

A

	# of FG Repeats	Plurality Motif	
<i>S. cerevisiae</i>	270	FXFG	16%
		GLFG	20%
		FXF	2%
		XFGX	62%
<i>H. sapiens</i>	205	FXFG	20%
		GLFG	4%
		XFGX	76%
<i>T. brucei</i>	286	FXFG	8%
		[A/V]FGQ	34%
		GGFG[A/Q]	22%
		[F]GFG	7%
		VFGT	6%
		[S/T]FG	2%
		XFGX	21%

B

Nucleoporin class	Number of Nups		
	<i>S. cerevisiae</i>	<i>H. sapiens</i>	<i>T. brucei</i>
Structural scaffold	12	14	10
FG Nups	12	9	9
Transmembrane	3	2	not found
Other	3	3	3
Total	30	28	22

Fold Type/Domain	Number of Folds/Domains		
	<i>S. cerevisiae</i>	<i>H. sapiens</i>	<i>T. brucei</i>
α -solenoid	10	10	9
β -propeller	8	10	5
FG repeat	12	11	10
Autoproteolytic	3	1	1