

Structure-based tailoring of compound libraries for  
high-throughput screening: Discovery of novel  
EphB4 kinase inhibitors  
**Supplementary Material**

Peter Kolb<sup>1,†</sup>, Catherine Berset Kipouros<sup>2</sup>, Danzhi Huang<sup>1</sup>,  
Amedeo Caflich<sup>1,\*</sup>

<sup>1</sup>Department of Biochemistry  
University of Zurich  
Winterthurerstrasse 190  
CH-8057 Zurich, Switzerland

<sup>2</sup>Oncalis AG  
Biotech Center Zurich  
Wagistrasse 21  
CH-8952 Schlieren, Switzerland

\*Corresponding author  
Phone: (41 44) 635 55 21  
FAX: (41 44) 635 68 62  
email: [caflisch@bioc.uzh.ch](mailto:caflisch@bioc.uzh.ch)

Keywords: EphB4, tyrosine kinases, angiogenesis, cancer,  
docking, high-throughput screening

<sup>†</sup>Present address: Department of Pharmaceutical Chemistry  
University of California, San Francisco  
1700 4<sup>th</sup> Street  
San Francisco, CA 94158, USA

October 25, 2007

## Contents

Compound ranking 3

Enzymatic assays 4

References 5

**Compound ranking.** It is necessary to discard poses with unlikely binding modes through the application of filters before calculating the binding energies with LIECE.<sup>1</sup> The quotient of the van der Waals energy and the molecular weight (the “van der Waals efficiency”) is a very useful filter. Compounds **1-3** were identified in rankings using it as a filter. The three applied filter/ranking combinations were: (i) ranking according to the vdW-efficiency; (ii) vdW-efficiency  $\leq -0.116 \text{ kcal}\cdot\text{g}^{-1}$ , then ranking according to  $\Delta G^L$  (free energy of binding predicted by the LIECE model); (iii) vdW-efficiency  $\leq -0.1 \text{ kcal}\cdot\text{g}^{-1}$ , number of rotatable bonds  $\leq 5$ , and van der Waals interaction energy  $\leq -35 \text{ kcal}\cdot\text{mol}^{-1}$ , then ranking according to  $\Delta G^L$ . The cutoffs were obtained from the distributions of the respective values. Compounds **1-3** and the five weak EphB4 inhibitors with different anchors were among the best 300 poses in at least one of these rankings. Compounds **4-8** were not present in the original library of 728202 molecules.

**FRET-based enzymatic assay.** The EphB4 FRET enzymatic assay (Z'Lyte Tyr kinase assay, Invitrogen, USA) was performed according to the manufacturer's protocol. Briefly, EphB4 kinase, at a concentration of 3.5 ng per reaction well, was incubated for one hour with FRET-peptide substrate in the presence of ATP at a concentration of 30  $\mu$ M, which is near its  $K_m$ . Reactions were developed and stopped and fluorescence was measured at 440 nm and 520 nm emission with excitation at 400 nm. IC<sub>50</sub> values were calculated with GraphPad Prism 5.0 (GraphPad Software).

**FRET-based enzymatic assay at Cerep.** The compounds showing the strongest inhibition effects were verified in the FRET-based assay<sup>2</sup> offered by Cerep (France). Briefly, compounds were measured in duplicates at eight different concentrations ranging from 10 nM to 20  $\mu$ M. The concentrations of EphB4 and ATP in the assay were 0.2  $\mu$ g/ml and 0.75  $\mu$ M (which corresponds to its  $K_m$ ), respectively.

## References

- [1] Huang D, Caflisch A. Efficient evaluation of binding free energy. *J Med Chem* 2004; 47:5791–5797.
- [2] Mathis G. Probing molecular interactions with homogeneous techniques based on rare earth cryptates and fluorescence energy transfer. *Clin Chem* 2003; 41:1391–1397.