

[S-5]

A ubiquitin-proteasome system as a determination factor involved in methylmercury toxicity

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The methylmercury (MeHg) is a toxic environmental pollutant, causing serious neurological and developmental effects in humans. Recent epidemiological studies have indicated that ingestion of MeHg in fish during pregnancy can result in neuroethological effects in the offspring. However, the mechanism underlying the MeHg-toxicity is not fully understood.

To elucidate the mechanisms of toxicity of MeHg and of defense against MeHg, we searched for factors that determine the sensitivity of yeast cells to MeHg, and found that overexpression of Cdc34, a ubiquitin-conjugating enzyme (E2) that is a component of the ubiquitin-proteasome (UP) system, induces a resistance to MeHg toxicity in both yeast and human cells. The UP system is involved in the intracellular degradation of proteins. When Cdc34 is overexpressed in cells, ubiquitination reactions are activated and the degradation of certain proteins by the UP system is enhanced. Therefore, it seems likely that certain as-yet-unidentified proteins that increase MeHg toxicity might exist in cells and that toxicity might be reduced by the enhanced degradation of such proteins, mediated by the UP system, when Cdc34 is overexpressed.

SCF ubiquitin-ligase is a component of UP system and consists of Skp1, the scaffold protein Cdc53, the RING-finger protein Hrt1, and one member of the family of F-box proteins. The F-box proteins directly bind to the substrates and are the determinants of substrate specificity of SCF. Therefore, we searched for the F-box protein that confers resistance to MeHg, and found that overexpression of Hrt3 or Yil224w induced resistance to MeHg toxicity in yeast cells. Since the protein(s) that enhance toxicity of MeHg might plausibly be induced in substrates of both F-box proteins, we next searched for substrate proteins that are recognized by Hrt3 or Ylr224w using two-hybrid screen. We found that Did3 or Grs1 interacts with Hrt3; and Eno2 interacts with Yir224w. The yeast cells that overexpressed each those proteins showed hypersensitivity to MeHg, respectively, indicating that those proteins enhance the MeHg toxicity. Both Dld3 and Eno2 are proteins involved in the synthesis of pyruvate, and


overexpression of both proteins might induce increase in interacellular levels of pyruvate. Deletion of Yil006w that transports pyruvate into the mitochondria induced a resistance to MeHg. These results suggest that the promotion of the pyruvate inflow into the mitochondria might enhance MeHg toxicity. This study provides important key for the elucidation of the molecular mechanism of MeHg toxicity.

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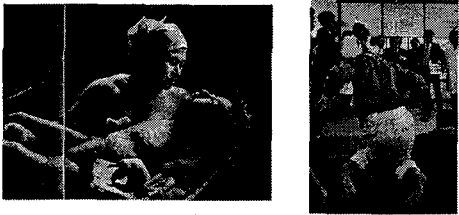

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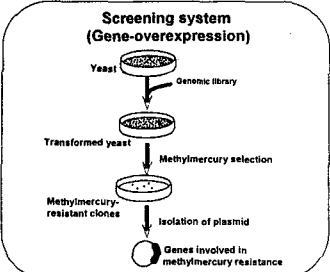
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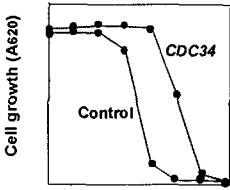
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Model organism : Yeast (*Saccharomyces cerevisiae*)
Powerful genetic approaches can be used to elucidate fundamental but complex eukaryotic processes

Screening system (Gene-overexpression)



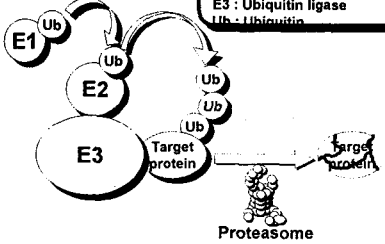
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Effects of overexpression of Cdc34 on sensitivity of yeast to methylmercury (MeHg)

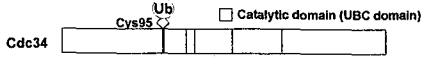
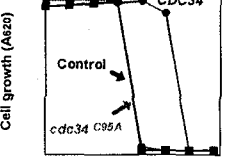
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E1 : Ubiquitin-activating enzyme
E2 : Ubiquitin-conjugating enzyme
E3 : Ubiquitin ligase
Ub : Ubiquitin

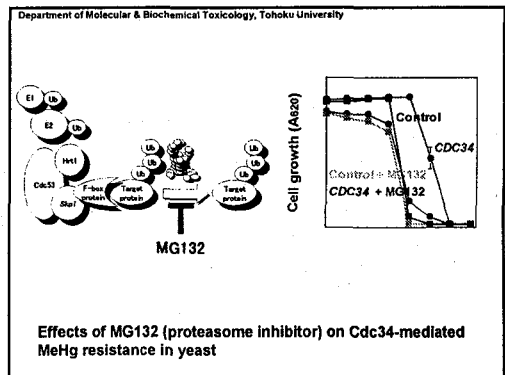
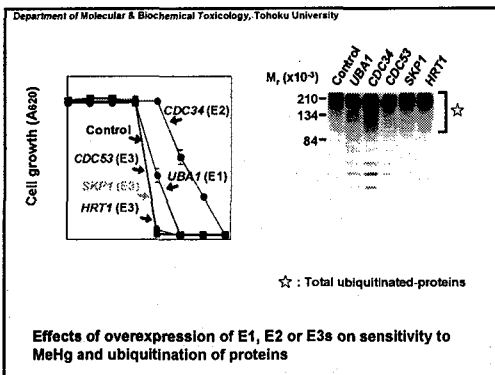
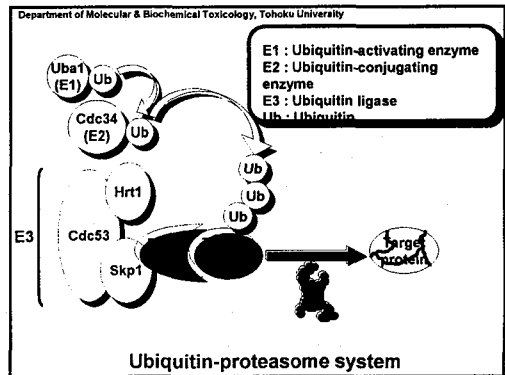
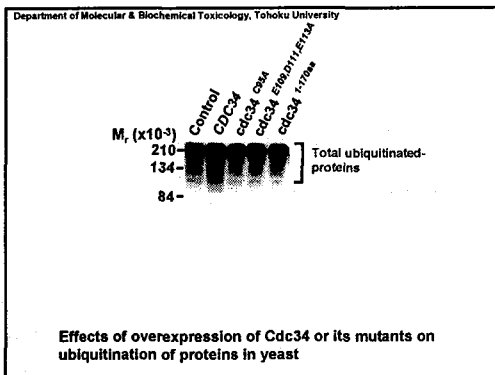
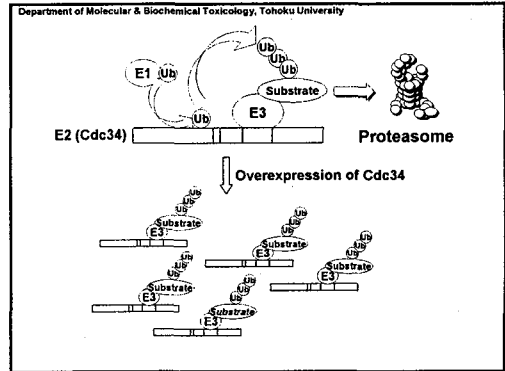
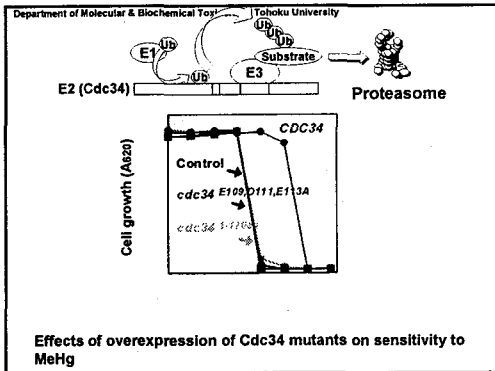


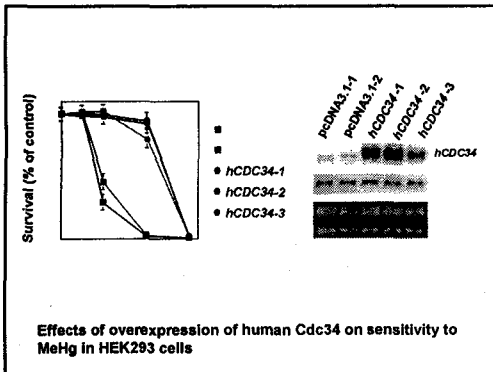
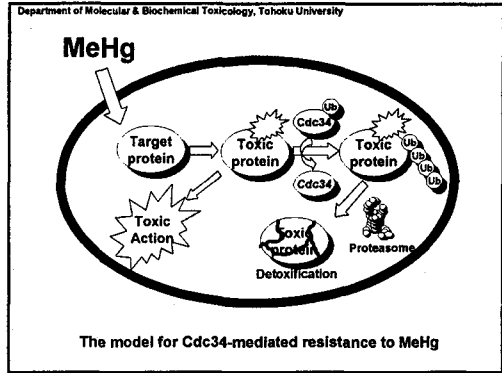
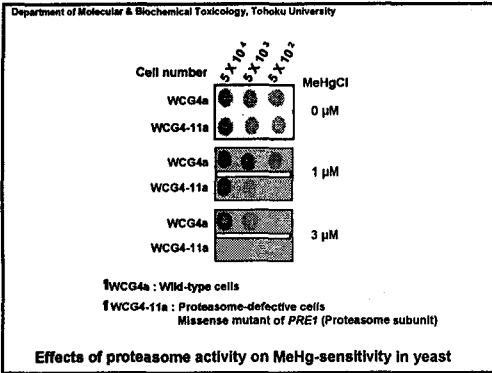
Ubiquitin-proteasome system

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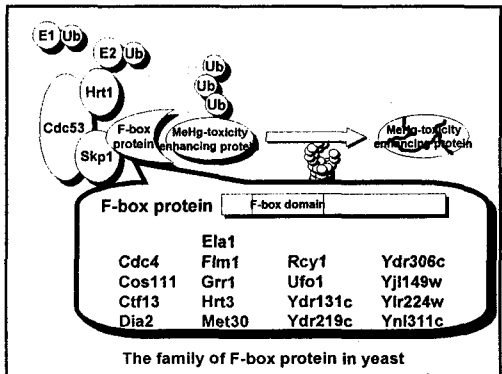
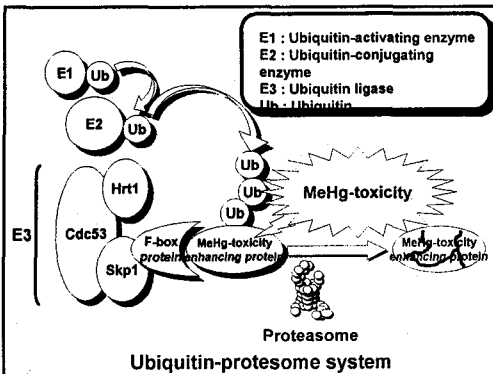



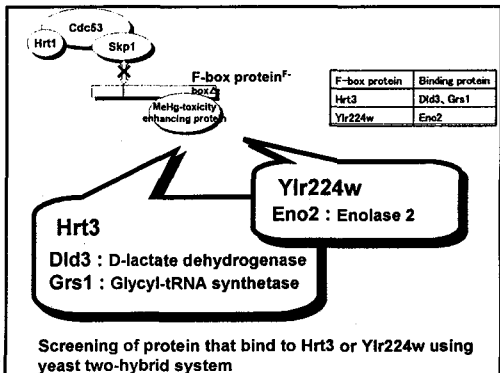
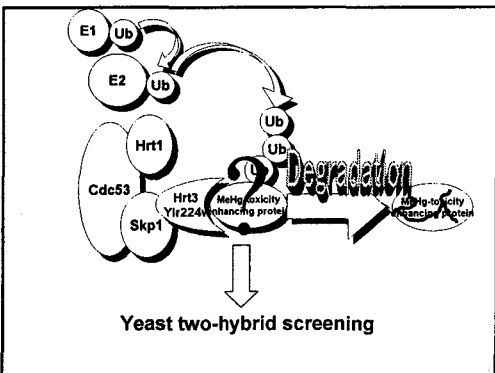
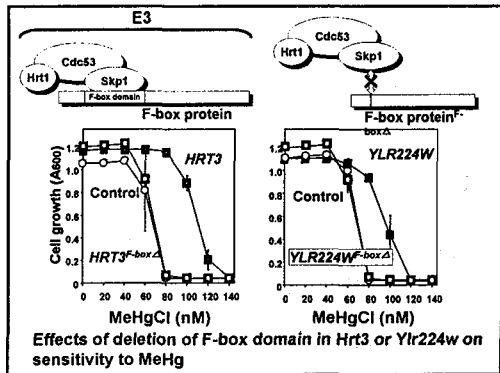
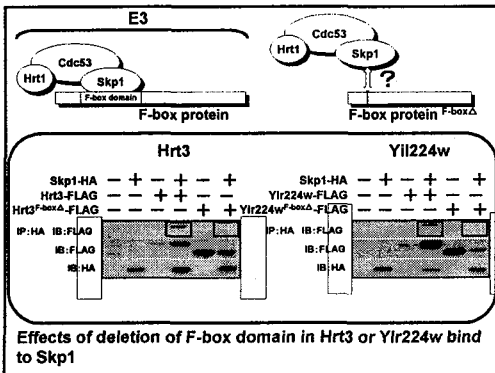
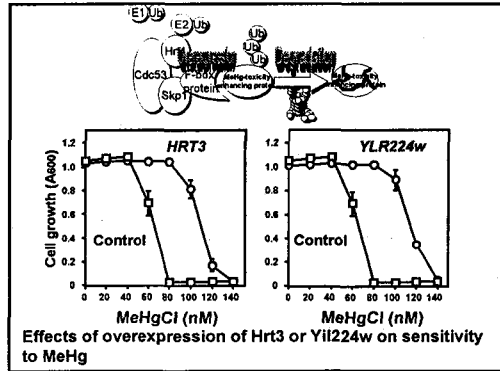
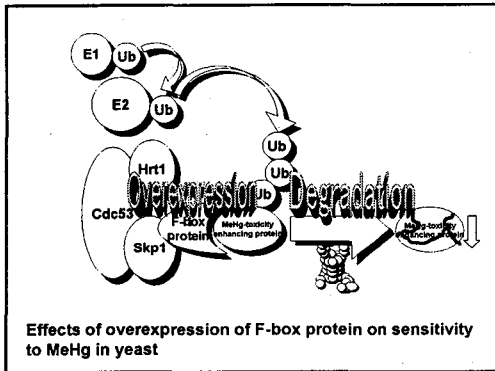
Effects of overexpression of Cdc34 mutant (point mutation of ubiquitin-binding site) on sensitivity to MeHg

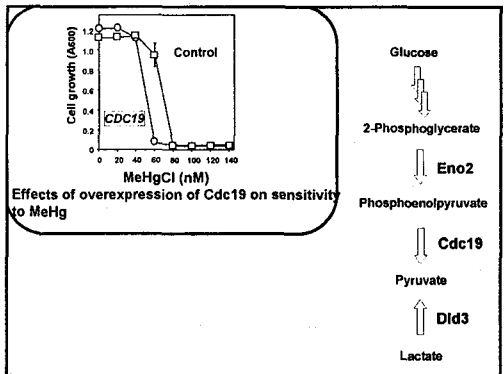
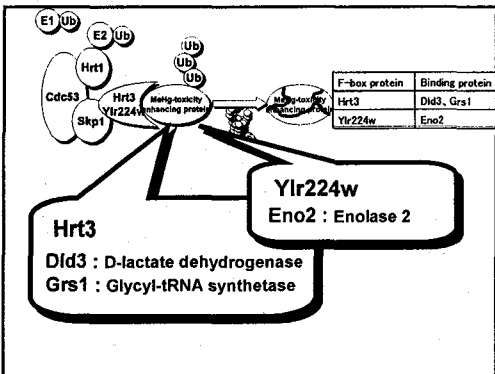
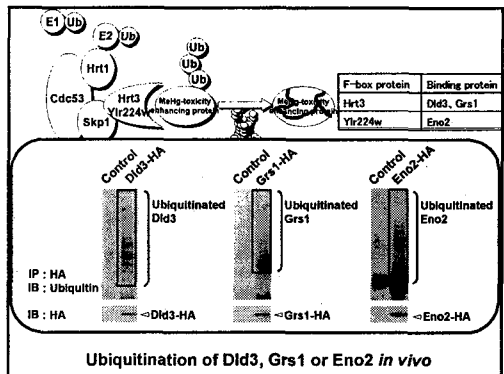
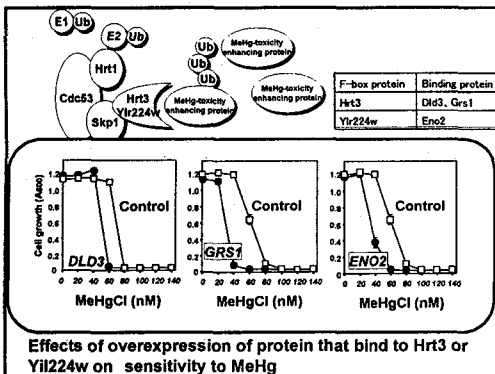
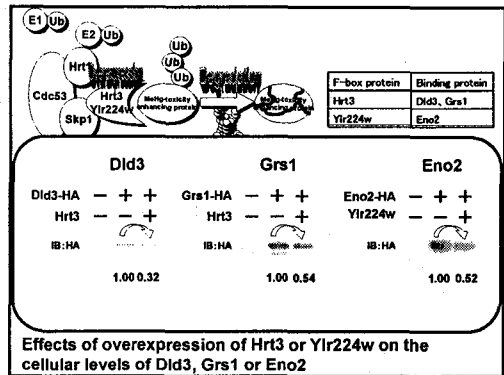
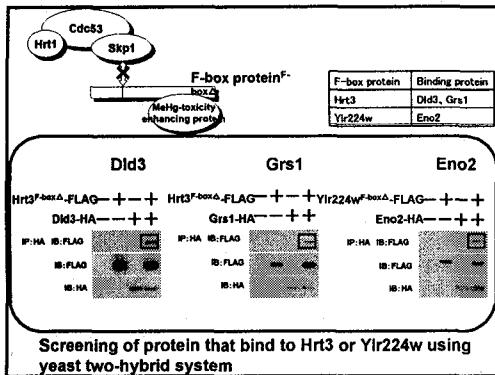


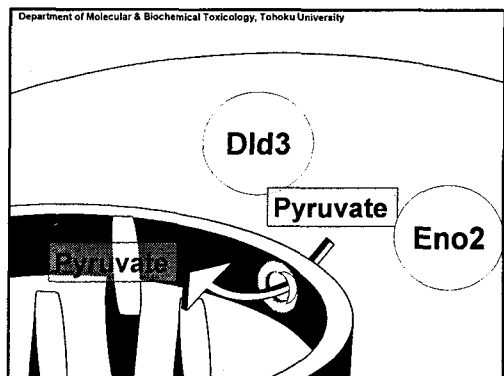
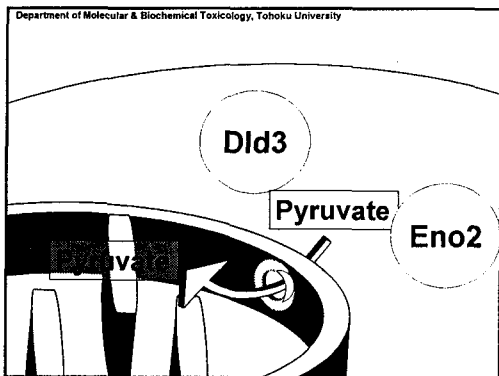
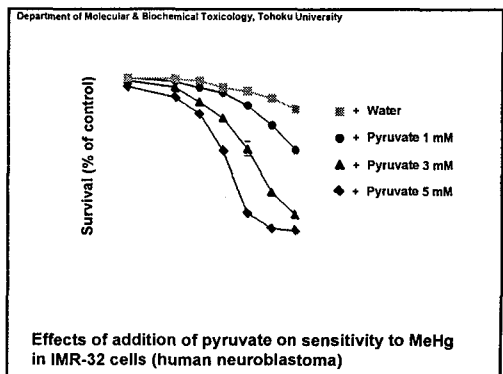
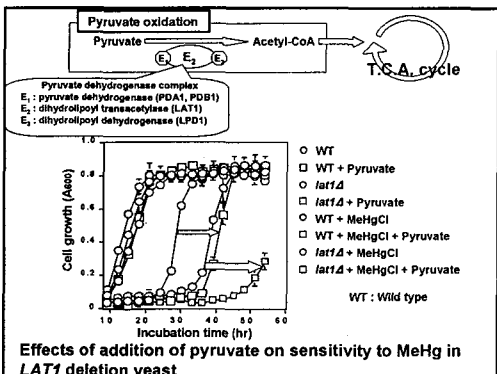
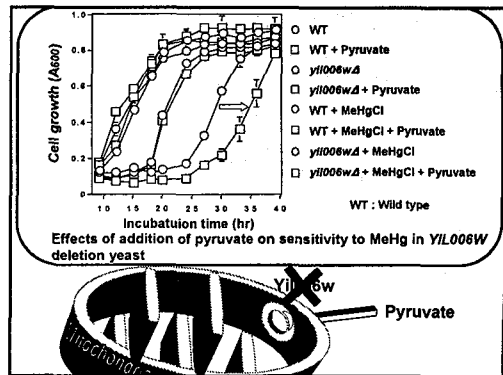
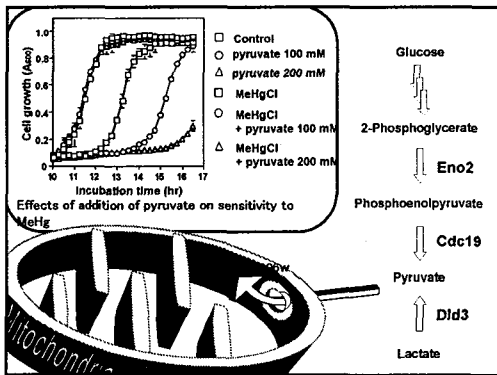



Screening of MeHg-toxicity enhancing protein











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