

Supplementary Material

Article Title Transcription factors in fungi: TFome dynamics, three major families, and dual-specificity TFs

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Table S2. Major taxonomic assignments of the TF-like DBD families found in fungi and Microsporidia. **A.** TFDFs found in fungi and microsporidia in trace amounts (1-5 species). The second column of the section A shows the taxonomic assignment and number of species (in the parentheses) possessing the found TFDFs. Abbreviations: A - ascomycetes, B - Basidiomycetes, Z - Zygomycetes, C - Chitrids, M - Microsporidia. For details, consult Suppl. Table S3. **B.** TFDFs reliably detected in fungal and microsporidial genomes (>5 TFs per species).

IPR ID	Domain name		Found in*		
A. Rare DBDFs (found in 1-5 species)					
IPR004210	BESS motif	A(4),B(1)	Metazoa		
IPR011539	Rel homology	A(3),B(2)	Metazoa		
IPR013019	MAD homology, MH1	A(2)	Metazoa		
	Transcription factor AP-2, C-terminal	B(1)	Metazoa		
IPR001628	Zinc finger, nuclear hormone receptor-type	Z(2)	Metazoa		
IPR013524	Acute myeloid leukemia 1 (AML 1)/Runt	B(1)	Metazoa		
IPR009508	Churchill	C(1)	Metazoa		
IPR009638	Fez1	A(1)	Metazoa		
IPR003350	Homeodomain protein CUT	B(1)	Metazoa		
IPR002546	Myogenic basic muscle-specific protein	B(1)	Metazoa		
IPR000770	SAND	B(1)	Metazoa+Viridiplantae		
IPR005735	Zinc finger, LSD1-type	A(4)	Viridiplantae, plant-spec		
IPR003441	No apical meristem (NAM) protein	A(2),B(3)	Viridiplantae		
IPR003340	Transcriptional factor B3	A(1)	Viridiplantae, plant-spec		
IPR007050	HTH_10, Bacterio-opsin activator	A(1)	Archea		
IPR002831	Transcriptional regulator TrmB	A(1)	Bacteria/ Archea		
IPR006199	LexA DNA-binding region	A(3),B(2)	Bacteria		
IPR000524	Bacterial regulatory protein GntR, HTH	A(1),Z(1),M(2)	Bacteria		
IPR000847	Bacterial regulatory protein, LysR	A(2),Z(1),M(1)	Bacteria		
IPR002481	Ferric-uptake regulator	A(3),Z(1)	Bacteria		
IPR007985	Haemolysin expression modulating, HHA	B(2),Z(2)	Bacteria		
IPR013199	HTH Mga	B(2),Z(1)	Bacteria		
IPR013196	Helix-turn-helix, type 11	A(1),Z(2)	Bacteria		
IPR007911	Flagellar transcriptional activator	A(1),Z(1)	Bacteria		
IPR007737	M trans-acting positive regulator	A(1),Z(1)	Bacteria		
IPR000831	Trp repressor	A(1),B(1)	Bacteria		
IPR002577	Helix-turn-helix, HxIR type	A(1),M(1)	Bacteria		
IPR005569	Arc-like DNA binding	Z(1)	Bacteria		
IPR010330	Competence-induced protein CoiA-like	A(1)	Bacteria		
IPR013198	GTP-sensing helix-turn-helix, CodY, C-terminal	A(1)	Bacteria		

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	GTP-sensing transcriptional pleiotropic repressor	A (4)	Destaria
IPR010312		A(1)	Bacteria
IPR000281	HTH_6, Helix-turn-helix protein RpiR	A(1)	Bacteria
	LytTr DNA-binding region	A(1)	Bacteria
IPR004356	P pili regulatory PapB protein	Z(1)	Bacteria
IPR005650	Penicillinase repressor	A(1)	Bacteria
IPR001867	Signal transduction response regulator, C-terminal	A(2),Z(1),M(1)	Bacteria
IPR009300	Transcriptional activator RinB	A(1)	Bacteria / Viruses
IPR003176	Viral DNA-binding protein	A(3),B(1),Z(1)	Viruses
	Alpha trans-inducing protein (Alpha-TIF)	B(2),Z(2)	Viruses
	TF MotA, C-terminal, bacteriophage	B(2),C(1)	Viruses
IPR007031	Poxvirus late transcription factor VLTF3 like	C(2)	Viruses
-	VITF-3 subunit protein	A(1)	Viruses
		Γ.(1)	VIIUSES
B. DBDFs t	ypically found in fungi (in >5 species)		
			FUNGAL+Various eukar groups, 1-
IPR001138	Zn_cluster		5 species in each
IPR003163			FUNGAL
	Copper fist DNA-binding		FUNGAL
IPR006856	Mating-type protein MAT alpha 1		FUNGAL
			ALL but not true fungi so far: Found
	Nucleic acid-binding, OB-fold		in Opisthokonta incertae sedis
	Pathogenesis-related transcriptional factor and ERF, DNA-binding		ALL but not metazoans
	Helix-turn-helix type 3	ALL but mainly bacteria	
	Helix-turn-helix, Psq		ALL (eukar+bact)
	Homeodomain-like		ALL
	Putative DNA binding	ALL	
	Cold-shock protein, DNA-binding	ALL	
IPR018004	KilA, N-terminal/APSES-type HTH, DNA-binding		ALL
IPR009044	LeeDNIA binding transprintional regulator		
	ssDNA-binding transcriptional regulator		All eukar (plants)
IPR009044 IPR008895	YL1 nuclear		All eukar (Metazoa)
IPR008895	YL1 nuclear		All eukar (Metazoa) Ichthyosporea, Nucleariidae and
IPR008895 IPR007604	YL1 nuclear CP2 transcription factor		All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida
IPR008895 IPR007604 IPR012294	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ?
IPR008895 IPR007604 IPR012294 IPR005011	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar+Archea
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR000814	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar+Archea Eukar+Archea+(Viruses only 3)
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR000814 IPR004827	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various Eukar - various
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289 IPR007196	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various Eukar - various Eukar - various
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289 IPR007196 IPR003150	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289 IPR001289 IPR007196 IPR003150 IPR016177	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding, integrase-type	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289 IPR001289 IPR007196 IPR003150 IPR016177 IPR001878	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-kArchea Eukar+Archea Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar+viruses (only 13)
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289 IPR001289 IPR007196 IPR003150 IPR016177 IPR001878 IPR025659	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type Tubby C-terminal-like domain	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea+(Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar+viruses (only 13) All eukar (mostly plants)
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IPR008895 IPR007604 IPR012294 IPR005011 IPR003958 IPR003958 IPR004827 IPR001289 IPR007196 IPR007196 IPR0016177 IPR001878 IPR025659 IPR007087 IPR000571	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type Tubby C-terminal-like domain Zinc finger, C2H2-type Zinc finger, CCCH-type	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar+viruses (only 13) All eukar All eukar
IPR008895 IPR007604 IPR012294 IPR005011 IPR003958 IPR003958 IPR004827 IPR001289 IPR007196 IPR0016177 IPR001878 IPR025659 IPR007087 IPR000571 IPR000679	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type Tubby C-terminal-like domain Zinc finger, C2H2-type Zinc finger, CCCH-type Zinc finger, GATA-type	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar +viruses (only 13) All eukar All eukar All eukar
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IPR008895 IPR007604 IPR012294 IPR012294 IPR010770 IPR003958 IPR003958 IPR004827 IPR004827 IPR001289 IPR007196 IPR001878 IPR025659 IPR007087 IPR000571 IPR004823 IPR004823 IPR00481 IPR004823 IPR004823 IPR004181 IPR00571 IPR004823 IPR004823 IPR004823 IPR004181 IPR013932 IPR013932 IPR013932 IPR013656 IPR004198 IPR013921	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type Tubby C-terminal-like domain Zinc finger, CCH-type Zinc finger, GATA-type TATA box binding protein associated factor (TAF) Zinc finger, MIZ-type TATA-binding protein interacting (TIP20) Zinc finger, GRF-type Zinc finger, GRF-type Zinc finger, C5HC2-type TATA-binding related factor	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar+Archea Eukar+Archea (Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar - various All eukar All eukar
IPR008895 IPR007604 IPR012294 IPR012294 IPR005011 IPR010770 IPR003958 IPR004827 IPR004827 IPR001289 IPR007196 IPR001878 IPR0016177 IPR001878 IPR00571 IPR000571 IPR0004823 IPR004181 IPR000967 IPR010666 IPR003656 IPR004198 IPR013921 IPR000007	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type Tubby C-terminal-like domain Zinc finger, CCH-type Zinc finger, GATA-type TATA box binding protein associated factor (TAF) Zinc finger, MIZ-type TATA-binding protein interacting (TIP20) Zinc finger, GRF-type Zinc finger, GF-type Zinc finger, C5HC2-type TATA-binding related factor Tubby, C-terminal	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-Archea Eukar+Archea+(Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar - various All eukar (mostly plants) All eukar All eukar
IPR008895 IPR007604 IPR012294 IPR005011 IPR010770 IPR003958 IPR00814 IPR004827 IPR001289 IPR001289 IPR001289 IPR001289 IPR001289 IPR001289 IPR001878 IPR001878 IPR00571 IPR004823 IPR004856 IPR00485	YL1 nuclear CP2 transcription factor Transcription factor TFIID, C-terminal/DNA glycosyla SART-1 protein SGT1 Transcription factor CBF/NF-Y/archaeal histone TATA-box binding Basic-leucine zipper (bZIP) transcription factor CCAAT-binding transcription factor, subunit B CCR4-Not complex component, Not1 DNA-binding RFX DNA-binding, integrase-type Zinc finger, CCHC-type Tubby C-terminal-like domain Zinc finger, CCH2-type Zinc finger, GATA-type TATA box binding protein associated factor (TAF) Zinc finger, MIZ-type TATA-binding protein interacting (TIP20) Zinc finger, GRF-type Zinc finger, GF-type Zinc finger, C5HC2-type TATA-binding related factor Tubby, C-terminal Zinc finger, A20-type	se, N-terminal	All eukar (Metazoa) Ichthyosporea, Nucleariidae and Fonticulagroup, Choanoflagellida ? Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar-ubiquitous Eukar+Archea Eukar+Archea (Viruses only 3) Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various Eukar - various All eukar - various All eukar All eukar
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IPR000232	Heat shock factor (HSF)-type, DNA-binding	All eukar
	Transcription factor, MADS-box	All eukar
	Helix-loop-helix DNA-binding	All eukar
IPR004022		All eukar
	GCN5-like 1	All eukar
	Transcription factor E2F/dimerisation partner (TDP)	All eukar
		Metazoa+Fungi (some other eukar
IPR000818	TEA/ATTS	in trace amounts)
IPR024061	NDT80 DNA-binding domain	Fungi+Metazoa
IPR001356		Metazoa
IPR008967	p53-like transcription factor, DNA-binding	metazoa
IPR008917	Eukaryotic transcription factor, Skn-1-like	Metazoa
IPR015988	STAT transcription factor, coiled coil	Metazoa
IPR000418		Metazoa
	Transcriptional regulator, GCM-like	metazoa
	Paired box protein, N-terminal	Metazoa
	DM DNA-binding	Metazoa
IPR010919		Metazoa
	Transcription factor, T-box	Metazoa
	Maf transcription factor	Metazoa
	POU-specific	Metazoa
	YABBY protein	Plants+Fungi
	Transcription factor, SBP-box	Plants only
	DNA-binding WRKY	Plant
	DNA-binding protein Tfx	archea
	Transcriptional regulator, Rrf2	Bacteria+ <mark>archea</mark>
	Ribbon-helix-helix	Bacteria+archea
	Helix-turn-helix, AraC type	Bacteria
	Lambda repressor-like, DNA-binding	Bacteria
IPR002197		Bacteria
	Bacterial regulatory protein, LuxR	Bacteria
	Bacterial regulatory protein, Lacl	Bacteria
	Signal transduction response regulator, C-terminal effector	Bacteria
	Bacterial regulatory protein, MarR	Bacteria
	Trp repressor/replication initiator	Bacteria
	Bacterial regulatory protein, ArsR	Bacteria
	Bacterial regulatory protein, MerR	Bacteria
	Bacterial regulatory protein, DeoR N-terminal	Bacteria
IPR001808	Bacterial regulatory protein, Crp	Bacteria

* - in addition to fungi.