

## Determination of Association ( $k_{on}$ ) and Dissociation ( $k_{off}$ ) Rates of inhibitors of the PI3K $\delta$ isoform using the LanthaScreen® technology

B. Oliosi<sup>1</sup>, C. Griffante<sup>1</sup>, F. Visentini<sup>1</sup>, S. Bertolini<sup>2</sup>, M. Biagetti<sup>2</sup>, S. Capacchi<sup>2</sup>, A. Capelli<sup>2</sup>, L. Carzaniga<sup>2</sup>, V. Mileo<sup>2</sup>, M. Corsi<sup>1</sup>. <sup>1</sup>Center for Drug Discovery and Development, Aptuit, Verona, Italy, <sup>2</sup>Chiesi Farmaceutici, Parma, Italy.

**Introduction:** There is an increasing evidence of potential advantages on duration of pharmacodynamics and target selectivity for kinase inhibitors with long residence time. Obtaining kinetic parameters at early stage of drug discovery may allow improved predictions of *in vivo* drug efficacy (1). PI3K kinase inhibitors have been studied in a variety of inflammatory processes and hematologic malignancies (2).

**Method:** In order to obtain  $k_{on}$  and  $k_{off}$  values of inhibitors at human PI3K $\delta$  kinase isoform, a simple non-radioactive method was setup and the binding kinetics of reference inhibitors CAL-101 (3), IPI-145 (4) and SW13 (5) were determined by using Motulsky and Mahan method (6). The LanthaScreen® (ThermoFisher) binding of Tracer-314 to PI3K $\delta$ -His enzyme coupled to Eu-anti-His antibody produced an increase of Fluorescence Resonance Energy Transfer (FRET), which was measured with the Envision plate-reader. PI3K $\delta$ -Eu kinase and tracer titration curves were measured after 1h incubation at 23°C. The tracer saturation curve gave an average  $K_D$  value of 8.1nM ( $pK_D=8.09\pm0.04$ , average $\pm$ StDev, n=3). Increasing tracer concentrations were associated to a fixed amount of PI3K $\delta$ -Eu.  $K_{on}$  was calculated as  $5.7E7M^{-1}min^{-1}$  and  $k_{off}$   $0.47min^{-1}$  and the  $K_D$  calculated as from  $k_{off}/k_{on}$  was 8.1nM. Tracer  $k_{off}$  average value from dissociation experiments was  $0.41min^{-1}$  (n=3), corresponding to a 2min dissociation half-life ( $t_{1/2off}$ ). Tracer  $k_{on}$  was also calculated from dissociation and saturation experiments, ( $k_{off}/K_D$ ,  $5.1E7M^{-1}$ ).

**Results:** The kinetics binding properties of the three PI3K $\delta$  reference inhibitors were analyzed. Tracer was associated to PI3K $\delta$ -Eu with increasing inhibitor concentrations. Data were analyzed using GraphPad v5 software. The  $t_{1/2off}$  for CAL-101, IPI-145 and SW13 was 2, 6 and 12min, respectively. The inhibitor affinities ( $K_D$ ) were calculated as  $k_{off}/k_{on}$  ratio and were compared with the potency values obtained in a functional ADP-Glo enzymatic assay (Promega). A good correspondence between inhibitor affinities and potencies was found

(Table 1).

Compound	PI3K $\delta$ binding						PI3K $\delta$ functional		
	$k_{on}$ $M^{-1}min^{-1}$	$k_{off}$ $min^{-1}$	$t_{1/2off}$ min	$pK_D$	$pK_D$ stdev	n	$pK_i$	$pK_i$ stdev	n
CAL-101	3.7E+08	0.46	2	8.87	0.14	3	8.84	0.04	316
IPI-145	2.3E+08	0.13	6	9.26	0.13	3	9.18	0.35	4
SW13	1.5E+08	0.06	12	9.37	0.22	3	9.15	0.08	3

**Conclusions:** The LanthaScreen® technology allowed the determination of kinetics properties of the three PI3K $\delta$  inhibitors. CAL-101 was the faster dissociating PI3K $\delta$  kinase inhibitor whereas SW13 the slowest. These results may have an implication on their pharmacodynamics.

### References:

1. Copeland R (2006). *Nature Rev Drug Discov.* **5**: 730-739.
2. Wang X et al. (2015). *Acta Pharmacol Sin.* **36** (10): 1170-1176.
3. Somoza JR et al. (2015). *J Biol Chem.* **290** (13): 8439-8446.
4. Winkler DG et al. (2013). *Chem Biol.* **20**: 1364-1374.
5. Williams O et al. (2010). *Chem Biol.* **17** (2): 123-144.
6. Motulsky HJ and Mahan LC (1984). *Mol Pharmacol.* **25**: 1-9.