



Research paper

Ynamide Click chemistry in development of triazole VEGFR2 TK modulators



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ABSTRACT

Structure novelty, chemical stability and synthetic feasibility attracted us to design 1,2,3-triazole compounds as potential inhibitors of VEGFR2 tyrosine kinase. Novel triazoles **T1–T7** were proposed by oxazole (**AAZ** from PDB: 1Y6A)/1,2,3-triazole isosteric replacement, molecular modelling and docking. In order to enable synthesis of **T1–T7** we developed a methodology for preparation of ynamide **22**. Compound **22** was used for all Click chemistry reactions leading to triazoles **T1–T3** and **T6–T7**. Among the obtained products, **T1**, **T3** and **T7** specifically bind VEGFR2 TK and modulate its activity by concentration dependent manner. Moreover predicted binding poses of **T1–T7** in VEGFR2 TK were similar to the one known for the oxazole inhibitor **AAZ** (PDB: 1Y6A). Unfortunately the VEGFR2 inhibition by triazoles e.g. **T3** and **T7** is lower than that determined for their oxazole bioisosters **T3-ox** and **AAZ**, resp. Different electronic properties of 1,2,3-triazole/oxazole heterocyclic rings were proposed to be the main reason for the diminished affinity of **T1–T3**, **T6** and **T7** to an oxazole **AAZ** inhibitor binding site in VEGFR2 TK (PDB: 1Y6A or 1Y6B). Moreover **T1–T3** and **T6** were screened on cytotoxic activity against two human hepatocellular carcinoma cell lines. Selective cytotoxic activity of **T2** against aggressive Mahlavu cells has been discovered indicating possible affinity of **T2** to Mahlavu constitutionally active PI3K/Akt pathway.

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1. Introduction

Anti-angiogenesis agents that target malignant vasculature are of considerable interest due to their perceived potential to target tumour resistance towards chemo- and radiotherapy [1,2]. Vascular endothelial growth factors (VEGFs) and their corresponding family of receptor tyrosine kinases (VEGFRs) are the key proteins

modulating angiogenesis, the formation of new vasculature from an existing blood network. These include VEGFR1 (Flt1), VEGFR2 (Kinase Insert Domain Receptor (KDR) or Flk1). The last one, VEGFR3 is specialized for lymphangiogenesis [3]. VEGFR2 is the major positive signal transducer for endothelial cells proliferation and differentiation [4]. There has been considerable evidence, including clinical observations, that the abnormal angiogenesis is implicated in a number of diseases including rheumatoid arthritis, inflammation, degenerative eye conditions and cancer [5,6].

Cancer stem cells (CSCs) represent a small but the most tumourigenic subpopulation from the tumour cells responsible for metastasis, tumour recurrence and drug resistance. CSCs, also called “the roots of cancer”, are considered to be a new promising therapeutic target [7]. VEGFR2 is regarded as an endothelial cell protein but evidences suggest that VEGFRs may be expressed also by cancer cells. Glioblastoma multiforme (GBM) is characterized by

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florid vascularisation and aberrantly elevated VEGF. Antiangiogenic therapy with bevacizumab reduces GBM tumour growth. VEGFR2 is expressed on CD133 + human glioma cancer stem cells (GCSCs). VEGF-VEGFR2-Neuropilin1 (NRP1) axis influences GCSCs viability, self-renewal, and tumourigenicity. GCSCs viability was attenuated by direct inhibition of VEGFR2 TK activity and/or knockdown of VEGFR2 or NRP1. VEGFR2 inhibitors may block the VEGF-VEGFR2-NRP1 pathway [8,9]. VEGF via VEGFR2 stimulates proliferation of glioblastoma multiform CSCs. VEGF stimulates GCSCs tumourigenesis and angiogenesis. Suppression of VEGFR2 signalling is therefore a potential therapeutic strategy in GBM [10]. VEGFR2 plays a key role in ability of glioblastoma CSCs to vasculogenic mimicry (VM) formation, neovascularisation and tumour initiation. Knockdown of VEGFR2 in GCSCs markedly reduced their self-renewal, forming tubules, initiating xenograft tumours, promoting vascularisation and the establishment of VM. VEGFR2 is an essential molecule to sustain the “stemness” of GCSCs, their capacity to initiate tumour vasculature, and direct initiation of tumours [11]. Therefore VEGFR2 inhibitors are important compounds reducing angiogenesis and promising compounds to interfere with CSCs resistance.

The PDB database contains VEGFR2 TK complex 1Y6A possessing N-aryl-5-aryloxazol-2-amine ligand **AAZ** (Fig. 1) determined as a powerful VEGFR2 inhibitor (IC_{50} : 22 nM). Ligand **AAZ** was prepared in five steps in a low overall 10% yield mostly due to the problematic oxazole-2-amine fragment formation [12]. Low yields of oxazole-2-amine formation (1–58 %) have been also described [13,14]. Moreover **AAZ** contains N-aryloxazol-2-amine part that uses to liberate from connection with oxazole by influence of nucleophilic reagent (e.g. amines, alkoxides etc.). Nucleophilic attack on C(2) of **AAZ** oxazole ring resulting in toxic aniline **26** (Scheme 1). (unpublished results) Low yielding synthesis of N-aryloxazol-2-amines, their problematic stability and potential toxicity resulting from releasing aniline inspired us to develop novel, stable and synthetically more feasible VEGFR2 modulators based on the oxazole/1,2,3-triazole isosteric replacement. (Fig. 1) Replacing the heterocyclic ring in the structure of some inhibitors can provide a novel compounds with improved properties [15].

1,4-Disubstituted 1,2,3-triazoles are stable compounds easily obtainable in high yields from organic alkynes and azides by Cu(I) catalyzed reaction (CuACC Click chemistry) [16]. Click reaction allows rapid preparation of different triazoles that is especially advantageous if one of the reactants is the same in all performed reactions (e.g. ynamide **22a** in our case, Schemes 8 and 11). Additionally, Click reaction can selectively provide also 1,5-regioisomeric compounds via Ru (II) catalyzed cycloaddition [17] (Fig. 1).

Only few active 1,2,3-triazole containing VEGFR2 TK inhibitors are described in the literature: 6 inhibitors **1–6** possessing

$IC_{50} < 50$ nM, (Fig. 2) 10 substances with $IC_{50} < 100$ nM and 19 compounds having $IC_{50} \leq 200$ nM [18].

Imidazopyridazine **1** is the most active 1,2,3-triazole containing VEGFR2 inhibitor (IC_{50} : 1.2 nM) [19]. Other active inhibitors are e.g. benzoxazolamine **2** (IC_{50} : 8 nM) [20], quinazolinamines **3** and **4** (IC_{50} : 10 and 30 nM, resp.) [21,22], indazolurea **5** (IC_{50} : 45 nM) [23] and ureidopyridine-N-oxide **6** (IC_{50} : 49 nM) [24]. All of these VEGFR2 inhibitors (Fig. 2) possess a terminal 1,2,3-triazole fragment. In these cases, triazole group need not contribute to the inhibitor target affinity. Triazole group can be used to improve the ligand pharmacokinetic properties. E.g. the most potent inhibitor **1** (IC_{50} : 1.2 nM) was designed by pyrazole/triazole isosteric replacement in the structure of ligand from PDB complex 3VO3. In case of PDB: 3VO3 a pyrazole ring does not bind the VEGFR2 TK directly. It is exposed towards the solvent accessible part of the protein [19]. Therefore triazole core that replaces a pyrazole ring can have the same position. Compound **7** is a close analogue of sunitinib that is an active base of the drug Sutent (Pfizer Inc.). Sunitinib inhibitory activity $IC_{50} = 39$ nM (VEGFR2 TK) was determined [25]. The structure **7** possesses N-1,2,3-triazolyethyl group instead of N,N-diethylaminoethyl group present in sunitinib (Fig. 3). Recently an X-ray structure of VEGFR2 TK/sunitinib complex (PDB: 4AGD) appeared [26]. From its analysis is clear that the N,N-diethylamino group in sunitinib is exposed out of the VEGFR2 protein and represents only a group improving the ligand pharmacokinetic profile.

Although X-ray structure of complex **7** with VEGFR2 kinase is not known, an analogous function can be expected also for the triazole group in **7** (Fig. 3).

Only few VEGFR2 TK inhibitors possessing internal 1,2,3-triazole core were found in the literature. Compounds **8–10** were described as VEGFR2 activity modulators ($K_i > 915$ nM (VEGFR2)) [27]. The staurosporin-like inhibitor **11** moderately influences the VEGFR2 kinase activity (IC_{50} : 200 nM) [28] (Fig. 4).

Kiselyov et al. described the most active VEGFR2 inhibitors **12–14** (IC_{50} : 51–87 nM) possessing an internal 1,2,3-triazole core [29] (Fig. 5).

Our docking results proposed that the triazole fragment in **12–14** directly contributes the binding with VEGFR2 TK. The predicted intermolecular interactions of **12** and **14** are depicted on Fig. 6. Moreover proposed positions of the above triazole inhibitors in VEGFR2 ATP active site is similar to the poses of ligands from PDB complexes 2P2I (IC_{50} : 38 nM), and 3EFL (IC_{50} : 3 nM). (not shown) Both VEGFR2 kinase conformers (PDB: 2P2I and 3EFL) used in docking experiments are VEGFR2 inactive (DFG-out) kinases originally accommodating Type II inhibitor [14].

Considering the data mentioned above, we decided to prepare VEGFR2 TK modulators **T1–T7** possessing an internal 1,2,3-triazole core and determine their VEGFR2 TK inhibition potential.

2. Results and discussion

2.1. Interaction analysis of AAZ conformers

Oxazole VEGFR2 TK inhibitor **AAZ** was developed by GlaxoSmithKline [12]. The intermolecular interactions for both **AAZ** conformers present in VEGFR2 TK complex PDB: 1Y6A are depicted on Fig. 7.

Based on the above analysis, we decided to keep the pharmacophorically interesting 5-(ethylsulfonyl)-2-methoxyphenylamine fragment from **AAZ** ligand in all predicted triazole structures **T1–T7**. (Fig. 10) The proposed intermolecular interactions of triazoles **T1–T7** and their poses in VEGFR2 kinase were similar to those known for **AAZ** ligand from PDB complex 1Y6A. (e.g. **T3** from Fig. 12).

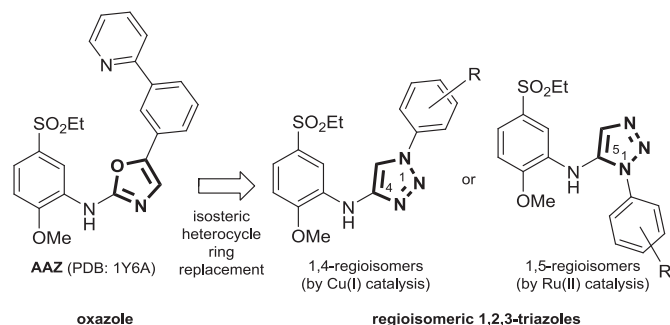
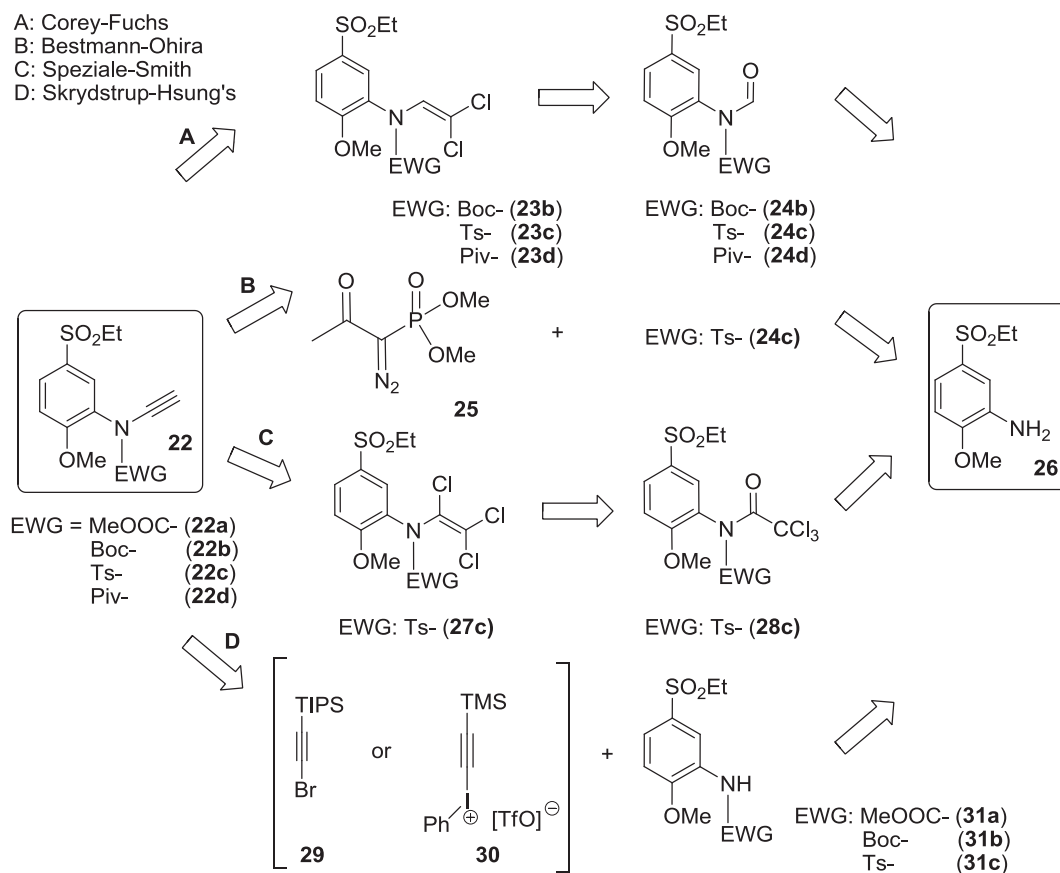


Fig. 1. The isosteric oxazole/1,2,3-triazole replacement. Click reaction can produce selectively one of the two triazole regioisomers based on different catalytic conditions.



Scheme 1. Proposed retrosynthetic approach to ynamides 22a–d.

2.2. In Silico predictions

An interaction analyse, molecular modelling and docking were used to identify the skeletons of 1,2,3-triazole derivatives **T1–T7** (Fig. 10). A reduced success of *in Silico* predictions is often associated with the ligand based target induced fit [30–32]. Therefore, for our docking experiments we selected the kinase variants of VEGFR2 from PDB complexes 1Y6A and 1Y6B possessing structurally the most relative oxazole ligands (**AAZ** and **AAX**, resp.). These ligands

represent Type I, ATP competitive VEGFR2 inhibitors that bind to an exceptional inactive VEGFR2 tyrosine kinase possessing an opened activation loop as was discovered by us recently [14]. Because the structure similarity between **AAZ** (**AAX**) and proposed triazoles (e.g. **T7**, **T6**, **T3** and **T5**) we did not expect strong influence of triazole ligand based induced fit. On the other hand, it was not easy to find triazole structures possessing **AAZ**-like score and pose by docking. The initial *in Silico* experiments were performed by an older DOCK software version with a kinase taken from PDB complex 1Y6B [33].

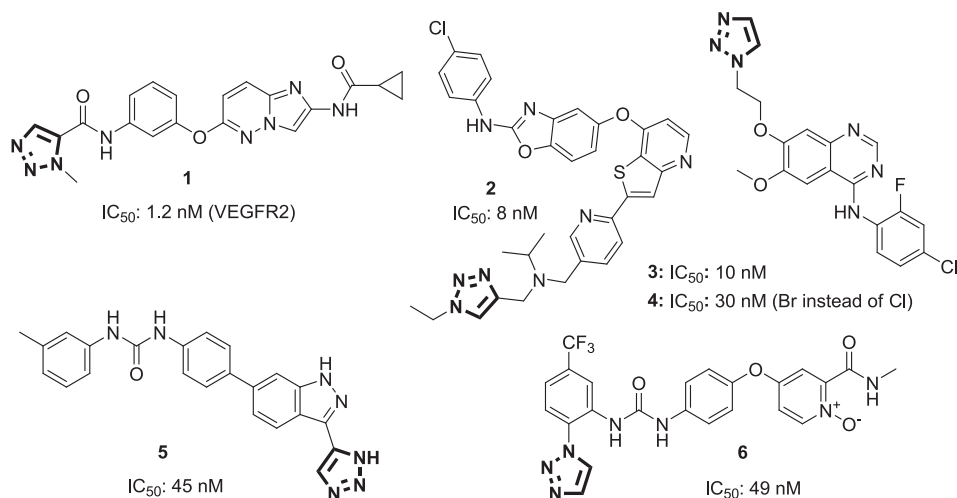


Fig. 2. The structures of VEGFR2 TK inhibitors 1–6.



Fig. 3. On the Left: The structure of sunitinib and its VEGFR2 TK inhibitory activity. In the middle: The PDB: 4AGD complex of VEGFR2 TK with sunitinib ligand that projects $Et_2NCH_2CH_2-$ group out of the protein into the solvent accessible area. (juxtamembrane part Leu802–Pro812, part of VEGFR2 activation loop: Asp1046–Asp1056 and water molecules were omitted for clarity). On the right: The structure of 1,2,3-triazole containing inhibitor 7 (mimicking sunitinib).

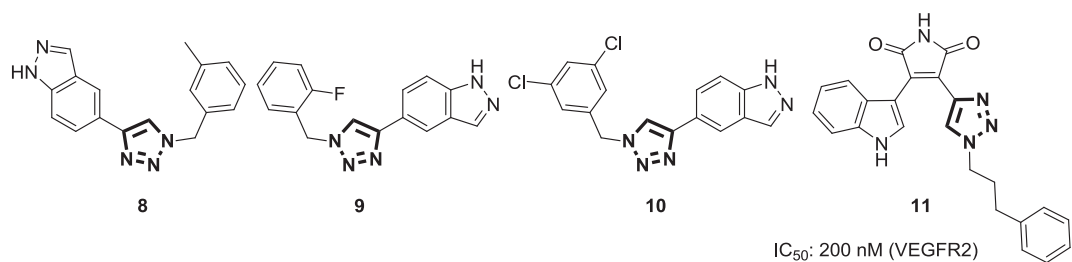


Fig. 4. VEGFR2 modulators 8–11 possessing an internal 1,2,3-triazole core.

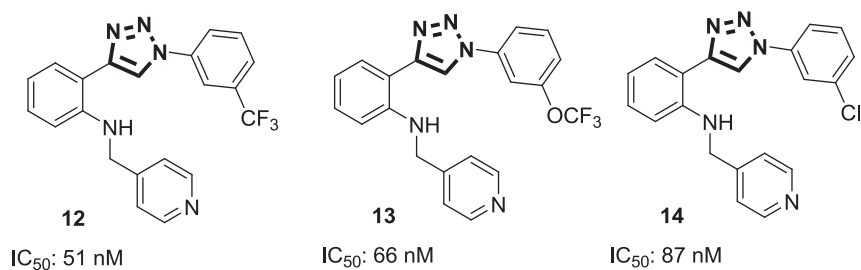


Fig. 5. Potent VEGFR2 inhibitors 12–14 with an internal 1,2,3-triazole ring.

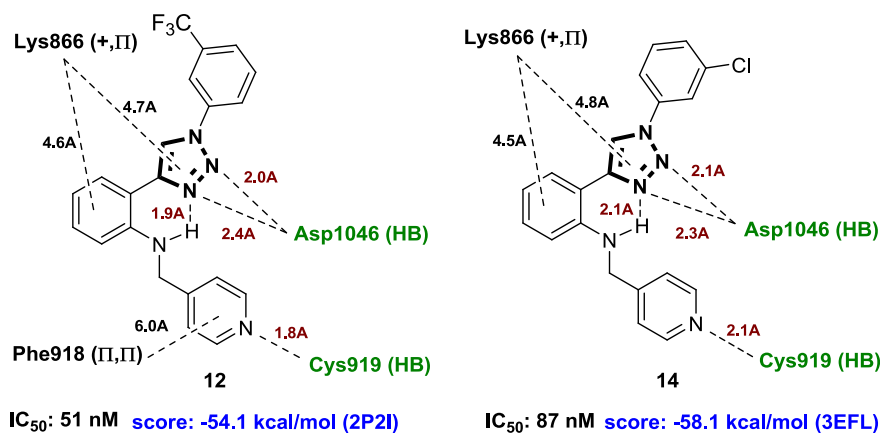


Fig. 6. Structures, VEGFR2 TK IC_{50} activities, predicted intermolecular interactions and docking scores of 12 and 14 obtained in VEGFR2 TK variants from PDB: 2P2I and 3EFL, resp. In both cases a triazole fragment in 12 and 14 directly contributes to the ligand/receptor binding (+ Π : induced dipole, HB: hydrogen bond, Π : stacked interaction). However the position of the triazole ring in 12 and 14 is conformationally blocked by an intermolecular HB with neighbouring NH group.

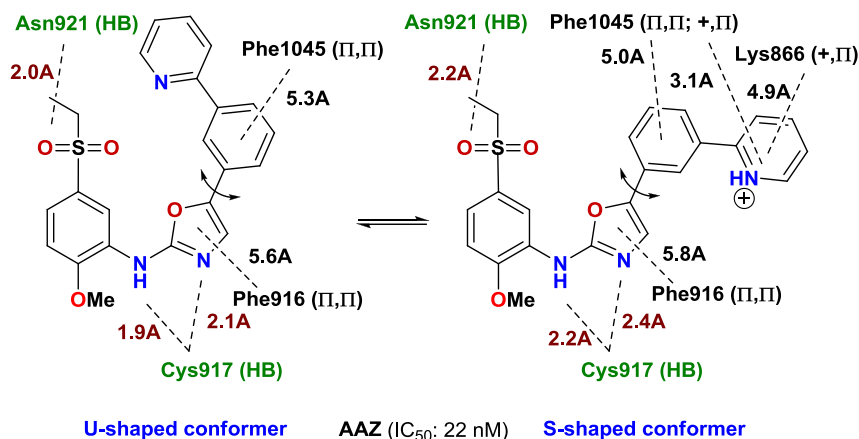


Fig. 7. The intermolecular interactions for both AAZ conformers present in VEGFR2 TK complex PDB: 1Y6A.

Some 1,4-disubstituted triazole skeletons that fulfilled both required score (AAZ: −53.6 kcal/mol) and pose conditions were selected: e.g. **T3** (score −57.2), **T5** (−52.9), **T6** (−51.3), **T2** (−51.3), **T4** (−49.2). The triazole core of **T1–T7** retains predicted hydrogen bond interaction with the kinase backbone amino acid residue Cys917 an important interaction known from AAZ oxazole ring in PDB: 1Y6A (Fig. 7). Pharmacokinetic properties of triazoles **T1–T7** predicted by Molinspiration toolkit supported their drug-likeness [34–38] (see supporting material). Proposed properties for 1,4-regioisomeric triazoles **T1–T7** were promising. Therefore we decided to perform their synthesis. 1,5-Regioisomeric triazole analogues (Fig. 1) of **T1–T7** were also docked by the same conditions. All these compounds were less advantageous possessing lower docking scores and several of them did not retain expected AAZ-like pose in VEGFR2 TK. (not shown) Therefore 1,5-regioisomers were omitted.

For the synthesis of all 1,4-disubstituted 1,2,3-triazoles **T1–T7** the pharmacophoric ynamide **22** (in modifications **22a–22d**) was required. (Schemes 8, 9 and 11) Its synthesis was developed starting from aniline derivative **26**. (Scheme 1).

2.3. Synthesis

In order to obtain compounds **T1–T7** (Fig. 10), we started synthesis their precursors azides **15–21** and ynamides **22a–22d** (Fig. 8).

2.3.1. Preparation of ynamides **22a,b**

Ynamides are stabilized equivalents of the corresponding reactive ynamines [39,40]. Diminishing the electron-donating ability of ynamine nitrogen by its protection with an electron-withdrawing group (EWG) leads to more stable ynamides. Since the first synthesis of ynamides reported by Viehe in 1972 [41], different methods have been published and the possible pathways towards required ynamides **22a,d** are summarized in Scheme 1. Starting 5-(ethylsulfonyl)-2-methoxyaniline (**26**) was prepared from commercially available 4-methoxybenzenesulfonyl chloride according to a recently described procedure in four steps and 59% overall yield [42].

Corey–Fuchs alkyne alkylation proved to be ineffective in our hands. (Scheme 1, Path A) In fact, application of the Brückner procedure [43] on protected formamides **24**, which have to be protected with an EWG protecting group prior to their formulation, was successful only in the case of the corresponding dichlorovinyl intermediate **23c** (tosyl protecting group) in a very modest yield (9%), and therefore this path was abandoned. The Bestmann–Ohira

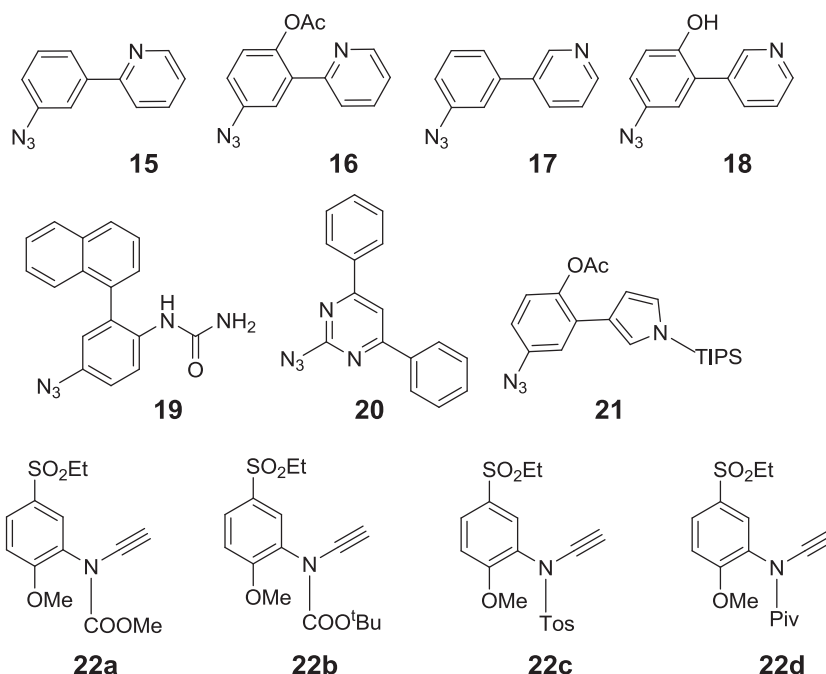
alkynylation (Scheme 1, Path B) was ineffective with **24c** and only starting material was recovered [44]. A transformation of *N*-trichloroacetate **28c** to the corresponding ynamide **22c** according to the methodology of Speziale and Smith has been tested [45]. (Scheme 1, Path C and Scheme 2) Treatment of *N*-trichloroacetate **28c** obtained from *N*-tosyl-aniline **31c** with PPh₃ in refluxing toluene led to a mixture of uncharacterized products. (Scheme 2).

An exciting expansion in ynamide chemistry [46,47] has been initiated by the pioneer work of Stang and Kitamura who prepared ynamides by reaction of alkynyl iodonium triflates [48] or tosylates [49] with lithium amides. Unfortunately, this methodology applied to amides **31a,b** and iodonium triflate **30** failed. (Scheme 1, Path D) Inspired by Buchwald's copper-catalyzed *N*-alkynylations of amides [50], practical cross-coupling between amides and alkyne bromides have been developed using copper salts (CuSO₄ · 5H₂O [51], CuI, Cu₂O, Cu(OAc)₂) or simple copper powder. Later on, W. Tam [52] significantly improved the yield of ynamides by using modified reaction conditions (0.20–0.30 eq of CuI, 0.22–0.36 eq of the 1,10-phenanthroline ligand and adding 1.20 eq of the base KHMDS slowly over 3–4 h in toluene at 90 °C). In 2008, Skrydstrup et al. [53] published the Hsung's second generation protocol where potassium phosphate or carbonate was used as the mild base instead of KHMDS. In this case, anhydrous K₃PO₄ provided higher ynamide yields (52–91%). The latter conditions brought positive results and ynamides **22a,b** were prepared from carbamates **31a,b** and alkyne bromide **29** in 94 and 59% yield respectively over two steps (Scheme 1, Path D and Scheme 3).

2.3.2. Preparation of azides **15–21**

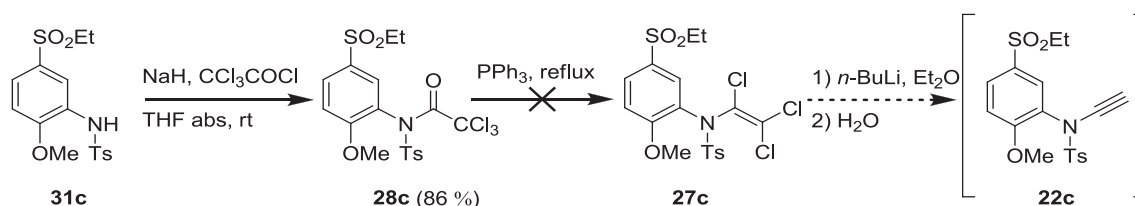
The synthesis of azides **15–19** and **21** was performed by a Suzuki–Miyaura cross coupling conditions followed by an aromatic substitution (Scheme 4).

According to this strategy, azides **15**, **16**, **17** and **19** were prepared in 57 and 45% yield over two steps for **15** and **17**, 33% over three steps for **16** and 37% yield over seven steps for **19** starting from *p*-nitroaniline **37**. (Scheme 5) Palladium catalyzed coupling of 2-bromopyridine with boronic acid **33** in the presence of sodium carbonate in a mixture of water/ethanol and DME at 75 °C within 16 h [54] afforded biaryl bromide **34** which was treated with sodium azide and copper(I) iodide according to the procedure of Liang and co-workers to deliver azide **15** [55]. Subsequently a palladium-catalyzed acetoxylation of arene C–H bond at *ortho* position to the pyridine-2-yl substituent has been performed by phenyliodine diacetate (PhI(OAc)₂) in acetic anhydride [56]. These conditions led regioselectively to azide **16** in 57% yield. Biaryl bromide **36** was obtained from the Suzuki–Miyaura cross coupling

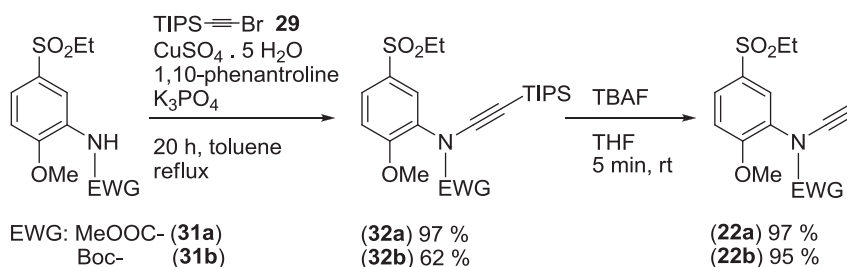


TIPS: $i\text{-Pr}_3\text{Si}$; Tos: tosyl; Piv: pivaloyl

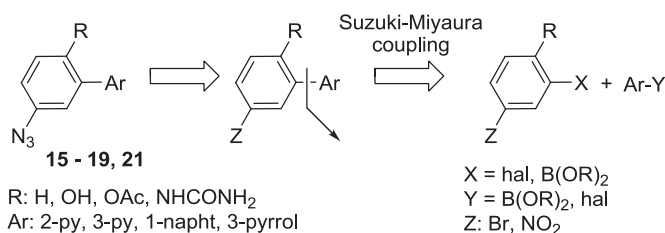
Fig. 8. Structures of required azides **15**–**21** and ynamides **22a**–**22d**.



Scheme 2. An attempt for preparation of ynamide **22c** via trichloroacetate **28c**.

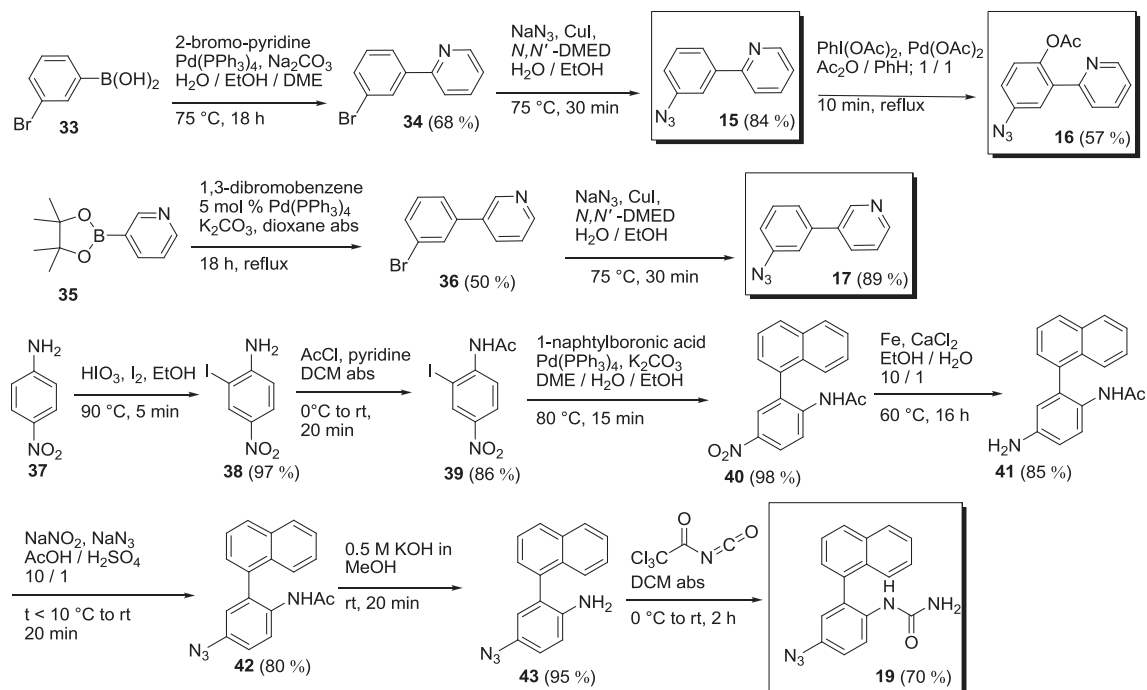


Scheme 3. Successful synthesis of ynamides **22a,b** via copper-mediated amide alkynylation.



Scheme 4. General strategy towards azides **15**–**19** and **21**.

[54] between pinacolboronic ester **35** [57] and 1,3-dibromobenzene and converted to azide **17** using the Liang's protocol. (Scheme 5) Preparation of azide **19** started with *ortho*-iodination [58] of *p*-nitroaniline **37**, followed by protection of amino group of the resulting iodide **38** as acetamide **39** [59] and subsequent Suzuki–Miyaura coupling with 1-naphthylboronic acid [60] to deliver biaryl compound **40** in 82% yield over three steps. The transformation of nitro derivative **40** to the corresponding azide **42** was performed in 2 steps via aniline **41** [61,62]. Deacetylation of **42** was performed under basic conditions. Finally, urea derivative **19** was prepared from unstable amino azide **43** by trichloroacetyl

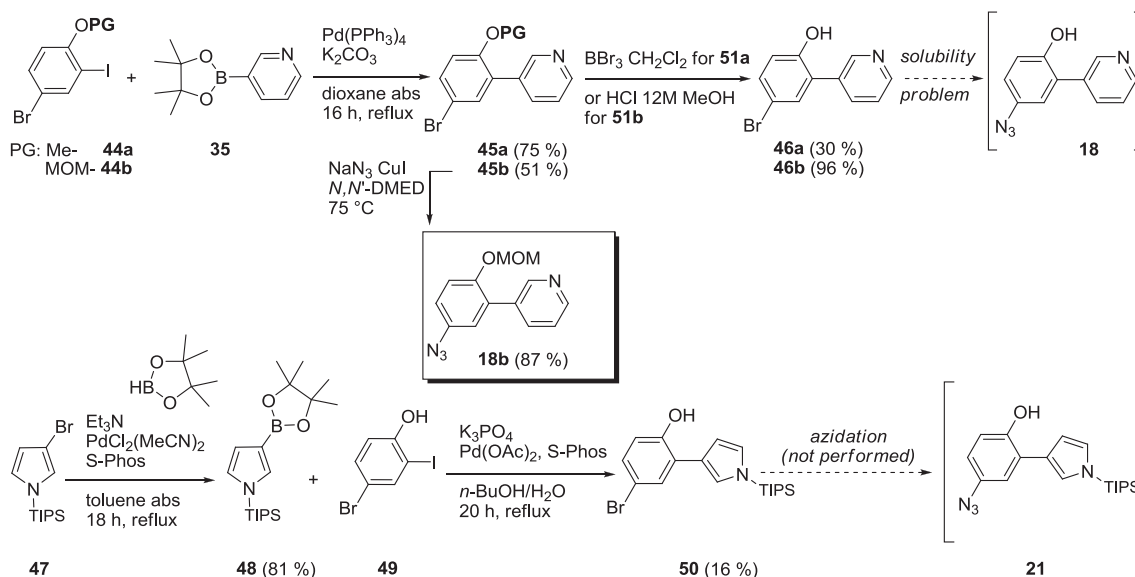
Scheme 5. Synthesis of azides **15**–**17** and **19**.

isocyanate in dry dichloromethane followed by basic treatment [63] (Scheme 5).

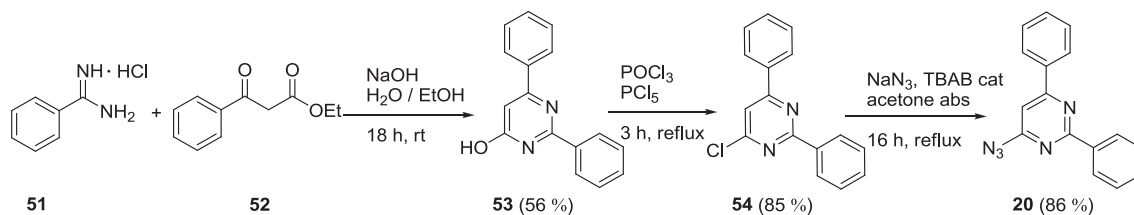
Preparation of azides **18** and **21** following the same pathway proved to be unsuccessful with exception of **18b** that was prepared in 44% yield over two steps (Scheme 6).

Biaryllic bromophenols **46a,b** were obtained by coupling reaction of pinacolboronic ester **35** with protected iodophenols **44a,b**, prepared from *para*-bromoanisole respectively in one (98% yield) or three steps (87% overall yield), and subsequent deprotection of the resulted phenol ethers **45a,b** in 23 and 49% two step yield, respectively. Unfortunately zwitterionic bromophenol **46** was insoluble and its transformation into the azide **18** was impossible.

For this reason we decided to perform the Huisgen cycloaddition prior HO- deprotection step in order to prepare triazole **T5** (Scheme 8). Methoxymethyl protected phenol **18b** was obtained in 44% yield through two steps from **44b** (Scheme 6). Preparation of pyrrole boronic ester **48** from commercially available 3-bromo-N-triisopropylsilylpyrrole **47** has been performed using pinacolborane in the presence of a catalytic amount of bis(acetonitrile)palladium dichloride and S-Phos (dicyclohexyl(2',6'-dimethoxybiphenyl-2-yl) phosphine) [64]. The Suzuki–Miyaura cross coupling [63,65,66] between **48** and the dihalogenated phenol **49** led only to 16% yield of biaryllic compound **50** (Scheme 6). This low yield led us to consider another pathway and to perform the coupling reaction



Scheme 6. Attempts for the synthesis of azides **18** and **21**. Preparation of **18b**.

Scheme 7. Preparation of pyrimidine azide **20**.

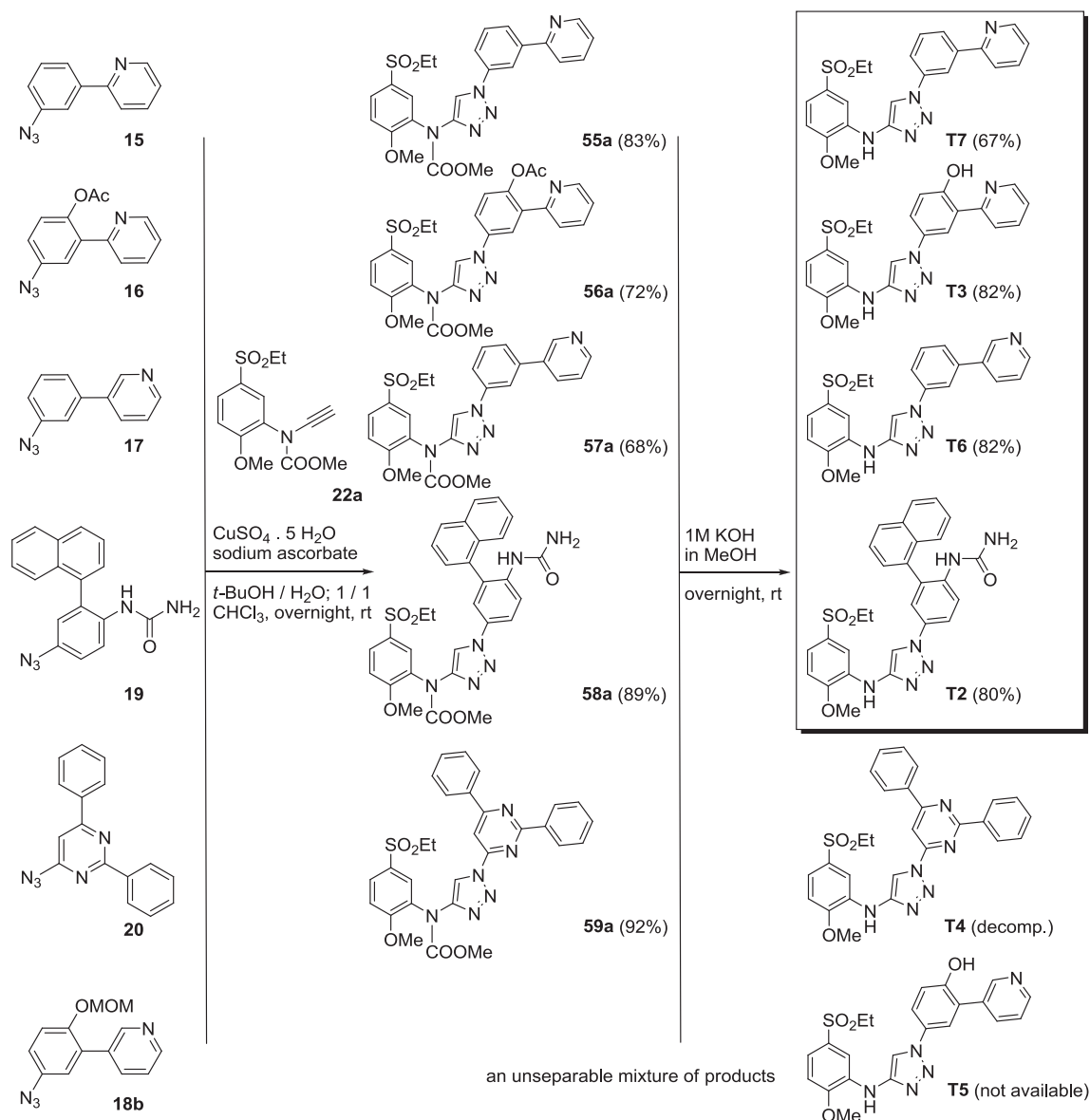
after the Click cycloaddition in order to prepare triazole **T1** (Scheme 11).

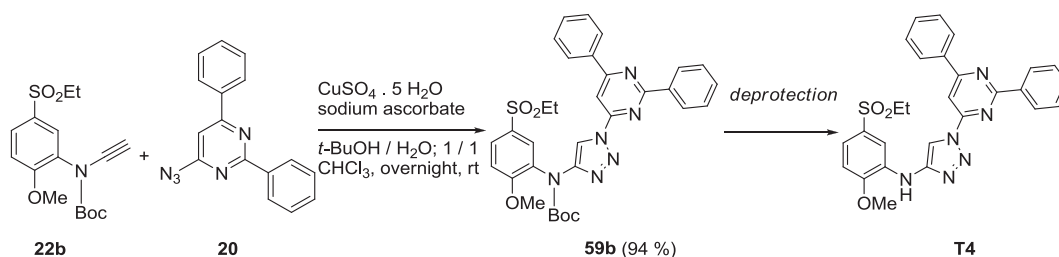
The pyrimidine azide **20** was prepared through pyrimidine core construction in 41% yield over three steps. Compound **53** was obtained from condensation of benzamidine hydrochloride **51** with β -ketoester **52** in 56% yield according to the literature [67] (Scheme 7). Treatment of **53** in refluxing POCl_3 and PCl_5 within 3 h delivered **54** in 85% yield. Transformation of chloride **54** to azide **20** was performed using sodium azide and catalytic amount of tetra-*n*-

butylammonium bromide (TBAB) [68] (Scheme 7).

2.3.3. Click reactions and preparation of target compounds

We tried to perform the synthesis of predicted 1,2,3-triazoles **T2** – **T7** via cycloaddition between azides **15**–**20** and ynamide **22a** (Scheme 8). The Click chemistry concept was introduced by Sharpless and co-workers in 2001 [69,70]. All Click reactions were performed under mild conditions using *in situ* prepared catalytic amount of copper(I) to control the 1,4-regioselectivity (5 mol %

Scheme 8. Preparation of target molecules **T2**, **T3**, **T6** and **T7** via Click chemistry approach.

Scheme 9. Synthesis of **T4** via deprotection of **59b**.

$\text{CuSO}_4 \cdot 5\text{H}_2\text{O}$ /10 mol % sodium ascorbate) in a mixture of solvents (*t*-BuOH/ H_2O / CHCl_3) at room temperature. Deprotection of the resulting methyl carbamates **55a**–**59a** was performed in 1 M methanolic solution of KOH to give triazoles **T2**, **T3**, **T6**, **T7** in good yields (67–82%). Unfortunately pyrimidine triazole **T4** was unstable in these conditions and only decomposition products were observed. Cycloaddition of **18b** with **22a** and its crude mixture deprotection resulted in a mixture of products containing the expected triazole **T5** which was impossible to isolate (Scheme 8).

In order to get pyrimidine triazole **T4**, different conditions for deprotection of **59a** have been tested (1 M KOH in MeOH or 0.5 M KOH ethylene glycol both at rt or by reflux) and gave only products of decomposition. In order to circumvent these difficulties we prepared also *N*-Boc protected triazole **59b** by Click reaction of ynamide **22b** and azide **20**. Then we tried different conditions for deprotection of cycloadduct **59b** (Scheme 9) and found that 12 M HCl in EA or TFA in THF led to **T4** with 80 and 90% conversion [71,72] (Scheme 9). Unfortunately, we were unable to isolate the pure triazole **T4** in reasonable amounts as it was unstable on silica or alumina gel. Trituration or crystallization of the crude mixture was also unsuccessful in our hands.

Since the preparation of azide **18** failed due to the high insolubility of its precursor **46** (Scheme 6), we tried a Suzuki–Miyaura coupling between triazole derivative **60** and pinacol boronic ester **35** [54] or the boronic acid **61** [73,74] but also in these cases only insoluble material has been obtained (Scheme 10).

Finally, the triazole **T1** has been prepared by the Click reaction prior to the biaryl formation (Scheme 11). The synthesis started with the preparation of the functionalized azides **65** and **68** by nitration of *o*-iodo phenol **62** using a 70% solution of nitric acid giving nitrophenol **63**. Part of **63** was acetylated to ester **66** and a subsequent reduction of both nitro compounds **63** and **66** using SnCl_2 provided anilines **64** and **67**, resp. [75] that were transformed into azides **65** and **68** by diazotation and reaction with NaN_3 . Cycloaddition between ynamide **22a** and azides **65** or **68** using the Click chemistry conditions furnished triazole derivatives **69** or **70** in 80 and 92% yield respectively. The latter were submitted to Suzuki–Miyaura coupling with pinacolboronic ester **54** to afford *N*-protected triazole **71**. Compound **71** was deprotected under basic

conditions to give triazole **T1** in 5% yield over 6 steps (via azide **65**) or 8% yield over 7 steps (via azide **68**) both started from 2-iodophenol **62** (Scheme 11).

2.4. Inhibition of VEGFR2 TK activity

Prepared compounds **T1**–**T3** and **T6**–**T7** were screened on their ability to inhibit VEGFR2 kinase activity. Their IC_{50} values were determined by radiometric protein kinase assay in 10 semi-logarithmic concentrations [25]. Compounds **T1**, **T3** and **T7** bind specifically to VEGFR2 tyrosine kinase and resulting typical concentration dependent enzymatic activity sigmoid curves. (e.g. Fig. 9).

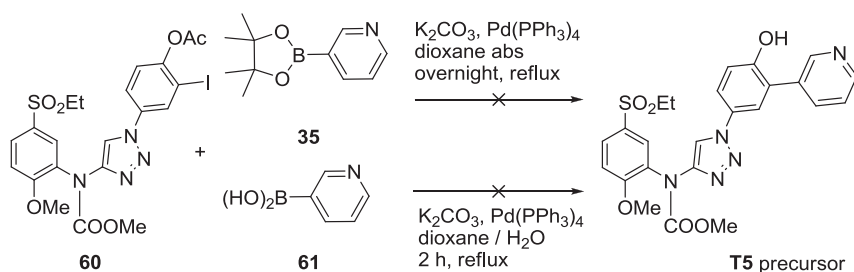
The structures of **AAZ** and 1,2,3-triazoles **T1**–**T7** together with their docking score [76] and obtained biological activity (IC_{50} , VEGFR2 TK) are depicted on Fig. 10.

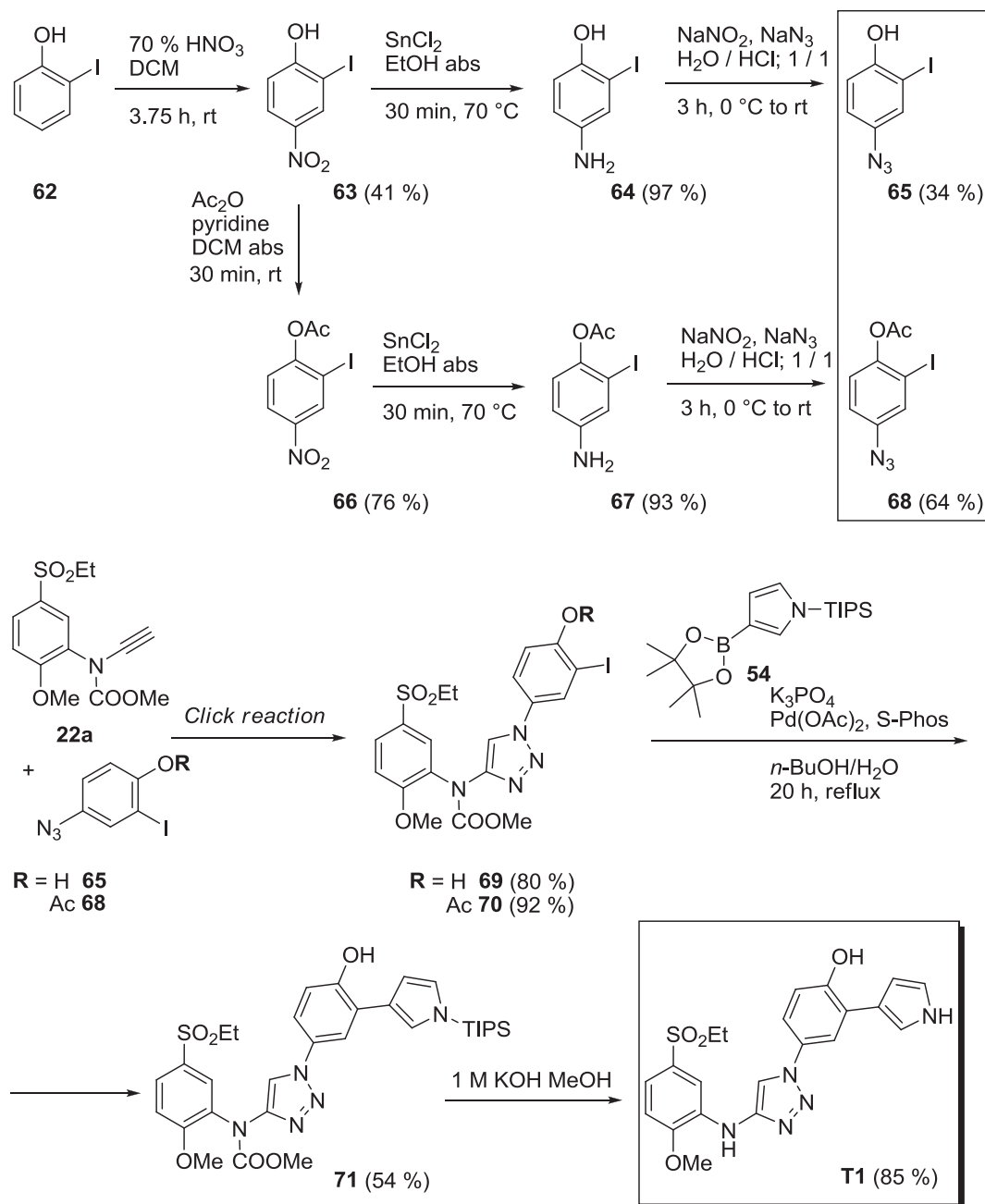
The compounds **T1**–**T3**, **T6**–**T7** exhibited different inhibitory properties: from inactive compounds **T2** and **T6** ($\text{IC}_{50} > 1\text{E-4 M}$), through weakly active **T7**, **T1** (IC_{50} : 42 and 40 μM , resp.) to moderately active inhibitor **T3** (IC_{50} : 6.96 μM). Triazole compounds **T1**–**T3** and **T6**–**T7** possess the same pharmacophoric 5-(ethylsulfonyl)-2-methoxyphenylamine moiety therefore the observed activities are dependent on the remaining aryltriazole fragments.

2.5. Re-docking experiments

In initially performed docking experiments (an older DOCK software) triazoles **T2**–**T6** showed better or similar scores as their oxazole precursor **AAZ** (vide supra, see part 2.2 In Silico predictions). These results do not correlate well with obtained IC_{50} activities determined from the biological assay. Therefore an additional docking experiment was performed with newer version of the docking software (DOCK 3.6) on more calibration reliable VEGFR2 kinase taken from PDB: 1Y6B [76]. By these conditions all triazole structures although still retaining **AAZ**-like pose in VEGFR2 TK, showed worse score values compared to their oxazole analogue **AAZ** (Figs. 10 and 11).

A similar result, disfavoring 1,2,3-triazoles compare to oxazoles, was obtained from docking experiment of isosteric pairs

Scheme 10. Attempts for the preparation of **T5** precursor.

Scheme 11. Synthesis of triazole **T1** from iodophenol **62**.

derived from **T1**–**T7** (values not shown).

Predicted VEGFR2 TK interactions of the most active 1,2,3-triazole **T3** and its oxazole bioisostere **T3-ox** are depicted on Fig. 12. The activity of **T3-ox** inhibitor (IC_{50} : 12.8 nM, VEGFR2 TK) was published by us recently [77].

2.6. Influence of isosteric oxazole/triazole replacement

The above re-docking experiment of 1,2,3-triazoles **T1** – **T7** resulted lower score than that obtained for their oxazole analogues. Triazoles **T3** and **T7** exhibit much less inhibitory activity against VEGFR2 kinase compared to their known oxazole bioisosters **T3-ox** and **AAZ** (**T3/T3-ox**, IC_{50} : 6950 nM/12.8 nM and **T7/AAZ**, IC_{50} : 42 400 nM/22 nM). Therefore we proposed that different electronic properties of oxazole/triazole core (the size and dipole moment

orientation) could be responsible for disfavouring of 1,2,3-triazole derivatives in VEGFR2 TK **AAZ** binding site. The structures and dipole moments depicted on Fig. 13 were performed by Discovery Studio 3.5 Visualizer [78].

VEGFR2 kinase surrounds the oxazole core of **AAZ** ligand (PDB: 1Y6A) by seven nonpolar amino acid residues: Phe916, Val914, Val897, Ala864, Leu838, Leu1033, and Cys917. This lipophilic pocket is less favourable to accommodate more polar 1,2,3-triazole ring. Therefore triazoles **T1**–**T7** are less suitable to inhibit VEGFR2 kinase compare to their bioisosteric oxazole analogues.

2.7. Cytotoxic activity of triazoles on hepatocellular carcinoma cell lines

Four of developed 1,4-triazole compounds **T1**–**T3** and **T6** were

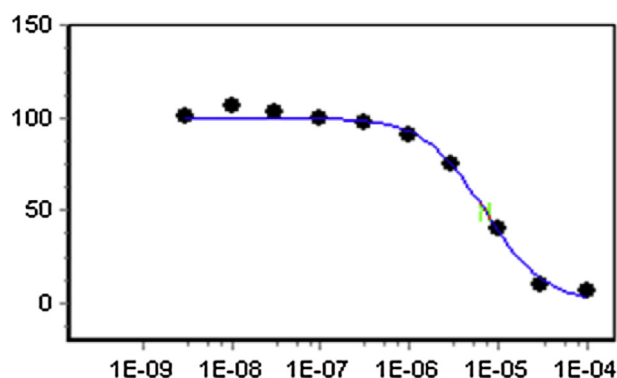


Fig. 9. An example of diminishing VEGFR2 TK activity (y axis, %) by increasing concentration of triazole **T3** (x axis, logarithmic scale) and determining **T3** activity (IC_{50} : 6.96 μ M).

screened on their cytotoxic activity against well and poorly-differentiated aggressive human hepatocellular carcinoma cell lines (Huh-7 and Mahlavu). Huh-7 is a well-differentiated (epithelial-like) hepatocellular carcinoma cell line commonly used studying liver cancer potential therapies. Huh7 expresses mutated but functional p53 protein. Mahlavu is a poorly-differentiated (mesenchymal-like) hepatocellular carcinoma cell line. This cell line is associated with the loss of PTEN protein expression leading to the constitutive activation of PI3K/Akt pathway involved in cell survival and anti-apoptotic signalling. Comparative analysis of both cell lines with lower IC_{50} values for Mahlavu cells are preferred for the reason that the inhibitor may acts on its hyperactive PI3K/Akt pathway. Additionally selective inhibitor can be further analysed for its possible specific target in PI3K/Akt pathway.

Results indicate that three of four tested triazole molecules were able to inhibit growth of both tumour cell lines by half in indicated

concentrations. (Table 1).

All triazoles **T1–T3** and **T6** display lower IC_{50} values in mesenchymal-like Mahlavu cells. Even though no enzymatic modulation of **T2** on VEGFR2 TK was obtained, triazole **T2** is highly selective inhibitor of growth of Mahlavu cells. The mechanism of its action as possible inhibitor of PI3K/Akt pathway could be further investigated at molecular level.

3. Conclusions

Seven 1,2,3-triazole compounds **T1–T7** derived from oxazole VEGFR2 TK inhibitor **AAZ** were designed. A methodology for the synthesis of the pharmacophoric ynamides **22a,b** was developed. Ynamide **22a** was used as a joined precursor for the synthesis of all proposed compounds **T1–T7**. Cu(I) catalyzed Click reaction performed from **22a** with different azides (yields: 68–92 %) confirmed the synthetic reliability of this still “exotic” ynamide reagent. Five triazole compounds **T1–T3**, **T6–T7** were prepared (Schemes 8 and 11) and screened in the radiometric VEGFR2 kinase assay. Within concentration limits of used biological test, two compounds **T2**, **T6** showed to be inactive ($IC_{50} > 1E-4$ M) and other three triazoles modulate VEGFR2 tyrosine kinase activity: **T7** (IC_{50} : 42.0 μ M), **T1** (IC_{50} : 40.1 μ M) and **T3** (IC_{50} : 6.96 μ M). (Fig. 10) The activities of new compounds were significantly lower than the ones obtained for their oxazole bioisosters (e.g. **T3/T3-ox** in Fig. 12 and **T7/AAZ** in Fig. 10, resp.). Despite the diminished activity, the triazole modulators **T1**, **T3** and **T7** inhibit VEGFR2 kinase by concentration dependent manner. Concerning the identical substructure, the inhibitory activity of 1,4-triazoles **T1–T3**, **T6–T7** is depending on the decoration of their aryl part joined to N(1) of the triazole core. (Fig. 1) The different electronic properties of 1,2,3-triazole and oxazole fragments (size and orientation of the dipole moments, Fig. 13) were proposed to be responsible for low activities of triazoles **T1–T3**, **T6–T7**. The more polar triazole core binds less readily into VEGFR2 TK lipophilic oxazole binding pocket known

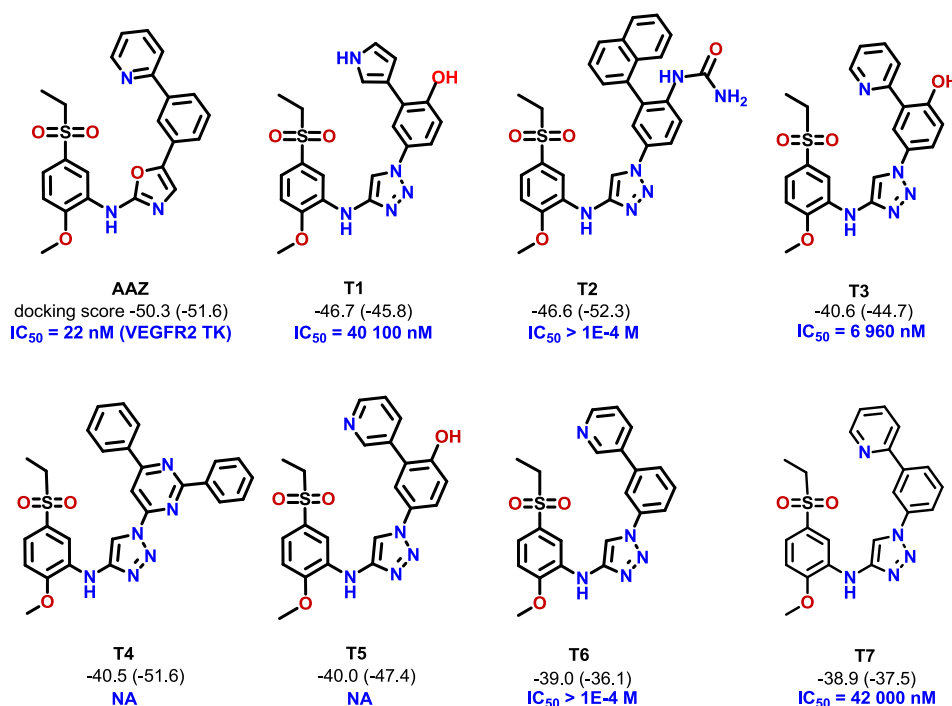


Fig. 10. The structures of **AAZ** and **T1–T7**, their docking score (software DOCK3.6, [76] VEGFR2 TK conformer from PDB: 1Y6B (1Y6A), resp.) and obtained IC_{50} activity (VEGFR2 TK), if not otherwise stated. NA: the compound was not available.

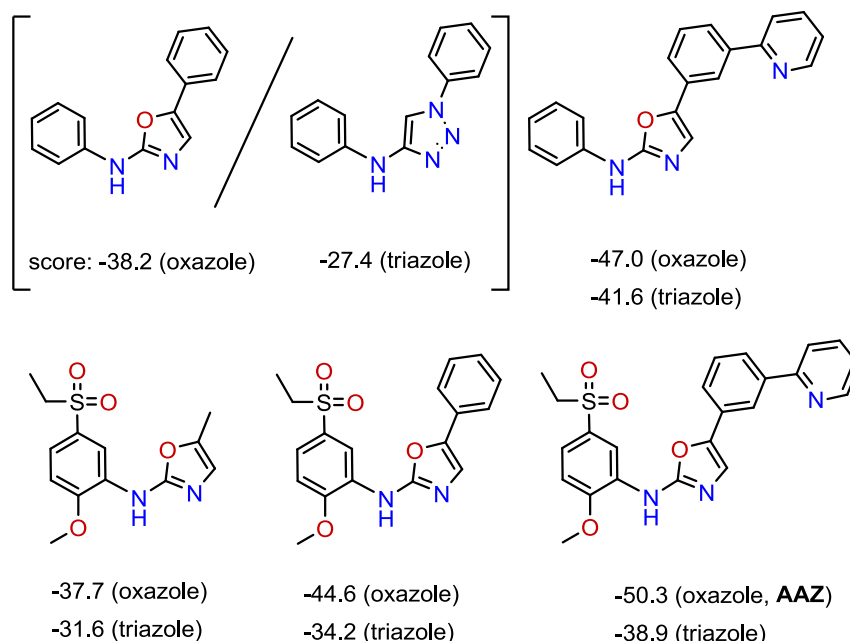


Fig. 11. The structures of isosteric oxazole/triazole pairs and relative binding energies obtained after their docking into VEGFR2 TK variant from PDB: 1Y6B. (software DOCK 3.6) All depicted structures are predicted to occupy **AAZ**-like binding site in VEGFR2 kinase.

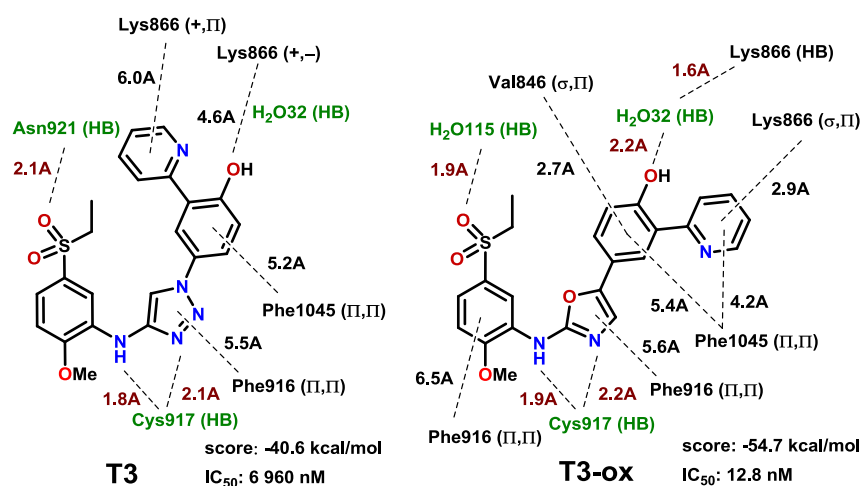


Fig. 12. The structures, docking scores, IC_{50} activities, predicted conformations and interactions (DOCK 3.6) of **T3** and **T3-ox** isosteres.

from PDB: 1Y6A or 1Y6B. We can conclude that the 1,2,3-triazole compounds **T1**, **T3** and **T7** are weaker VEGFR2 inhibitors than their oxazole analogues.

In addition **T1–T3** and **T6** were screened on their cytotoxic activity against two human hepatocellular carcinoma cell lines (Huh-7 and Mahlavu). Mahlavu is aggressive hepatocellular carcinoma

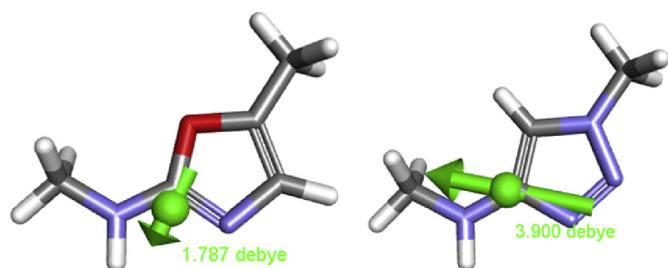


Fig. 13. Calculated dipole moment (value and orientation) for simplified N,5-dimethyloxazol-2-amine (left structure) and its 1,2,3-triazole isostere in the right.

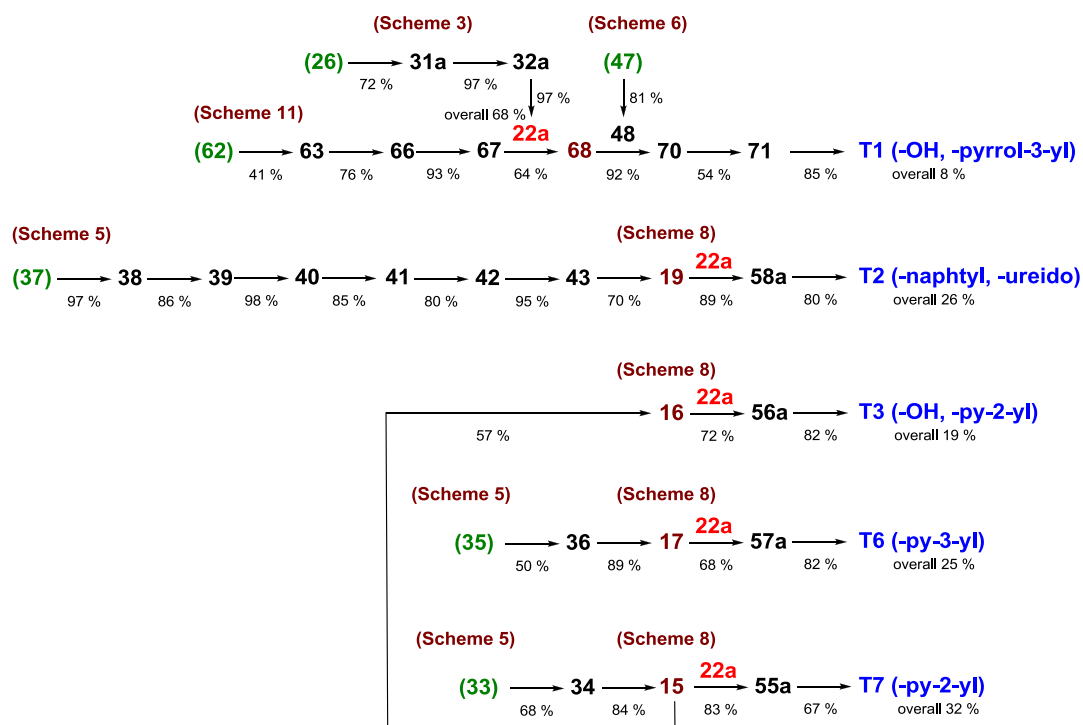
Table 1

IC_{50} growth inhibition values of compounds **T1–T3** and **T6** in Huh7 and Mahlavu human cancer hepatocellular cell lines.

Compound	Tumour cell lines IC_{50} [uM]	
	Huh7	Mahlavu
T6	12.8	11.8
T1	18.6	11.9
T3	17.7	13.8
T2	NA ^a	27.9

Data in the table are sorted according to activity against aggressive Mahlavu tumor cells.

^a NA: not active.



Scheme 12. The syntheses of 1,2,3-triazoles **T1** – **T3**, **T6** – **T7** (marked in blue), their intermediates and yields. Compounds in parenthesis are commercially available starting materials (marked green). Ynamide **22a** and azides **15–17** (marked in red and brown, resp.) The schemes in parenthesis present over the compound(s) number(s) (marked in brown) describe synthetic pathway and its reaction conditions. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

with constitutive active PI3K/Akt anti-apoptotic signalling. All screened triazole compounds performed low IC_{50} activity (12–28 μ M) against Mamlu cells (Table 1). Even though compounds **T6** and **T2** were not active in VEGFR TK assay, both of them showed interesting cytotoxic activity against Mamlu carcinoma. The high selectivity of triazole **T2** to Mamlu over Huh-7 cell lines indicates its possible affinity to PI3K/Akt pathway that can be further investigated.

4. Experimental

4.1. Molecular docking

Docking experiments were accomplished according the expert self-assessing system DOCK Blaster (University of California, San Francisco) described in the literature [79]. The calculations were performed by an older DOCK version and later on with the DOCK 3.6 version of UCSF DOCK software [33,76].

4.2. Chemistry

All reactions and compounds leading to prepared 1,2,3-triazole products **T1** – **T3**, **T6** and **T7** are completely described in Supplementary Material in order as they are depicted in the Scheme 12. Beside other compound characteristics the Supplementary material contains also 1H and ^{13}C NMR spectral graphical abstracts.

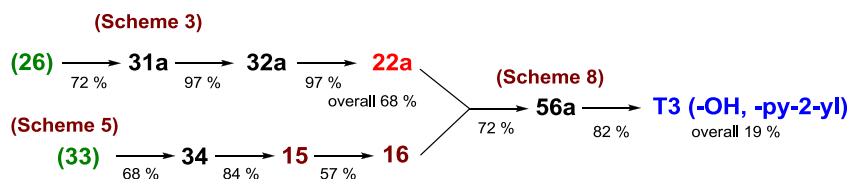
For simplicity, only the synthetic pathway leading to the most active triazole **T3** is described here (Scheme 13).

4.2.1. General procedures

THF and Et_2O were dried over and distilled from Na/benzophenone under Ar atmosphere. DCM and Et_3N were dried over calcium

hydride or KOH pellets, resp. and distilled. Commercially available chemicals and solvents were purchased from Sigma–Aldrich company and were used without further purification. The course of the reactions was followed by TLC analysis (Merck Silica gel 60 F₂₅₄). UV lamp (254 nm) and iodine vapours were used for visualization of TLC spots. Flash column chromatography was performed on silica gel (40–60 mesh). Melting points were determined on a Büchi B-540 apparatus (Büchi Labortechnik, Flawil, Switzerland) and were uncorrected. All 1H -NMR and ^{13}C -NMR spectra were recorded on Bruker instruments (500, 300 MHz for hydrogen and 100 MHz for carbon, Bruker Bioscience, Billerica, MA, USA) with $CDCl_3$, $DMSO-d_6$ or acetone- d_6 as solvent. Chemical shifts are given in parts per million (ppm). IR spectra were acquired on FT-IR-ATR REACT IR 1000 (ASI Applied Systems) with diamond probe and MTS detector. ESI Mass spectral data were obtained by Esquire-LC-00075 spectrometer (Bruker Bioscience). Other mass spectra were performed on LC-MS (Agilent Technologies 1200 Series equipped with Mass spectrometer Agilent Technologies 6100 Quadrupole LC-MS). Elemental analyses for carbon, hydrogen and nitrogen were performed with an Eager 300 analyzer. Used abbreviations: EA: ethyl acetate, Boc: tert-butylloxycarbonate, Cy: cyclohexane, DCM: dichloromethane, EWG: electron withdrawing group (e.g. –COOMe, –Boc, –Ts etc.), FLC: Flash liquid chromatography, N,N' -DMED: N,N' -dimethylethylenediamine, RT: room temperature, S -Phos: 2-dicyclohexylphosphino-2',6'-dimethoxybiphenyl [CAS 657408-07-6], TIPS: triisopropylsilyl, TLC: thin layer chromatography (SiO_2/UV_{254}). Elemental analyses indicated in the experimental part by the symbols of the elements were within $\pm 0.4\%$ of the theoretical values.

4.2.1.1. A: N -alkynylation of protected anilines with bromoalkyne 29 (Skydrup's protocol). 1-Bromoalkyne (1.1 mol eq) **29** (1.0 M solution in toluene) was added to a mixture of an amide (1.0 mol eq) (EWG protected aniline) **31a–c**, (3.0 mol eq) K_3PO_4 , (0.2 mol eq)



Scheme 13. The syntheses of triazole **T3** (marked in blue), its intermediates and yields. Starting compounds are marked green, ynamide **22a** and azides (**15–16**) are marked in red and brown, resp. (For interpretation of the references to colour in this figure legend, the reader is referred to the web version of this article.)

CuSO₄ · 5H₂O and (0.4 mol eq) 1,10-phenantroline. The reaction mixture was heated at 65–75 °C (oil bath temperature) for 20 h. Upon completion, detected by TLC, the reaction mixture was cooled to RT, diluted with EA, filtered through Celite and the filtrate concentrated under reduced pressure. The crude product was purified by flash chromatography on silica gel (Scheme 3).

4.2.1.2. B: TIPS group deprotection. A (1.0 mol eq) 1.0 M solution of TBAF in THF was added dropwise to a solution of TIPS protected ynamide (1.0 mol eq) **32a–b** in dry THF. After 5 min stirring at RT, the volatile parts were evaporated under reduced pressure. The concentrated reaction mixture was partitioned in a mixture of EA and brine (1:1). Collected organic layers were dried over anhydrous MgSO₄, filtered and evaporated leading to a desired product (Scheme 3).

4.2.1.3. C: substitution of an aromatic halide to azide. The solution of (1.0 mol eq) aryl bromide, (2.0 mol eq) NaN₃, (0.05 mol eq) sodium ascorbate, (0.1 mol eq) CuI and (0.15 mol eq) *N,N'*-DMED in 4 mL of mixture EtOH : H₂O (7 : 3) was refluxed under Ar atmosphere. Progress of the reaction was monitored by TLC. When the aryl bromide was completely consumed, or when the progress of the reaction had stopped, the reaction mixture was cooled to RT and the crude product purified either by extraction and/or FLC chromatography giving the desired aryl azide (Scheme 5).

4.2.1.4. D: copper-catalyzed Click chemistry reaction. Prepared (1.0 mol eq) ynamide **22a** and (1.0 mol eq) of corresponding azide were suspended in 6 mL of mixture *tert*-butanol : H₂O (1 : 1) and 5 mL CHCl₃. Consequently, premixed solution of sodium ascorbate (10 mol %) and CuSO₄ · 5H₂O (5 mol %) in 3 mL of H₂O was added. After stirring overnight the reaction mixture was diluted with 10 mL H₂O and extracted with EA (3 × 10 mL). The combined organic layers were washed with 10 mL of NH₄OH (3% solution in brine), treated over MgSO₄, filtered, evaporated and dried in HV. Purification using FLC chromatography afforded 1,4-disubstituted 1,2,3-triazole product (Scheme 8).

4.2.1.5. E: deprotection of methoxycarbonyl group. Protected aminotriazole derivative (2.0 mmol, 1.0 mol eq) was stirred in 5 mL of 1 M solution of KOH in MeOH at RT overnight. The progress of reaction was monitored by LCMS or ¹H NMR spectroscopy. When the reaction was accomplished, the mixture was neutralized with 1 M HCl and extracted with EA (3 × 8 mL). The combined organic layers were washed with water, treated over MgSO₄, filtered, evaporated under reduced pressure and dried with HV. Obtained crude product was purified by FLC (Scheme 8).

4.3. Synthesis of triazole **T3** (–OH, –py-2-yl)

4.3.1. Synthesis of methyl 5-(ethylsulfonyl)-2-methoxyphenylcarbamate (**31a**)

Pyridine 1.7 mL (21.2 mmol, 1.1 mol eq) was added to solution of 4.15 g (9.3 mmol, 1.0 mol eq) 5-(ethylsulfonyl)-2-methoxyaniline

(**26**) in 35 mL of dry DCM. ClCOOMe 1.6 mL (21.2 mmol, 1.1 mol eq) was added to the reaction mixture dropwise at 0 °C. The mixture was stirred at RT for 2.5 h and consumption of starting material was confirmed by TLC. The reaction was quenched with brine (2 × 30 mL) and the organic layer was separated, dried over anhydrous MgSO₄ and concentrated under reduced pressure. The crude product was purified by crystallization from Et₂O with charcoal to yield 3.15 g (13.9 mmol, 72%) **31a** in form of white crystals. **M.p.** 121.0–123.0 °C [Et₂O].

¹H-NMR (300 MHz, CDCl₃): δ 8.54 (br s, 1H, –NH–), 7.53 (dd, 1H, *J*(3,4) = 8.6 Hz, *J*(4,6) = 2.3 Hz, H–C(4)), 7.19 (d, 1H, *J*(4,6) = 2.3 Hz, H–C(6)), 6.90 (d, 1H, *J*(3,4) = 8.6 Hz, H–C(3)), 3.89 (s, 3H, –OCH₃), 3.74 (s, 3H, –COOCH₃), 3.05 (2H, q, *J*(CH₂,CH₃) = 7.4 Hz, –SO₂CH₂CH₃), 1.21 (3H, t, *J*(CH₂,CH₃) = 7.4 Hz, –SO₂CH₂CH₃).

¹³C-NMR (100 MHz, CDCl₃): δ 153.6 (s, –NHCOO–), 151.2 (s, C(2)), 130.9 (d, C(4)), 128.5 (s, C(5)), 123.7 (d, C(6)), 117.6 (s, C(1)), 109.7 (d, C(3)), 56.2 (q, CH₃O–), 52.6 (t, –SO₂CH₂–), 50.6 (q, –COOCH₃), 7.6 (q, –CH₂CH₃).

IR *v* (solid): 3416, 1721, 1595, 1535, 1304, 1275, 1264, 1239, 1127, 1064, 766 cm^{–1}.

Anal. Calcd for C₁₁H₁₅NO₅S (273.31): C, 48.34; H, 5.53; N, 5.12. Found: C, 48.30; H, 5.47; N, 5.02.

4.3.2. Synthesis of methyl 5-(ethylsulfonyl)-2-methoxyphenyl((triisopropylsilyl)ethynyl)carbamate (**32a**)

Compound **32a** was prepared according to the general procedure **A**. Yield: (97%). Purification: filtration through a silica gel pad (eluent: Cy/EA, 1/1). **M.p.** 42.3–45.1 °C [Cy/EA], pale yellow solid.

¹H-NMR (300 MHz, CDCl₃): δ 7.82 (d, 1H, *J*(4,6) = 2.3 Hz, H–C(6)), 7.81 (dd, 1H, *J*(3,4) = 8.2 Hz, *J*(4,6) = 2.3 Hz, H–C(4)), 7.04 (d, 1H, *J*(3,4) = 8.2 Hz, H–C(3)), 3.88 (s, 3H, ArOCH₃), 3.77 (s, 3H, –COOCH₃), 3.02 (q, 2H, *J*(CH₂,CH₃) = 7.5 Hz, –SO₂CH₂–), 1.20 (t, 3H, *J*(CH₂,CH₃) = 7.5 Hz, –SO₂CH₂CH₃), 0.98 (br s, 21H, all CH and –CH₃ from TIPS: –Si(CH(CH₃)₂)₃).

¹³C-NMR (75 MHz, CDCl₃): δ 158.6 (s, C(2)), 157.6 (s, N(C=O)), 2 × 130.3 (s and d, C(4) and C(5)), 128.8 and 128.7 (s and d, C(1) and C(6)), 112.4 (d, C(3)), 96.2 and 77.2 (2 × s from C≡C), 56.4 (q, ArOCH₃), 54.5 (q, –COOCH₃), 50.9 (–SO₂CH₂CH₃), 18.6 (q, 6 × Me from TIPS), 11.3 (d, 3 × CH from TIPS), 7.6 (q, –SO₂CH₂CH₃).

IR *v* (solid): 2941, 2864, 2180, 1744, 1440, 1290, 1132, 730 cm^{–1}.

Anal. Calcd for C₂₂H₃₅NO₅SSi (453.67): C, 58.24; H, 7.78; N, 3.09. Found: C, 58.04; H, 7.80; N, 2.91.

4.3.3. Synthesis of methyl 5-(ethylsulfonyl)-2-methoxyphenyl(ethynyl)carbamate (**22a**)

Compound **22a** was prepared according to the general procedure **B**. Yield: 97%. Purification: filtration through a silica gel pad (eluent: Cy/EA, 1/2). **M.p.** 158.1–163.2 °C [Cy/EA], pale yellow solid.

¹H-NMR (300 MHz, CDCl₃): δ 7.83 (dd, 1H, *J*(3,4) = 8.5 Hz, *J*(4,6) = 2.3 Hz, H–C(4)), 7.81 (d, 1H, *J*(4,6) = 2.3 Hz, H–C(6)), 7.05 (d, 1H, *J*(3,4) = 8.5 Hz, H–C(3)), 3.90 (s, 3H, ArOCH₃), 3.79 (s, 3H, –COOCH₃), 3.04 (q, 2H, *J*(CH₂,CH₃) = 7.5 Hz, –SO₂CH₂CH₃), 2.74 (s, 1H, –C≡CH), 1.23 (t, 3H, *J*(CH₂,CH₃) = 7.5 Hz, –SO₂CH₂CH₃).

¹³C-NMR (75 MHz, CDCl₃): δ 158.8 (s, –C(=O)OCH₃), 154.6 (s,

C(2)), 130.8 and 130.6 ((d, C(4)) and (s, C(5))), 129.0 (d, C(6)), 128.2 (s, C(1)), 112.5 (d, C(3)), 75.8 (s, $\text{C}\equiv\text{CH}$), 57.5 (d, $\text{C}\equiv\text{CH}$), 56.5 (t, $-\text{SO}_2\text{CH}_2\text{CH}_3$), 54.7 (q, ArOCH_3), 50.9 (q, $-\text{COOCH}_3$), 7.5 (q, $-\text{SO}_2\text{CH}_2\text{CH}_3$).

IR ν (solid): 3267, 2152, 1726, 1500, 1443, 1315, 1293, 1130, 1089, 827, 730 cm^{-1} .

Anal. Calcd for $\text{C}_{13}\text{H}_{15}\text{NO}_5\text{S}$ (297.33): C, 52.51; H, 5.09; N, 4.71. Found: C, 52.62; H, 5.16; N, 4.39.

4.3.4. Synthesis of 2-(3-bromophenyl)pyridine (**34**)

A degassed mixture of 2.4 mL (24.9 mmol, 1.00 mol eq) 2-bromopyridine, 5.75 g (54.8 mmol, 2.20 mol eq) Na_2CO_3 , 28 mL H_2O , 20 mL EtOH, 63 mL 1,2-dimethoxyethane, 5.0 g (24.9 mmol, 1.00 mol eq) 3-bromophenylboronic acid (**33**) and 288.0 mg (0.25 mmol, 0.01 mol eq) $\text{Pd}(\text{PPh}_3)_4$ was refluxed for 18 h. The reaction mixture was filtered through a short Celite pad. Organic layer was separated and water layer extracted with EA (2×35 mL). Collected organic layers were washed with water (50 mL), dried over anhydrous MgSO_4 , filtered and concentrated in vacuum. The crude product was purified by FLC column chromatography (SiO_2 , eluent: Cy/EA, 1/9) to afford 3.96 g (16.9 mmol, 68%) of **34** as colourless oil. The analytical data corresponded to the literature [82].

¹H-NMR (300 MHz, CDCl_3): δ 8.67 (ddd, 1H, $J(\text{B}_5, \text{B}_6) = 5.0$ Hz, $J(\text{B}_4, \text{B}_6) = 1.8$ Hz, $J(\text{B}_3, \text{B}_6) = 1.0$ Hz, H-C_B(6)), 8.18 (dd, 1H, $J(\text{A}_2, \text{A}_4) = J(\text{A}_2, \text{A}_6) = 1.8$ Hz, H-C_A(2)), 7.88 (ddd, 1H, $J(\text{A}_5, \text{A}_6) = 7.9$, $J(\text{A}_2, \text{A}_6) = 1.8$ Hz, $J(\text{A}_4, \text{A}_6) = 1.3$ Hz, H-C_A(6)), 7.69 (ddd, 1H, $J(\text{B}_3, \text{B}_4) = 7.9$ Hz, $J(\text{B}_4, \text{B}_5) = 7.2$ Hz, $J(\text{B}_4, \text{B}_6) = 1.8$ Hz, H-C_B(4)), 7.63 (ddd, 1H, $J(\text{B}_3, \text{B}_4) = 7.9$ Hz, $J(\text{B}_3, \text{B}_5) = 1.3$ Hz, $J(\text{B}_3, \text{B}_6) = 1.0$ Hz, H-C_B(3)), 7.51 (ddd, 1H, $J(\text{A}_4, \text{A}_5) = 8.0$ Hz, $J(\text{A}_2, \text{A}_4) = 1.8$ Hz, $J(\text{A}_4, \text{A}_6) = 1.3$ Hz, H-C_A(4)), 7.30 (dd, 1H, $J(\text{A}_4, \text{A}_5) = 8.0$ Hz, $J(\text{A}_5, \text{A}_6) = 7.9$ Hz, H-C_A(5)), 7.20 (ddd, 1H, $J(\text{B}_4, \text{B}_5) = 7.2$ Hz, $J(\text{B}_5, \text{B}_6) = 5.0$ Hz, $J(\text{B}_3, \text{B}_5) = 1.3$ Hz, H-C_B(5)).

¹³C-NMR (75 MHz, CDCl_3): δ 155.8 (s, C_B(2)), 149.8 (d, C_B(6)), 141.4 (s, C_A(1)), 136.9 (d, C_B(4)), 131.9, 130.0, 130.3 and 125.4 ($4 \times$ d, C_A(2), C_A(4), C_A(5) and C_A(6)), 123.1 (s, C_A(3)), 122.7 (d, C_B(5)), 120.6 (d, C_B(3)).

Anal. Calcd for $\text{C}_{11}\text{H}_8\text{BrN}$ (234.09): C, 56.44; H, 3.44; N, 5.98. Found: C, 56.28; H, 3.48; N, 5.62.

4.3.5. Synthesis of 2-(3-azidophenyl)pyridine (**15**)

Compound **15** was prepared according to the general procedure **C**. Yield: 84%. Purification: filtration through silica gel (eluent: Cy/EA, 1/1) Colourless liquid.

¹H-NMR (300 MHz, CDCl_3): δ 8.62 (ddd, 1H, $J(\text{B}_5, \text{B}_6) = 4.7$ Hz, $J(\text{B}_4, \text{B}_6) = 1.8$ Hz, $J(\text{B}_3, \text{B}_6) = 1.0$ Hz, H-C_B(6)), 7.71–7.60 (m, 4H, H-C_A(2), H-C_A(6), H-C_B(3) and H-C_B(4)), 7.36 (dd, 1H, $J(\text{A}_4, \text{A}_5) = J(\text{A}_5, \text{A}_6) = 7.9$ Hz, H-C_A(5)), 7.17 (ddd, 1H, $J(\text{B}_4, \text{B}_5) = 7.2$ Hz, $J(\text{B}_5, \text{B}_6) = 4.7$ Hz, $J(\text{B}_3, \text{B}_5) = 1.3$ Hz, H-C_B(5)), 6.98 (ddd, 1H, $J(\text{A}_4, \text{A}_5) = 7.9$ Hz, $J(\text{A}_2, \text{A}_4) = 2.3$ Hz, $J(\text{A}_4, \text{A}_6) = 1.2$ Hz, H-C_A(4)).

¹³C-NMR (75 MHz, CDCl_3): δ 156.1 (s, C_B(2)), 149.7 (d, C_B(6)), 141.1 (s, C_A(1)), 140.6 (d, C_B(4)), 136.8 and 130.0 ($2 \times$ d, C_A(2) and C_A(5)), 123.3 and 122.6 ($2 \times$ d, C_A(4) and C_A(6)), 120.5 (s, C_B(3)), 119.5 (d, C_B(5)), 117.5 (d, C_A(3)).

IR ν (solid): 3053, 2095, 1578, 1564, 1463, 1449, 1414, 1295, 1270, 1260, 991, 879, 765, 736, 666 cm^{-1} .

Anal. Calcd for $\text{C}_{11}\text{H}_8\text{N}_4$ (196.21): C, 67.34; H, 4.11; N, 28.55. Found: C, 67.41; H, 4.17; N, 28.25.

4.3.6. Synthesis of 4-azido-2-(pyridin-2-yl)phenyl acetate (**16**)

Azide 675.0 mg (3.4 mmol, 1.0 mol eq) **15**, 1.22 g (3.8 mmol, 1.1 mol eq) $\text{PhI}(\text{OAc})_2$ and 38.6 mg (0.17 mmol, 0.05 mol eq) $\text{Pd}(\text{OAc})_2$ was suspended in a mixture of 8 mL benzene and 8 mL Ac_2O . The reaction vial was sealed with a Teflon cap and heated at 100 °C for 1.5 h. Then the volatile parts were evaporated under

reduced pressure, and the resulting oil purified by FLC chromatography (SiO_2 , eluent: Cy/EA, 3/1). Product 446 mg (1.9 mmol, 57%) **16** was isolated in form of pale yellow oil.

¹H-NMR (300 MHz, CDCl_3): δ 8.70 (ddd, 1H, $J(\text{B}_5, \text{B}_6) = 4.9$ Hz, $J(\text{B}_4, \text{B}_6) = 1.8$ Hz, $J(\text{B}_3, \text{B}_6) = 1.0$ Hz, H-C_B(6)), 7.69 (ddd, 1H, $J(\text{B}_3, \text{B}_4) = 7.9$ Hz, $J(\text{B}_4, \text{B}_5) = 7.8$ Hz, $J(\text{B}_4, \text{B}_6) = 1.8$ Hz, H-C_B(4)), 7.54 (ddd, 1H, $J(\text{B}_3, \text{B}_4) = 7.9$ Hz, $J(\text{B}_3, \text{B}_5) = 1.1$ Hz, $J(\text{B}_3, \text{B}_6) = 1.0$ Hz, H-C_B(3)), 7.41 (d, 1H, $J(\text{A}_2, \text{A}_4) = 2.7$ Hz, H-C_A(2)), 7.26 (ddd, 1H, $J(\text{B}_4, \text{B}_5) = 7.8$ Hz, $J(\text{B}_5, \text{B}_6) = 4.9$ Hz, $J(\text{B}_3, \text{B}_5) = 1.1$ Hz, H-C_B(5)), 7.15 (d, 1H, $J(\text{A}_4, \text{A}_5) = 8.6$ Hz, H-C_A(5)), 7.06 (dd, 1H, $J(\text{A}_4, \text{A}_5) = 8.6$ Hz, $J(\text{A}_2, \text{A}_4) = 2.7$ Hz, H-C_A(4)), 2.17 (s, 3H, $-\text{OCOCH}_3$).

¹³C-NMR (75 MHz, CDCl_3): δ 169.3 (s, $-\text{OC}(=\text{O})$), 154.6 (s, C_B(2)), 149.7 (d, C_B(6)), 145.0 (s, C_A(6)), 138.0 (s, C_A(3)), 136.4 (d, C_B(4)), 134.5 (s, C_A(1)), 124.8 and 123.6 ($2 \times$ d, C_A(2) and C_B(5)), 122.7, 121.0 and 120.1 ($3 \times$ d, C_A(4), C_A(5) and C_B(3)), 20.9 (q, $\text{CH}_3\text{COO}-$).

IR ν (solid): 2958, 2928, 2107, 1724, 1595, 1486, 1463, 1287, 1269, 1240, 1128, 1072, 887, 781, 738, 719, 663 cm^{-1} .

Anal. calcd for $\text{C}_{13}\text{H}_{10}\text{N}_4\text{O}_2$ (254.24): C, 61.41; H, 3.96; N, 22.04. Found: C, 61.09; H, 4.07; N, 22.14.

4.3.7. Synthesis of 4-(4-(5-(ethylsulfonyl)-2-methoxyphenyl)(methoxycarbonyl)amino)-1H-1,2,3-triazol-1-yl)-2-(pyridin-2-yl)phenyl acetate (**56a**)

Compound **56a** was prepared according to the general procedure **D**. Yield: 72%. Purification: Flash chromatography (SiO_2 , eluent: Cy/EA, 1/3). **M.p.** 155.0–157.5 °C. Pale yellow foam.

¹H-NMR (300 MHz, CDCl_3): δ 8.72 (ddd, 1H, $J(\text{D}_5, \text{D}_6) = 4.8$ Hz, $J(\text{D}_4, \text{D}_6) = 1.8$ Hz, $J(\text{D}_3, \text{D}_6) = 0.8$ Hz, H-C_D(6)), 8.46 (br s, 1H, H-C_B(5)), 8.10 (d, 1H, $J(\text{A}_4, \text{A}_6) = 2.7$ Hz, H-C_A(6)), 7.92 (dd, 1H, $J(\text{C}_5, \text{C}_6) = 8.6$ Hz, $J(\text{C}_2, \text{C}_6) = 2.2$ Hz, H-C_C(6)), 7.88 (d, 1H, $J(\text{C}_2, \text{C}_6) = 2.2$ Hz, H-C_C(2)), 7.84 (dd, 1H, $J(\text{A}_3, \text{A}_4) = 8.7$ Hz, $J(\text{A}_4, \text{A}_6) = 2.7$ Hz, H-C_A(4)), 7.77 (ddd, 1H, $J(\text{D}_3, \text{D}_4) = 7.9$ Hz, $J(\text{D}_4, \text{D}_5) = 7.7$ Hz, $J(\text{D}_4, \text{D}_6) = 1.8$ Hz, H-C_D(4)), 7.61 (ddd, 1H, $J(\text{D}_3, \text{D}_4) = 7.9$ Hz, $J(\text{D}_3, \text{D}_5) = 1.1$ Hz, $J(\text{D}_3, \text{D}_6) = 0.8$ Hz, H-C_D(3)), 7.33 (d, 1H, $J(\text{C}_5, \text{C}_6) = 8.6$ Hz, H-C_C(5)), 7.29 (ddd, 1H, $J(\text{D}_4, \text{D}_5) = 7.7$ Hz, $J(\text{D}_5, \text{D}_6) = 4.8$ Hz, $J(\text{D}_3, \text{D}_5) = 1.1$ Hz, H-C_D(5)), 7.15 (d, 1H, $J(\text{A}_3, \text{A}_4) = 8.7$ Hz, H-C_A(3)), 3.88 (s, 1H, C_A(2) OCH_3), 3.77 (s, 3H, $-\text{NCOOCH}_3$), 3.13 (q, 2H, $J(\text{CH}_2, \text{CH}_3) = 7.5$ Hz, $-\text{SO}_2\text{CH}_2\text{CH}_3$), 2.21 (s, 3H, $-\text{COCH}_3$), 1.31 (t, 3H, $J(\text{CH}_2, \text{CH}_3) = 7.5$ Hz, $-\text{SO}_2\text{CH}_2\text{CH}_3$).

¹³C-NMR (75 MHz, CDCl_3): δ 169.0 (s, $-\text{COCH}_3$), 154.2 and 153.6 ($2 \times$ s, $-\text{NCOOCH}_3$ and C_D(2)), 159.8 (s, C_A(2)), 149.9 (d, C_D(6)), 148.0 (s, C_C(4)), 136.6 (d, C_D(4)), 135.2, 134.5, 131.1, $2 \times$ 130.6, 130.5 ($5 \times$ s and d, C_A(1), C_A(5), C_B(4), C_C(1), C_C(3) and C_B(5)), 124.9, 123.7, 123.0, $2 \times$ 122.7, 121.9 and 121.5 ($7 \times$ d, C_A(4), C_A(6), C_C(2), C_C(5–6), C_D(3) and C_D(5)), 112.3 (d, C_A(3)), 56.4 (q, C_A(2) OCH_3), 53.8 (t, $-\text{SO}_2\text{CH}_2\text{CH}_3$), 51.1 (q, $-\text{COOCH}_3$), 21.0 (q, $-\text{COCH}_3$), 7.6 (q, $-\text{SO}_2\text{CH}_2\text{CH}_3$).

IR ν (solid): 2955, 1764, 1725, 1565, 1500, 1443, 1371, 1313, 1182, 1132, 1091, 1039, 735, 532 cm^{-1} .

Anal. calcd for $\text{C}_{26}\text{H}_{25}\text{N}_5\text{O}_7\text{S}$ (551.57): C, 56.62; H, 4.57; N, 12.70. Found: C, 56.72; H, 4.97; N, 12.40.

4.3.8. Synthesis of 4-(4-(5-(ethylsulfonyl)-2-methoxyphenylamino)-1H-1,2,3-triazol-1-yl)-2-(pyridin-2-yl)phenol (**T3**)

Compound **T3** was prepared according to the general procedure **E**. Yield: 82%. Purification: Flash chromatography (SiO_2 , eluent: Cy/EA, 1/4). **M.p.** 148.2–149.1 °C. Pale yellow foam.

¹H-NMR (600 MHz, CDCl_3): δ 8.56 (ddd, 1H, $J(\text{D}_5, \text{D}_6) = 5.1$ Hz, $J(\text{D}_4, \text{D}_6) = 1.8$ Hz, $J(\text{D}_3, \text{D}_6) = 0.9$ Hz, H-C_D(6)), 8.24 (d, 1H, $J(\text{C}_2, \text{C}_6) = 2.6$ Hz, H-C_C(2)), 8.03 (ddd, 1H, $J(\text{D}_3, \text{D}_4) = 8.3$ Hz, $J(\text{D}_3, \text{D}_5) = 1.1$ Hz, $J(\text{D}_3, \text{D}_6) = 0.9$ Hz, H-C_D(3)), 7.92 (ddd, 1H, $J(\text{D}_3, \text{D}_4) = 8.3$ Hz, $J(\text{D}_4, \text{D}_5) = 7.6$ Hz, $J(\text{D}_4, \text{D}_6) = 1.8$ Hz, H-C_D(4)), 7.84 (s, 1H, H-C_B(5)), 7.64 (d, 1H, $J(\text{A}_2, \text{A}_6) = 2.2$ Hz, H-C_A(2)), 7.55 (dd, 1H, $J(\text{C}_5, \text{C}_6) = 8.8$ Hz, $J(\text{C}_2, \text{C}_6) = 2.6$ Hz, H-C_C(6)), 7.41 (dd, 1H,

$J(A_3, A_4) = 8.4$ Hz, $J(A_4, A_6) = 2.2$ Hz, H- $C_A(4)$), 7.34 (ddd, 1H, $J(D_4, D_5) = 7.6$ Hz, $J(D_5, D_6) = 5.1$ Hz, $J(D_3, D_5) = 1.1$ Hz, H- $C_D(5)$), 7.15 (d, 1H, $J(C_5, C_6) = 8.8$ Hz, H- $C_C(5)$), 6.97 (d, $J(A_3, A_4) = 8.4$, H- $C_A(3)$), 6.92 (br s, 1H, –OH), 4.00 (s, 3H, $C_A(2)OCH_3$), 3.10 (q, 2H, $J(CH_2, CH_3) = 7.4$ Hz, – $SO_2CH_2CH_3$), 1.27 (t, 3H, $J(CH_2, CH_3) = 7.4$ Hz, – $SO_2CH_2CH_3$).

^{13}C -NMR (75 MHz, $CDCl_3$, **2D-NMR: HSQC and HMBC** were used for more exact carbon assignment): δ 160.5 (s, $C_C(4)$), 156.5 (s, $C_D(2)$), 150.7 ($C_A(2)$), 145.9 (d, $C_D(6)$), 138.3 (s, $C_B(4)$), 133.4 and 130.6 ($2 \times$ s, $C_A(1)$ and $C_A(5)$), 128.9 (s, $C_C(3)$), $2 \times$ 123.7 (s and d, $C_C(1)$ and $C_C(6)$), 122.5 (d, $C_D(5)$), 120.4 (d, $C_A(4)$), 119.6 and 119.5 ($2 \times$ d, $C_C(5)$ and $C_D(3)$), 119.3 (d, $C_D(4)$), 119.0 (d, $C_C(2)$), 111.0 (d, $C_A(6)$), 109.6 and 109.5 ($2 \times$ d, $C_A(3)$ and $C_B(5)$), 56.2 (q, $C_A(2)OCH_3$), 50.8 (t, – $SO_2CH_2CH_3$), 7.6 (q, – $SO_2CH_2CH_3$).

IR ν (solid): 3354, 2939, 1597, 1566, 1509, 1428, 1302, 1260, 1142, 1123, 792, 735 cm^{-1} .

Anal. calcd for $C_{22}H_{21}N_5O_4S$ (451.50): C, 58.52; H, 4.69; N, 15.51. Found: C, 58.62; H, 4.62; N, 15.40.

4.4. *In vitro* VEGFR-2 kinase assay

A radiometric protein kinase assay (^{33}P anQinase® Activity Assay) was used for measuring the activity of VEGFR2 protein kinase. VEGFR2 tyrosine kinase was expressed in Sf9 insect cells as human recombinant GST-fusion protein. The kinase was purified by affinity chromatography using GSH-agarose. The purity of the kinase was checked by SDS-PAGE/silver staining and the identity of the kinase was verified by mass spectroscopy. IC_{50} values were measured by testing 10 concentrations from $1E-4$ to $1E-9$ M of each compound at 1 μ M ATP conc. The measurements were performed by ProQinase GmbH, Freiburg, Germany [25].

4.5. *In vitro* cytotoxic tumour cell lines evaluation

4.5.1. Cell culture

Huh7 and Mahlavu [80], human Hepatocellular Carcinoma (HCC) cells (ATCC) were maintained in Dulbecco's Modified Eagle's Medium (DMEM) (Invitrogen GIBCO), with 10% fetal bovine serum (FBS) (Invitrogen GIBCO), 2 mM L-glutamine, 0.1 mM nonessential amino acids, 100 units/mL penicillin and 100 g/mL streptomycin at 37 °C in a humidified incubator under 5% CO_2 . Both Huh7 and Mahlavu cells are tested their authentication by STR analysis.

4.5.2. NCI-60 Sulforhodamine B (SRB) cytotoxicity assay

SRB is anionic dye that can bind to proteins. The SRB assay measures the cellular protein content in order to determine cell density since cell proliferation is directly proportional to total protein synthesis [81].

Method: Huh7 (2000 cell/well) and Mahlavu (1000 cell/well) cells were inoculated into 96 well plates (150 μ l/well). 24 h later, molecules of interest and DMSO control were applied in concentrations 40 μ M–2.5 μ M in serial dilutions. After 72 h of treatment, cells were fixed by cold 10% (w/v) trichloroacetic acid (MERCK) for an hour. Then the wells were washed with ddH_2O and dried. 50 μ l of 0.4% SRB dye (Sigma–Aldrich) was applied to each well and incubated at RT for 10 min. Then wells were washed with 1% acetic acid and left for air-drying. SRB dye was solubilized in a 100 μ l 10 mM Tris-Base solution and the absorbance was measured at 515 nm. The experiment was performed in triplicates and the absorbance values were normalized to DMSO controls. Standard deviations were less than 10%.

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Appendix Supplementary data

Supplementary data associated with this article can be found in the online version, at <http://dx.doi.org/10.1016/j.ejmech.2015.08.012>. These data include physico-chemical characteristics, spectra and spectra graphical abstracts of prepared compounds depicted in Scheme 12 (also those not included in this printed version), MOL files and InChIKeys of the most important compounds described in this article.

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