

## Poster Presentation

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### *Brazzein and structurally similar proteins: structural/functional comparisons*

K. Nagata<sup>1</sup>, N. Hongo<sup>1</sup>, Y. Kameda<sup>1</sup>, A. Yamamura<sup>1</sup>, H. Sasaki<sup>1</sup>, W. Lee<sup>1</sup>, K. Ishikawa<sup>2</sup>, E. Suzuki<sup>2</sup>, M. Tanokura<sup>1</sup>

<sup>1</sup>The University of Tokyo, Graduate School of Agricultural and Life Sciences, Tokyo, Japan, <sup>2</sup>Ajinomoto Co. Inc., Institute for Innovation, Kanagawa, Japan

Brazzein, a 6.5-kDa protein consisting of 54 amino acids and four disulfide bonds, is the smallest sweet-tasting protein yet isolated from the wild African plant *Pentadiplandra brazzeana*. Brazzein has various desirable properties for use as a low-calorie sweetener in the diets of individuals suffering from diabetes, obesity, and metabolic syndrome. For example, brazzein has a high water solubility and a high thermostability. In addition, brazzein is 2000-times sweeter than sucrose on a weight basis. Both the solution and crystal structures of brazzein have been reported. In the crystal structure [1], brazzein has a defensin-like fold containing two  $\alpha$ -helices and a three-stranded antiparallel  $\beta$ -sheet. Defensins are small cysteine-rich cationic proteins found in both animals and plants, which function by binding to the microbial cell membrane, and, once embedded, forming pore-like membrane defects that allow efflux of essential ions and nutrients. In fact, Yount and Yeaman reported that brazzein has antimicrobial activity against Gram positive (*Bacillus subtilis* and *Staphylococcus aureus*) and negative (*Escherichia coli*) bacteria and a fungus (*Candida albicans*) at pH 7.5 rather than pH 5.5 [2]. A search for proteins with a similar backbone fold to brazzein using the DALI server shows that structurally similar proteins to brazzein include plant defensins, scorpion neurotoxins (K<sup>+</sup> channel blockers), arthropod defensins, mollusc defensins, mold defensins, and a plant trypsin inhibitor. These proteins commonly have a  $\gamma$ -core sequence. Here we compare their sequences, structures and functions, which has led to a conclusion that the C-terminal half of brazzein is important for its antimicrobial activity, brazzein will not have a neurotoxin activity, and it will not act as a trypsin inhibitor.

[1] Nagata, K. et al. *Acta Crystallogr. D* 69, 642-647 (2013)., [2] Yount, N. Y. and Yeaman, M. R. *Proc. Natl. Acad. Sci. USA* 101, 7363-7368 (2004).

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