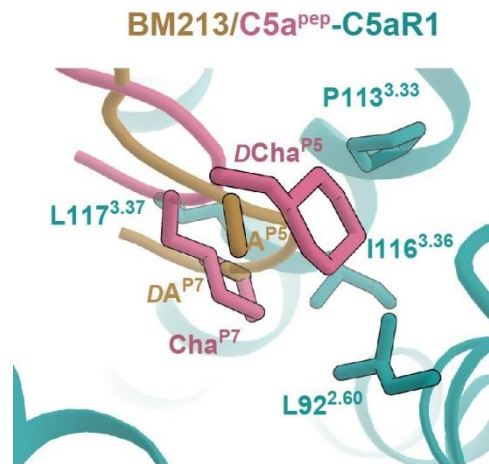


**Supplementary information, Fig. S16**



**Fig. S16. Structural comparison of BM213 and C5a<sup>pep</sup> in the orthosteric binding pocket of C5aR1.** *D*Cha<sup>P5</sup> and Cha<sup>P7</sup> of C5a<sup>pep</sup> forms more extensive hydrophobic interactions with C5aR1, when compared to BM213.