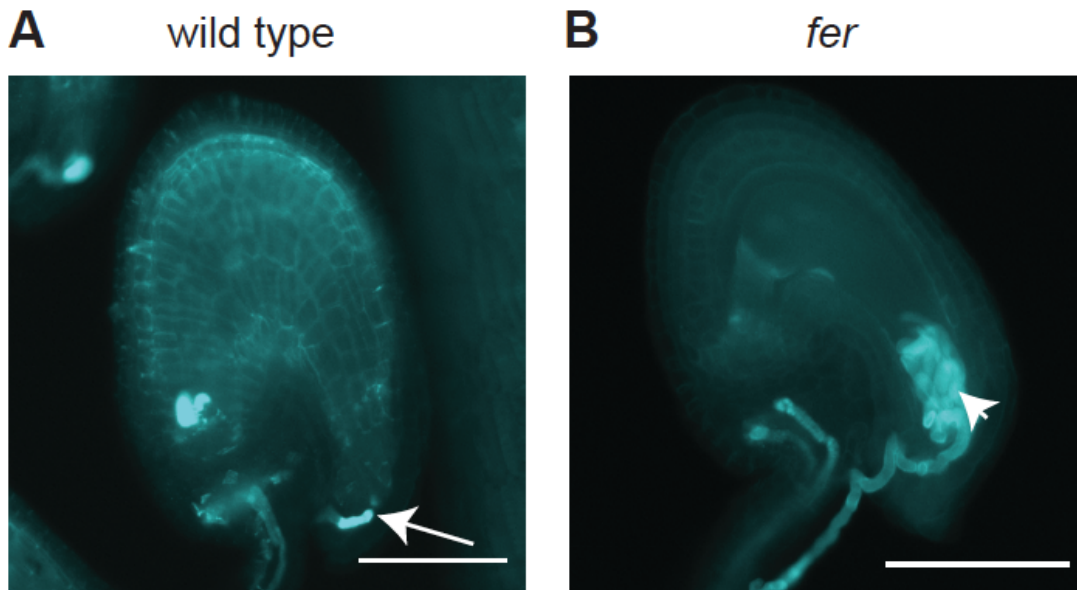


## SUPPLEMENTARY INFORMATION

### **Functional analysis of related CrRLK1L receptor-like kinases in pollen tube reception**

Sharon A Kessler, Heike Lindner, Daniel S Jones & Ueli Grossniklaus

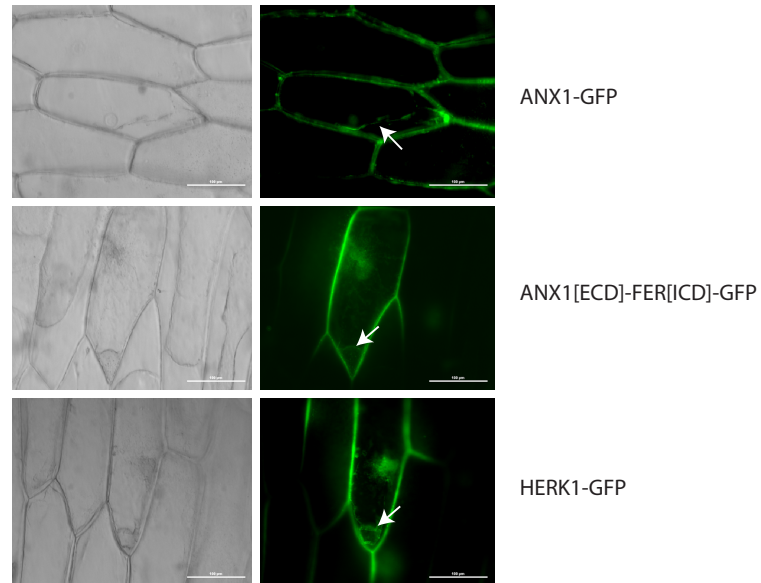
### **Supplementary Figures**



#### **Supplementary Figure S1. PT reception is disrupted in *fer* mutants.**

Aniline blue staining of PTs in wild-type (A) and *fer-1* (B) ovules. Arrow indicates normal PT reception and arrowhead indicates PT overgrowth. Staining was performed according to [1].





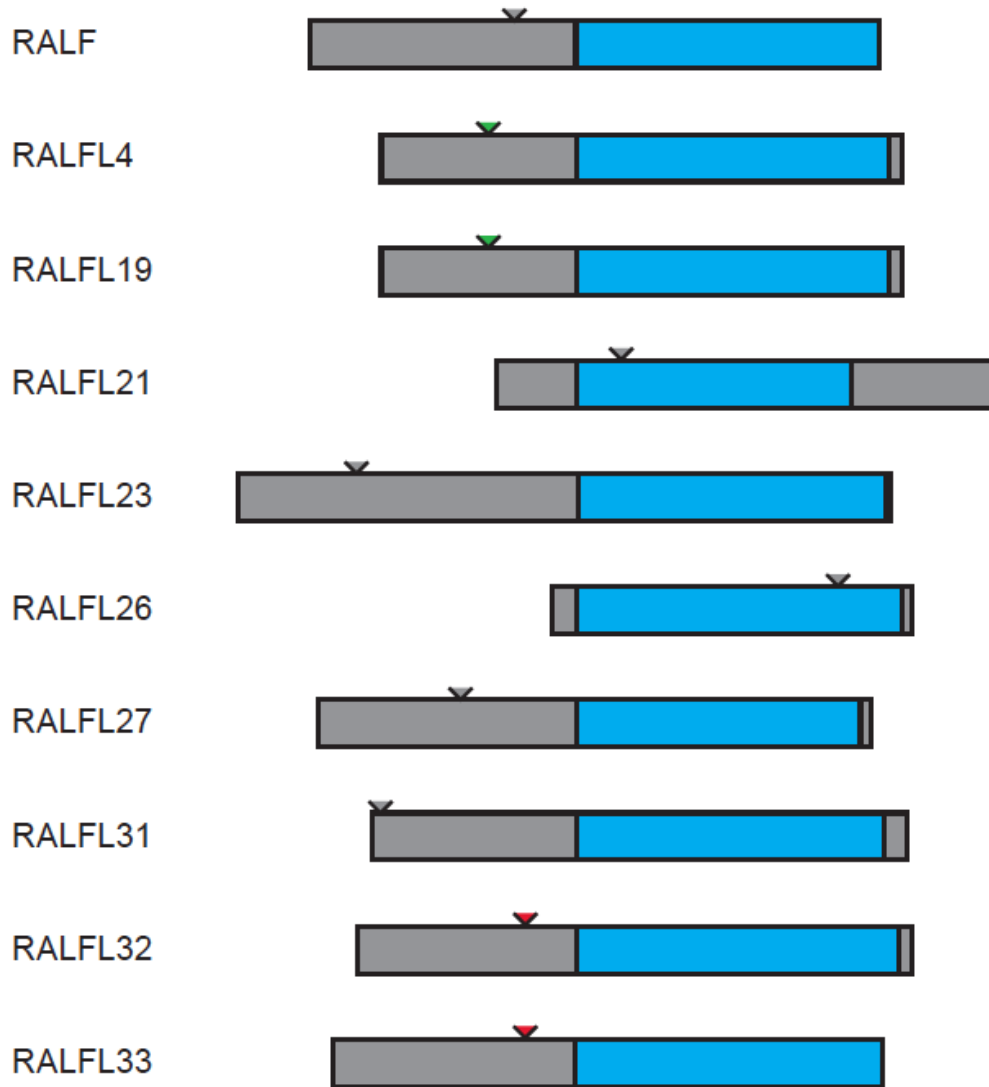
**Supplementary Figure S3. Membrane localization of GFP fusion proteins in onion epidermal cells.**

Left panels, DIC images of onion epidermal cells plasmolyzed in 30% sucrose. Right panels, transient GFP fusion protein expression. Arrows indicate GFP signal in plasma membrane that has retracted from the cell wall upon plasmolysis. Transient onion infections were done with *Agrobacterium tumefaciens* strains carrying the GFP constructs described in Materials and Methods according to the protocol described in [2].

	Subdomain I	II	III
AtCLV1   692-751	LKEENII	<b>GKGGAGI</b> VYRGSMPNNV-DVAIKRLVGRG-TGRS-DHGFTA	EIQTLGRIRRRHIVR-
AtFLS2   870-929	FNSANI	<b>IGSSSL</b> STVYKQLEDGT-VIAVKVLNLKEFSAES-DKWFYTEAKT	LSQLKHRNLV--
AtACR4   512-571	FKEESI	<b>VGKGSF</b> SCVYKGLRDGT-TVAVKRAIMSSDKQKN-SNEFRTELD	LLSRLNHAHLL--
AtBRI1   883-942	FHNSLI	<b>IGSGG</b> FDVYKAILKDGS-AVAIKKLIHVS--GQG-DREFMAEMETIG	KIKHRNLVPL
AtANX1   517-576	FDDSNV	<b>IGVGG</b> FGKVYKGVIDG-TTKVAVKKNPNS--EQG-LNEFETE	IELLSRLRHKHLVSL
AtFER   536-595	FDESRV	<b>LGVGG</b> FGKVYRGEIDGGTTKVAIKRGNPMS--EQG-VHEFQTE	IEMLSKLRRHRLVS-
AtHERK1   485-544	FDESRN	<b>IGVGG</b> FGKVYKGEINDGT-KVAVKRGPNKS--QQG-LAEFRTE	IEMLSQFRHRLVSL
DmCDK1   4-63	FEKIEKI	<b>IGEGTY</b> GVVYKGRNRLTGQIVAMKKIRLES-DEGVPSTAI	REISLLKELKHENI---
ScSNF1   55-114	YQIVKTL	<b>IGEGS</b> FGKVKLAYHTTGGQVALKI INKKVLAKSDMQGRIERE	ISYLRLLRHPH----
	.    : * . . * .	: : * :	* . : : * .

**Supplementary Figure S4. FER, HERK1, and ANX1 have typical kinase domains.**

Amino acid alignment of the first 60 amino acids of kinase domains from *Arabidopsis* RLKs, and *Drosophila* CDK1, and yeast SNF1 kinases. Subdomains are defined based on [3]. The subdomain II critical K of the highly conserved ATP-binding site is highlighted in green, the nearly conserved glycine loop in subdomain I of kinases is shown in yellow, and the conserved E in subdomain II, which forms a salt bridge with the critical K [3], is highlighted in blue. The K565 residue was changed to R, A, and E in FER and all three changes complemented the *fer* phenotype (Fig 4B). A K911E change in BRI1 abolished kinase activity and resulted in a dwarf phenotype [4] and K560L, M, and W changes in ACR4 all led to loss of kinase activity [5-7]. Amino acid sequences and locations of predicted kinase domains were downloaded from the Universal Protein Resource (uniProt) database (CLV1, [Q9SYQ8](#); FLS2, [Q9FL28](#); ACR4, [Q9LX29](#); BRI1, [O22476](#); ANX1, [Q9SR05](#); FER, [Q9SCZ4](#); HERK1, [Q9LX66](#); CDK1, [P23572](#); SNF1, [P06782](#)).



**Supplementary Figure S5. Potential glycosylation sites in RALF and RALFL-like peptides.**

RALF and RALFL peptides were analyzed for potential glycosylation sites with the NetNGlyc 1.0 Server (<http://www.cbs.dtu.dk/services/NetNGlyc/>). The 10 peptides with predicted glycosylation sites are shown with the RALF domain (predicted by PFAM analysis) in blue. Conserved glycosylation sites are red and green, while non-conserved sites are in grey. Peptides shown are before potential processing by a protease, which may remove some of the putative glycosylation sites. RALF and RALFL23 were shown to be processed by a Site-1 protease (S1P) [8,9]. Pollen expressed RALFL4 and RALFL26 do not have the motif for S1P recognition, but might be processed by other proteases.

## Supplementary Tables

**Supplementary Table S1: Seed count data from domain swap complementation experiment (see also Figure 3)**

Construct	Plant number	genotype	unfertilized ovules	fertilized ovules	% unfertilized
Ler untransformed control	Ler-1	<i>FER/FER</i>	11	355	3.0
	Ler-2	<i>FER/FER</i>	13	338	3.7
	Ler-3	<i>FER/FER</i>	4	357	1.1
	Ler-4	<i>FER/FER</i>	4	363	1.1
	Ler-5	<i>FER/FER</i>	5	469	1.1
	Ler-6	<i>FER/FER</i>	21	442	4.5
	Ler-7	<i>FER/FER</i>	7	457	1.5
<i>fer-1/FER</i> untransformed control	SAK543-1	<i>fer-1/FER</i>	171	234	42.2
	SAK543-2	<i>fer-1/FER</i>	135	221	37.9
	SAK543-3	<i>fer-1/FER</i>	169	203	45.4
	SAK543-4	<i>fer-1/FER</i>	176	224	44.0
	SAK543-5	<i>fer-1/FER</i>	178	207	46.2
	SAK543-6	<i>fer-1/FER</i>	187	234	44.4
	SAK543-7	<i>fer-1/FER</i>	192	225	46.0
	SAK543-8	<i>fer-1/FER</i>	147	208	41.4
<i>fer-1/fer-1</i> untransformed control	SAK534-1	<i>fer-1/fer-1</i>	267	53	83.4
	SAK534-2	<i>fer-1/fer-1</i>	262	50	84.0
	SAK534-3	<i>fer-1/fer-1</i>	200	41	83.0
	SAK534-4	<i>fer-1/fer-1</i>	203	50	80.2
<i>pFER::FER-GFP/-</i> (complements)	SAK836-31	<i>fer-1/FER</i>	66	220	23.1
	SAK836-30	<i>fer-1/FER</i>	75	241	23.7
	SAK836-35	<i>fer-1/FER</i>	62	231	21.2
	SAK836-42	<i>fer-1/FER</i>	66	243	21.3
	SAK836-44	<i>fer-1/FER</i>	69	211	24.6
	SAK836-39	<i>fer-1/fer-1</i>	101	112	47.4
	SAK836-40	<i>fer-1/fer-1</i>	72	100	41.9
	SAK836-45	<i>fer-1/fer-1</i>	103	135	43.3
<i>pFER::ANX1-GFP/-</i>	SAK540-2	<i>FER/FER</i>	8	407	1.9

(does not complement)

SAK540-3	<i>FER/FER</i>	11	281	3.8
SAK540-7	<i>FER/FER</i>	19	319	5.6
SAK540-10	<i>FER/FER</i>	12	422	2.8
SAK540-11	<i>FER/FER</i>	15	425	3.4
SAK540-12	<i>FER/FER</i>	16	342	4.5
SAK540-13	<i>FER/FER</i>	22	382	5.4
SAK540-19	<i>FER/FER</i>	25	382	6.1
SAK540-24	<i>FER/FER</i>	20	312	6.0
SAK540-26	<i>FER/FER</i>	12	375	3.1
SAK540-30	<i>FER/FER</i>	14	407	3.3
SAK540-1	<i>fer-1/FER</i>	141	210	40.2
SAK540-8	<i>fer-1/FER</i>	148	169	46.7
SAK540-15	<i>fer-1/FER</i>	159	226	41.3
SAK540-16	<i>fer-1/FER</i>	152	220	40.9
SAK540-20	<i>fer-1/FER</i>	152	206	42.5
SAK540-23	<i>fer-1/FER</i>	171	220	43.7
SAK540-25	<i>fer-1/FER</i>	170	238	41.7
SAK540-28	<i>fer-1/FER</i>	162	235	40.8

*pFER::ANX1[ECD]-  
FER[ICD]-GFP/-*  
(does not complement)

SAK537-1	<i>FER/FER</i>	11	347	3.1
SAK537-3	<i>FER/FER</i>	10	373	2.6
SAK537-4	<i>FER/FER</i>	21	337	5.9
SAK537-5	<i>FER/FER</i>	21	331	6.0
SAK537-8	<i>FER/FER</i>	12	388	3.0
SAK537-9	<i>FER/FER</i>	21	388	5.1
SAK537-10	<i>FER/FER</i>	24	408	5.6
SAK537-2	<i>fer-1/FER</i>	94	121	43.7
SAK537-6	<i>fer-1/FER</i>	184	240	43.4
SAK537-7	<i>fer-1/FER</i>	175	200	46.7
SAK537-26	<i>fer-1/FER</i>	113	161	41.2
SAK537-27	<i>fer-1/FER</i>	122	165	42.5
SAK537-30	<i>fer-1/FER</i>	98	132	42.6
SAK537-38	<i>fer-1/FER</i>	79	126	38.5

*pFER::FER[ECD]-  
ANX1[ICD]-GFP/-*  
(complements)

SAK538-1	<i>FER/FER</i>	35	307	10.2
SAK538-7	<i>FER/FER</i>	22	386	5.4
SAK538-12	<i>FER/FER</i>	11	368	2.9
SAK538-18	<i>FER/FER</i>	15	408	3.5
SAK538-19	<i>FER/FER</i>	22	401	5.2
SAK538-20	<i>FER/FER</i>	12	420	2.8

SAK538-23	<i>FER/FER</i>	18	436	4.0
SAK538-2	<i>fer-1/FER</i>	89	317	21.9
SAK538-3	<i>fer-1/FER</i>	101	257	28.2
SAK538-13	<i>fer-1/FER</i>	103	332	23.7
SAK538-24	<i>fer-1/FER</i>	111	301	26.9
SAK538-27	<i>fer-1/FER</i>	95	263	26.5
SAK538-4	<i>fer-1/fer-1</i>	172	231	42.7
SAK538-8	<i>fer-1/fer-1</i>	151	181	45.5
SAK538-17	<i>fer-1/fer-1</i>	153	224	40.6

*pFER::HERK1-GFP/-  
(does not complement)*

SAK541-1	<i>FER/FER</i>	19	324	5.5
SAK541-3	<i>FER/FER</i>	16	388	4.0
SAK541-4	<i>FER/FER</i>	16	395	3.9
SAK541-9	<i>FER/FER</i>	19	334	5.4
SAK541-13	<i>FER/FER</i>	15	304	4.7
SAK541-24	<i>FER/FER</i>	16	376	4.1
SAK541-26	<i>FER/FER</i>	19	389	4.7
SAK541-6	<i>fer-1/FER</i>	155	253	38.0
SAK541-7	<i>fer-1/FER</i>	166	224	42.6
SAK541-10	<i>fer-1/FER</i>	153	212	41.9
SAK541-16	<i>fer-1/FER</i>	180	220	45.0
SAK541-23	<i>fer-1/FER</i>	152	228	40.0
SAK541-25	<i>fer-1/FER</i>	165	209	44.1

*pFER::FER[ECD]-  
HERK1[ICD]-GFP/-  
(complements)*

SAK539-3	<i>FER/FER</i>	14	416	3.3
SAK539-5	<i>FER/FER</i>	34	392	8.0
SAK539-7	<i>FER/FER</i>	25	386	6.1
SAK539-10	<i>FER/FER</i>	66	345	16.1
SAK539-11	<i>FER/FER</i>	20	413	4.6
SAK539-12	<i>FER/FER</i>	33	364	8.3
SAK539-15	<i>FER/FER</i>	23	396	5.5
SAK539-16	<i>FER/FER</i>	15	420	3.4
SAK539-2	<i>fer-1/FER</i>	94	252	27.2
SAK539-6	<i>fer-1/FER</i>	83	272	23.4
SAK539-9	<i>fer-1/FER</i>	91	270	25.2
SAK539-27	<i>fer-1/FER</i>	63	277	18.5
SAK539-28	<i>fer-1/FER</i>	85	171	33.2
SAK539-29	<i>fer-1/FER</i>	81	242	25.1
SAK539-33	<i>fer-1/FER</i>	70	291	19.4
SAK539-39	<i>fer-1/FER</i>	69	182	27.5
SAK539-40	<i>fer-1/FER</i>	38	205	15.6



SAK539-44	<i>fer-1/FER</i>	59	146	28.8
SAK539-45	<i>fer-1/FER</i>	48	166	22.4
SAK539-46	<i>fer-1/FER</i>	125	373	25.1
SAK539-8	<i>fer-1/fer-1</i>	143	178	44.5
SAK539-9	<i>fer-1/fer-1</i>	160	157	50.5
SAK539-14	<i>fer-1/fer-1</i>	143	147	49.3
SAK539-19	<i>fer-1/fer-1</i>	127	196	39.3
SAK539-21	<i>fer-1/fer-1</i>	154	180	46.1

<i>pFER::FERΔICD-GFP/-</i> (does not complement)	SAK542-3	<i>FER/FER</i>	21	349	5.7
	SAK542-4	<i>FER/FER</i>	28	319	8.1
	SAK542-6	<i>FER/FER</i>	6	271	2.2
	SAK542-7	<i>FER/FER</i>	11	271	3.9
	SAK542-9	<i>FER/FER</i>	5	270	1.8
	SAK542-1	<i>fer-1/FER</i>	151	222	40.5
	SAK542-2	<i>fer-1/FER</i>	172	217	44.2
	SAK542-3	<i>fer-1/FER</i>	133	147	47.5
	SAK542-4	<i>fer-1/FER</i>	145	181	44.5
	SAK542-5	<i>fer-1/FER</i>	175	134	56.6
	SAK542-7	<i>fer-1/FER</i>	196	217	47.5
	SAK542-10	<i>fer-1/FER</i>	91	158	36.5
	SAK542-11	<i>fer-1/FER</i>	131	175	42.8
	SAK542-15	<i>fer-1/FER</i>	121	178	40.5
	SAK542-18	<i>fer-1/FER</i>	75	72	51.0
	SAK542-20	<i>fer-1/FER</i>	124	199	38.4
	SAK542-21	<i>fer-1/FER</i>	120	166	42.0
	SAK542-23	<i>fer-1/FER</i>	93	164	36.2
	SAK542-24	<i>fer-1/FER</i>	99	173	36.4

<i>pFER::FER[ECD]-BRI1[ICD]-GFP/-</i> (does not complement)	SAK837-3	<i>FER/FER</i>	30	211	12.4
	SAK837-17	<i>FER/FER</i>	19	241	7.3
	SAK837-13	<i>FER/FER</i>	17	238	6.7
	SAK837-1	<i>fer-1/FER</i>	94	164	36.4
	SAK837-4	<i>fer-1/FER</i>	102	167	37.9
	SAK837-5	<i>fer-1/FER</i>	107	143	42.8
	SAK837-7	<i>fer-1/FER</i>	111	143	43.7
	SAK837-8	<i>fer-1/FER</i>	94	163	36.6
	SAK837-9	<i>fer-1/FER</i>	107	169	38.8
	SAK837-11	<i>fer-1/FER</i>	94	158	37.3

**Supplementary Table S2: Seed count data from FER point mutation complementation experiments (see also Figure 4).**

Construct	line number	genotype	unfertilized ovules	fertilized ovules	% unfertilized
Ler untransformed control	Ler-1	<i>FER/FER</i>	14	330	4.1
	Ler-2	<i>FER/FER</i>	7	315	2.2
	Ler-3	<i>FER/FER</i>	15	322	4.5
<i>fer-1/FER</i> untransformed control	SAK433-1	<i>fer-1/FER</i>	149	232	39.1
	SAK433-2	<i>fer-1/FER</i>	157	221	41.5
	SAK433-3	<i>fer-1/FER</i>	155	220	41.3
	SAK433-4	<i>fer-1/FER</i>	154	230	40.1
	SAK433-5	<i>fer-1/FER</i>	159	245	39.4
	SAK433-6	<i>fer-1/FER</i>	151	256	37.1
	SAK433-7	<i>fer-1/FER</i>	126	198	38.9
<i>fer-1/fer-1</i> untransformed control	SAK429-1	<i>fer-1/fer-1</i>	141	33	81.0
	SAK429-2	<i>fer-1/fer-1</i>	153	39	79.7
	SAK429-3	<i>fer-1/fer-1</i>	138	39	78.0
	SAK429-4	<i>fer-1/fer-1</i>	159	33	82.8
	SAK429-5	<i>fer-1/fer-1</i>	168	39	81.2
<i>pFER::FER[K-R]-GFP/-</i> (complements)	SAK91-3	<i>FER/FER</i>	14	336	4.0
	SAK91-6	<i>FER/FER</i>	12	387	3.0
	SAK91-7	<i>FER/FER</i>	16	302	5.0
	SAK91-2	<i>fer-1/FER</i>	103	310	24.9
	SAK91-5	<i>fer-1/FER</i>	84	338	19.9
	SAK91-9	<i>fer-1/FER</i>	78	268	22.5
	SAK116-2	<i>fer-1/fer-1</i>	91	130	41.2
	SAK116-3	<i>fer-1/fer-1</i>	81	102	44.3
	SAK118-5	<i>fer-1/fer-1</i>	86	129	40.0
	SAK118-6	<i>fer-1/fer-1</i>	88	128	40.7
<i>pFER::FER[K-A]-GFP/-</i> (complements)	SAK839-5	<i>fer-1/FER</i>	73	167	30.4
	SAK839-9	<i>fer-1/FER</i>	75	214	26.0
	SAK839-12	<i>fer-1/FER</i>	54	226	19.3
	SAK839-15	<i>fer-1/FER</i>	77	213	26.6
	SAK839-20	<i>fer-1/FER</i>	73	222	24.7
	SAK839-10	<i>fer-1/FER</i>	64	228	21.9

	SAK839-11	<i>fer-1/fer-1</i>	126	160	44.1
	SAK839-27	<i>fer-1/fer-1</i>	100	156	39.1
<i>pFER::FER[K-E]-GFP/- (complements)</i>	SAK840-6	<i>fer-1/FER</i>	59	182	24.5
	SAK840-19	<i>fer-1/FER</i>	70	185	27.5
	SAK840-23	<i>fer-1/FER</i>	64	205	23.8
	SAK840-27	<i>fer-1/FER</i>	75	206	26.7
	SAK840-10	<i>fer-1/fer-1</i>	100	88	53.2
<i>pFER::FER[STS-AAA]-GFP/- (complements)</i>	SAK309-3	<i>fer-1/FER</i>	50	117	29.9
	SAK309-5	<i>fer-1/FER</i>	50	168	22.9
	SAK393-1	<i>fer-1/FER</i>	103	279	27.0
	SAK393-2	<i>fer-1/FER</i>	74	323	18.6
	SAK393-3	<i>fer-1/FER</i>	75	255	22.7
	SAK393-4	<i>fer-1/FER</i>	74	239	23.6
<i>pFER::FER[STS-DDD]-GFP/- (does not complement)</i>	SAK92-2	<i>FER/FER</i>	12	288	4.0
	SAK92-6	<i>FER/FER</i>	5	190	2.6
	SAK92-7	<i>FER/FER</i>	6	203	2.9
	SAK92-8	<i>FER/FER</i>	9	180	4.8
	SAK92-12	<i>FER/FER</i>	9	221	3.9
	SAK446-2	<i>fer-1/FER</i>	159	220	42.0
	SAK446-4	<i>fer-1/FER</i>	146	208	41.2
	SAK446-11	<i>fer-1/FER</i>	151	190	44.3
	SAK446-14	<i>fer-1/FER</i>	150	221	40.4
	SAK446-15	<i>fer-1/FER</i>	144	201	41.7
	SAK446-16	<i>fer-1/FER</i>	142	222	39.0
	SAK446-17	<i>fer-1/FER</i>	164	224	42.3
	SAK446-18	<i>fer-1/FER</i>	156	218	41.7
	SAK446-20	<i>fer-1/FER</i>	141	231	37.9
<i>pFER::FER[S695A]-GFP/- (complements)</i>	SAK305-1	<i>fer-1/FER</i>	38	118	24.4
	SAK305-2	<i>fer-1/FER</i>	36	117	23.5
	SAK305-4	<i>fer-1/FER</i>	43	127	25.3
<i>pFER::FER[T696A]-GFP/- (complements)</i>	SAK306-3	<i>fer-1/FER</i>	47	103	31.3
	SAK306-4	<i>fer-1/FER</i>	53	118	31.0
	SAK306-5	<i>fer-1/FER</i>	44	125	26.0
<i>pFER::FER[S701A]-GFP/- (complements)</i>	SAK308-1	<i>fer-1/FER</i>	44	168	20.8
	SAK308-3	<i>fer-1/FER</i>	38	145	20.8

<i>pFER::FER[S695A-T696A]-GFP/-</i> (complements)	SAK307-3	<i>fer-1/FER</i>	34	167	16.9
	SAK307-4	<i>fer-1/FER</i>	36	194	15.7
<i>pFER::FER[S695D]-GFP/-</i> (complements)	SAK96-1	<i>fer-1/fer-1</i>	74	81	47.7
	SAK96-4	<i>fer-1/fer-1</i>	69	73	48.6
	SAK96-5	<i>fer-1/FER</i>	56	164	25.5
<i>pFER::FER[T696D]-GFP/-</i> (complements)	SAK123-1	<i>fer-1/fer-1</i>	71	80	47.0
	SAK123-2	<i>fer-1/fer-1</i>	56	71	44.1
<i>pFER::FER[S701D]-GFP/-</i> (complements)	SAK124-1	<i>fer-1/FER</i>	52	179	22.5
	SAK124-7	<i>fer-1/FER</i>	42	141	23.0
	SAK124-28	<i>fer-1/fer-1</i>	46	58	44.2
<i>pFER::FER[S695D-T696D]-GFP/-</i> (complements)	SAK122-2	<i>fer-1/fer-1</i>	71	74	49.0
	SAK122-5	<i>fer-1/fer-1</i>	76	76	50.0
	SAK122-6	<i>fer-1/fer-1</i>	67	77	46.5

**Supplementary Table S3: Primers used to generate complementation constructs**

Primer name	Sequence
SKpromFER-attB1	ggggacaagttgtacaaaaaagcaggctggtaagcttcgatttaagcg
promFER-R	cgatcaagagcacttctccg
FERIC-F	gctattattgcaggcgcagc
attB2FERIC-R	ggggaccactttgtacaagaaagctgggtacgtcccttggattcatgatc
FEREC-R	cgtattgcttttcgatttct
ANX1FER-F	cggagaagtgccttgatcgatgagcgggaaaactcggattc
ANX1FER-R	actggctgcgcctgcaataatagcgtgtctctttcgttcttg
FERANX1-F	aggaaatcgaaaagcaatacggcttcatcattggttcggc
attB2ANX1-R	ggggaccactttgtacaagaaagctgggttcgtccttgggatttacaatc
HERK1FER-F	cggagaagtgccttgatcgatgggtattgaaaagttgaaac
HERK1FER-R	actggctgcgcctgcaataatagcccgagattactcttactgc
FERHERK1-F	aggaaatcgaaaagcaatacgtgattgtaggttcagccat
attB2HERK1-R	ggggaccactttgtacaagaaagctgggttcttccttcagatttcaccag
SKFER-17FattB1	aaaaagcaggcttcggaagcttcgatttaagcg
SKFER-15RattB2	agaaagctgggtgaccaccaacaccaagcacc
SKFERcdBRI-F	aggaaatcgaaaagcaatacggctggtagtggtggcggatgg
SKBRI1-attB2	ggggaccactttgtacaagaaagctgggttaattttccttcaggaaacttctttatac

**Supplementary Table S4: Primer combinations used for fusion constructs**

<b>Construct</b>	<b>Primers</b>
<b><i>pFER::ANX1-GFP</i></b>	SKpromFER-attB1
	promFER-R
	ANX1FER-F
	attB2ANX1-R
<b><i>pFER::HERK1-GFP</i></b>	SKpromFER-attB1
	promFER-R
	HERK1FER-F
<b><i>pFER::ANX1[ECD]-FER[ICD]-GFP</i></b>	SKpromFER-attB1
	promFER-R
	ANX1FER-F
	ANX1FER-R
	FERIC-F
	attB2FERIC-R
<b><i>pFER::FER[ECD]-ANX1[ICD]-GFP</i></b>	SKpromFER-attB1
	FEREC-R
	FERANX1-F
	attB2ANX1-R
<b><i>pFER::FER[ECD]-HERK1[ICD]-GFP</i></b>	SKpromFER-attB1
	FEREC-R
	FERHERK1-F
	attB2HERK1-R
<b><i>pFER::FER<math>\Delta</math>ICD-GFP</i></b>	SKFER-17FattB1
	SKFER-15RattB2
<b><i>pFER::FER[ECD]-BRI1[ICD]-GFP</i></b>	SKpromFER-attB1
	FEREC-R
	SKFERecdBRI-F
	SKBRI1-attB2

**Supplementary Table S5: Primers used for site-directed mutagenesis**

SKFER-5F	tcacgaattccgggtgcttggtgtgg
SKFER-6R	tcacggatccacagaacatctccattg
SKferS695DFor	gtagacacagttgtgaaaggaag
SKferS695DRev	tcacaactgtgtctacgtgtgtgtggtctagtg
SKferS695Afor	gtagccacagttgtgaaaggaag
SKferS695Arev	tcacaactgtggctacgtgtgtgtggtctagtg
SKferT696Dfor	gtaagcgatgttggtgaaaggaag
SKferT696Drev	tcacaacatcgcttacgtgtgtgtggtctagtg
SKferT696Afor	gtaagcgagttgtgaaaggaag
SKferT696Arev	tcacaactgcgcttacgtgtgtgtggtctagtg
SKferSTDDfor	gtagacgatgttggtgaaaggaag
SKferSTDDrev	tcacaacatcgcttacgtgtgtgtggtctagtg
SKferSTAAfor	gtagccgcagttgtgaaaggaag
SKferSTAArev	tcacaactgcgcttacgtgtgtgtggtctagtg
SKferS701Dfor	gagatttcggttatcttgaccag
SKferS701Drev	gataaccgaaatctcctttcacaactgtgcttac
SKferS701Afor	gagctttcggttatcttgaccag
SKferS701Arev	gataaccgaaagctcctttcacaactgtgcttac
SKferSTSDDDfor	gttgtgaaaggagatttcgg
SKferSTSDDDrev	atctcctttcacaacatcgcttacg
SKferSTSAAAfor	gttgtgaaaggagcttcgg
SKferSTSAAArev	agctcctttcacaactgcggttacg
SKFERKA-F	aaggtagccatcgagaggcaaccaatgtcc
SKFERKA-R	cgcgatggctacctttgtagttc
SKFERKE-F	aaggtagccatcgagaggcaaccaatgtcc
SKFERKE-R	ctcgatggctacctttgtagttc

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