

wCfeT proteins		Top Blastp hit								
Protein	Size	Accession	Size	Annotation	Taxon	Max, tot.	Cov.	E value	%ID	
WP_168464803	309	APR98706	311	Patatin-like phospholipase	wFol	514, 514	100%	0.0	78.14%	
WP_168464804	72	AZU37351	74	ParD-like family antidote	wBta	107, 107	97%	7e-29	72.86%	
WP_168464805	96	OAM00647	92	RelE/ParE family toxin	wDacA	154, 154	95%	5e-47	80.43%	
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WP_168464806	262	APR98707	257	PHA03095 (Ank repeats)	wFol	321, 321	98%	1e-107	60.31%	
WP_168464807	82	APR99034	86	XRE family transcriptional regulator	wFol	111, 111	97%	5e-30	68.75%	
WP_168464808	648	APR98618	651	DNA ligase (NAD(+)) LigA	wFol	1007, 1007	99%	0.0	73.92%	
WP_168464809	79	APR97852	96	XRE family transcriptional regulator	wFol	139, 139	98%	4e-41	89.74%	
WP_168464414	545	WP_143688845	493	Group II intron RevTranscriptase/maturase	wStr	992, 992	89%	0.0	99.59%	
WP_168464810	310	APR98606	312	Recombination-promoting nuclease/put. tnp	wFol	506, 506	100%	3e-179	79.35%	
WP_168464811	301	APR98615	302	Hypothetical protein (put. rhopty protein)	wFol	421, 421	100%	9e-146	74.17%	
WP_168464812	226	APR98945	217	DNA repair protein RadC	wFol	367, 367	96%	7e-127	84.86%	

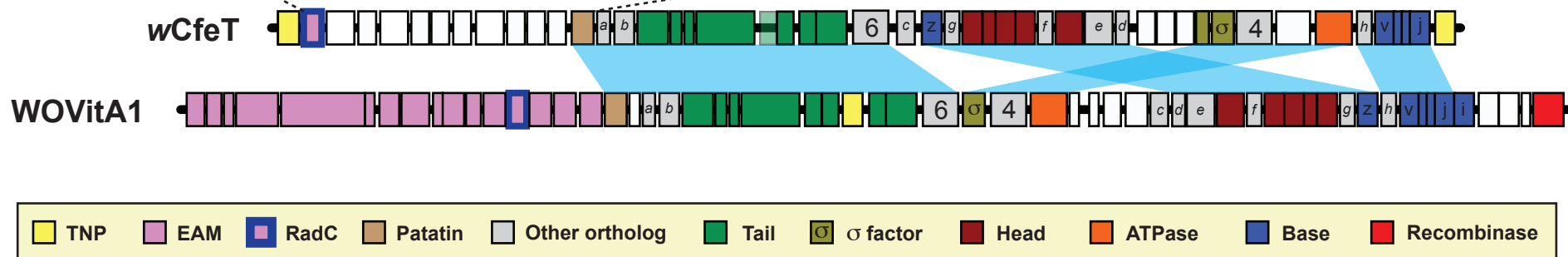


Fig. S2