

5th World Congress on **Biotechnology**

June 25-27, 2014 Valencia Conference Centre, Valencia, Spain

Enzymatic glycosylation of epothilone A

Jae Kyung Sohng¹, Prakash Parajuli¹, Ramesh Prasad Pandey¹ and Yeo Joon Yoon²

¹SunMoon University, Republic of Korea

²EwhaWomans University, Republic of Korea

Many natural products and antibiotics have been modified by tailoring enzymes through the process of glycosylation which usually improves their biological activities. Epothilone having epoxide, thiazole, and ketone moieties are extremely cytotoxic agents with a similar mode of action to taxol. Nevertheless, Epothilones represent a novel structural class of compounds with equipotent kinetic similarity to taxol. These antifungal and cytotoxic epothilones are produced by a few strains of myxobacterium, *Sorangium cellulosum*. In *S. cellulosum*, 8 analogs of 16-membered macrolide epothilones (A to H) containing 29 variants have been described. Epothilone A and B are the major products having potential applications in therapy and cytotoxicity against different tumor cell lines. Thus, it could be interesting to elucidate and search for glycosylated analogues of epothilones. Here we report an enzymatic glycosylation study of epothilone A with the help of GT (YjiC) from *Bacillus licheniformis* DSM 13. UDP-D-glucose and TDP-2 deoxyD-glucose were used as a sugar donor whereas epothilone A as an acceptor substrate for YjiC. The reaction products were analyzed by HPLC and high resolution LC-QTOF-ESI/MS which revealed the presence of glycosylated products. Although 3-hydroxyl and 7-hydroxyl positions were prominent in epothilone A for glycosylation; reactions with UDP-glucose and TDP-2-deoxy glucose, mono glucosides were detected which was confirmed by mass analysis. Exact glycosylation position is yet to be elucidated.

sohng@sunmoon.ac.kr