

Table S1. All sequences with EFFDA-E or any of 127 combinations of allowed residues.

TABLE S1A. Motifs starting DF...

1 & 2 E/D–F/Y	3 F or Y	4 D or E	5 A or C	7 D/E/S/T	count (6)	notes
DF	DFF	DFFD	DFFDA	DFFDA-D		
				DFFDA-E		
				DFFDA-S		
				DFFDA-T		
		DFFC	DFFC	EFFDC-D		
				EFFDC-E	1	KSL10 (rice)
				EFFDC-S	1	CHCH-1 (mouse)
				EFFDC-T	1	Atg1-kinase (fungus)
	DFFE	DFFEA	DFFEA	DFFEA-D		
				DFFEA-E	1	Rhophilin-1 (mouse)
				DFFEA-S	1	Lim domain kinase-2 (bovine)
				DFFEA-T		
		DFFEC	DFFEC	DFFEC-D		
				DFFEC-E		
				DFFEC-S		
				DFFEC-T		
DFY	DFYD	DFYDA	DFYDA	DFYDA-D		
				DFYDA-E		
				DFYDA-S		
				DFYDA-T		
	DFYDC	DFYDC	DFYDC	DFYDC-D		
				DFYDC-E		
				DFYDC-S		
				DFYDC-T	1	Hir3 (fungus)
	DFYE	DFYE	DFYE	DFYE-A-D		
				DFYE-A-E		
				DFYE-A-S		
				DFYE-A-T		
	DFYEC	DFYEC	DFYEC	DFYEC-D		
				DFYEC-E		
				DFYEC-S		
				DFYEC-T		

A. All eukaryotic sequences containing each of the 32 variants of the FFAT-like motif starting DF...

TABLE S1A. Motifs starting DY...

1 & 2 E/D-F/Y	3 F or Y	4 D or E	5 A or C	7 D/E/S/T	count (11)	notes
DY	DYF	DYFD	DYFDA	DYFDA-D		
				DYFDA-E	1	Slx1 endonuclease (fungal)
				DYFDA-S		
				DYFDA-T		
		DYFE	DYFEA	DYFDC-D	1	arylsuphatase (sea urchin)
				DYFDC-E		
				DYFDC-S	1	Mating pheromone 2 (Euplotes)
				DYFDC-T		
		DYYD	DYYDA	DYFEA-D		
				DYFEA-E	1	Ycf1 plastid protein
				DYFEA-S	1	Wit2 (Arabidopsis)
				DYFEA-T		
	DYY	DYYD	DYYDA	DYFEC-D		
				DYFEC-E		
				DYFEC-S		
				DYFEC-T		
		DYYE	DYYDC	DYYDA-D		
				DYYDA-E		
				DYYDA-S	2	CQ072 (Hs unknown function)[2]
				DYYDA-T	2	Hemocyanin (octopus)[2]
		DYYE	DYYEA	DYYDC-D		
				DYYDC-E		
				DYYDC-S		
				DYYDC-T		
		DYYE	DYYEC	DYYEA-D		
				DYYEA-E		
				DYYEA-S		
				DYYEA-T		
		DYYE	DYYEC	DYYEC-D		
				DYYEC-E	1	Cation-Cl ⁻ cotransporter 1 (Arabidopsis)
				DYYEC-S	1	CHD-1 (Bombyx)
				DYYEC-T		

B. All eukaryotic sequences containing each of the 32 variants of the FFAT-like motif starting DY... (see footnote)

TABLE S1C. Motifs starting EF...

1 & 2 E/D–F/Y	3 F or Y	4 D or E	5 A or C	7 D/E/S/T	count (60)	notes
EF	EFF	EFFD	EFFDA	EFFDA-D	2	Vps13C (mouse) Dicer-like-2 (fungal)
				EFFDA-E	29	C[8] O[10] RB[7] R3[1] R11[2] Opi1
				EFFDA-S		
				EFFDA-T	2	O[2]
			EFFC	EFFDC-D	2	RB R3
				EFFDC-E	1	Mth-like 7 (fly)
				EFFDC-S		
				EFFDC-T		
		EFFE	EFFEA	EFFEA-D	1	Myosin IB light chain (Dicty.)
				EFFEA-E	2	Nop2[2]
				EFFEA-S		
				EFFEA-T		
		EFFY	EFFEC	EFFEC-D		
				EFFEC-E		
				EFFEC-S	5	
				EFFEC-T		
EFY	EFYD	EFYDA	EFYDA	EFYDA-D		
				EFYDA-E	10	O[5] Flipt1[2] R11[2] R07G3.7(worm)
				EFYDA-S	4	O[3] Rad2 DNA repair (yeast)
				EFYDA-T		
		EFYDC	EFYDC	EFYDC-D	1	nuclear phosphoprotein p8 (bovine)
				EFYDC-E		
				EFYDC-S		
				EFYDC-T		
	EFYE	EFYEA	EFYEA	EFYEA-D	2	alpha-1 acid glycoprotein (Hs lumenal)
				EFYEA-E	1	Formin-like 5 (rice)
				EFYEA-S		
				EFYEA-T	1	alpha-1 acid glycoprotein (lumenal)
		EFYEC	EFYEC	EFYEC-D	3	THEMIS1[3]
				EFYEC-E		
				EFYEC-S		
				EFYEC-T		

C. All eukaryotic sequences containing each of the 32 variants of the FFAT-like motif starting EF... (see footnote)

TABLE S1D. Motifs starting EY...

1 & 2 E/D-F/Y	3 F or Y	4 D or E	5 A or C	7 D/E/S/T	count (17)	notes
EY	EYF	EYFD	EYFDA	EYFDA-D	2	Vps13C (Hs) Mfh1 helicase (S. pombe)
				EYFDA-E	2	O[2]
				EYFDA-S		
				EYFDA-T		
			EYFC	EYFDC-D	1	Torso (tyrosine-kinase receptor, fly)
				EYFDC-E		
				EYFDC-S		
				EYFDC-T		
	EYFE	EYFEA	EYFEA-D			
			EYFEA-E	2	ALMT[2]	
			EYFEA-S	5	Orc2[4] Mrt4 (fungal)	
			EYFEA-T	1	CD46 (guinea pig)	
		EYFEC	EYFEC-D			
			EYFEC-E			
			EYFEC-S			
			EYFEC-T			
EYY	EYYD	EYYDA	EYYDA-D	2	Ferm/PDZ comtaining-1[2]	
			EYYDA-E			
			EYYDA-S			
			EYYDA-T	1	beta-glucan synthesis-assoc pr.	
		EYYDC	EYYDC-D			
			EYYDC-E			
			EYYDC-S			
			EYYDC-T			
	EYYE	EYYEA	EYYEA-D	1	COG-4 (worm)	
			EYYEA-E			
			EYYEA-S			
			EYYEA-T			
		EYYEC	EYYEC-D			
			EYYEC-E			
			EYYEC-S			
			EYYEC-T			

D. All eukaryotic sequences containing each of the 32 variants of FFAT-like motif starting EY...

Footnote: Numbers in column 6 indicate the number of proteins with a specific motif. In column 7, C, O, RB, R3 and R11 refer to homologues of CERT, OSBP, rdgB, Rab3GAP1 and rabphilin11 respectively, and numbers in square brackets indicate where the list contains multiple homologues in the same family.