

especially the properties regarding its carbohydrate recognition domain structure. As a consequence, it is absolutely essential to understand the structure of TM, in order to get into more functional details of its regulation in the aforementioned properties. Thrombomodulin (TM) forms a 1:1 complex with thrombin. Whereas thrombin alone cleaves fibrinogen to make the fibrin clot, the thrombin-TM complex cleaves protein C to initiate the anticoagulant pathway. Until present, the so-far available structures, either through NMR or through X-ray analyses, can not shed lights into the decent structural-functional interpretations for TM regulations. Crystallographic investigations of the complex between thrombin and TM-EGF456 did not show any changes in the thrombin active site. Therefore, research has focused recently on how TM may provide a docking site for the protein C substrate with different  $\text{Ca}^{2+}$  concentration. Previous work, however, showed that when the thrombin active site was occupied by substrate analogues labeled with fluorophores, the fluorophores responded differently to active (TMEGF1-6) versus inactive (TMEGF56) fragments of TM.

**Keywords:** Thrombomodulin, Structural analysis, Calcium-induced dimerization.

#### FA1-MS05-P20

##### Crystal structure of 6- Methoxy- 4-

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Coumarins are a class of naturally occurring oxygen heterocycles which have been found to exhibit wide ranging biological activities [1-3] through its innumerable derivatives. Structural studies on coumarins have been focused on their solid state photochemical dimerization [4], hydrogen bonding [5], mode of packing [6], molecular self assembling [7] and photophysical properties [8]. Introduction of bromine has resulted in formation of hydrates, intermolecular hydrogen bonding, eclipsed conformation observed in 3-bromocoumarin [9], 6-bromo-3-acetylcoumarin [10] and 3-bromoacetylcoumarin [11] respectively. 3-Bromophenyl-6-acetoxymethyl-coumarin-3-carboxylates have been found to exhibit potential anticancer and antitumour activity [12].

Crystals suitable for diffraction studies were grown by slow evaporation technique using acetic acid as a solvent. The crystals of the compound crystallize in Monoclinic with space group  $P2_1/n$  having 8 molecules in the unit cell of dimensions crystal system:  $a = 4.3573(3)$ ,  $b = 9.2859(6)$ ,  $c = 25.2677(17)$  Å and  $\beta = 91.927(3)^\circ$ . The three dimensional intensity data was collected using a crystal of size  $0.25 \times 0.15 \times 0.10$  mm mounted on Bruker axis kappa apex2 ccd diffractometer with  $\text{MoK}_\alpha$  radiation. The data was collected using  $\omega$  and  $\phi$  scan mode. 9957 measured reflections of which 2130 independent reflections and 1502 reflections with  $I > 2\sigma(I)$ . With absorption correction: multi-scan.

The structure was solved using wingx software package and the model was refined by the full-matrix least-square method. The refinement was continued till the final  $R = 0.0449$ ,  $R_w = 0.1103$ .

The title compound is cyclic and planar but non-aromatic in nature due to the continuous delocalization of electrons around the coumarin ring. Skeleton is not possible. There is a significance deviation in bond angle at  $01\text{-C1-C2}$  ( $117.2(3)^\circ$ ) due to the electronic repulsion of oxygen (O2) atom which is present at C1 carbon atom.

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**Keywords:** x-ray, single crystal, coumarin.