

Plant sphingolipids, from cytokinesis to membrane dynamics

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Sphingolipids were found in yeast and mammals to be essentials in several basic cellular functions such as endocytosis, protein transport, apoptosis and stress responses. A key step of the sphingolipid biosynthetic pathway is the acylation of long chain bases (LCBs) catalyzed by the sphingoid base N-acyl transferase or ceramide synthase. Three ceramide synthases homologues (named LOHs) have been identified in the model plant *Arabidopsis* with different acyl chain specificities (Markham et al. 2011). Pharmacological and genetic approaches demonstrated that very-long-acyl-chain (VLCFA)-sphingolipids are essentials for plant development. Acyl chain length is thought to be critical for biophysical properties of membrane in particular during cell division when active vesicular fusions are necessary. *In vivo* imaging of endomembrane and plasma membrane markers of VLCFA-depleted showed that sub-cellular trafficking and plasma membrane dynamics were altered during cytokinesis. *In vitro* analysis of the involvement of VLCFA-sphingolipid in vesicle fusion and aggregation will also be presented. Finally, a model presenting the specific role of sphingolipid acyl chain length in plant membranes will be discussed.