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. 4 Setting the SERIAL Communication Device on the GSM and PC side. The Developer Assistant (DA).4 Setting the Programming Port on the Server Unit. (page 20). Pl707 V4 Programming Software 13 - DOWNLOAD d77fe87ee0 documents for Schneider Electric PLC Programming Software 4. (page 20).4 Setting the Communication Device on the GSM and PC side. User Manual. (page 12). In the menu start up menu for Schneider Electric PL7 series "Programming" select the software named "PL7-07".The crystal structure of activated ferulic acid esterase-1 from Neurospora crassa. The structure of ferulic acid esterase-1 (FAE-1) from Neurospora crassa (NcFAE-1) was determined by X-ray crystallography. The protein belongs to the α/β class of hydrolytic enzymes. NcFAE-1 consists of a characteristic α/β catalytic domain and a β-sandwich domain, which is structurally similar to fungal lipase/esterase A from Aspergillus niger and the A. oryzae norskeolase (Afn1). The enzyme consists of four subdomains; α/β, α/β' and α/α' subdomains and an α-helical sandwich region. In the active site, the catalytic triad of Cys12-His199-Asp66 was observed to form a distorted tetrahedral arrangement, and the location of the catalytic histidine residue in the active site may be conserved between NcFAE-1 and feruloyl esterase from Aspergillus niger (AnFE). A disulfide linkage was observed between Cys14 and Cys199 in NcFAE-1. Activated NcFAE-1 was thought to be the same as the chemically modified form of FAE-1 or its C-terminal truncated mutant (NcΔN-FAE-1). A C-terminal helix was observed in activated NcFAE-1 structure, while the corresponding C-terminal helix in NcΔN-FAE-1 consists of only the following 24 residues. This structure provides a basis for chemical modification of the C-terminal region of FAE-1 and for the stabilization of catalytically active FAE-

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