

Identification of a chemical chaperone for preventing protein aggregation and proteotoxicity under endoplasmic reticulum stress

Keisuke Kitakaze^{1,2}, Shusuke Taniuchi¹, Eri Kawano¹, Yoshimasa Hamada¹, Masato Miyake¹, Miho Oyadomari¹, Hirotatsu Kojima³, Hidetaka Kosako⁴, Tomoko Kuribara⁵, Suguru Yoshida⁵, Takamitsu Hosoya⁵, Seiichi Oyadomari¹

¹*Div. of Mol. Biol., Inst. of Adv. Med. Sci., Tokushima Univ.*, ²*Dept. of Pharmacol., Kawasaki Med. Sch.*, ³*DDI, The Univ. of Tokyo*, ⁴*Fujii Memorial Inst. of Med. Sci., Inst. of Adv. Med. Sci., Tokushima Univ.*, ⁵*Lab. of Chem. Biosci., Inst. of Biomater. and Bioeng., TMDU*

Endoplasmic reticulum (ER) is responsible for protein biosynthesis and folding, but accumulation of unfolded proteins leads to disturbance of ER proteostasis and subsequent clinical pathologies including diabetes, neurodegenerative disease and cancer. Chemical chaperones are chemical compounds that help protein folding and suppress aggregation, and receiving increased attention as potential therapeutic approaches for ER stress-related diseases. In this study, we established a novel ER stress reporter cell line and identified compound X as a chemical chaperone from the 217,765-compound chemical library. Compound X directly binds to secreted or membrane proteins and inhibits protein aggregation during tunicamycin induced ER stress. Furthermore, compound X significantly prevented cell death caused by chemically induced ER stress and by an aggregation-prone mutant prion protein. These results show the therapeutic potential of compound X as a chemical chaperone that can ameliorate ER stress-related diseases.