

# *Defining D-Arabinose Metabolism in Leishmania major and Crithidia fasciculata*

Elda Iljazi and Michael A. J. Ferguson

Division of Biological Chemistry & Drug Discovery School of Life Sciences

The metabolism of D-Arabinose (D-Ara) in eukaryotes is poorly understood. Arabinose (Ara) is one of the “rare” aldopentose sugars distributed in Nature principally as a component of cell wall structures in plants and bacteria. Arabinose exists naturally in both pyranose and furanose conformations and D- and L- configurations. The most abundant form of arabinose is L-arabinose which is present in the arabinogalactans of plants. D-arabinofuranose (D-Araf) is found mainly in the arabinomannans, arabinogalactans, lipoarabinomannans and mycolylarabinogalactan-peptidogalactans of mycobacterial cell walls. However, D-arabinopyranose (D-Arap) is found, uniquely, in cell surface glycoconjugate structures of certain trypanosomatid parasites: *Leishmania major* lipophosphoglycan (LPG) *Crithidia fasciculata* lipoarabinogalactan (LAG) and *Endotrypanum schaudinni* glycoinositol phospholipids (GIPLs). The activated donor molecule of D-Arap has been identified in *L. major* as GDP- $\alpha$ -D-Arap. However, the source of the GDP-Arap is not fully understood. So far it is known that both *L. major* and *C. fasciculata* have a salvage pathway allowing the parasites to internalize D-Ara from the extracellular medium or the lumen of the insect guts and convert it to GDP- $\alpha$ -D-Arap via an arabinose-1-kinase/pyrophosphorylase. A *de novo* pathway, whereby D-Glucose (D-Glc) is converted to D-Arap via loss of the Glc C-1 carbon atom has been postulated but many details are missing. Many gram-negative bacteria have an Arabinose-5-phosphate isomerase (APIs) enzyme. In bacteria API enzymes catalyse the interconversion of D-ribulose-5-phosphate (Ru5P), the product of the oxidative phase of the pentose phosphate pathway, and D-arabinose-5-phosphate (A5P). A5P is a precursor to 3-deoxy-D-manno-octulosonic acid (KDO) that is a component of the bacterial capsular polysaccharides and lipopolysaccharides (LPS). KDO is an essential component of the cell envelope of gram-

negative bacteria. We speculate that trypanosomatids may also convert D-Glc to D-Arap via Ru5P and its isomerisation to A5P followed by dephosphorylation to D-Arap. Apart from cell surface incorporation by *L. major* and *C. fasciculata*, it is possible that D-Arap may be used by all the kinetoplastids to make D-erythroascorbate, a C5 ascorbate analogue similar in structure and physicochemical properties to ascorbate (Vitamin C). In animals and plants, L-ascorbate is involved in cellular defence against oxidative stress. The D-erythroascorbate analogue of ascorbate appears in yeast and other fungi, but this antioxidant role is not well understood. The biosynthesis of D-erythroascorbic acid starts from D-Arap, which is oxidized by NAD(P)<sup>+</sup> specific dehydrogenases to D-arabino-1,4-lactone, which is further oxidized to D-erythroascorbic acid by D-arabino-1,4-lactone oxidase. The source of the original D-Arap in yeast and fungi is, however, unknown. In summary, central to the problem of eukaryotic D-Arap metabolism is the bioconversion from D-Glc to D-Arap.