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A: The file is not really compressed but it's encoded with unicode, so that you should be able to use the following regular expression: /[\x00-\x1F\x7F]{1,4}/ Yeast copper, zinc-superoxide dismutase: a potential biomarker for oxidative stress caused by genotoxicants. Cytosolic Cu,Zn-superoxide dismutase (Cu,Zn-SOD) is implicated in the regulation of cellular ROS concentrations. The ability to measure levels of this enzyme may aid in the determination of cellular susceptibility to redox stress. We demonstrate that Cu,Zn-SOD is expressed in Saccharomyces cerevisiae as a single band of approximately 57 kDa on immunoblots. Substantial levels of enzyme were found in cytosolic extracts of yeast cells regardless of the strain background. Levels of enzyme declined following exposure to hydrogen peroxide or the alkylating agent methyl methanesulfonate (MMS). In contrast, only a small amount of enzyme was found in a mitochondrial fraction following homogenization of cells in the presence of protease inhibitors. These results are consistent with the possibility that Cu,Zn-SOD is located within the cytosol in yeast. We observed that levels of this enzyme decreased in an SOD1-dependent manner following exposure of yeast to hydrogen peroxide. However, the levels of SOD1 mRNA and protein were not substantially altered following treatment with MMS. These results suggest that some component of the cellular ROS detoxification system other than Cu,Zn-SOD is responsible for the down-regulation of enzyme levels.We have previously published in this journal the analysis of the M4 consensus sequences from the ALK tyrosine kinase and from the mesenchymal-epithelial transition (MET) receptor. Both receptors belong to the type of RTKs that are activated by their ligands, which are themselves transmembrane proteins. In contrast, their intracellular domains have similar sequences and their ligands (ALK and HGF respectively) are normally expressed in a specific tissue or cell type, therefore it is not surprising that the mechanisms leading to activation are specific and that the recognition domain involved is also tissue specific. This has led to the choice of the extracellular domain as a target for the development of drugs for the treatment of different forms of cancer. However, the intrace 2d92ce491b