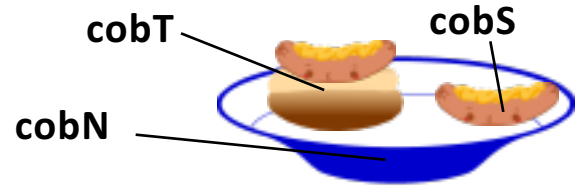


Mg-chelatase chlIDH



Co-chelatase cobNST

	Production of Mg-/Co-chelatases	Genotype(s) [<i>main taxa</i>] (reference)
A	<p>Mg-chelatase</p>	<p>chlH, chlD, chlI</p> <p><i>Proteobacteria</i></p> <p>Mg-chelatase: (Gibson et al. 1995)</p>
B	<p>Mg-chelatase</p> <p>Co-chelatase</p>	<p>cobN, cobT, cobS, chlH, chlD, chlI</p> <p><i>Proteobacteria</i></p> <p>Mg-chelatase: (Gibson et al. 1995) Co-chelatase: (Debussche et al. 1992)</p>
C	<p>Mg-chelatase</p> <p>Co-chelatase</p>	<p>cobN, chlH, chlD, chlI cobN, 2xchlH, chlD, chlI</p> <p><i>Cyanobacteria</i></p> <p>Mg-chelatase: (Gibson et al. 1995) Co-chelatase: (Rodionov et al. 2003)</p>
D	<p>Co-chelatase</p>	<p>cobN, chlD, chlI</p> <p><i>Actinobacteria</i></p> <p>Co-chelatase: (Rodionov et al. 2003)</p>
E	<p>Co-chelatase</p>	<p>cobN, fs-chlD</p> <p><i>Proteobacteria, Actinobacteria, Archaea</i></p> <p>Co-chelatase: (this study)</p>
F	<p>Co-chelatase</p>	<p>cobN, chlD</p> <p><i>Proteobacteria, Actinobacteria, Archaea</i></p> <p>Co-chelatase: (this study)</p>

- *chlD* gene with a frameshifting signal

- *fs-chlD* gene with a frameshifting signal and a frameshift mutation