ACS Medicinal Chemistry Letters

Letter

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ACS Med. Chem. Lett., Just Accepted Manuscript • DOI: 10.1021/acsmedchemlett.6b00149 • Publication Date (Web): 10 Jun 2016 Downloaded from http://pubs.acs.org on June 12, 2016

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CZ415, a highly selective mTOR inhibitor showing *in vivo* efficacy in a collagen induced arthritis (CIA) model

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KEYWORDS. mTOR, kinase inhibitor, inflammation, collagen induced arthritis model.

ABSTRACT: CZ415, a potent ATP-competitive mTOR inhibitor with unprecedented selectivity over any other kinase is described. In addition to a comprehensive characterization of its activities *in vitro*, *in vitro* ADME and *in vivo* pharmacokinetic data are reported. The suitability of this inhibitor for studying *in vivo* mTOR biology is demonstrated in a mechanistic mouse model monitoring mTOR proximal downstream phosphorylation signalling. Furthermore, the compound reported here is the first ATP-competitive mTOR inhibitor described to show efficacy in a semi-therapeutic collagen induced arthritis (CIA) mouse model.

The mammalian target of rapamycin (mTOR/FRAP1) is a serine/threonine protein kinase, which belongs to a family of phosphoinositide 3-kinase-related kinases (PIKK), which also includes ATR, ATM, DNAPK and SMG-1. FRAP1 is the catalytic subunit of two signaling complexes called mTORC1 and mTORC2, which play a central role in regulation of cell metabolism, proliferation, survival and migration.¹ It has been reported that mTOR is a key modulator in ageing, and dysregulated mTOR signaling can promote metabolic diseases, cancer and is often associated with inflammation.^{2,3} The broad interest in mTOR as a drug target is reflected by the growing number of mTOR inhibitors that have already been approved or are under clinical development.^{4–9}

To define further potential therapeutic uses and guide the development of safe and efficacious drugs it is crucial to fully understand the biological role of mTOR and the effects of its pharmacological inhibition.¹⁰ Ideally, target validation studies using small molecules should be conducted with multiple chemical probes to increase the confidence of target dependency of the observed phenotype and reduce the risk of misinterpretation of potential off-target effects.¹¹

Quality criteria such as high selectivity, well characterized mode of action and a favorable pharmacokinetic profile are a requirement to generate meaningful biological data and enable translation of *in vitro* findings to *in vivo* models.¹²

We report here a highly selective drug-like small molecule mTOR inhibitor from a novel chemical series,¹³ which shows efficacy in *in vivo* models.

The morpholine substituted pyrimidine core is a common structural motif that has been extensively explored for the development of ATP competitive mTOR inhibitors.¹⁴ In previous efforts on discovering mTOR inhibitors, we have also worked on this compound class and identified **CZ830** (1) and **CZ109** (2) as leads from two different chemical series (Figure 1).^{15,16} Both inhibitors showed moderate submicromolar affinity for mTOR and exhibited high selectivity of more than 100 fold over other lipid kinases from the same family, namely phosphatidylinositide 3-kinases (PI3K) $\alpha/\beta/\gamma/\delta$ and DNA-dependent protein kinase (DNAPK).

In order to discover more potent mTOR inhibitors, we combined both series, incorporating the sulfone moiety of compound 1 into the fused pyrimide core of .





In addition to what was previously reported on cyclic sulfones,¹⁷ we broadly explored methyl-substitution on the 5membered ring. Compared to other 5-membered sulfones, the introduction of a dimethyl-substituent was advantageous in terms of potency, selectivity or solubility and gave beneficial PK profile as compared to the 6-membered sulfopyrimidines. The morpholine substituent was found to be crucial for maintaining high potency and selectivity, whereas multiple groups were tolerated on the urea moiety. This is consistent with reported mTOR crystal structure and models on close analogues showing morpholine making an important hinge binding interaction.^{14,17,18} Exploring this chemical series, our efforts culminated in the discovery of **CZ415 (3)**, **Figure 1**. The synthesis of **3** and related cyclic sulfones has been previously reported¹³ and is described in the supporting information.

The potency and selectivity of **3** was assessed using Cellzome's chemoproteomic platform. Competition binding experiments combined with a proteomic readout are a powerful tool to determine the selectivity of small molecules against a large portion of the proteome in a single experiment.^{19,20} With this approach, the compound's binding affinities were measured for approximately 285 protein kinases, including the family of lipid and atypical kinases. The apparent dissociation constant pK_d^{app} for **3** against mTOR was found to be 8.2, while any other kinase identified in the experiments, including other PI3K family members – found to be frequent off-targets of other mTOR inhibitors – did not show any affinity within 1000 fold. The remarkable selectivity of **3** is depicted in **Figure 2** (detailed list of proteins is provided in Supporting Information).



Figure 2. MS-based kinase-binding profile of **3** across a set of protein kinases identified from mixed human cell-line lysates (285 kinases identified). The bars indicate pK_d^{app} values defined as the concentration of drug at which half-maximal competition of binding is observed, corrected by the influence of the immobilized ligand on the binding equilibrium using the Cheng-Prusoff equation.²¹ The dose dependent binding curve for mTOR is also depicted.

To verify that binding of 3 to mTOR results in inhibition of both mTORC1 and mTORC2 dependent signaling in the cellular context, 3 was tested in a cellular mechanistic assay. Inhibition of mTORC1 and mTORC2 protein complex activity was assessed by monitoring the phosphorylation of downstream targets, S6 Ribosomal Protein (S6RP, downstream of mTORC1) and Protein Kinase B (=Akt, mTORC2 substrate). Phosphorylation levels for both proteins, pS6RP (S240/244) and pAkt (S473) were assessed in dose response format from HEK293T cells after treatment with **3**. As depicted in Figure 3, inhibition of phosphorylation for both downstream targets resulted in 14.5 nM IC₅₀ for pS6RP and 14.8 nM for pAKT (Western Blot analysis and Caspase 3/7 activity assay are reported in the supporting information). Retaining high potency against mTOR in the cellular system proves functional inhibition of and the target indicates very

good cell permeability ($K_d^{app} = 6.9$ nM determined by chemoproteomic binding assay from cellular lysate, see above). Furthermore, interferon gamma (IFN γ) release in whole blood was used to test mTOR inhibition in primary cells. Treatment of whole blood with monoclonal antibodies binding to cluster of differentiation 3 and 28 (α CD3/ α CD28) mimics antigen receptor signalling and activates proliferation of Tlymphocytes. Co-stimulation with Interleukin 2 (IL-2) then induces interferon gamma gene (IFNG) expression and IFN γ is released.²² The immunosuppressive effect of **3** was measured by detecting secreted IFN γ after 18 hours in stimulated human whole blood and the resulting IC₅₀ was 226 nM (**Figure 3**).



Figure 3. Activity of **3** in cellular assays: A: Dose dependent inhibition of S6RP phosphorylation in HEK293T after 2 hours treatment of **3**, normalized to total S6RP levels. $IC_{50} = 14.5 \text{ nM} (95\% \text{ CI } 11.5 \text{ to } 18.3 \text{ nM}, n = 4)$. **B**: Dose dependent inhibition of Akt phosphorylation in HEK293T after 2 hours treatment of **3**, normalized to total Akt levels. $IC_{50} = 14.8 \text{ nM} (95\% \text{ CI } 10.4 \text{ to } 21.0 \text{ nM}, n = 4)$. **C**: Dose dependent inhibition of IFN γ release in stimulated human whole blood after 18 hours treatment of **3**. $IC_{50} = 226 \text{ nM} (95\% \text{ CI } 169 \text{ to } 303 \text{ nM}, n = 4)$.

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59 60 We have further investigated the properties of **3** in dedicated in vitro assays for early drug safety prediction. The cytochromes P450 (CYPs) are a family of enzymes that play a major role in drug metabolism and interaction with them is related to potential drug-drug interactions and adverse drug reactions. In human microsomes, no inhibition of the main P450 isoforms CYP1A, CYP2C8, CYP2C9, CYP2C19, CYP2D6 and CYP3A4 was observed within a 1000 fold window of mTOR affinity. Further tests for cytochrome CYP3A4 also showed no induction at 10 μ M and no time dependent inhibition (TDI) at a concentration as high as 50 μ M.

As a predictor for cardiotoxicity, the activity of **3** against the human cardiac ion channel hERG was assessed in a whole-cell patch-clamp assay in HEK293 cells resulting in an IC₅₀ of 48 μ M. We were further pleased that **3** showed no genotoxic potential. It was neither mutagenic in a bacterial mutation assay (Ames test) nor did it show genotoxicity in the mouse lymphoma assay (MLA), in either the presence or absence of rat-liver S9 mix. Given the data reported above for 3, there were no safety related liabilities predicted for this compound. Next to potency/selectivity assessment and early safety prediction, physicochemical properties are important for the successful development of drugs as well as for high quality probes, that can be used *in vivo*.^{23,24} Compound **3** has a moderate molecular weight (MW 460) and polar surface area (PSA 114), both in the desirable chemical space for bioavailable drugs. Despite a relatively high distribution coefficient (logD_{pH74} 4.31), **3** showed good kinetic solubility (CLND 149.5 µM) as well as solubility in biorelevant media, which is important for oral administration (FaSSIF 11.1 µg/mL, FeSSIF 60.1 µg/mL). The permeability measured in a Caco-2 assay, predictive for in vivo absorption of drugs across the gut wall, was moderate (Papp (A-B) 0.47 nm/s, Papp (B-A) 1.02 nm/s). A table summarizing physicochemical properties of 3 is provided in the supporting information.

The pharmacokinetic (DMPK) properties of a compound determine its ability to act over time on the molecular target. A low rate of drug metabolism is desirable for long acting compounds to maintain sufficient compound levels in plasma. As in vitro determination of the metabolic stability can help predicting in vivo PK, stability of 3 was assessed in microsomes and hepatocytes from different species. Low to moderate intrinsic clearance was measured, resulting in a calculated Clint <0.6/<1.1/1.5 µL/min/g liver in microsomes and Cl_{int} n.d./<0.8/≤0.5 µL/min/g liver in hepatocytes (rat/dog/human). In addition to the metabolic stability, measurement of plasma protein binding (PPB) enables estimation of the free fraction of compound in the bloodstream. PPB of 3 in rat, mouse and human plasma was determined to be 69/89/81% (r/m/h), indicating a sufficient free fraction of the compound being available for binding to the target.

For full characterization of **3** and to enable improved dose predictions, the pharmacokinetic (PK) profile was assessed in rat. PK and oral bioavailability were determined after of 1 mg/kg intravenous (iv) bolus and 3 mg/kg oral (po) administration (**Figure 4**). The observed plasma clearance, corresponding to 45% liver blood flow, suggested that sufficient levels of free compound were circulating in the animal over time. The oral uptake was rapid with a T_{max} of 0.5 h and bioavailability F = 44% indicated very good absorption from the gut. The plasma concentration peaked at 0.43 μ M which was above IC₅₀ determined in cellular assays for **3**.



Figure 4. Time dependent plasma concentration of 3 after intravenous bolus (iv, circle) and oral solution (po, square) administration to rats. Rats were dosed at 1 mg/kg (iv, n = 3) or 3 mg/kg (po, n = 3). Vehicle: 5% DMSO / 95% (10% Kleptose).

To demonstrate the efficacy of 3 in vivo, modulation of mTOR proximal downstream signaling has been monitored in a mechanistic mouse model. In vivo treatment with anti-CD3 antibody specifically stimulates T cells upon binding to the CD3 receptor and subsequent downstream activation of the PI3K/mTOR signalling pathway.²⁵ To determine the effects of **3** on its pharmacological target, dose-dependent changes in phosphorylation levels of S6 Ribosomal Protein and Akt both downstream targets of mTOR - were assessed. The inhibitor was administered orally at 1, 3 and 10 mg/kg to mice 1 h before anti-CD3 stimulus. 15 min after stimulation, spleens were dissected and analyzed for pS6RP and pAKT levels. A dose related significant inhibition of phosphorylation of both S6RP and Akt was observed after compound administration (Figure 5, A and C). In particular, 1 and 3 mg/kg 3 could fully inhibit S6RP phosphorylation induced by anti-CD3 stimulation and 10 mg/kg additionally decreased the constitutive phosphorylation levels as measured in the control group. While phosphorylation of Akt was not increased upon anti-CD3 stimulation (Figure 5, C), 3 - already at the lowest dose of 1mg/kg - significantly reduced pAkt constitutive levels, as measured in the control group. The PK/PD relationship was determined following measurement of compound concentration in blood and the exposure-response correlation is depicted in Figure 5, B and D. An EC₅₀ of 0.22 µM was measured for pS6RP and 0.055 μ M for pAkt, which correlate well with the nanomolar potency of 3 in the cell based assays. These results showed 3 to be a potent mTOR inhibitor which retained its activity in vivo

In vivo efficacy of **3** was then assessed in an inflammatory context. In the collagen induced arthritis (CIA) model, mice reliably develop polyarthritis when immunized against bovine type II collagen and this model can be utilized to monitor antiinflammatory effects of the tested compounds. It has been shown, that treatment with Rapalogs can inhibit the clinical severity of arthritis in mice,^{26–28} but the effect of ATP-competitive mTOR inhibitors has not previously been reported. A so called semi-established study design was utilized for testing **3** at 10 mg/kg twice daily. Mice were challenged with two injections of collagen, the first at day 0 and a boost injection at day 21.



Figure 5. Compound **3** in anti-CD3 mouse model. **A:** pS6RP levels (normalized to total S6RP) measured in spleens of compound treated as compared to disease vehicle group (p<0.01 for 1 mg/kg of **3**, p<0.001 for 3 and 10 mg/kg of **3**-; one outlier removed in normal control and disease vehicle group). **B:** Exposure response fit: pS6RP levels at terminal exposure. EC₅₀ 0.22 μ M (95% CI 0.15 to 0.32 μ M). **C:** pAkt levels (normalized to total Akt) measured in spleens of compound treated as compared to disease vehicle group (p<0.001 for 1, 3 and 10 mg/kg of **3**). **D:** Exposure response fit: pAkt levels at terminal exposure. EC₅₀ 0.055 μ M (95% CI 0.048 to 0.065 μ M). **Legend:** White – normal controls (n = 4). Black – anti-CD3 / vehicle 5% DMSO / 95% (10% Kleptose) (n = 6). Grey – anti-CD3 / **3** (1, 3 or 10 mg/kg, of **3** PO) (n = 6).

Administration of **3** was started at day 22, immediately before the onset of the disease and continued until day 34 at termination of the study. The development of arthritis, which reached 100% disease incidence in the vehicle group, was determined by evaluation of clinical arthritis scores (from 0 to 5) that were measured by inspection of the paws for erythema and swelling. The daily clinical arthritis scores of the treated group differed from disease control group over time with significant reductions (p<0.05) at day 33-34 (**Figure 6**, **A**). Clinical arthritis scores expressed as area under the curve (AUC) were reduced in the group treated with **3** by 59% as compared to the disease vehicle group (data not shown). No significant difference in body weight loss was observed compared to vehicle-treated, disease-induced mice at end of the study (**Figure 6**, **B**). Measurement of compound concentrations in blood, determined on day 31, indicated that up to 8 hours after dosing the levels of circulating **3** were above EC_{50} as determined both *in vitro* in the whole blood assay and *in vivo* in the PK/PD mouse model (**Figure 6**, **C**). Unlike many anti-inflammatory molecules that require prophylactic dosing in the CIA, this data demonstrates that **3** is active in a situation more akin to human disease.



Figure 6. Compound **3** in a mouse CIA model. **A:** Clinical arthritis score – all paws (Scored 0-5). Normal controls (white squares): significance ($p \le 0.05$) from disease vehicle group (black squares) observed at day 27 through day 34; Compound **3** at 10 mg/kg, bid (grey dots): significant reduction as compared to disease vehicle group at day 33-34 ($p \le 0.05$). **B:** Body weight over time. Significant difference of normal controls at day 26 through day 34 as compared to disease vehicle group, and of 10 mg/kg **3** at day 26 ($p \le 0.05$) to disease vehicle). **C:** Drug levels of **3** in blood on study day 31. Exposure: AUC 14 μ M×h, C_{max} 1.7 μ M. Legend: Normal controls (n = 4), Disease vehicle (5%DMSO/95% (10% kleptose), n = 10), Compound **3** (10 mg/kg, PO, n = 10).

We have presented here a highly potent mTOR inhibitor **CZ415 (3)** with exquisite selectivity over other lipid and protein kinases. It has a defined mechanism of action, which was demonstrated in mechanistic and phenotypic cell based assays, including whole blood assays. Beside favorable physicochemical properties, early *in vitro* safety assessment did not exhibit any liabilities and pharmacokinetic properties of moderate

clearance and good oral bioavailability showed suitability of **3** for progression to *in vivo* studies. In an anti-CD3 mouse model **3** efficiently inhibited mTOR downstream signaling and in a CIA mouse model showed significant anti-inflammatory effects. With its extraordinary selectivity, drug-like properties and proven efficacy *in vivo*, **3** represents an ideal molecule for

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the pharmacological investigation of mTOR pathophysiological role *in vivo*.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge on the ACS Publications website.

NMR and MS spectra, detailed selectivity profiling data, protocols for cellular assays, DMPK and supplemental information on animal studies (PDF)

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Author Contributions

G. Bergamini and A.J.R. wrote the manuscript, T.L. and M.S. designed and participated to the execution of the work, A.D.C., R.L, K.E. and J.F. participated to the chemical design and synthesis, A.L.N. designed and performed the cellular assays, A.L. supervised and analyzed the DMPK studies, G. Bennett supervised and analyzed the selectivity profiling work, A.P.W. and O.R. contributed to the design of the studies, G. Bergamini designed and supervised the work reported. All authors have given approval to the final version of the manuscript.

Notes

The human biological samples were sourced ethically and their research use was in accord with the terms of the informed consents. All animal studies were ethically reviewed and carried out in accordance with European legislation and guidelines for animal welfare.

ACKNOWLEDGMENT

We thank Cellzome chemistry, biology, pharmacology, biochemistry, informatics and mass spectrometry departments, the leadership team and the whole mTOR program team for support and advice.

ABBREVIATIONS

Akt, Protein Kinase B; ATM, Ataxia telangiectasia mutated kinase; ATR, ataxia telangiectasia and Rad3-related protein; α CD3/ α CD28, monoclonal antibodies binding to cluster of differentiation 3/28; CIA, collagen induced arthritis; DNAPK, DNAdependent protein kinase; IFN γ , interferon gamma; IFNG, interferon gamma gene; IL-2, Interleukin 2; K_d^{app}, apparent dissociation constant; mTOR, mammalian target of rapamycin; PI3K, phosphatidylinositide 3-kinases; PIKK, family of phosphoinositide 3-kinase-related kinases; PK/PD, pharmacokinetic/pharmacodynamic; SMG-1, nonsense mediated mRNA decay associated PI3K related kinase; S6RP, S6 Ribosomal Protein.

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