CORRECTION

Correction: Mechanisms of action of *Coxiella burnetii* effectors inferred from host-pathogen protein interactions

The *PLOS ONE* Staff

A second version of Table 6 incorrectly replaces [Table 7](#). Please see [Table 7](#) here. The publisher apologizes for the error.

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Table 7. Unknown secreted effector characterization based on host-protein interaction data.

Locus	Name	Pair testing (% identified)	Y2H protein interaction data						
			N _{host}	Top locations (%)				Individual interactions	Pathways
CBU0794	coxCC4	22	22	Nucleus 50	ER/Golg 27	Vesicle 22	Memb 10	indeterminate	focal adhesion; NF-κB immune-response
CBU0881	coxCC5	41	71	Nucleus 44	Exosome 24	Centro 13	ER/Golg 11	ESCRT; centrosome; transcription	endosomal sorting; focal adhesion; NF-κB immune-response; mitochondrial matrix
CBU1724	cetCb6	65	50	Nucleus 42	ER/Golg 30	Exosome 28	Mito 18	pleiotropic; cholesterol transport	endosomal sorting; focal adhesion; Fc receptor pathway; cell cycle phase transition; mitochondrial matrix
CBU2078	coxFIC1	67	38	Nucleus 50	Exosome 21	ER/Golg 18	Memb 18	endosomal sorting; ubiquitin processing	endosomal sorting; focal adhesion

Centro, centrosome; ER/Golg, endoplasmic reticulum/Golgi apparatus; ESCRT, *endosomal sorting complexes required for transport*; Memb, plasma membrane; Mito, mitochondria; N_{host}, number of host-pathogen protein interactions; Y2H, yeast two-hybrid.

<https://doi.org/10.1371/journal.pone.0196373.t001>

Reference

1. Wallqvist A, Wang H, Zavaljevski N, Memišević V, Kwon K, Pieper R, et al. (2017) Mechanisms of action of *Coxiella burnetii* effectors inferred from host-pathogen protein interactions. PLoS ONE 12(11): e0188071. <https://doi.org/10.1371/journal.pone.0188071> PMID: 29176882