

	IBAQ (log2)	
Protein	GFP-CTD _{AmyB}	SP-GFP
AmyB	24,28	NaN
PorV	22,54	15,67
SprA	20,88	15,66
Ppi	19,90	13,78

	IBAQ (log2)	
Protein	GFP-CTD _{RemA}	SP-GFP
RemA	28,71	17,41
PorV	24,14	15,18
SprA	20,78	14,17
Ppi	19,60	NaN

	IBAQ (log2)	
Protein	GFP-CTD _{SprB}	SP-GFP
SprB	23,47	17,97
SprF	19,25	16,43
SprA	19,28	14,17
Ppi	17,05	NaN

	IBAQ (log2)	
Protein	GFP-CTD ₃₉₅₂	SP-GFP
Fjoh_3952	28,47	14,65
Fjoh_3951	29,05	15,86
SprA	19,01	15,66
Ppi	17,98	13,78

	IBAQ (log2)	
Protein	GFP-CTD ₁₁₂₃	SP-GFP
Fjoh_1123	26,55	13,90
Fjoh_1677	17,60	NaN
SprA	14,55	15,66

	IBAQ (log2)	
Protein	GFP-CTD ₁₇₂₀	SP-GFP
Fjoh_1720	25,65	15,65
Fjoh_1677	20,56	NaN
SprA	12,66	13,21

	IBAQ (log2)	
Protein	GFP-CTD ₄₉₃₄	SP-GFP
Fjoh_4934	27,45	15,89
Fjoh_1677	24,71	13,75
SprA	13,42	12,17

Figure S2. Translocon complex proteins abundance for each sfGFP-CTD strain relative to the sfGFP_{peri} strain based on IBAQ.