



### University of Dundee

# C11orf70 Mutations Disrupting the Intraflagellar Transport-Dependent Assembly of Multiple Axonemal Dyneins Cause Primary Ciliary Dyskinesia

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Homo	1 MATGELGDLGGYYFRFLPQKTFQSLSSKEITSRLRQWSMLGRIKAQAFG	FDQ
Bos	1 MAAGEPGDIGVYSFRFLPQKTFQSLSTPQTTSRLRQWSMLGRIEAQAFG	FDQ
Mus	1 MAAGEPRDGGGYYFRFLPHRTFSSLSAREITSRLRQWSMLGRIQAQAFS	FDQ
Rattus	1 MATGEPRDRGGYYFRFLPQRTFSSLSAREITNRLRQWSMLGRIQAQAFG	FDQ
Xenopus	1 MSAGSFSTEAAKFSFSPILNKTFGFLINRDTRELIMKWSMNGRITAQAFR	Υ <mark>D</mark> E
Danio	1 MVTTTIGIMAEEKLSFEQTENENLLSTKSENFQEDPKTSRLLMKWSMLGRITAQAEN	FDQ
Chlamydomonas	1 MTAFVPVSLPSTSALNDAYVKSQLTKWD LRN RCVAVR	ΥTK
Paramecium	1 MQIESDNQVTNQSYSFFRQE-NIIDDKKFMEIDQKWGDQHSIKVSTFL	FDI
TT		
HOIIIO Dan	53 TEQSIRCDEF WAFFECPENTICULAUS SSGQWIILGIEVENIEAINVPCIQUSMS	
BOS	53 TFQAIRKDDFVTAFFKDPNVIPNLKLLSFSSGEWLTLGTEVKKIEAINVPCTQLSMS	EEN
Mus	53 TFQPYQKDDFVMAFFKDPNVIPNLQLLSDSSGQWTTLGBEVKKILAINVPCTQLSMS	ΓFQ
Rattus	53 TFQPYQKDDFVTAFFKDPNVIPHLQLLAFSSGQWTTLGTEVKRIEAINVPCTQLSMS	ΕΕQ
Xenopus	54 CFQPYQKNDFVWAFFQDPDV SHIK VSENSGQWVILGTKVKKV VQE LOSQLSMS	LED
Danio	61 SEQPYRSNDFAWNFFQDPCVKHNLNVLD-PTGSWTRLC-DLTHVNVEVVPCLKVSVD	IED
Chlamydomonas	43 Y HKLOGQ L ADLF DEKVQEAFQVURK-GGAWGQLCGPVTKV ATL AS LTRND	LFD
Paramecium	52 KEDHL PNOF LDLENSKDVRGSLHYVSF-K-QNVL SQIKFQP TCKS K D	LFD
Homo	113 RLYDEDIVRDSGHIVKCLDSFCDPFLISDELRRVLLVEDSEKYEIFSOPDREEFL	FCL
Bos	13 RIYDE – AUVRONGYTVKCHOSECOPETTSDEHRKVILVEDSEKYEVES	FCL
Mus	13 RLYDE NTVRESGHTVKCLDSFCDPFLTSDELRKVLLMEDSEKYEVFS	FCL
Rattus	13 RLYDE NIVRESGHIVKCLDSFCDPFLISDELRKVLIMEDSEKYEVFSPVEREEFI	FCL
Xenopus	14 CLYSEGUVRESGHICKCLDEYLDDFTTSDELRKVLLLDDCEKHDVFSOSDREOFL	DT.D
Danio	19 PUYSNGURPSGHUVKOYHETYPDPDEURM IJEADSPYHHUTSPSDBODEU	FRT
Chlamydomonas	102 KUTPTSPPTVBSNGDIGKOMPDNREGDOVSDOLBEL UVEPSEHAALESEAPROPIL	R
Paramecium	106 KUTED KUVV-KGHUKOOFECOFENTOUADEURKAUVUEDSEOMOVENEADBOETU	PKT
r ar anno or an		
Homo	171 FKHLCLGGALCQYEDVI <mark>S</mark> PYLETTKLIYKDLVSVRK <mark>N</mark> PQTK <mark>K</mark> IQITSSVFKVSAYDS	-AG
Bos	171 FKHLCLGGALCQYEDVI <mark>N</mark> PYLETTKLIYKDLVSVRK <mark>N</mark> PQTK <mark>K</mark> IQITSSIFKVTAYDS	-VG
Mus	171 FKHLCLGG <mark>SLCQYEDV K</mark> PYLET <mark>AKLIYKDLVSVRK</mark> HPRTKEIQITSSVFKVKAYDS	-VG
Rattus	171 FKHLCLGG <mark>SLCQYEDVIK</mark> PYLET <mark>AKLIYKDLVSVRKH</mark> PRTKEIQITSSVFKVKAYDS	-LG
Xenopus	172 FKHLCLGGAICQFEDTIGPYLETTKSIYKDLISVQKDPETKQIRIISTVFKVSAYDE	-NG
Danio	174 FKHYVLGGELCQYEDVIDPYLETVKIMYKDLVSVQKDTDTKEINVVTTVLKVSAYDH	-SG
Chlamydomonas	162 FEHVVIGGACCOFEDKVEPYVETSKRIYKEIVCAQKDPATGKVQTVSAVYKINSIQG	DSG
Paramecium	163 FQI <mark>IVLGGQLCQYED</mark> EIQA <mark>YLDWTK</mark> YI <mark>YK</mark> NT <mark>VNARK</mark> YADKDETYIDSYAYDIRKL	
Homo	230 WOYPSAKNHEQTESYFIVDPIRRHIHVLYHCYGVGDMS	
Bos	2.30 V <b>OYPS</b> TKS <b>HEQTESYE</b> T <b>VDPTKRHV</b> HVDYHCYGMCEVS	
Mus	230 VCYPSPKEHEQTFSYFVVDPIKRHVNVLYHCYGVCHMA	
Rattus	230 VOYESPKEHEQTESYFVVDPIKRHVNVLYHCYGVCEMA	
Xenopus	231 MCYPSGRPHQOTEANL VDP KRHVYVLYHC GGCA-F	
Danio	233 CYPSATENKOTEANLCEDECKRHVYVLEHSEGECEFSGN	
Chlamydomonas	222 PLELYPSRSRONDCYAAVDEVERIVK LYHAYVPYW	
Paramecium	218 - ENSYSSDHPONVMYVVVNESLRIVNI ENQALKVW	

Figure S1. Cross species protein alignment for C11orf70



Figure S2. Quantification of TEM analysis of outer and inner dynein arm loss in affected individuals with *C11orf70* mutations



**Figure S3. Successful RNAi ablation of** *C11orf70* **shown in** *Paramecium* **transformed with GFP-tagged C11orf70.** *ND7-* (control) and *C11orf70-* silenced Paramecia were fixed and visualised for C11orf70-GFP protein expression after 72h of RNAi. Scale bars, 10 µm.



**Figure S4. Depletion of** *C11orf70* **in** *Paramecium* **does not affect cilia number and length**. *ND7* (control) and *C11orf70* silenced Paramecia were fixed and stained for cilia after 72h of RNAi. Cilia were stained using anti - polyglutamylated tubulin antibodies to indicate the cilia (PolyE). Scale bars, 20 µm.



Figure S5. *Chlamydomonas* FBB5 is expressed at similar levels in a number of mutant *Chlamydomonas* strains. Amido black stained gels shows equivalent protein in strains with mutations in proteins shown in Figure 5, main text.

Primer ID	(5'- Sequence -3')						
Sanger sequencing for c.776A>G mutation							
Fwd	GATCCTATCAGGCGTCACCTT						
Rev	ATCCCCGGTTAGTATTCCACAC						
Sanger sequencing for c.154C>T mutation							
Fwd	GCCAGCCCCCAGATGATTT						
Rev	GAGACCCAGGGAACTCTCCG						
Sanger sequencing for c.361C>T mutation							
Fwd	GTTCCTTGCACACAGCTTTCAA						
Rev	TCCTGAAATAAAATTCCACTGCGA						
q(RT-PCR) for expression profiling of C11orf70 in ALI culture							
C11orf70-Fwd	TGGGTGGCTACTACTTCAGGT						
C11orf70-Rev	GCCTTGATTCTGCCCAGCAT						
DNAH5-Fwd	TGCAGATGCCATGGTTCACT						
DNAH5-Rev	ATGAAGCCAACCTCGTCAGG						
GAPDH-Fwd	TGCACCACCAACTGCTTAGC						
GAPDH-Rev	GGCATGGACTGTGGTCATGAG						
561bp segment of Paramec	ium GSPATG00011350001 for RNAi design						
Fwd	TAGAACTAGTATGTAAATTGAATCAGACAATTAAGTTAC						
Rev	TTTTTTCTCGAGCTTTGTCCAATCTAAATAGGCTTG						
q(RT-PCR) for GSPATG000 <sup>2</sup>	11350001 knockdown in <i>Paramecium</i>						
GSPATG00011350001-F	TGCTCGAAAATATGCAGATAAGGA						
GSPATG00011350001-R	TGATGGGTTCACAACGACAT						
GAPDH-Fwd	GAGAGCCGGAAGAGCTGCTA						
GAPDH-Rev	TGGTGGAACTCTGAAGGCCATA						
Amplification of Paramecium GSPATG00011350001 (C11orf70) for GFP expression vector cloning							
Fwd	TAGAACTAGTATGTAAATTGAATCAGACAATTAAGTTAC						
Rev	TTTTTCTCGAGTCCTCCTCCCCATACTTTTAACCATTAGTTTTCT						
Amplification of Paramecium GSPATG00024708001 (IFT46) for GFP expression vector cloning							
Regulator-Fwd	ATACTTGGATCCTATTTAATATAATTAAATAAATACTGTTATTC						
Regulator-Rev	GACAAAACAATGATGAAATTTAAGGCATGCAAGTAT						
GSPATG00024708001-F	ATACTTGGTACCTGAATATTCAATAATATAACATTATTTC						
GSPATG00024708001-R	GGAATATTTATTATAATTATTAA						

Table S1. Primers

# used in the study

Antigen	Antibody	Host	Source	Application
		species		
DNAH5	HPA037470	Rabbit	Sigma-Aldrich	IF (1:800)
DNALI1	HPA053129	Rabbit	Sigma-Aldrich	IF (1:200)
RSPH4A	HPA031196	Rabbit	Sigma-Aldrich	IF (1:400)
GAS8	HPA041311	Rabbit	Sigma-Aldrich	IF (1:500)
Acetylated Tubulin	T7451	Mouse	Sigma-Aldrich	IF (1:500)
Acetylated Tubulin	YF488	Mouse	Proteintech	IF (1:500)
Polyglutamylated Tubulin	PolyE	Rabbit	A gift from J. Cohen	IF (1:500)
GFP	IgG	Rabbit	Interchim, France	IF (1:500)
НА	McAb 3F10	Rat	Roche, Indianapolis	IB (1:1000)
Chlamydomonas IFT46C	α-IFT46C404	Guinea Pig	A gift from G. Witman	IB (1:10000)
Chlamydomonas IFT46N	α-IFT46-17601	Rabbit	A gift from H. Qin	IB (1:5000)
Chlamydomonas IFT81	81.3	Mouse	A gift from D. Cole	IB (1:10000)
Chlamydomonas IFT139	139.1	Mouse	A gift from D. Cole	IB (1:10000)
Chlamydomonas IC2	C11.4	Mouse	DRM lab	IB (1:2000)
Chlamydomonas ODA16	α-ODA16-927	Rabbit	DRM lab	IB (1:100)

## Table S2: Primary antibodies used in the study

Antibody	Host Species	Source	Application
Alexa 488, 594 – conjugated anti-rabbit IgG (H+L)	Goat	Invitrogen (Molecular Probes)	IF (1:1000)
Alexa 488, 594 – conjugated Anti-mouse IgG1	Goat	Invitrogen (Molecular Probes)	IF (1:1000)
HRP-conjugated Anti-mouse IgG	Goat	BioRad	IB (1:2000)
HRP-conjugated Anti-rabbit IgG	Goat	Sigma-Aldrich	IB (1:2000)
HRP-conjugated Anti-guinea Pig IgG	Goat	Rockland	IB (1:1000)
HRP-conjugated Anti-rat IgG	Goat	Sigma-Aldrich	IB (1:5000)

Table S3: Secondary antibodies used in the study