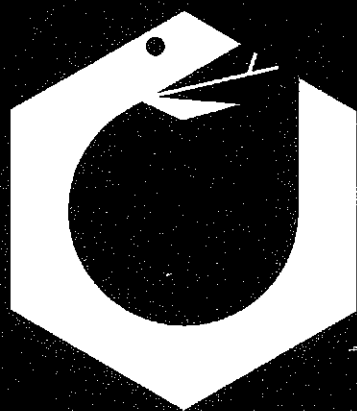
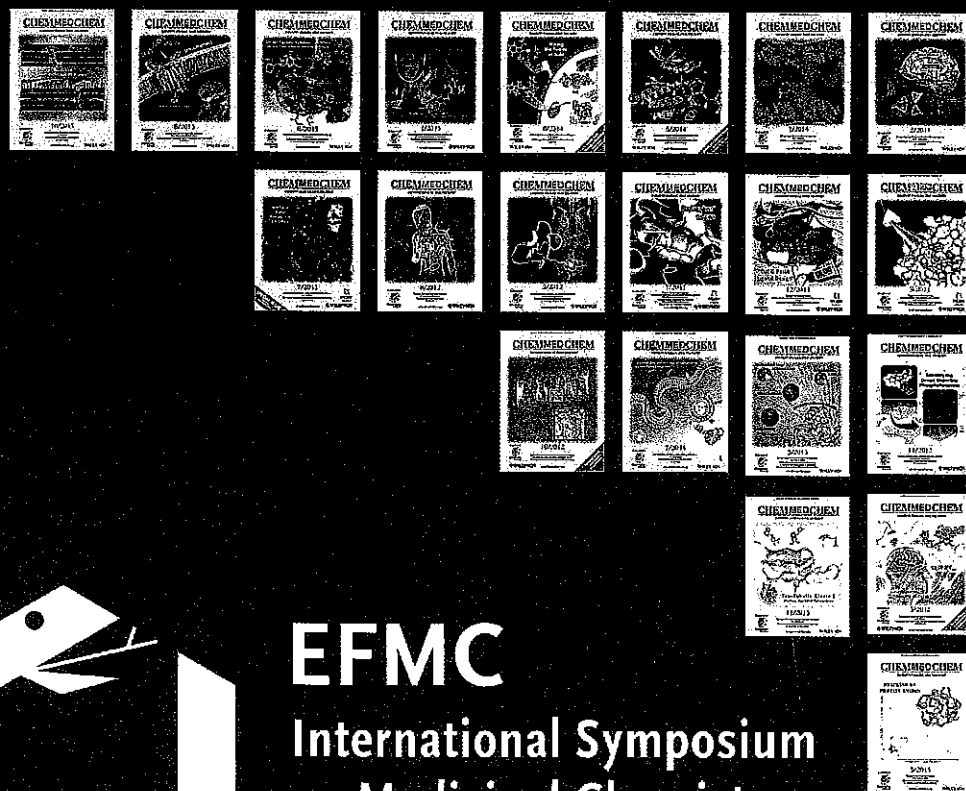


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ABSTRACTS**

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R026 | Synthesis, Molecular Docking and Biological Evaluation of New 1-Aryl-3-[3-(thieno[3,2-*b*]pyridin-7-ylthio)phenyl]ureas as Potent Type II VEGFR-2 Tyrosine Kinase Inhibitors

Vera A. Machado,^(1,2) Daniela Peixoto,⁽¹⁾ Raquel Costa,⁽²⁾ Ricardo C. Calhela,^(1,3) Rui M. V. Abreu,⁽³⁾ Isabel C.F.R. Ferreira,⁽³⁾ Raquel Soares,⁽²⁾ Maria-João R.P. Queiroz⁽¹⁾

1) Departamento/Centro de Química (U686-FCT), Escola de Ciências, Universidade do Minho, Campus de Gualtar 4710-057Braga, Portugal; E-mail: mjrpq@quimica.uminho.pt

2) Departamento de Bioquímica (U38-FCT), Faculdade de Medicina, Universidade do Porto, 4200-319 Porto, Portugal

3) CIMO (U690-FCT)-ESA, Instituto Politécnico de Bragança, Campus de Sta. Apolónia, Apartado 1172, 5301-855Bragança, Portugal

The vascular endothelial growth factor receptor 2 (VEGFR-2) is a tyrosine kinase receptor, expressed primarily in endothelial cells, and is activated by the specific binding of VEGF to the VEGFR-2 extracellular regulatory domain. Once activated, VEGFR-2 undergoes autophosphorylation, triggering signaling pathways leading to endothelial cell proliferation and subsequent angiogenesis.^[1] Small molecules may act as inhibitors by competing for the ATP-binding site of the VEGFR-2 intracellular tyrosine kinase domain, thereby preventing the intracellular signaling that leads to angiogenesis.^[2]

Here, we present the synthesis of new 1-aryl-3-[3-(thieno[3,2-*b*]pyridin-7-ylthio)phenyl]ureas **1a–c**, as potent type II VEGFR-2 inhibitors based on molecular docking (Figure A) and biological evaluation including enzymatic assays using the VEGFR-2 tyrosine kinase domain (IC₅₀=10–28 nM) and studies in human umbilical vein endothelial cells (HUVECs). The latter included cell viability (MTS), proliferation (BrdU) and Western blot for total and phosphorylated VEGFR-2 (Figure B).

The predicted docked poses were analyzed in detail and a plausible explanation for compounds **1** potency was obtained based on the simultaneous presence of a *S*-linker and the arylurea moiety in the *meta* position as a new substitution pattern for the type II VEGFR-2 inhibitors. These chemical features place the thieno[3,2-*b*]pyridine and the terminal aryl ring in close superimposition to a pyrrolo[3,2-*d*]pyrimidine derivative. The presence of hydrophobic substituents (F and Me) in the terminal aryl ring is also important. For these compounds a significant inhibition in HUVECs proliferation upon VEGF stimulation was observed at low concentrations (0.5–1.0 μM) without affecting cell viability. Western blot analysis demonstrated that compounds **1** significantly inhibited the VEGFR-2 phosphorylation at 1.0 μM, thus confirming their anti-angiogenic potential.

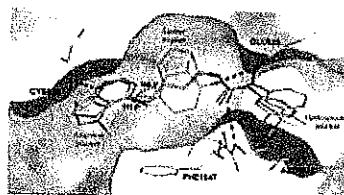
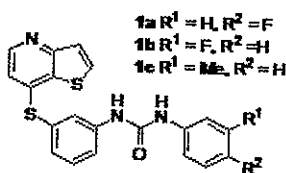


Fig. A. Docking pose superimposition at the VEGFR-2 kinase binding site for compounds **1a** and **1c** with a known type II inhibitor (pyrrolo[3,2-*d*]pyrimidine derivative)

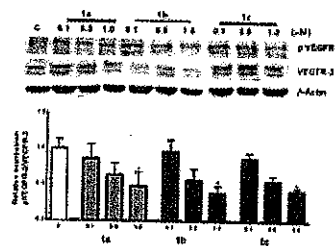


Fig. B. Western blot for total and phosphorylated VEGFR-2. **p* < 0.05 vs control (DMSO).

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References:

- [1] L.M. Strawn et al. *Cancer Res.* **1996**, *56*, 3540–3545.
 [2] S. Baka, A.R. Clamp, G.C. Jayson, *Expert Opin. Ther. Targets* **2006**, *10*, 867–876.

R027 | Novel Pyrimido-Oxazepinones as Potent and Selective mTOR Inhibitors

Gilbert Marciniak,⁽¹⁾ Alain Braun,⁽²⁾ Yann Foricher,⁽³⁾ Nicolas Muzet,⁽¹⁾ Eric Nicolaï,⁽²⁾ Cécile Pascal,⁽²⁾ Sukhvinder Sidhu,⁽³⁾ Bertrand Vivet,⁽¹⁾ Fabrice Viviani,^(2,4) Axel Ganzhorn⁽¹⁾

1) Sanofi R&D, DPU Early to Candidate, 16 rue d'Ankara, 67080 Strasbourg, France

2) Sanofi R&D, DPU Early to Candidate, 1 avenue Pierre Brosolette, 91385 Chilly-Mazarin, France

3) Sanofi R&D, Oncology Division, 13 quai Jules Guesde, 94403 Vitry-Sur-Seine, France

4) Present address: Laboratoire Glaxo Smith Kline, 25 avenue du Québec, 91951 Les Ulis, France

Mammalian target of rapamycin (mTOR), a 289 kDa serine/threonine kinase of the phosphoinositide 3-kinase-like kinase family, is a central regulator of cell growth and proliferation. Mutations and dysregulation of the PI3K/mTOR pathway (amplification of RTKs, loss of