

	PCF (Spectrum count)	Reservosome fraction (# unique peptides)		
		B1TU	B1MTU	B1TG
<b>Vertommen et al - mitochondrial proteins</b>				
2-Oxoglutarate dehydrogenase E1 component	60	0	0	0
l-Threonine 3-dehydrogenase	42	1	1	1
2-Oxoglutarate dehydrogenase subunit	37	0	0	0
2-Amino-3-ketobutyrate coenzyme A ligase	35	0	1	0
delta-1-pyrroline-5-carboxylate dehydrogenase	34	2	2	2
2-Oxoglutarate dehydrogenase, E2 component, dihydrolipoamide succinyltransferase	16	0	0	0
ATP synthase beta chain, mitochondrial precursor	10	0	0	0
Dihydrolipoyl dehydrogenase	16	0	0	0
Glutamate dehydrogenase	16	7	9	2
Mitochondrial processing peptidase, beta subunit	14	2	7	0
<b>Vertommen et al - glycosomal proteins</b>				
fructose-bisphosphate aldolase	438	2	2	0
glyceraldehyde 3-phosphate dehydrogenase	530	0	0	0
hexokinase	305	2	3	0
phosphofructokinase	249	0	0	0
glycerol-3-phosphate dehydrogenase	241	0	1	0
glycerol kinase	196	0	0	0
glucose-6-phosphate isomerase	122	0	0	0
Hypoxanthine-guanine phosphoribosyltransferase	85	2	2	1
Glycosomal phosphoenolpyruvate carboxykinase	57	0	0	0
Arginine kinase	54	0	0	0
Triosephosphate isomerase	34	0	0	0
<b>Colasante et al - glycosomal proteins</b>				
pyruvate phosphate dikinase	73	14	23	2
phosphoenolpyruvate carboxykinase	73	9	9	4
glycerol-3-phosphate dehydrogenase	66	0	1	0
glycerol kinase	64	0	0	0
glycosomal malate dehydrogenase	57	0	0	0
fructose-bisphosphate aldolase	53	2	2	0
glyceraldehyde 3-phosphate dehydrogenase	55	0	0	0
triosephosphate isomerase	50	0	0	0
TIM, triosephosphate isomerase	50	0	0	0
beta tubulin	46	18	0	8
<b>Broadhead et al - flagellar proteins</b>				
paraflagellar rod component	38	0	0	0
flagellar radial spoke component	20	0	0	0
dynein heavy chain	50	1	0	0
microtubule-associated protein	10	0	0	0
paraflagellar rod protein 1	43	0	0	0
paraflagellar rod protein 2	58	0	0	0
paraflagellar rod component PAR4	33	0	0	0
dynein arm light chain	24	0	0	0
flagellar radial spoke component	20	0	0	0
adenylate kinase	17	0	0	0