

### Effect of intracellular loop 1 pepducins on ligand-receptor binding kinetics at the CXCR4 receptor

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**Introduction.** Pepducins are a class of lipopeptide ligands derived from the intracellular loops of a GPCR and have been reported to act as allosteric modulators, although their exact mechanism remains unclear.<sup>1</sup> Pepducins for the chemokine receptor, CXCR4, have been described to act as positive or negative allosteric modulators and can exhibit agonist activity in the absence of the endogenous ligand CXCL12.<sup>2</sup> In this study, we investigated the interaction and mechanism of intracellular loop 1 pepducins with CXCR4.

**Method.** Experiments were performed in HEK293 cells stably expressing human CXCR4 tagged with the luciferase, NanoLuc, on its N-terminus (NL-CXCR4) or C-terminus (CXCR4-NL). The binding and displacement of fluorescently labelled CXCR4 ligands was studied in a NanoBRET assay<sup>3</sup> at equilibrium or as kinetic experiments reading once every minute using NL-CXCR4 or CXCR4-NL cells or membranes quantified on a PHERAstar FS plate reader (BMG).

**Results.** The affinity of fluorescent CXCL12 (CXCL12-red) was determined in NL-CXCR4 cells and membranes using NanoBRET saturation binding ( $pK_d(\text{Cells})=7.15\pm 0.04$ ,  $n=5$ ;  $pK_d(\text{Membranes})=7.61\pm 0.10$ ,  $n=5$ ) or alternatively the association binding kinetics in cells using 1.5625 to 100nM ( $pK_d=7.10\pm 0.18$ ,  $n=7$ ) were analysed. Competition binding experiments in membranes showed that CXCL12-red binding was inhibited by pre-incubation with small molecules AMD3100 ( $pK_i=7.03\pm 0.04$ ,  $n=5$ ), IT1t ( $pK_i=8.07\pm 0.02$ ,  $n=5$ ) and CXCR4 pepducin ATI-2341 ( $pK_i<6$ , maximal inhibition  $58.3\pm 2.2\%$  ( $10\mu\text{M}$ ),  $n=5$ ). Control pepducins missing a lipid tail or the last three amino acids did not displace CXCL12-red at concentrations up to  $10\mu\text{M}$ . Displacement of CXCL12-red by ATI-2341 was followed in a kinetic assay in cells. ATI-2341 ( $10\mu\text{M}$ ) shows displacement of CXCL12-red which reaches a plateau of  $53.47\pm 0.98\%$  of the initial signal within 10 minutes ( $k_{\text{off}}=0.43\pm 0.04\text{ min}^{-1}$ ,  $n=5$ ). ATI-2341f, a TAMRA-labelled ATI-2341, was used in the membrane binding assay to detect interaction with the Nluc-CXCR4 receptor. ATI-2341f showed no displaceable binding towards the N-terminal labelled NL-CXCR4, but did interact with the C-terminal CXCR4-NL. In CXCR4-NL cells, binding of ATI-2341f reached saturation after 15 min and displacement with high concentrations of ATI-2341 was possible within 10 minutes. AMD3100, IT1t and CXCL12 are not able to displace ATI-2341f binding at concentrations up to 100 or  $1\mu\text{M}$ , respectively.

**Conclusion.** These data suggest that ATI-2341 interacts with CXCR4 from an allosteric site on the inner leaflet of the cell membrane. Moreover, ATI-2341 decreases the binding of CXCL12 to CXCR4 suggesting an influence on the orthosteric binding site.

### References

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- 3.Stoddart et al.(2015). Nature Methods 12:661 -663.