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INTRODUCTION

PI3K (Phosphoinositide 3-kinase) is the family of lipid kinases that participates in many key cellular processes like proliferation, growing up, migration, cytokines production, and apoptosis. These proteins are highly expressed in many human cells, such as leukocytes, which have implications for the body's immune response. They are involved in metabolism regulation, normal embryogenesis, or maintaining glucose homeostasis. Inhibition of PI3K (especially the first class which contains four subunits: α , β , γ , δ) is regarded to be very attractive as a potential mechanism for the treatment of many diseases like SLE (Systemic Lupus Erythematosus), MS (Multiple Sclerosis), IBD (Inflammatory Bowel Disease) and other inflammatory and autoimmune diseases [1-4]. In this work, we present a series of new, very potent, active, and selective small molecule inhibitors based on the structure of appropriately substituted 5-(2-difluoromethylbenzimidazo-1-yl)pyrazolo[1,5-*a*]pyrimidine [5] as highly potent drug candidates.

SYNTHESIS

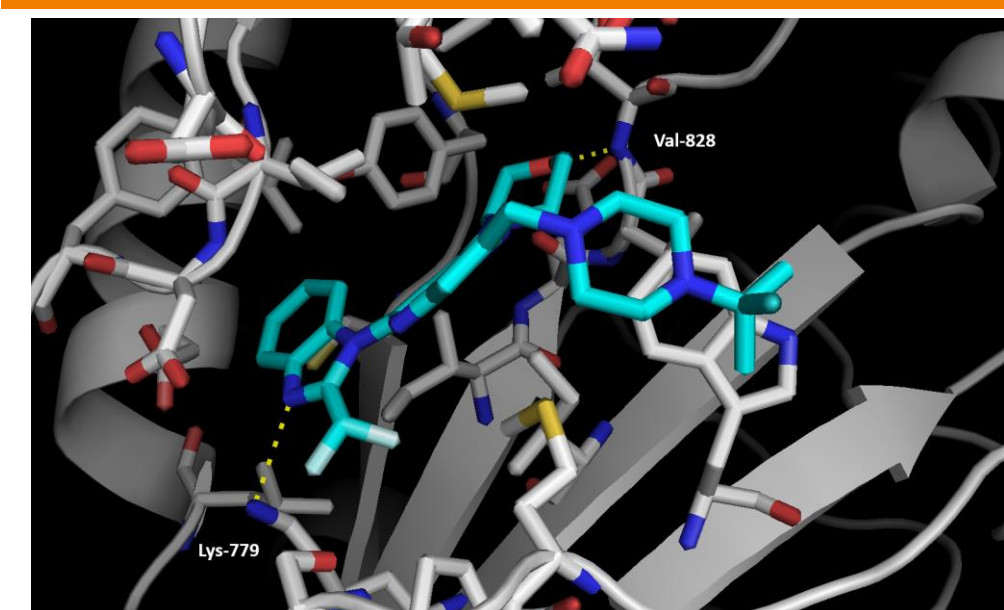


Figure 1. Example of 3D modeling binding mode of CPL302415 – binding with amino acids.

In our research, it was reported that modifications of benzimidazoles groups and amine subunits play a crucial role in the activity and selectivity of PI3K δ inhibitors. We focused on the pyrazolo[1,5-*a*]pyrimidine core. It was investigated that there are two very important interactions. The first one is the binding between the oxygen atom of the morpholine ring and Val-828. The second bond is formed between the nitrogen atom at the third position of benzimidazole derivative (especially 2-(difluoromethyl)-1*H*-benzimidazole) and Lys-779. (Fig.1).

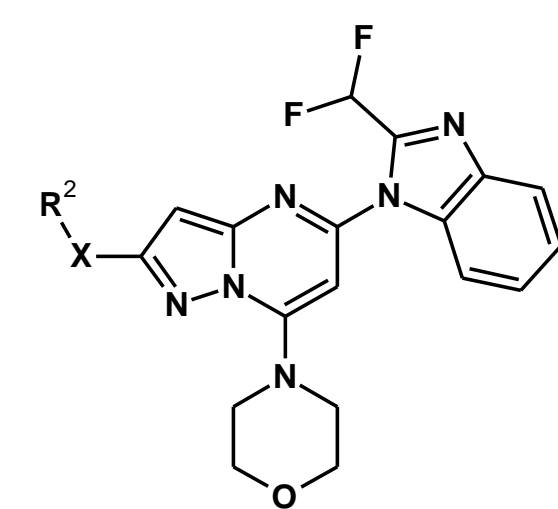
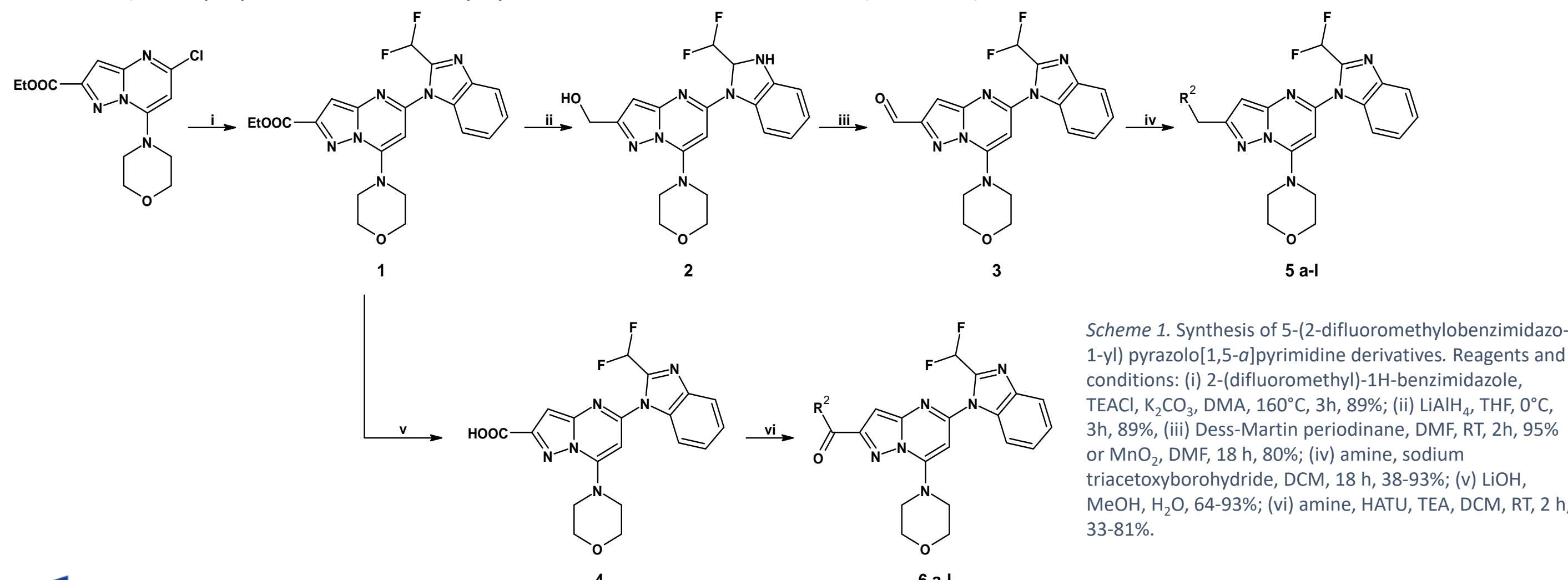


Figure 2. General structure of pyrazolo[1,5-*a*]pyrimidine derivatives.

A new family of substituted pyrazolo[1,5-*a*]pyrimidines (Fig.2) was prepared in multi-step synthesis included the hydrolysis reaction, oxygenation reaction, or reductive amination. Compounds (5a – 5l) were obtained after four-step synthesis from 5-chloro-7-(morpholin-4-yl)pyrazolo[1,5-*a*]pyrimidine-2-carboxylate. Compounds (6a – 6l) (replacement of the (-CH₂) group with a (-CO) group in position no.2 of pyrazolo[1,5-*a*]pyrimidine derivatives) were prepared after three-step synthesis from the same substrate (Scheme 1).



Scheme 1. Synthesis of 5-(2-difluoromethylbenzimidazo-1-yl)pyrazolo[1,5-*a*]pyrimidine derivatives. Reagents and conditions: (i) 2-(difluoromethyl)-1*H*-benzimidazole, TEACl, K₂CO₃, DMA, 160°C, 3h, 89%; (ii) LiAlH₄, THF, 0°C, 3h, 89%; (iii) Dess-Martin periodinane, DMF, RT, 2h, 95% or MnO₂, DMF, 18 h, 80%; (iv) amine, sodium triacetoxyborohydride, DCM, 18 h, 38-93%; (v) LiOH, MeOH, H₂O, 64-93%; (vi) amine, HATU, TEA, DCM, RT, 2 h, 33-81%.

RESULTS

Compound	R ²	X	IC ₅₀ PI3K δ [nM]	IC ₅₀ PI3K α [nM]	IC ₅₀ PI3K γ [nM]	α/δ	γ/δ
5 a		CH ₂	18	1 428	16 904	79	939
6 a		CO	84	12 481	48 777	148	580
5 b		CH ₂	52	1 729	6 347	33	122
6 b		CO	101	4 863	2 483	48	25
5 c		CH ₂	31	624	2 197	20	71
6 c		CO	74	n/a	3 593	n/a	52
5 d		CH ₂	24	73	156	3.0	6.5
6 d		CO	63	n/a	3 831	0.0	61
5 e		CH ₂	135	498	1 380	3.7	10
6 e		CO	226	2 385	8 536	11	38
5 f		CH ₂	443	721	1 709	1.6	3.6
6 f		CO	801	4 561	35 750	5.7	45
5 g		CH ₂	240	1 259	1 855	5.2	7.7
6 g		CO	314	3 127	2 793	10	8.9
5 h		CH ₂	615	1 852	7 553	3.0	12
6 h		CO	213	5 408	1 818	25	8.5
5 i		CH ₂	593	1 988	8 318	3.4	14
6 i		CO	90	2 354	1 782	26	20
5 j		CH ₂	561	1 570	4 317	2.8	7.7
6 j		CO	195	797	2 081	4.1	10.7
5 k		CH ₂	445	n/a	1 503	0.0	3.4
6 k		CO	141	1 476	1 689	10	12
5 l		CH ₂	38	n/a	13 330	0.0	350
6 l		CO	768	n/a	11 397	n/a	15

The library of active compounds was obtained. It was observed that the most promising are piperazine- and piperidine- derivatives (position R₂) (Fig.2). Compounds with satisfactory parameters were selected from the entire library. Nine final structures with an IC₅₀ value of less than 100 nM were obtained (Tab.1). The most promising compound (CPL302415) is 1-{2-[(4-*tert*-butylpiperazin-1-yl)methyl]-7-(morpholin-4-yl)pyrazolo[1,5-*a*]pyrimidin-5-yl}-2-(difluoromethyl)-1*H*-benzimidazole (Fig.3) with potent activity (IC₅₀ = 18 nM), selectivity (PI3K α /PI3K δ = 79; PI3K γ /PI3K δ = 939) and promising others parameters (Tab.2.)

Solubility*	>500 μ M	PAMPA	13.3 x 10 ⁻⁶ cm/s
MiceLM t _{1/2}	378 min	PPB Human	79 %
MiceLM CL	3.7 mL x min ⁻¹ x mg ⁻¹	PPB Monkey	81 %
HumanLM t _{1/2}	145 min	PPB Mice	83 %
HumanLM CL	9.6 mL x min ⁻¹ x mg ⁻¹	PPB Rat	82 %

Table 2. CPL302415 parameters, * KINETIC solubility pH 7.4.

Table 1. Obtained, selected PI3K inhibitors – pyrazolo[1,5-*a*]pyrimidine derivatives; n/a – data not available.

CONCLUSIONS

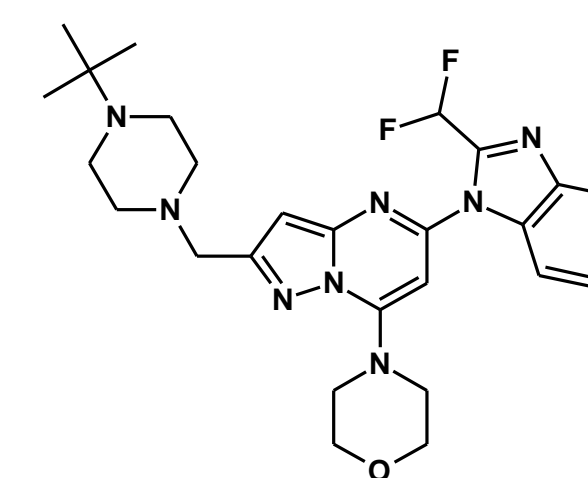


Figure 3. Structure of CPL302415.

In our work, we synthesized a library of very active benzimidazole derivatives (PI3K δ IC₅₀ = 0.02 – 0.80 μ M). Many substituted pyrazolo[1,5-*a*]pyrimidines were prepared with different amine groups in multi-step synthesis. Then we developed the process of lead compound(s) synthesis to select the preclinical candidate for SLE. Molecular modeling studies explain the interaction of active structures in the PI3K δ ATP binding site. Pyrazolo[1,5-*a*]pyrimidine with morpholine at position 7, 2-(difluoromethyl)benzimidazole at position 5 and *N*-*tert*-butylamine as amine proved to be the most potent structure (IC₅₀ = 18 nm). Moreover, this structure shows good selectivity to other PI3K isoforms (PI3K α /PI3K δ = 79; PI3K γ /PI3K δ = 939) and promising physical properties. That lead compound (CPL302415) (Fig.3) is subjected to further biological, toxicological, and pharmacokinetic studies.

REFERENCES

- [1] – John G. Foster, Matthew D. Blunt, Edward Carter, Stephen G. Ward; "Inhibition of PI3K Signaling Spurs New Therapeutic Opportunities in Inflammatory/Autoimmune Diseases and Hematological Malignancies"; Pharmacol Rev.; 2012 Oct.; 64(4):1027-54; doi: 10.1124/pr.110.004051.
- [2] – Edward Banham-Hall, Menna R. Clatworthy nad Klaus Okkenhaug; "The Therapeutic Potential for PI3K Inhibitors in Autoimmune Rheumatic Diseases"; Open Rheumatol J.; 2012; 6:245-58.; doi: 10.2174/1874312901206010245.
- [3] – Anne-Katrien Stark, Srividya Sriskantharajah, Edith M. Hessel, Klaus Okkenhaug; "PI3K inhibitors in inflammation, autoimmunity and cancer"; Curr Opin Pharmacol.; 2015 Aug; 23:82-91.; doi: 10.1016/j.coph.2015.05.017.
- [4] – Peter K. Vogt, Jonathan R. Hart, Marco Gymnopoulos, Hao Jiang, Sohye Kang, Andreas G. Bader, Li Zhao, Adam Denley; "Phosphatidylinositol 3-kinase (PI3K): The Oncoprotein"; Curr. Top Microbiol. Immunol.; 2010; 347:79-104.; doi: 10.1007/82_2010_80.
- [5] – EP3277687B1; "7-(Morpholin-4-yl)pyrazolo[1,5-*a*]pyrimidine derivatives which are useful for the treatment of immune or inflammatory diseases or cancer".