells were transfected with siRNAs as indicated and expression of Dbl3 was analyzed by immunoblotting and RT-PCR (G). (H) The effect of Dbl depletion on total levels of active Cdc42 (shown are means  $\pm$  SD,  $n = 4$ ). Bars, 20  $\mu$ m.



**Figure S3. Expression of Dbl isoforms and ezrin analysis.** (A–D) Myc-tagged Dbl isoforms were transiently expressed in MDCK and Caco-2 cells and localized by immunofluorescence using antibodies against the myc tag (A, C, and D) or Dbl (B). (E and F) Caco-2 cells were transfected with siRNAs and processed for immunofluorescence as indicated. Bars: (A, C, and D) 5  $\mu$ m; (B) 10  $\mu$ m; (E and F) 20  $\mu$ m.



**Figure S4. Dbl is required for enrichment of active Cdc42 at apical cell-cell contacts.** (A) Total levels of apical FRET activity after transfection of siRNAs and FRET Cdc42 biosensor (shown are means  $\pm$  SD,  $n \geq 12$  imaged fields). (B) Caco-2 cells were transfected with siRNAs and processed for immunofluorescence using anti-Cdc42-GTP and ZO-1 antibodies. (C and D) Caco-2 cells were transiently transfected with plasmids encoding the indicated constructs. The cells were then extracted for immunoblotting (C) and determination of total levels of active Cdc42 by G-LISA assay (D; shown are means  $\pm$  SD,  $n = 5$ ). Bars, 20  $\mu$ m.



**Figure S5. *Dbl3 activates the Par3–Par6– $\alpha$ PKC pathway.*** (A–C) Conditional MDCK cell lines stably expressing active or mutant Dbl3-myc were analyzed by phase-contrast microscopy [A], immunofluorescence followed by confocal microscopy [B; shown are confocal xy sections taken at the apical–lateral border], or immunoblotting [C]. (D and E) Dbl3-myc MDCK cells were transfected with siRNAs and incubated with tetracycline as indicated before analysis by immunoblotting and immunofluorescence combined with confocal microscopy [xy sections were taken at the apical–lateral border]. (E2) A z scan and the corresponding xy section of an area that contains ezrin-expressing and -depleted cells. Bars: (A) 30  $\mu$ m; (B, D, and E) 10  $\mu$ m.

**Table S1. Alignment of CRAL-TRIO domains.** The CRAL-TRIO domains of DOME and the indicated proteins were aligned using the Eukaryotic Linear Motif (<http://elm.eu.org>). The leucine residue mutated to inactivate the CRAL-TRIO domain of Dbl3 is labelled with an asterisk.

Db13	---QDIAFLS GGR-----G	KDNAWIITFP	-----E NCNFR <i>C</i> IPPE VAKVITYLT SIARQ.----	-----NGSDS
AAD25756/1-0	<b>EDLATAAYMN</b> G-----V	<b>R</b> ESHPVCYNV	<b>H</b> SEELY...Q TIGSEKNRE <i>K</i> FWRWFQLM <i>E</i> KGIQK.----	-LNLKPGGV
A49678/1-0	YYDIAKRQIV EVA----G	<b>D</b> KRKKIIVFS	<b>A</b> CRMPP.----S HOLDHS <i>K</i> LIGYLKHTLD QYVES.----	-DYTLT
00939/1-0	<b>E</b> ATGKIVRS SC-----	<b>V</b> D	<b>P</b> T <sub>*</sub> ASIALFT ARLH.----P PKSAQHV VIQYLVCF <i>L</i> RAVDS.----	-LPPGE
O35239/1-0	<b>E</b> ILSGKFTIL NVR-----	<b>D</b> P	<b>T</b> PT <sub>*</sub> ASIALFT ARLH.----P PKSAQHV VIQYLVCF <i>L</i> RAVDS.----	-FETQRN
CAA15638/1-0	KRTFDYADIL PLP-----GLT	<b>P</b> DCKVWSLYC	<b>F</b> RE <i>E</i> .----S ASKMHHTE DTRAFFMVSD CRFVT.----PD DLAKPDVLSE	
AAF25952/1-0	LLQERVVFLT GGR-----	<b>R</b> RGPPLCFP	<b>T</b> RRDRLKPE DRLRLSYLI SIPSD.----	-AAKNL
NIP2_HUMAN/1-0	YKVVSHGGY YGD-----	<b>G</b> LNAIVVFAVC	<b>F</b> MPE.----S OPNYRVLMDN LFKYVIGTLE LLVAE.----	-NYMIV
CAB55301/1-0	-----AFLS GGR-----G	KDNAWIITFP	-----E NCNFR <i>C</i> IPPE VAKVITYLT SIARQ.----	-----NGSDS
YUQP_CAEEL/1-0	<b>R</b> RKHAPIDII GPQ----RKE	DGRVLVWDR	<b>R</b> AKRDV.----S GLMKSVQPTF YIHEMFRSF <i>E</i> EIQR.----LM KMEAETGVQC	
Q9ZW6/1-0	<b>E</b> LEEVLYQYP OGYH----G	<b>K</b> E	<b>R</b> PVYIER LGKAHP.----S KLMRITTD <i>R</i> YIYHVQE <i>E</i> RALQE. KFPA CSIAAKRRIC	
Q91870/1-0	<b>E</b> LLSGKFTVL PGR-----	<b>E</b> N	<b>A</b> ALALFT ARLH.----P DVDTTHKA VIQAIILYQLD KAIE.----	-VETQRN
O43304/1-0	PPALLEEFYA GGWH----YQ	<b>I</b> D	<b>R</b> PYLILR LGQMDT.----K GLMKAVGEEA LRHVLVSNE EGQKR. CEG STRQLGRPIS	
190854/1-0	<b>T</b> IEAGYPGVL SSR-----	<b>K</b> Y	<b>R</b> VVMLFN IENQ.----S SQEITFDE ILQAYCF <i>L</i> KILEN.----	-EETQIN
AAF02821/1-0	<b>D</b> LDDLQFFTL QGL-----	<b>R</b> S	<b>N</b> IRIFRIV GKYP.----S ARVUSAER LKXYIQS <b>K</b> IS NQCP.----	-GPLCLVY
CAB51563/1-0	<b>E</b> NATGKVLL G-----YD	<b>K</b> D	<b>R</b> PCPLYLY PAR.----Q NTKTSP <i>L</i> QRHLYFSL <i>E</i> CAIDL.----	-MPPGVE
KARI_RAT/1-0	<b>I</b> LKEVKAFVFS GGR-----	<b>K</b> D	<b>G</b> RPILP.----ART.----NHDRIQRE DLRKVLNLD SVPSE.----	-DVCKR
O48940/1-0	<b>E</b> LADQKVFMQ G-----HO	<b>K</b> I	<b>R</b> PILMFV GGR <i>H</i> .----S QNKDGLDE FKRKVYVVLD KVCAS.----	-MPPGQE
O23217/1-0	<b>E</b> GETGKASRA SF-----HO	<b>R</b> O	<b>R</b> VVIMR PAM.----S QNSTSQEG NTRHLVYLL NAIIN.----	-LPKGQK
A53057/1-0	<b>P</b> PDV1QKFMT GGDV----GH	<b>K</b> D	<b>S</b> VLVRIEP WGGLDM.----K GLIMYSCKKD LEKSKLLQ <b>C</b> E KHLKD. LEA QSEKVGKPC	
4437/1-0	<b>E</b> KPLIAKFY P QYH----KT	<b>K</b> D	<b>R</b> PVYFEE LGAVNL.----H EMNKVTSER M KMLNVWEY <i>E</i> SVVQY. RLPA CSRAAGHLVE	
Q23130/1-0	<b>F</b> NDISAHEII QVIA----DG	<b>R</b> V	<b>R</b> PIVVVY AYRIP.----S SKEDIDHAR LQYQLVQIID KIVDQ.----	-DYTV
O48939/1-0	<b>P</b> ADLYRAVRD SQLIGLSGY	<b>R</b> E	<b>L</b> LPVFAIG VGLST.----S FDKASVHY YVQSHIQINE YRERI. ILPS ASKKQGRPIT	
SC14_SCHPO/1-0	<b>E</b> KEAVSKYYP QFYH----KT	<b>I</b> D	<b>R</b> PVYFEE LGNDL.----K KLYQITTPER MMQNLYVYE <i>E</i> MLALK. RFPA CSRKAGGLIE	
CAB63153/1-0	<b>D</b> LEGKAVAYMR G-----YD	<b>K</b> E	<b>R</b> WCYNA YGVEKE(5)R VFGEDEKLNK F RWRVQVLE RGVM.----	-LHFPGGVN
Q22824/1-0	<b>N</b> MSAVAAYFP GGI-----GKS	<b>K</b> R	<b>D</b> VVYMQA MAKAHP.----K TLVKAGPTSQ LFQLCSET <i>E</i> MSFK.----IR QTEQETERKM	
Q17910/1-0	<b>N</b> PIFQKRLMP RGEI-LEKT	<b>N</b> Q	<b>R</b> LLNWYIE YATITV.----E SIASHRSSE ACKFQFLQF <i>E</i> YMLRK.----VM EQEERTGCLS	
DBS_HUMAN/1-0	<b>Q</b> LKQKRFAYLS GGR-----	<b>G</b> Q	<b>D</b> SPVITF DYP.----S AFSEIPDK EFQNVMTLYT SIPSL.----	-QDAGI-
CAB62036/1-0	<b>E</b> TWSSLFLFWH G-----YD	<b>K</b> N	<b>R</b> QPCPLIVR LQHFL.----S KLPSPHERPR FAQAIISQVE HGVLH.----	-LLTPENS
AAF18525/1-0	<b>G</b> SEFEKLVFT H-----GVO	<b>K</b> Q	<b>H</b> VVYISS YGEFON.----KE IFSDKEKLSK FKWRIQFQ <i>E</i> KCVRS.----	-LDFSPPEAKS
O24659/1-0	<b>D</b> LDDDKVYMG G-----AO	<b>R</b> T	<b>R</b> PILLGF PVKNF.----S SAKRDMPK FSKYCVYLLD SICAR.----	-IPRGQE
TPPA_RAT/1-0	<b>L</b> LKAGYHGVL RSR-----	<b>P</b> T	<b>S</b> RVLLYR ISYND.----S PKVETAYD VFRVSLITS LIVQE.----	-VETORN
NF1_DROME	<b>E</b> FKTLKSMNI FYQA----GTS	<b>K</b> S	<b>Y</b> VPVYIY ARRKY.----S IGETNGD LIYHVILTL KPFCH.----	-SPFEVVID
AAD23650/1-0	<b>E</b> IDDEVLKYYP QCYH----G	<b>K</b> D	<b>R</b> PVYIILER LGQVDA.----T KLMQVTTIDR Y KYHVREF <i>E</i> KTFNI. KLPA CSIAAKHHID	
Q93546/1-0	<b>D</b> HPVHHHWKT GLTG--ESGI	<b>I</b> P	<b>N</b> PTIVNIEQ TGSNDY.----W GLMLHSVPTNE IIARHVHDLE SMLKA.----VM DLEKKTNQCC	
Q22467/1-0	<b>D</b> CSVPLRYLP GSLI-----GL	<b>H</b> E	<b>N</b> ENVVSLQM IGHADA.----A GLMPATRNSD LYRMRIAES GVMI.----IR KMEEQGKPL	
Q06705/1-0	<b>N</b> LELOKATIQ G-----YD	<b>N</b> D	<b>M</b> RPVILVR PRLLH.----S SDQTE <i>Q</i> LEKFSLLV <b>E</b> QSKLF.----	-FKENYPA
081806/1-0	<b>L</b> IEKEIFKI HGR-----	<b>K</b> R	<b>K</b> RKILRII GKKF.----S ARFLSLDV LKLYKEEKIF PRLGR.----	-KPFAVLY
AAF19259/1-0	<b>P</b> PEVIQLYDS GGLC-----GY	<b>Y</b> E	<b>C</b> PVYFN1 IGSLLDP.----K GLLLSASKQD MRKRIKVC <i>E</i> LLLHE. CEL QTQKLGRKIE	
Q9Z1J8/1-0	<b>P</b> PEVIQKYMP GGLC-----GY	<b>R</b> D	<b>C</b> PVWYDI IGPPLDP.----K GLLFSTVKQD LKTKMRDC <i>E</i> RILHE. CDL QTERLGRKIE	
076636/1-0	<b>D</b> HPLIKKYFP LGLVG----ETG	<b>K</b> D	<b>Q</b> NLQLLVIC <i>E</i> AGRIDL.----M GILKSVLHSD FIQIQRFFQ <i>E</i> KMLAA.----MN EMERKYGTQ	
O15915/1-0	<b>I</b> REIGSAGCV YVN-----KR	<b>K</b> K	<b>R</b> PIIIFAV PRNDT.----S LKNVPSEL KFKNLVYVWL <i>E</i> QGFSR.----	-MDEPKGIE
YNX1_YEAST/1-0	<b>E</b> NESGKQVIL G-----YE	<b>N</b> D	<b>A</b> RPILYLYK PGR.----S QNTKTSHR QVQHLVFM <i>E</i> RVIDF.----	-MPAGQD
Q01324/1-0	<b>E</b> DEELNKYVPE IDIVG-QNHQ	<b>D</b> D	<b>N</b> DKVNLMPER TGKID <i>E</i> S GLVDNVLHM FMQIKLKM <i>E</i> GVHQK.----VV AAERKTGRQS	
BAA90344/1-0	<b>E</b> AETGKIVRA DY-----K	<b>K</b> H	<b>R</b> TVLVLV PGL.----S ENTSGKG QKLYVYCYCLE KAIMS.----	-LTEDQE
Q9ZW01/1-0	<b>E</b> LAEGLAYVA G-----D	<b>D</b> C	<b>R</b> CPVLFVFR IKQDY.----S QKLHTQKQ LTRLVVFTE L VAIST.----	-MSRNVE
BAA90976/1-0	<b>D</b> IASLAKALYQ T-----GV	<b>N</b> C	<b>R</b> RTVMMVV GRN <i>E</i> .----VTLIDDM <i>D</i> A LYF1HVM <i>D</i> HIAVK.----EYVLV	
061528	<b>V</b> LRDGIAVLP GGR-----	<b>R</b> A	<b>Q</b> AVIVCP SRE <i>E</i> Q.----S VNQDNLN RFLYLF <i>E</i> VT KMAR.----	-K
YN02_CAEEL/1-0	<b>R</b> PVIKQYFP GCWH----NS	<b>K</b> A	<b>R</b> PMYILR FGQLDT.----K GLMLRSCGVEN LVKLTISICE DGLQR. AAE ATRKLGRPIS	
YJ05_YEAST/1-0	<b>N</b> TELQNVGIL TFDA----NG	<b>A</b> N	<b>K</b> AKAVTWNL YGQIV.----K KKELFQNVDK FVYR <i>E</i> GLM <i>E</i> KGLSL.----	-LDFTSSDNN
YQF6_CAEEL/1-0	<b>S</b> IAAPPFLQFIA SSRL(4)WS	<b>D</b> S	<b>H</b> NAFLFVER AWSQP.----K EFIKTFKTS <i>D</i> YLHCFGYS <i>E</i> MLQQLLIRRE KKQSADKGPV	
O17907/1-0	<b>P</b> PECLEKYYCG YGLL-----GD	<b>T</b> E	<b>R</b> PILMSL LGNVDV.----E GLLRSVASLD YIKFSIAAI <i>E</i> KGML. CEE KAKESGRPFE	
5824562/1-0	<b>N</b> HPIHKHWKY GITG--PSGH	<b>M</b> D	<b>N</b> DIVIVNIEQ CGKTDY.----T GMMETYSILE VMRARMDL <i>E</i> QMLHH.----VM ELEAKTGKQA	
P78783/1-0	<b>N</b> FKVAKSMYFI WGO-----	<b>K</b> K	<b>R</b> AIYVFLN LHNP <i>E</i> IP.----S PKNTKDM <i>E</i> LKALILYAME NARLF.----LDSEQNAAK	
YN04_YEAST/1-0	<b>E</b> NETGKQVIL GF-----D	<b>N</b> A	<b>R</b> KRPLYYMK NGR.----S QNTESSFR QVQELVYMM <i>E</i> TATTV.----	-APQGVE
YAS2_SCHPO/1-0	<b>D</b> DDFVRQLRI GKCFI-FGE	<b>K</b> H	<b>N</b> RPVCYIR ARLH.----S VGDVSPE VERLTWWM <i>E</i> TARLI.----	-LKPPIE
NF1_HUMAN	<b>E</b> FKALKTLSI FYQA----GTS	<b>K</b> A	<b>N</b> PIFIYVV ARRFK.----S TGQINGD LIYHVLLTL KPYYA.----	-KPYEI-
Q9XVU9/1-0	<b>G</b> KDCPERFH SQLN-YEVT SKNLHVVFQK MEGIDI.----K EILKVMPLSY VHHSYFML <i>E</i> NFSRA. MA HTERKTGKTS			
Consensus/60%	pbchhpbbb. sh.....D +pGpslbbb. hspb.....pshs.pc b1-bb1bbbe phbp. ....s..bp			

Db13 RFTI<sub>1</sub>LD<sub>2</sub>RRL DT----... - --WSSLKIS LQK<sub>1</sub>SA .SPE GNLHLV<sub>1</sub>LVR PTSF--LQRT FTDIGFWESQ EDFMLKLPVV  
 AAD25756/1-0 SLLQ<sub>1</sub>HD<sub>2</sub>LKN APG<sub>1</sub>VSR... - TEIWVG<sub>1</sub>IKKV IET<sub>1</sub>QD.NYP EFVSRNIFIN VFW--FYA MRAVLSPELT QRTKS<sub>1</sub>KFVVA  
 A49678/1-0 YLHHLG<sub>1</sub>TS<sub>2</sub>DN KPS---- - --LSWL<sub>1</sub>RDA YREF<sub>1</sub>DR.KYK KNIAKALYIVH PTM---FIK<sub>1</sub> LLLIFKPLIS FKFG<sub>1</sub>KIFYV  
 T00939/1-0 QMVWM<sub>1</sub>ID<sub>2</sub>FHG YSIAN---- - VSLRTT<sub>1</sub>KET AHV<sub>1</sub>QE.HY<sub>1</sub> ERLAFAVLYN PCK---FFEP FWKVARP<sub>1</sub>ELE PKTRNKVKFV  
 O35239/1-0 GLVFIY<sub>1</sub>DMCG SNYAN---- - FELDLG<sub>1</sub>KKV LNL<sub>1</sub>KG.AFP ARLK<sub>1</sub>VLIVR AIIW---FRV PY<sub>1</sub>SII<sub>1</sub>LLK DKVRERIPIL  
 CAA15638/1-0 GEVO<sub>1</sub>IF<sub>2</sub>DMKG TTMRHI..S RLTISTL<sub>1</sub>RAY IKF<sub>1</sub>QL.AFP VRLRAIHMIN CQT---YLDR IVSVVKP<sub>1</sub>EIS DEVFKLIRFH  
 AAF25952/1-0 GFTVI<sub>1</sub>DMRG NG----... - NCSTNAKTI LKV<sub>1</sub>QE.HFS ANIHN<sub>1</sub>VIIK PDN----F WQKQRASISS HKY<sub>1</sub>KETTTV  
 NIP2\_HUMAN/1-0 YLN<sub>1</sub>GATT<sub>2</sub>RK MPS---- - --LGWL<sub>1</sub>RKC YQQ<sub>1</sub>DR.RLR KNL<sub>1</sub>KSLIIIVH PSW---FIR<sub>1</sub> LLAVTRE<sub>1</sub>EIS SKFSC<sub>1</sub>KIRV  
 CAB55301/1-0 RFTI<sub>1</sub>LD<sub>2</sub>RRL DT----... - --WSSLKIS LQK<sub>1</sub>SA .SPE GNLHLV<sub>1</sub>LVR PTSF--LQRT FTDIGFWESQ EDFMLKLPVV  
 YUQP\_CAEEL/1-0 YM<sub>1</sub>H<sub>2</sub>YF<sub>3</sub>DL<sub>4</sub>EA LNF<sub>1</sub>DPT..LL GVNVGP<sub>2</sub>FRVS T WOL<sub>1</sub>QG.HY<sub>1</sub> ERL<sub>1</sub>DF<sub>2</sub>IVIN S<sub>3</sub>S---YINV LWSALSPELP EQSKQ<sub>1</sub>RIVFA  
 Q9ZWD6/1-0 STT<sub>1</sub>LD<sub>2</sub>DVQG LG<sub>1</sub>KNF..T PTAANLVAA<sub>1</sub>M SKIDNS.YYP ET<sub>1</sub>HLRMYIVN AGTG---FKKM LWPA<sub>1</sub>AQ<sub>2</sub>KELD AKTIA<sub>1</sub>KHVL  
 Q91870/1-0 GLIFIY<sub>1</sub>DMIN SSYGN---- - FDYELCV<sub>1</sub>KI LNL<sub>1</sub>KG.AFP ARLK<sub>1</sub>C<sub>2</sub>FI<sub>3</sub>VS SLW---FRA PFAVRLF<sub>2</sub>VR EKL<sub>1</sub>RR<sub>2</sub>VRTV  
 O43304/1-0 SWTC<sub>1</sub>LD<sub>2</sub>DL<sub>3</sub>EG LNMRH<sub>1</sub>..W RPGV<sub>1</sub>KALLRM IEV<sub>1</sub>WED.NYP ET<sub>1</sub>LG<sub>2</sub>FL<sub>3</sub>IP<sub>4</sub> FVPP<sub>1</sub> LWL<sub>1</sub>LI<sub>2</sub>SP<sub>3</sub>EIN ENTRRK<sub>2</sub>FLIY  
 190854/1-0 GFCI<sub>1</sub>EN<sub>2</sub>FKG FT<sub>3</sub>MQA...A SLRTSD<sub>1</sub>RKL VDML<sub>1</sub>QD.SPF ARFKAHF<sub>1</sub>H QW---YFTT TYNVVKP<sub>2</sub>ELK SKLLE<sub>1</sub>RVFVH  
 AAF02821/1-0 MHSTVQ<sub>1</sub>KDDN SPG---- - --ITIL<sub>1</sub>RWI YEDL<sub>1</sub>PS.DIK DRLQLVY<sub>1</sub>FH PG<sub>1</sub>RLR-SRLV IATLGR<sub>1</sub>LLS GGLYWK<sub>1</sub>KYV  
 CAB51563/1-0 TLAL<sub>1</sub>Y<sub>2</sub>INF<sub>3</sub>S SSNRS---- - NP<sub>1</sub>SVGQ<sub>2</sub>GKEV<sub>3</sub> LN<sub>1</sub>IQ<sub>2</sub>T.HYC ERLG<sub>1</sub>RAL<sub>2</sub>VIN IEM---AVWG FF<sub>1</sub>KL<sub>2</sub>ISE<sub>3</sub>PID PITREK<sub>1</sub>KFN  
 KARI\_RAT/1-0 GFTVI<sub>1</sub>DMRG SK----... - --WDL<sub>1</sub>IKPL LKT<sub>1</sub>QE.AFP AEI<sub>1</sub>HVAL<sub>2</sub>IK PDN----F WQKQ<sub>1</sub>KTN<sub>2</sub>EGS SKF<sub>1</sub>IFETSMV  
 O48940/1-0 KFGV<sub>1</sub>AEL<sub>2</sub>KG WG<sub>3</sub>YSN---- - SDVG<sub>1</sub>YLSA LS<sub>1</sub>IQ<sub>2</sub>OD.NYP ERLG<sub>1</sub>KLF<sub>2</sub>IVN AY<sub>3</sub>--IFM<sub>1</sub> VWQ<sub>1</sub>VY<sub>2</sub>PEID NKT<sub>1</sub>KK<sub>2</sub>I<sub>3</sub>VFV  
 O23217/1-0 QMSWL<sub>1</sub>ID<sub>2</sub>FTG WSM<sub>3</sub>AV---- - NPM<sub>1</sub>MKT<sub>2</sub>TREI IH<sub>1</sub>IQ<sub>2</sub>ON.YYP ERLG<sub>1</sub>IA<sub>2</sub>FLY<sub>3</sub>N PER---LFQA VYRAAKY<sub>2</sub>ELD PRTAEK<sub>1</sub>WKFV  
 A53057/1-0 GLTV<sub>1</sub>Y<sub>2</sub>DM<sub>3</sub>N HG<sub>4</sub>SKHM..W KPG<sub>1</sub>LDM<sub>2</sub>YL VQ<sub>1</sub>LED.NYP EMMK<sub>1</sub>RL<sub>2</sub>FV<sub>3</sub>N ACT---LFPV LYK<sub>1</sub>IV<sub>2</sub>KP<sub>3</sub>LLS EDM<sub>1</sub>KNK<sub>2</sub>I<sub>3</sub>FV  
 4437/1-0 TSCT<sub>1</sub>IM<sub>2</sub>DL<sub>3</sub>KG IS<sub>4</sub>I<sub>1</sub>SSA... - YSVM<sub>1</sub>SY<sub>2</sub>RE<sub>3</sub> AY<sub>4</sub>SI<sub>1</sub>Q<sub>2</sub>SON.YYP ERMGK<sub>1</sub>FY<sub>2</sub>IVN AEPG---FST AFR<sub>1</sub>LF<sub>2</sub>KE<sub>3</sub>ELD PVT<sub>1</sub>VS<sub>2</sub>K<sub>3</sub>I<sub>4</sub>FIL  
 Q23130/1-0 YFHYGLRS<sub>1</sub>HN K----... - PPVRWLFQ<sub>1</sub>A YKQ<sub>1</sub>DR.RPK KN<sub>1</sub>KL<sub>2</sub>ALYVHH PTR---FIR<sub>1</sub> IFS<sub>2</sub>LF<sub>3</sub>K<sub>4</sub>EIS SKF<sub>1</sub>EN<sub>2</sub>K<sub>3</sub>FHYV  
 O48939/1-0 TC<sub>1</sub>KV<sub>2</sub>LD<sub>3</sub>MTG LK<sub>4</sub>LSAL.. - - NQ<sub>1</sub>IKL<sub>2</sub>TII SS<sub>1</sub>DDL<sub>2</sub>ND<sub>3</sub>NYP EKTNT<sub>1</sub>YY<sub>2</sub>IVN AY<sub>3</sub>--IF<sub>1</sub>DSA CWKVVKP<sub>2</sub>LLQ ERT<sub>1</sub>RR<sub>2</sub>K<sub>3</sub>IQL  
 SC14\_SCHPO/1-0 TSCT<sub>1</sub>IM<sub>2</sub>DL<sub>3</sub>KG VG<sub>4</sub>ITSI... - HSV<sub>1</sub>SY<sub>2</sub>IR<sub>3</sub> AY<sub>4</sub>SI<sub>1</sub>Q<sub>2</sub>OD.NYP ERLG<sub>1</sub>IA<sub>2</sub>FLY<sub>3</sub>N AEWG---FSS AFNL<sub>1</sub>KG<sub>2</sub>ELD EATVKK<sub>3</sub>I<sub>4</sub>HIL  
 CAB63153/1-0 SI<sub>1</sub>IQ<sub>2</sub>TD<sub>3</sub>L<sub>4</sub>K MPK---- - RELRV<sub>1</sub>AS<sub>2</sub>NQI LSL<sub>1</sub>QD.NYP ELV<sub>1</sub>AT<sub>2</sub>K<sub>3</sub>IF<sub>4</sub> VEW---YFSV IYSMFSPELT QRT<sub>1</sub>SK<sub>2</sub>FVMS  
 Q22824/1-0 GVII<sub>1</sub>IM<sub>2</sub>DL<sub>3</sub>DG FS<sub>4</sub>M<sub>1</sub>LL..Y TPTLK<sub>1</sub>Y<sub>2</sub>MSL LTM<sub>1</sub>Q<sub>2</sub>N.IFP D<sub>3</sub>F<sub>4</sub>PAR<sub>1</sub>Y<sub>2</sub>I<sub>3</sub>IN CGA---MMSA VYAMVSPVLS SQ<sub>1</sub>TR<sub>2</sub>E<sub>3</sub>KV<sub>4</sub>---  
 Q17910/1-0 SLRH<sub>1</sub>IV<sub>2</sub>NMD<sub>3</sub>G YE<sub>4</sub>INPFT.MV FV<sub>1</sub>TSG<sub>2</sub>TL<sub>3</sub>AYY S<sub>4</sub>QL<sub>1</sub>HF<sub>2</sub>EN<sub>3</sub>Y<sub>1</sub>P E<sub>2</sub>LT<sub>1</sub>VP<sub>2</sub>DMVN IA<sub>3</sub>K---I<sub>4</sub>HV PYX<sub>1</sub>IA<sub>2</sub>KAMMP TG<sub>3</sub>SE<sub>4</sub>KFRLH  
 DBS\_HUMAN/1-0 GFLV<sub>1</sub>Y<sub>2</sub>DR<sub>3</sub>R DK----... - --WTSV<sub>1</sub>KAS VL<sub>2</sub>RA<sub>3</sub>A<sub>4</sub> S<sub>1</sub>RL<sub>2</sub>G<sub>3</sub>CF<sub>4</sub>IR L<sub>1</sub>EP---VVRV ISQTF<sub>1</sub>Q<sub>2</sub>IL<sub>3</sub>R P<sub>4</sub>TR<sub>1</sub>KK<sub>2</sub>L<sub>3</sub>RIE  
 CAB62036/1-0 ELTV<sub>1</sub>LV<sub>2</sub>DC<sub>3</sub>RG LSPLR---- - -IPMQM<sub>1</sub>MR<sub>2</sub>C SS<sub>3</sub>IQ<sub>4</sub>OD.HFP NRLG<sub>1</sub>CL<sub>2</sub>FI<sub>3</sub>IR L<sub>4</sub>EP---VVRV ISQTF<sub>1</sub>Q<sub>2</sub>IL<sub>3</sub>R P<sub>4</sub>TR<sub>1</sub>KK<sub>2</sub>L<sub>3</sub>RIE  
 AAF18525/1-0 SFVF<sub>1</sub>W<sub>2</sub>DF<sub>3</sub>FN APGLQ<sub>4</sub>... - - RALQ<sub>1</sub>Q<sub>2</sub>FI<sub>3</sub>KRA V<sub>4</sub>KQ<sub>1</sub>FD<sub>2</sub>N Y<sub>1</sub>P E<sub>2</sub>F<sub>3</sub>V<sub>4</sub>AK<sub>1</sub>EL<sub>2</sub>IN VEW<sub>1</sub>WW--YIP<sub>2</sub> YK<sub>3</sub>T<sub>4</sub>FG<sub>1</sub>SI<sub>2</sub>TS P<sub>3</sub>TR<sub>1</sub>SK<sub>2</sub>MVLS  
 024659/1-0 KF<sub>1</sub>VC<sub>2</sub>VD<sub>3</sub>L<sub>4</sub>KG WG<sub>1</sub>YSN---- - -CD<sub>1</sub>RAY<sub>2</sub>IA IEIM<sub>1</sub>QN.YYP ERLG<sub>1</sub>KALM<sub>2</sub>H V<sub>3</sub>Y<sub>4</sub>--MFM<sub>1</sub> AWK<sub>2</sub>MI<sub>3</sub>Y<sub>4</sub>ED NVT<sub>1</sub>RD<sub>2</sub>K<sub>3</sub>FV<sub>4</sub>V  
 TTPA\_RAT/1-0 GVKA<sub>1</sub>IF<sub>2</sub>DL<sub>3</sub>E<sub>4</sub>G WQ<sub>1</sub>ISHA...F QITPSV<sub>2</sub>AK<sub>3</sub>KI AAV<sub>1</sub>TD.S<sub>2</sub>FP<sub>3</sub> LKVR<sub>1</sub>GH<sub>2</sub>HL<sub>3</sub>IN E<sub>4</sub>V---IF<sub>1</sub>HA VFSM<sub>2</sub>K<sub>3</sub>PELT EKK<sub>1</sub>GR<sub>2</sub>I<sub>3</sub>HLH  
 NF1<sub>1</sub>\_DROME FTHTC<sub>2</sub>SD<sub>3</sub>NR<sub>4</sub>F R----... - -TEFL<sub>1</sub>Q<sub>2</sub>KW FY<sub>1</sub>VP.T.VAY ENV<sub>1</sub>H<sub>2</sub>AV<sub>3</sub>Y<sub>4</sub>IN CNS---WVR EY<sub>1</sub>TK<sub>2</sub>F<sub>3</sub>DR<sub>4</sub>L APL<sub>1</sub>GNR<sub>2</sub>KLL  
 AAD23650/1-0 QST<sub>1</sub>IL<sub>2</sub>D<sub>3</sub>VQG V<sub>4</sub>GL<sub>1</sub>KS<sub>2</sub>F..S KAARD<sub>1</sub>LL<sub>2</sub>QRI<sub>3</sub> QK<sub>1</sub>SD<sub>2</sub>.NYP ET<sub>1</sub>LN<sub>2</sub>RM<sub>3</sub>FI<sub>4</sub>IN AGSG---FRL LW<sub>1</sub>STV<sub>2</sub>K<sub>3</sub>ELD P<sub>4</sub>TT<sub>1</sub>AK<sub>2</sub>H<sub>3</sub>V<sub>4</sub>L  
 Q93546/1-0 SVIY<sub>1</sub>IM<sub>2</sub>DL<sub>3</sub>TG IKED<sub>4</sub>DKR..T<sub>1</sub> TLT<sub>2</sub>GG<sub>3</sub>LSAI SAF<sub>1</sub>ME<sub>2</sub> HY<sub>3</sub> ELV<sub>1</sub>H<sub>2</sub>S<sub>3</sub>V<sub>4</sub>L VN<sub>1</sub>A---FISA IWT<sub>2</sub>IA<sub>3</sub>K<sub>4</sub>PLP ERT<sub>1</sub>R<sub>2</sub>N<sub>3</sub>CN<sub>4</sub>L  
 Q22467/1-0 GT<sub>1</sub>S<sub>2</sub>V<sub>3</sub>IF<sub>4</sub>DL<sub>1</sub>DG LSM<sub>2</sub>MQ<sub>3</sub>QI..D LAALK<sub>1</sub>V<sub>2</sub>V<sub>3</sub>TTM LSQ<sub>1</sub>Q<sub>2</sub>OE.MFP DV<sub>1</sub>IR<sub>2</sub>K<sub>3</sub>IF<sub>4</sub>VN T<sub>1</sub>T---F<sub>2</sub>IQV LWSM<sub>3</sub>IS<sub>4</sub>PC<sub>1</sub>LA KQTQ<sub>2</sub>Q<sub>3</sub>K<sub>4</sub>V<sub>1</sub>K<sub>2</sub>  
 Q06705/1-0 STT<sub>1</sub>IL<sub>2</sub>FD<sub>3</sub>LN<sub>4</sub>GS FMSN---- - -MDY<sub>1</sub>AP<sub>2</sub>V<sub>3</sub>KFL ITC<sub>1</sub>FEA.HY<sub>2</sub>P E<sub>3</sub>SL<sub>1</sub>GH<sub>2</sub>LL<sub>3</sub>I<sub>4</sub>HK AQW---IFNP INWN<sub>1</sub>IK<sub>2</sub>NWLD PVV<sub>3</sub>ASK<sub>4</sub>IVFT  
 081806/1-0 VHTGV<sub>1</sub>Q<sub>2</sub>RS<sub>3</sub>EN FP<sub>4</sub>G---- - -IS<sub>1</sub>AL<sub>2</sub>RA<sub>3</sub>I YD<sub>1</sub>PA<sub>2</sub>PN<sub>3</sub>V<sub>4</sub>NR DN<sub>1</sub>LQ<sub>2</sub>EY<sub>3</sub>V<sub>4</sub>F<sub>1</sub>H PG<sub>2</sub>Q---SRL<sub>3</sub> LAT<sub>1</sub>Q<sub>2</sub>RF<sub>3</sub>LFS GGLY<sub>1</sub>GK<sub>2</sub>LR<sub>3</sub>Y<sub>4</sub>I  
 AAF19259/1-0 MALM<sub>1</sub>Y<sub>2</sub>DM<sub>3</sub>EG LSL<sub>1</sub>KH<sub>2</sub>L..W KPAVEVY<sub>3</sub>Q<sub>4</sub>QF FS<sub>1</sub>IL<sub>2</sub>EA.NYP E<sub>2</sub>TL<sub>1</sub>KN<sub>2</sub>L<sub>3</sub>IV<sub>4</sub>R P<sub>1</sub>AK<sub>2</sub>LF<sub>3</sub>SEMS EBT<sub>1</sub>RR<sub>2</sub>K<sub>3</sub>I<sub>4</sub>VIL  
 Q9Z138/1-0 TIVM<sub>1</sub>Y<sub>2</sub>DF<sub>3</sub>CE<sub>4</sub>G L<sub>1</sub>G<sub>2</sub>IK<sub>3</sub>HF..W K<sub>4</sub>PL<sub>1</sub>VEVY<sub>2</sub>Q<sub>3</sub>EF FG<sub>1</sub>LL<sub>2</sub>BE.NYP E<sub>2</sub>TL<sub>1</sub>K<sub>2</sub>FL<sub>3</sub>IV<sub>4</sub>K<sub>1</sub>W<sub>2</sub>RL<sub>3</sub> APL<sub>1</sub>GNR<sub>2</sub>KLL  
 076636/1-0 SVIY<sub>1</sub>Y<sub>2</sub>LD<sub>3</sub>L<sub>4</sub>R L<sub>1</sub>K<sub>2</sub>F<sub>3</sub>DPA..L<sub>4</sub> SI<sub>1</sub>VTG<sub>2</sub>PY<sub>3</sub>RI<sub>4</sub> WAS<sub>1</sub>Y<sub>2</sub>T.Y<sub>3</sub>P EW<sub>1</sub>INT<sub>2</sub>FL<sub>3</sub>LIN<sub>4</sub> AS---FMTL LW<sub>1</sub>KA<sub>2</sub>IG<sub>3</sub>PL<sub>4</sub>L ERT<sub>1</sub>TR<sub>2</sub>N<sub>3</sub>K<sub>4</sub>RC  
 O15915/1-0 QFC<sub>1</sub>F<sub>2</sub>IV<sub>3</sub>Y<sub>4</sub>D<sub>1</sub>K FGSGN---- - -MDM<sub>1</sub>KTN<sub>2</sub>LEA M<sub>3</sub>HFLD.RCP ERMGQS<sub>1</sub>FL<sub>2</sub>PD<sub>3</sub>A P<sub>4</sub>A<sub>1</sub>W<sub>2</sub>L<sub>3</sub>FE<sub>4</sub>LN<sub>1</sub>EVT<sub>2</sub>LS<sub>3</sub>K<sub>4</sub>RF<sub>1</sub>I  
 YNX1\_YEAST/1-0 SL<sub>1</sub>LL<sub>2</sub>AI<sub>3</sub>DF<sub>4</sub>K YPD<sub>1</sub>VPK<sub>2</sub>(5)K IPP<sub>1</sub>IGVG<sub>2</sub>KEV<sub>3</sub> L<sub>4</sub>H<sub>1</sub>IQ<sub>2</sub>T.HY<sub>3</sub> ERLG<sub>1</sub>K<sub>2</sub>AL<sub>3</sub>LT<sub>4</sub>N IW<sub>1</sub>W<sub>2</sub>--LAW<sub>3</sub> FL<sub>1</sub>KL<sub>2</sub>I<sub>3</sub>H<sub>4</sub>PID PLT<sub>1</sub>RE<sub>2</sub>K<sub>3</sub>L<sub>4</sub>VFD  
 001324/1-0 GGLF<sub>1</sub>IM<sub>2</sub>DL<sub>3</sub>DG IS<sub>4</sub>E<sub>1</sub>SP<sub>2</sub>K..L<sub>3</sub> LI S<sub>1</sub>V<sub>2</sub>L<sub>3</sub>GP<sub>4</sub>Y<sub>1</sub>RIM WGT<sub>1</sub>FD<sub>2</sub>D<sub>3</sub>Y<sub>4</sub> QL<sub>1</sub>Q<sub>2</sub>KK<sub>3</sub>Y<sub>4</sub>IN A<sub>1</sub>PS<sub>2</sub> F<sub>3</sub>Q<sub>4</sub>Y<sub>1</sub>AC<sub>2</sub>PE<sub>3</sub>EL<sub>4</sub> P<sub>1</sub>EDY<sub>2</sub>KE<sub>3</sub>K<sub>4</sub>IVIT  
 BAA90344/1-0 KM<sub>1</sub>WL<sub>2</sub>U<sub>3</sub>DF<sub>4</sub>QS WT<sub>1</sub>LG<sub>2</sub>S<sub>3</sub>... - -T<sub>1</sub>PL<sub>2</sub>K<sub>3</sub>V<sub>4</sub>RET V<sub>1</sub>N<sub>2</sub>Q<sub>3</sub>D.C<sub>4</sub>YP ERL<sub>1</sub>GL<sub>2</sub>AI<sub>3</sub>LY<sub>4</sub>IN PER---IFES FW<sub>1</sub>K<sub>2</sub>V<sub>3</sub>K<sub>4</sub>PEL<sub>1</sub> HET<sub>2</sub>Y<sub>3</sub>KK<sub>4</sub>V<sub>1</sub>F<sub>2</sub>V<sub>3</sub>  
 Q9ZW01/1-0 QF<sub>1</sub>VL<sub>2</sub>IF<sub>3</sub>DA<sub>4</sub>A H----... - -AV<sub>1</sub>L<sub>2</sub>V<sub>3</sub>TL<sub>4</sub>IV L<sub>1</sub>F<sub>2</sub>S<sub>3</sub>FL<sub>4</sub>.NYP LL<sub>1</sub>LC<sub>2</sub>NST<sub>3</sub>HA<sub>4</sub> LF<sub>1</sub>F<sub>2</sub>FFFF FIY<sub>1</sub>KT<sub>2</sub>NP<sub>3</sub>LL QIS<sub>1</sub>SFH--  
 BAA90976/1-0 YF<sub>1</sub>HT<sub>2</sub>LT<sub>3</sub>SE<sub>4</sub>YN H----... - -LDSDF<sub>1</sub>L<sub>2</sub>K<sub>3</sub>KL<sub>4</sub> YD<sub>1</sub>V<sub>2</sub>UV<sub>3</sub>D<sub>4</sub>Y<sub>1</sub>K<sub>2</sub>Y<sub>3</sub> RNL<sub>1</sub>K<sub>2</sub>AV<sub>3</sub>Y<sub>4</sub>F<sub>1</sub>B<sub>2</sub>T<sub>3</sub>PSV S<sub>1</sub>G<sub>2</sub>LD<sub>3</sub>K<sub>4</sub>IHHV  
 061528 GFLV<sub>1</sub>Y<sub>2</sub>DL<sub>3</sub>RG L<sub>4</sub>LSMRH<sub>1</sub>..W RPGVQ<sub>2</sub>CL<sub>3</sub>KI IEI<sub>1</sub>VEA.NYP ET<sub>1</sub>STV<sub>2</sub>Q<sub>3</sub>FL<sub>4</sub>IX<sub>1</sub> PEK---F WE<sub>1</sub>Q<sub>2</sub>KAQMSL CTW<sub>3</sub>DE<sub>4</sub>PE<sub>1</sub>EMI  
 YN02\_CAEEL/1-0 SW<sub>1</sub>SL<sub>2</sub>V<sub>3</sub>DD<sub>4</sub>L<sub>1</sub>GS L<sub>2</sub>LSMRH<sub>1</sub>..W RPGVQ<sub>2</sub>CL<sub>3</sub>KI IEI<sub>1</sub>VEA.NYP ET<sub>1</sub>STV<sub>2</sub>Q<sub>3</sub>FL<sub>4</sub>IX<sub>1</sub> PEK---F WE<sub>1</sub>Q<sub>2</sub>KAQMSL CTW<sub>3</sub>DE<sub>4</sub>PE<sub>1</sub>EMI  
 YJ05\_YEAST/1-0 YMTQ<sub>1</sub>Y<sub>2</sub>HD<sub>3</sub>Y<sub>4</sub>K<sub>1</sub> G<sub>2</sub>SV<sub>3</sub>WRM<sub>4</sub>..D SDIKNC<sub>1</sub>NS<sub>2</sub>KT<sub>3</sub>IV<sub>4</sub> IGI<sub>1</sub>FQ<sub>2</sub>K<sub>3</sub>Y<sub>4</sub>E<sub>1</sub>P ELLY<sub>2</sub>Q<sub>3</sub>AK<sub>4</sub>Y<sub>1</sub>FV<sub>2</sub> V<sub>3</sub>T<sub>4</sub>--VFG<sub>1</sub> VY<sub>2</sub>DL<sub>3</sub>KK<sub>4</sub>V<sub>1</sub>V<sub>2</sub>V<sub>3</sub>  
 YQF6\_CAEEL/1-0 QF<sub>1</sub>IV<sub>2</sub>IF<sub>3</sub>DL<sub>4</sub>NT V<sub>1</sub>N<sub>2</sub>TD<sub>3</sub>V<sub>4</sub>NPM SGY<sub>1</sub>M<sub>2</sub>KL<sub>3</sub>W<sub>4</sub>Q<sub>1</sub> SEL<sub>2</sub>QD.W<sub>3</sub>F<sub>4</sub>P EMVQ<sub>1</sub>RI<sub>2</sub>Y<sub>3</sub>LT<sub>4</sub>N P<sub>1</sub>DR<sub>2</sub>--LL<sub>3</sub>GL<sub>4</sub> LW<sub>1</sub>K<sub>2</sub>VAR<sub>3</sub>VEL<sub>4</sub>S EENLKR<sub>1</sub>IEII  
 O17907/1-0 QMTL<sub>1</sub>Y<sub>2</sub>FD<sub>3</sub>LN<sub>4</sub> ITS<sub>1</sub>A<sub>2</sub>H..SC KQ<sub>1</sub>F<sub>2</sub>ASS<sub>3</sub>FT<sub>4</sub>TL V<sub>1</sub>S<sub>2</sub>L<sub>3</sub>QD.H<sub>4</sub>F<sub>1</sub> YFL<sub>2</sub>R<sub>3</sub>K<sub>4</sub>LI<sub>1</sub>IR A<sub>2</sub>E<sub>3</sub>M<sub>4</sub> ARIA YAS<sub>1</sub>ST<sub>2</sub>AI<sub>3</sub>L<sub>4</sub>QD P<sub>1</sub>TR<sub>2</sub>L<sub>3</sub>VE<sub>4</sub>MP<sub>1</sub>  
 5824562/1-0 WIL<sub>1</sub>Y<sub>2</sub>LM<sub>3</sub>W<sub>4</sub>D<sub>1</sub>IT<sub>2</sub>G LQY<sub>1</sub>NNK..L<sub>2</sub>Y DLT<sub>2</sub>V<sub>3</sub>GS<sub>4</sub>MS<sub>1</sub>K<sub>2</sub>L ADF<sub>1</sub>MA<sub>2</sub>D.H<sub>3</sub>V<sub>4</sub> EM<sub>1</sub>IK<sub>2</sub>VG<sub>3</sub>V<sub>4</sub>WN AQT<sub>1</sub>VN-V<sub>2</sub>Y<sub>3</sub>G KIHVV<sub>1</sub>TK<sub>2</sub>I<sub>3</sub>K<sub>4</sub>AGR<sub>1</sub>LR<sub>2</sub>WW<sub>3</sub>V<sub>4</sub>R  
 P78783/1-0 GVLG<sub>1</sub>Y<sub>2</sub>DL<sub>3</sub>ITY<sub>4</sub> FSKRN---- - -ID<sub>1</sub>DL<sub>2</sub>DF<sub>3</sub>AR<sub>4</sub>V<sub>1</sub>E<sub>2</sub>TF<sub>3</sub>Q<sub>4</sub>N.Y<sub>1</sub>P EILG<sub>2</sub>Q<sub>3</sub>AL<sub>4</sub>IV<sub>1</sub> SG<sub>2</sub>FR<sub>3</sub>MA<sub>4</sub>LE<sub>1</sub>F EG<sub>2</sub>W<sub>3</sub>SI<sub>4</sub>GK<sub>1</sub>Y<sub>2</sub>Q<sub>3</sub>EL<sub>4</sub> P<sub>1</sub>E<sub>2</sub>V<sub>3</sub>RS<sub>4</sub>K<sub>1</sub>V<sub>2</sub>TC  
 YN04\_YEAST/1-0 K<sub>1</sub>IT<sub>2</sub>VL<sub>3</sub>Y<sub>4</sub>DF<sub>1</sub>K<sub>2</sub>S<sub>3</sub> YKE<sub>2</sub>PG<sub>3</sub>II<sub>4</sub>TD<sub>1</sub> A<sub>2</sub>PP<sub>3</sub>IS<sub>4</sub>ARM<sub>1</sub> C<sub>1</sub>LN<sub>2</sub>VM<sub>3</sub>D.H<sub>4</sub>F<sub>1</sub> ERL<sub>2</sub>AK<sub>3</sub>CV<sub>4</sub>IL<sub>1</sub> IN<sub>2</sub>W<sub>3</sub>--FA<sub>4</sub>W<sub>1</sub> FL<sub>2</sub>KM<sub>3</sub>Y<sub>4</sub>PEL<sub>1</sub> P<sub>2</sub>AT<sub>3</sub>K<sub>4</sub>IV<sub>1</sub>F<sub>2</sub>L  
 YAS2\_SCHPO/1-0 TAT<sub>1</sub>TV<sub>2</sub>Y<sub>3</sub>DF<sub>4</sub>MTD FS<sub>1</sub>MSN---- - -MDY<sub>1</sub>Y<sub>2</sub>GL<sub>3</sub>PK<sub>4</sub> F<sub>1</sub>IC<sub>2</sub>FEA.H<sub>3</sub>Y<sub>4</sub>P ECL<sub>2</sub>GE<sub>3</sub>C<sub>4</sub>IV<sub>1</sub>HK A<sub>2</sub>Y<sub>3</sub>W<sub>4</sub>Q<sub>1</sub>TF<sub>2</sub>Y<sub>3</sub>HER<sub>4</sub>LL TGL<sub>1</sub>GS<sub>2</sub>K<sub>3</sub>RL<sub>4</sub>V  
 NF1<sub>1</sub>\_HUMAN ---V<sub>1</sub>V<sub>2</sub>DL<sub>3</sub>L<sub>4</sub>TH<sub>1</sub> T<sub>2</sub>GE<sub>3</sub>PSN---- - RFK<sub>1</sub>TDF<sub>2</sub>LS<sub>3</sub>K<sub>4</sub> FW<sub>1</sub>V<sub>2</sub>FG<sub>3</sub>.F<sub>4</sub>Y<sub>1</sub> DV<sub>2</sub>VS<sub>3</sub>AV<sub>4</sub>Y<sub>1</sub>S V<sub>2</sub>EL<sub>3</sub>Q<sub>4</sub>LAFL<sub>1</sub> P<sub>2</sub>N<sub>3</sub>T<sub>4</sub>V<sub>1</sub>LAFL<sub>2</sub>  
 Q9XU9/1-0 SV<sub>1</sub>VC<sub>2</sub>Y<sub>3</sub>LD<sub>4</sub>L<sub>1</sub>KG L<sub>2</sub>LN<sub>3</sub>MD<sub>4</sub>FM<sub>1</sub>N<sub>2</sub>LS<sub>3</sub>PA<sub>4</sub>QL<sub>1</sub>AR<sub>2</sub>LV<sub>3</sub>Y<sub>4</sub>S E<sub>1</sub>HL<sub>2</sub>C<sub>3</sub>LL<sub>4</sub>IN<sub>1</sub> P<sub>2</sub>E<sub>3</sub>II<sub>4</sub>S V<sub>1</sub>SV MW<sub>2</sub>Q<sub>3</sub>V<sub>4</sub>TK<sub>1</sub>RL<sub>2</sub>VD P<sub>1</sub>N<sub>2</sub>T<sub>3</sub>V<sub>4</sub>LAFL<sub>1</sub>  
 Consensus/60% .bshlbDbct bsh..... sbssb+ph bpblps.pap -plpphbllp sPb...bb.s baplhp<sub>1</sub>sFls pp<sub>2</sub>pcp<sub>3</sub>K1.bl

\*

Dbl3	ML----SSVS D <sup>L</sup> TYID..D KQ <sup>L</sup> TP <sup>L</sup> EL <sup>L</sup> GT LQ...
AAD25756/1-0	RP---AKVRE T <sup>L</sup> IKYIP..A DE <sup>L</sup> EVQY <sup>L</sup> GF KT...
A49678/1-0	NY----LS E <sup>L</sup> SEH <sup>L</sup> V <sup>L</sup> KLEQ LGI <sup>L</sup> RQVLKY DDFLK
T00939/1-0	YS.DDPNTKV IMEENFD..M EK <sup>L</sup> MELAF <sup>L</sup> GN DD...
O35239/1-0	T-....TS E <sup>L</sup> TQH <sup>L</sup> P..R EC <sup>L</sup> EKNL <sup>L</sup> GY VK...
CAA15638/1-0	T-....QSIN T <sup>L</sup> YEFVP..R EM <sup>L</sup> EEY <sup>L</sup> GG AG...
AAF25952/1-0	S-....IE S <sup>L</sup> NKIVB..S HQ <sup>L</sup> TGDFE <sup>L</sup> Q QL...
NIP2_HUMAN/1-0	F-...NLAEL AELVPME..Y VG <sup>L</sup> E <sup>L</sup> CIKVQ DQELN
CAB55301/1-0	ML----SSVS D <sup>L</sup> TYID..D KQ <sup>L</sup> TP <sup>L</sup> EL <sup>L</sup> GT LQ...
YUQP_CAEEL/1-0	G-...SNWKE E <sup>L</sup> LDIVD..K EC <sup>L</sup> ERY <sup>L</sup> GM IP...
Q9ZWD6/1-0	EP---KSLF K <sup>L</sup> BEVID..S SQ <sup>L</sup> EFL <sup>L</sup> GS CS...
Q91870/1-0	K-....AH E <sup>L</sup> VNHIP..K ES <sup>L</sup> PEH <sup>L</sup> GT SK...
O43304/1-0	SG.SNYQPG G <sup>L</sup> VDYID..R EVI <sup>L</sup> PDFL <sup>L</sup> GE SV...
190854/1-0	G-...DDLS GFYQEID..E NI <sup>L</sup> ESDF <sup>L</sup> GT LP...
AAF02821/1-0	SR-...LQ Y <sup>L</sup> WED <sup>L</sup> K..K GEVEIPDFVK NH...
CAB51563/1-0	EP-....D <sup>L</sup> RYVP..R DQ <sup>L</sup> DSNF <sup>L</sup> GS LH...
KARI_RAT/1-0	S-....VE G <sup>L</sup> TKLVD..P SQ <sup>L</sup> TEEF <sup>L</sup> GS LD...
O48940/1-0	EK--NKVK <sup>L</sup> S T <sup>L</sup> LEENB..E SQ <sup>L</sup> VEIF <sup>L</sup> GS LP...
O23217/1-0	YP.KDKASDE LM <sup>L</sup> TT <sup>L</sup> HD..V EN <sup>L</sup> EKEF <sup>L</sup> GE AT...
A53057/1-0	G-...G <sup>L</sup> DYKD T <sup>L</sup> LEYID..A EEL <sup>L</sup> PAYL <sup>L</sup> GT KS...
4437/1-0	G-...SSYQ <sup>L</sup> E <sup>L</sup> LQ <sup>L</sup> IP..A EN <sup>L</sup> EVKF <sup>L</sup> GT SE...
Q23130/1-0	MC-...IDEL ENALSVA..R LN <sup>L</sup> PSPIRDH DKSF <sup>L</sup> S
O48939/1-0	P-...GCGRD E <sup>L</sup> DTIMD..Y SS <sup>L</sup> CFHFCRRE GS...
SC14_SCHPO/1-0	G-...SNYKS A <sup>L</sup> LEQIP..A DN <sup>L</sup> PAKL <sup>L</sup> GN CQ...
CAB63153/1-0	KE--GNAAE T <sup>L</sup> YKFR..P EDI <sup>L</sup> EVQY <sup>L</sup> QL SR...
Q22824/1-0	---...FA I <sup>L</sup> SQYAD..S HK <sup>L</sup> EGDRTKI AV...
Q17910/1-0	DR-...HFIE T <sup>L</sup> TEDIN..I DD <sup>L</sup> I <sup>L</sup> VSL <sup>L</sup> GN DA...
DBS_HUMAN/1-0	ML---SSVP D <sup>L</sup> HGYID..S SQ <sup>L</sup> TEFL <sup>L</sup> GT LD...
CAB62036/1-0	G-...ETFH <sup>L</sup> R V <sup>L</sup> SEYLQ..-T <sup>L</sup> PSYL <sup>L</sup> SN CN...
AAF18525/1-0	GP-...SKSAE T <sup>L</sup> FKYVA..P EV <sup>L</sup> EVVKY <sup>L</sup> GL SK...
O24659/1-0	DD---KSLQ <sup>L</sup> E <sup>L</sup> HQEID..D SQ <sup>L</sup> IEDTL <sup>L</sup> CK LA...
TPPA_RAT/1-0	G-...NNYKS S <sup>L</sup> IQHFP...DIL <sup>L</sup> PLEY <sup>L</sup> GN ES...
NF1_DROME	FL---ESPN K <sup>L</sup> TD <sup>L</sup> DAEQ QK <sup>L</sup> EGATL <sup>L</sup> SL DE...
AAD23650/1-0	G-...N <sup>L</sup> KYQS K <sup>L</sup> LEI <sup>L</sup> ID..S NE <sup>L</sup> PEFL <sup>L</sup> GN CT...
Q93546/1-0	N-...SEWRV <sup>L</sup> E <sup>L</sup> V <sup>L</sup> KMAB..G SC <sup>L</sup> CESYWNDE ED...
Q22467/1-0	G-...NDWKQ H <sup>L</sup> KENIG..E EV <sup>L</sup> FERW <sup>L</sup> GT RK...
Q06705/1-0	KN-....ID E <sup>L</sup> PHKF <sup>L</sup> Q..P QY <sup>L</sup> ERYL <sup>L</sup> QE ND...
Q81806/1-0	SR-....VD Y <sup>L</sup> WEHVR..R NEIEMPEFVY DH...
AAF19259/1-0	GD-...NWQ <sup>L</sup> E <sup>L</sup> TKFIS..P DQ <sup>L</sup> EV <sup>L</sup> GT MT...
Q9Z108/1-0	G-...NSWKE G <sup>L</sup> IKLIS..P EE <sup>L</sup> PAHF <sup>L</sup> GT LT...
076636/1-0	SGN--SDWKT SVQKHAH..I DN <sup>L</sup> E <sup>L</sup> KHW <sup>L</sup> GT LV...
O15915/1-0	NS(5)KRTFA E <sup>L</sup> LEYID..I EN <sup>L</sup> EQNL <sup>L</sup> GN LD...
YNX1_YEAST/1-0	EP-....-FV <sup>L</sup> KYVP..I NE <sup>L</sup> DSLY <sup>L</sup> GT LK...
001324/1-0	S-...EPAIG AIQKHAD..K CF <sup>L</sup> PSDL <sup>L</sup> GT LE...
BAA90344/1-0	YS.SDKE <sup>L</sup> SQ <sup>L</sup> IMADVF <sup>L</sup> .L DR <sup>L</sup> DSAF <sup>L</sup> GR NP...
Q9ZW01/1-0	---...EHTC D <sup>L</sup> TQNSR..R GIRFFVVLST AT...
BAA90976/1-0	D-...SLHQL FSAISPB..Q ID <sup>L</sup> PPFVLEY DA...
O61528	S-....FE S <sup>L</sup> TIKI <sup>L</sup> ID..S SH <sup>L</sup> E <sup>L</sup> KTV <sup>L</sup> GS YP...
YN02_CAEEL/1-0	GG.SGGDLKE E <sup>L</sup> RKH <sup>L</sup> E..E KFI <sup>L</sup> DFL <sup>L</sup> GS CL...
YJ05_YEAST/1-0	TD-....GS K <sup>L</sup> Q <sup>L</sup> QY <sup>L</sup> IK..-DCP <sup>L</sup> PEGY <sup>L</sup> CK DK...
YQF6_CAEEL/1-0	S-....DKSDLAG..S SCR <sup>L</sup> GEF <sup>L</sup> VNT VP...
O17907/1-0	E-...SDWKW S <sup>L</sup> AQIVN..L DAW <sup>L</sup> MYW <sup>L</sup> QN LV...
5824562/1-0	G-...NRNFG L <sup>L</sup> GVFHSD..E EQVADFFIAK QV...
P78783/1-0	K-....PA QVSGY <sup>L</sup> ID..S KY <sup>L</sup> E <sup>L</sup> LSMH <sup>L</sup> Q FD...
YN04_YEAST/1-0	G-(6)SDWKE E <sup>L</sup> IKYID..P EE <sup>L</sup> EEL <sup>L</sup> GT LK...
YAS2_SCHPO/1-0	RN-....YR D <sup>L</sup> Q <sup>L</sup> QY <sup>L</sup> IN..P DN <sup>L</sup> ILKEF <sup>L</sup> GP NP...
NF1_HUMAN	FI-...DCPG K <sup>L</sup> A <sup>L</sup> H <sup>L</sup> EHEQ QK <sup>L</sup> EAATLAL EE...
Q9XVU9/1-0	SN-....VE D <sup>L</sup> IKY <sup>L</sup> E..P EA <sup>L</sup> VEY <sup>L</sup> GT YR...
Consensus/60%	s.....p_bp pLhcblc... ppLP.pbGGp .s...

Table S2. siRNA sequences of confirmed hits of the library screening

Gene	First round siRNAs		Second round siRNAs
	siGenome	ON-TARGETplus	
SPATA13	5'-GAAGAACUUGCCAUGUUA-3'	5'-CCACACAGGAACACGGUGA-3'	
SPATA13	5'-GGAGAGCAUCGACAAGAU-3'	5'-GUACCAUGCUCUAAAGCGA-3'	
SPATA13	5'-GGACCAACGUCAUCCGGGA-3'	5'-CUGCAACCUCAGCGUGAA-3'	
SPATA13	5'-GCCAGCUAGCCACUAAA-3'	5'-GGACCAACGUCAUCCGGGA-3'	
VAV3	5'-UAAAUGAUCUUACAGCAGUA-3'	5'-GAAACAUUAUCACGACUUU-3'	
VAV3	5'-GCAGAGACCGAACUUAUUA-3'	5'-GCAAUAGCAUUAAGUACA-3'	
VAV3	5'-GCAAGCACAUCAAGAUUU-3'	5'-AGUACAAGAUAGCCAUAUA-3'	
VAV3	5'-AGACCGAACUUAUUAUAG-3'	5'-AGACCGAACUUAUUAUAG-3'	
TRIO	5'-GAUAAGAGGUACAGAGAU-3'	5'-GUAAAAGAAGUGAAAGAUUC-3'	
TRIO	5'-GGAAGUCGCUCCUUGACAA-3'	5'-CGACCUAUCCGUAGCAUUA-3'	
TRIO	5'-CAACGGAGAGGUCAUGUUA-3'	5'-GGAAUACAACCACGAAGAA-3'	
TRIO	5'-GAACACCAACUUCAGAUUA-3'	5'-AGAACAGGGUAUUGCAUUA-3'	
OBSCN	5'-GCAGACAGCGACACCUUA-3'	5'-GAUGUGAGCUGCAGAUUA-3'	
OBSCN	5'-AGACAUACCGCGAACAGAUGA-3'	5'-CCAAGGUGGUGGUUUGCUAA-3'	
OBSCN	5'-GGAAGGACAUACACUCUCA-3'	5'-GCAGACAGCGACACCUUA-3'	
OBSCN	5'-CAGGAGAGAUCCAUAUUGU-3'	5'-GAUGGAGAGUUCCGUGAUU-3'	
MCF2	5'-GGUGUAACCGCAAGUUUG-3'	5'-GAACUGGGCAAGAUGAUAA-3'	
MCF2	5'-GGUAUCAUCUGUUGAAGAA-3'	5'-UAACAGAACGGGAUAAGUU-3'	
MCF2	5'-GGAAGAAGUUAUUAUUGUC-3'	5'-GAAGAAGGCACUCUGAUGC-3'	
MCF2	5'-GCAACAGGAUCAUAUUAACA-3'	5'-GAGCUCAGUUAGUGAUUUG-3'	
ECT2	5'-GAUAAGAGGAUGAUCUUGAA-3'	5'-GCACUCACCUUGUAGUUGA-3'	
ECT2	5'-GCACUCACCUUGUAGUUGA-3'	5'-CAGAGGAGAUUAAGACUAU-3'	
ECT2	5'-GAAGGGCUCUUAUGACAUC-3'	5'-UAACAGCAAUGCCAAACGA-3'	
ECT2	5'-CAGAGGAGAUUAAGACUAU-3'	5'-GUAAUGAGUCGUCUUUCUA-3'	
FARP1	5'-CAGAAAACUCGUGGUCCAU-3'	5'-CAGAAAACUCGUGGUCCAU-3'	
FARP1	5'-AAGAAGAACUCACAAGGU-3'	5'-CAGGAGGCAUUUGAAGUUC-3'	
FARP1	5'-GCACGUUGUUGUUAAGUUU-3'	5'-CCGGAAAAAUUCGGGGAUCA-3'	
FARP1	5'-CAGCGCAGCUCUUGAUU-3'	5'-CUUGGAACGUGGACAGAACG-3'	
NET1	5'-GGCAAUAUAUGAAAUGUCC-3'	5'-GGAGGAUGCUUAUUGAU-3'	
NET1	5'-GAAGUCGCCUAGUAAAUA-3'	5'-GAGUCUCCCUCAGUCGAA-3'	
NET1	5'-GGAGGAUGCUUAUUGAU-3'	5'-UCACGUCCUUGGCAAAUUU-3'	
NET1	5'-UCACGUCCUUGGCAAAUUU-3'	5'-UUGGUGAUCUGGACUCUUA-3'	

## Reference

Elbediwy, A., C. Zihni, S.J. Terry, P. Clark, K. Matter, and M.S. Balda. 2012. Epithelial junction formation requires confinement of Cdc42 activity by a novel SH3BP1 complex. *J. Cell Biol.* 198:677–693. <http://dx.doi.org/10.1083/jcb.201202094>