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Abstract. An attempt to alter protein surface charges through traditional protein engineering approaches often affects the native protein structure significantly and induces misfolding. This limitation is a major hindrance in modulating protein properties through surface charge variations. In this study, as a strategy to overcome such a limitation, we attempted to co-introduce stabilizing mutations that can neutralize the destabilizing effect of protein surface charge variation.

Modulation of protein stability and aggregation properties ...

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Modulation of Protein Stability: Targeting the VHL Pathway ... Abstract The post-translational modification of proteins by the covalent attachment of carbohydrates to specific side chains, or glycosylation, is emerging as a crucial process in modulating the function of proteins. In particular, the dynamic processing of the oligosaccharide can ...

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At the same time steroid substrates decrease the affinity of both CYP11B1 \square Adx and CYP11B2 \square Adx complexes, predominantly reducing their stability (4-7 fold). This finding reveals differentiation of protein-protein interactions within the mitochondrial pool of CYPs, which have the same electron donor.

Substrate-induced modulation of protein-protein ...

Abstract Inactivation of the von Hippel-Lindau tumor suppressor protein (pVHL) has been linked to a variety of tumors such as renal cell carcinoma, pheochromocytomas, and cerebellar hemangioblastomas. The best characterized of its many proposed functions is the ability to downregulate hypoxia-inducible factor- \square (HIF \square) subunits.

Modulation of Protein Stability: Targeting the VHL Pathway ...

Electrodynamic pressure modulation of protein stability in cosolvents. Damodaran S (1). Author information: (1)Department of Food Science, University of Wisconsin-Madison, Madison, Wisconsin 53706, United States. Cosolvents affect structural stability of proteins in aqueous solutions. A clear understanding of the mechanism by which cosolvents impact protein stability is critical to understanding protein folding in a biological milieu.

Electrodynamic pressure modulation of protein stability in ...

Modulation of Helix Stability to Investigate Protein-Protein Interactions in the Paramyxovirus Replication Complex . By and Victor H. Arias and Victor H. Arias. Abstract. The focus of this project is to develop a greater understanding of the protein-protein interaction between the

measles Nucleocapsid Binding Domain (NBD) protein and its ligand ...

Modulation of Helix Stability to Investigate Protein ...

To overcome this difficulty, we have established a novel approach, termed the intein-mediated modulation of protein stability (imPS), in which a destabilizing domain and part of a split intein are fused to the essential protein. The growth of the mutant virus can then be regulated by the degradation and splicing of the protein.

An intein-mediated modulation of protein stability system ...

Modulation of Protein Stability and Function by Cysteine Mutations and Signal Peptides . By Likhesh Sharma. Abstract. Chapter 1 gives a general introduction to the CXXC motif found in natural proteins. It then reviews the studies where disulphides were engineered in various proteins. The various strategies developed to engineer metal binding ...

Modulation of Protein Stability and Function by Cysteine ...

Regulation of mRNA stability and translation by RNA-binding proteins (RBPs). They mainly influence the fates of target mRNAs at the post-transcriptional levels. In the cytoplasm, stabilized mRNAs are protected from degradation leading to more protein levels. Destabilized mRNAs are driven to degradation machinery leading to lower protein levels.

Modulation of Gene Expression by RNA Binding Proteins ...

When the protein properties such as fluorescent activity, folding rate and kinetic stability were assessed, we found the possibility that the protein stability can be modulated independently of activity and folding by engineering protein surface charges. The aggregation properties of GFP could also be altered through the surface charge engineering.

Modulation of protein stability and aggregation properties ...

Direct control of protein level enables rapid and efficient analyses of gene functions in crops. Previously, we developed the RDDK-Shield1 (Shld1) system in the model plant *Arabidopsis thaliana* for direct modulation of protein stabilization using a synthetic small molecule. However, it was unclear whether this system is applicable to economically important crops.

Direct and tunable modulation of protein levels in rice ...

LSD1 regulates HIF α protein stability in a RACK1-dependent manner. The fact that LSD1 depletion blocks HIF α protein accumulation under hypoxia without affecting its mRNA expression (Figs 1 and EV1, and Appendix Fig S1) indicates that LSD1-mediated HIF α regulation is achieved at the post-transcriptional level. We therefore examined the molecular mechanism of LSD1 action by investigating its effects on the stability of HIF α protein.

Regulation of hypoxia responses by flavin adenine ...

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