Genomic Analyses of the Quinol Oxidases and/or Quinone Reductases Involved in Bacterial Extracellular Electron Transfer

YUHONG ZHONG AND LIANG SHI

Department of Biological Sciences and Technology, School of Environmental Studies, China University of Geosciences, Wuhan, Hubei, 430074, People's Republic of China. <u>liang.shi@cug.edu.cn</u>

Up to date, 3 different families of quinol oxidases and/or quinone reductases involved in bacterial extracellular electron transfer (EET) have been discovered. They are the CymA, CbcL/MtrH/MtoC and ImcH families of quinol oxidases and/or quinone reductases that are all multiheme *c*-type cytochromes (*c*-Cyts). To investigate to what extent they are distributed among microorganisms, we search the bacterial as well as archaeal genomes for the homologs of these *c*-Cyts. Search results reveal that the homologs of these *c*-Cvts are only found in the Domain Bacteria. Moreover, the CymA homologs are only found in the phylum of Proteobacteria. In contrast to CymA, CbcL/MtrH/MtoC homologs are found in 15 phyla, while ImcH homologs are found in 12 phyla. Moreover, CymA and CbcL/MtrH/MtoC homologs are suggested to catalyze both guinol-oxidizing and guinonereducing reactions. ImcH homologs are implied to be only the quinol oxidases. Finally, some bacteria have the homologs of 2 different families of *c*-Cyts, which may improve the bacterial capability to exchange electrons with extracellular substrates.