Iron-sulfur-protein biogenesis in eukaryotes

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Iron-sulfur (Fe-S) clusters (ISCs) are versatile, ancient co-factors of proteins that are involved in electron transport, enzyme catalysis and regulation of gene expression. The synthesis of ISCs and their insertion into apoproteins involves the function of complex cellular machineries. In eukaryotes, the mitochondrial ISC-assembly machinery is involved in the maturation of all cellular iron-sulfur proteins. A mitochondrial export machinery and a recently discovered cytosolic assembly system specifically participate in the maturation of cytosolic and nuclear iron-sulfur proteins. Of the ~20 assembly components, more than ten are encoded by essential genes, which indicates that the process is indispensable for life. Mutations in two of the assembly components lead to neurological diseases. The essential character of Fe-S-protein biogenesis in eukaryotes and its importance for human disease identifies this evolutionary ancient process as one of the most important biosynthetic pathways of life.

Introduction

Iron-sulfur (Fe-S) clusters (ISCs) were discovered by Helmut Beinert and others in the early 1960s [1-3]. Until the late 1990s, the complexity of the in vivo assembly process of the ISC from its constituents, Fe2+/3+ and sulfide (S^{2-}) , and its incorporation into an apoprotein was not anticipated. It is now known that bacteria possess three distinct biosynthetic systems to generate Fe-S proteins (for a review, see Ref. [4]). The first discovered system, termed nitrogen-fixation (NIF) machinery, is dedicated to the assembly of the complex Fe-S protein nitrogenase, which is responsible for the conversion of N₂ to NH₃ in nitrogen-fixing bacteria [5,6] (Figure 1a). The second system, the ISC-assembly machinery, is required for generation of the majority of cellular Fe-S proteins and, thus, might perform a general 'house-keeping' biosynthetic function [7]. The ISC-assembly machinery consists of approximately ten proteins. Finally, the SUF (sulfur-mobilization) machinery was discovered as an independent assemblage that might be used predominantly under iron-limiting or exidative-stress conditions [8-11]. Although many bacteria contain both the ISC and SUF machineries, others harbour just one of these systems (Figure 1a), Strikingly, the NIF system of the eukaryotic parasite Entamoeba histolytica can replace both the ISC and SUF systems in Escherichia coli under anaerobic conditions, which shows that, in principle, these systems are interchangeable [12].

In eukaryotes, Fe-S proteins are present in the mitochondria, the cytosol and the nucleus (Table 1).

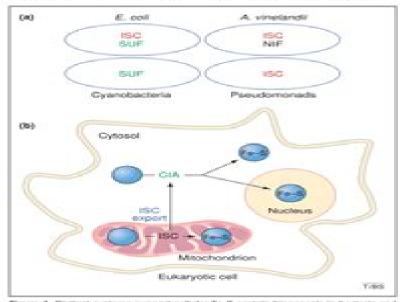


Figure 1. Distinct systems support cellular Fe-5 protein biogenesis in bacteria and autoroptes, (a) in bacteria, three systems have been identified that are capable of assembling Fe-S proteins. The ISC and SUF systems are responsible for general assembly of Fe-S proteins, whereas the NIF system present in nitrogen-foling bacteria in.g. Azotobacter vinelandili is specialized in assembling the ISCs of nitrogeness, a complex metalloprotein (5.6). Components of all these systems are usually encoded by operans. When both the ISC and SUF systems are present in one species, such as Eacherichia coil, the ISC system is responsible for biogenesis of house keeping Fe-S. proteins, in perticular, in low swygen concentrations. The SUF system is used preferentially under oxidative stress and iron limiting conditions III. Some of the SUF genes have been identified in archaelocterial genomes but, so far, nothing is known about Fe-S-protein biogenesis in Archeea. 86 In eukaryotes, such as the yeast Saccharomyces cerevisias, mitochondria sontain a machinery that shares similarity with the bacterial ISC system. This machinery is required for maturation of almost all Fe-S proteins in non-photosynthetic sukeryotes. The mitochondrial ISC-export machinery contributes an unknown factor to biogenesis of cytosolic and nuclear Fe-S proteins. The cytosolic CIA machinery functions downstream of mitochondria to accomplish Fe-S-protein assembly. Note that the situation in photosynthetic sukaryotes (plants and algae) might be similar but, in addition, they contain SUF proteins in their plastids, thus, resembling the situation in cyanobacteria. Fe-S. apoproteins are shown as blue circles.

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Albert A Gayle

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