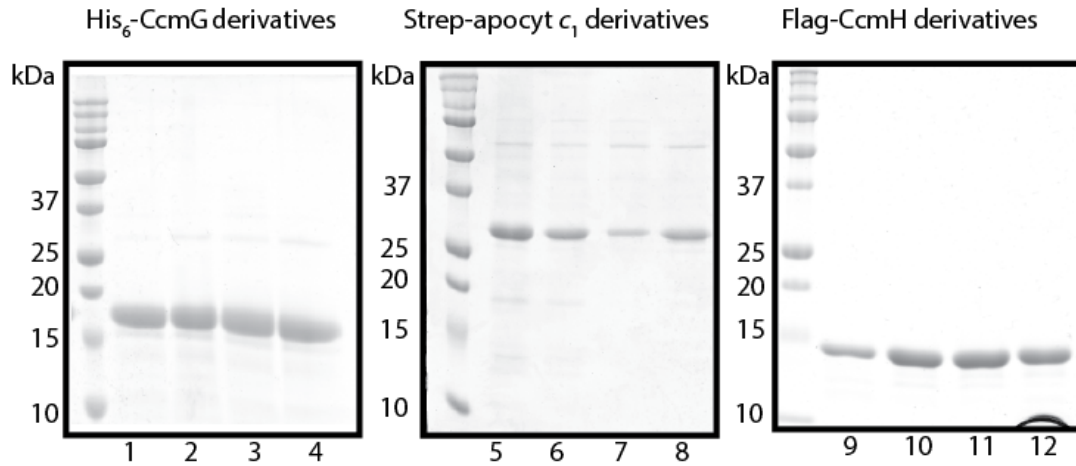


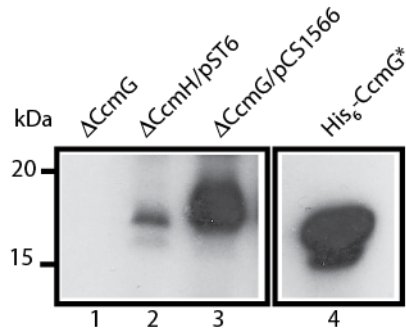
Verissimo *et al.*, Supplemental Table S1. nLC-MS/MS data used for identification of the protein bands seen in Fig. 5

Band MW	Peptide sequence	# Spectral matches	X <sub>corr</sub>	Protein	
13.5 kDa	HM*VQPDEVLSDPGLEAR	11	4.17	CcmH ( $\Sigma$ Cov 68.87)	
	FGEYVLFEPERR	9	3.48		
	SPVC*QGENIDESNAGVSR	6	4.98		
	LVAGDSDAQVIDYIK	6	3.97		
	QISQVLRSPVC*QGENIDESNAGVSR	5	2.51		
	LVAGDSDAQVIDYIKDR	2	3.86		
	ERLVAGDSDAQVIDYIKDR	1	3.37		
17.5 kDa	M*GLDWGVAGVPETFVVDGAGR	19	5.06	CcmG ( $\Sigma$ Cov 68.07)	
	VEHPNLIGLK	12	3.41		
	DTPDQAQGFLAEM*GSPYTR	12	4.54		
	LEPLGAEAPFTDADLR	12	3.59		
	KIDPLLAGTADR	8	3.42		
	IAGPLTEDVITK	7	3.78		
	LGADPGNKM*GLDWGVAGVPETFVVDGAGR	7	3.95		
	IAGPLTEDVITKK	5	2.74		
	LEPLGAEAPFTDADLRDGG	3	3.21		
	IDPLLAGTADR	3	3.27		
	QDGIEM*GVNWKDTPDQAQGFLAEM*GSPYTR	2	4.95		
	DTPDQAQGFLAEMGSPYTR	2	3.92		
	QDGIEMGVNWKDTPDQAQGFLAEM*GSPYTR	1	3.38		
	MGLDWGVAGVPETFVVDGAGR	1	2.65		
33 kDa	VEHPNLIGLK	12	3.56	CcmG ( $\Sigma$ Cov 68.07)	
	KIDPLLAGTADR	6	3.18		
	DTPDQAQGFLAEM*GSPYTR	6	4.53		
	QDGIEM*GVNWK	6	3.69		
	LEPLGAEAPFTDADLR	6	3.62		
	M*GLDWGVAGVPETFVVDGAGR	5	2.62		
	IAGPLTEDVITKK	4	3.09		
	IAGPLTEDVITK	3	3.36		
	LEPLGAEAPFTDADLRDGG	3	2.59		
	LGADPGNKM*GLDWGVAGVPETFVVDGAGR	3	5.45		
	SPVC*QGENIDESNAGVSR	6	5.21		CcmH ( $\Sigma$ Cov 60.38)
	HM*VQPDEVLSDPGLEAR	5	4.09		
	LVAGDSDAQVIDYIK	4	3.37		
	LVAGDSDAQVIDYIKDR	1	2.69		
FGEYVLFEPERR	1	3.42			

\*indicates oxidized methionine or carbamidomethylated cysteine; highlighted peptides are indicated on Fig. 5, and all MS/MS spectra are available upon request.

Verissimo *et al*, Supplemental Figure S1

**Figure S1. Purified mutant derivatives of His<sub>6</sub>-CcmG, Flag-CcmH and Strep-apocyt c<sub>1</sub>.**  
**Left panel:** Coomassie Blue stained SDS-PAGE of ~ 3 μg of Ni Sepharose HP purified His<sub>6</sub>-CcmG<sup>wt</sup> (lane 1), His<sub>6</sub>-CcmG<sup>C75</sup> (lane 2), His<sub>6</sub>-CcmG<sup>C78</sup> (lane 3) and His<sub>6</sub>-CcmG\* (lane 4).  
**Middle panel:** Coomassie Blue stained SDS-PAGE of ~ 8 μg of Strep-Tactin Sepharose purified Strep-apocyt c<sub>1</sub><sup>wt</sup> (lane 5), Strep-apocyt c<sub>1</sub><sup>C34</sup> (lane 6), Strep-apocyt c<sub>1</sub><sup>C37</sup> (lane 7) and Strep-apocyt c<sub>1</sub>\* (lane 8).  
**Right panel:** Coomassie Blue stained SDS-PAGE of ~ 3 μg of Anti-Flag (DYKDDDDK peptide) affinity gel purified Flag-CcmH<sup>wt</sup> (lane 9), Flag-CcmH<sup>C42</sup> (lane 10), Flag-CcmH<sup>C45</sup> (lane 11) and Flag-CcmH\* (lane 12).

Verissimo *et al*, Supplemental Figure S2

**Figure S2. Immunoblot analysis of *R. capsulatus* strains and purified CcmG protein for the validation of the specificity of CcmG polyclonal antibodies.** Immunoblot analysis of cell extracts from *R. capsulatus* mutants MD11 lacking CcmG ( $\Delta\text{CcmG}$ , lane 1), MD14/pST6 lacking CcmH and complemented with the plasmid pST6 carrying wild-type CcmH ( $\Delta\text{CcmH/pST6}$ , lane 2) and MD11/pCS1566 lacking CcmG and complemented with plasmid pCS1566 carrying wild-type CcmG ( $\Delta\text{CcmG/pCS1566}$ , lane 3) (**Table 1**) were used together with  $\sim 5$  ng of purified  $\text{His}_6\text{-CcmG}^*$  (a CcmG variant lacking its catalytic Cys residues) to validate that the newly raised rabbit anti-CcmG polyclonal antibodies are specific of CcmG. Note that no signal is recognized with extracts from a mutant lacking CcmG.