## LTB<sub>4</sub> 12-hydroxydehydrogenase/15-oxo-PG 13-reductase and Indomethacin Complex

<u>Tetsuya Hori</u><sup>a</sup>, Takehiko Yokomizo<sup>b,c,d</sup>, Hideo Ago<sup>a</sup>, Takao Shimizu<sup>b,c</sup>, Masashi Miyano<sup>a</sup>, <sup>a</sup>Structural Biophysics Laboratory, RIKEN Harima Institute at SPring-8. <sup>b</sup>Department of Biochemistry and Molecular Biology, Faculty of Medicine, The University of Tokyo. <sup>c</sup>CREST of JST. <sup>d</sup>PRESTO of JST. E-mail: thori@spring8.or.jp

The bi-functional leukotriene B<sub>4</sub> 12-hydroxydehydrogenase/15oxo-prostaglandin 13-reductase (LTB<sub>4</sub> 12-HD/PGR) is essential for eicosanoid inactivation. It catalyzes the first irreversible reactions in each eicosanoid inactivation and the activity may be regulated by a protein with SH3 domain. LTB<sub>4</sub> is oxidized to 12-oxo-LTB<sub>4</sub>, and 15oxo-PGE<sub>2</sub> are reduced to 13,14-dihydro 15-oxo-PGE<sub>2</sub>. Some none steroidal anti-inflammatory drugs inhibit LTB<sub>4</sub> 12-HD/PGR activity. Here we report the structure of LTB<sub>4</sub> 12-HD/PGR with NADP<sup>+</sup> and indomethacin. The indomethacin binds to the 15-oxo-PGE<sub>2</sub> binding site, indicating that the indomethacin competitively inhibits LTB<sub>4</sub> 12-HD/PGR. The chloro-benzene moiety of indomethacin enters into the hydrophobic pore that is the recognition site of  $\omega$ -chain of 15-oxo-PGE<sub>2</sub>, and the carboxyl group of indomethacin interacts with Arg56 and Tyr262. The architecture is identical to those of indomethacin and cyclooxygenase complexes. The result may be useful for further development of cyclooxygenase inhibitor.

Keywords: anti-inflammatory compounds, dehydrogenases, protein-inhibitor binding