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Asymmetric Synthesis of 2-Phenylpyrrolidines for SAR Study of Thienopyrimidine-based EGFR Kinase Inhibitors

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I hereby declare that the work done in this thesis is independent and in accordance with the exam regulations of the Norwegian University of Science and Technology.

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Acknowledgements

Abstract

The aim of this master's thesis was to carry out a preliminary structure-activity relationship (SAR) study of thieno[2,3-*d*]pyrimidines with racemic and enantioenriched 2-phenylpyrrolidine as a substituent on C-4. One objective was to investigate if secondary amines are essential substituents at C-4 by preparing thieno[2,3-*d*]pyrimidines with a tertiary heterocyclic amine. This study has also included compounds with pyrrolidine as a C-4 substituent, to study the contribution of the phenyl moiety on the amine on the EGFR-TK activity. All amines and thieno[2,3-*d*]pyrimidines have been synthesized as part of the project, and a second objective of this master's thesis was to investigate viable synthetic routes to obtain enantioenriched 2-phenylpyrrolidine with high chemical and enantiopurity.

Enantioenriched 2-phenylpyrrolidines were prepared in three synthetic steps, starting with Boc protection of pyrrolidine. Palladium-catalyzed α-arylation gave enantioenriched, Boc-protected intermediates, followed by Boc deprotection to achieve enantioenriched amines with 81-96 % ee. Enantionenriched amine was also obtained from racemic 2-phenyl-pyrrolidine via chiral resolution, employing a chiral acid. Racemic 2-phenylpyrrolidine was prepared in two synthetic steps, starting with a two-step, one-pot synthesis from pyrrolidin-2-one, followed by reduction of a conjugated double bond. Access to the target thieno[2,3-*d*]pyrimidines required further two to three synthetic steps, including nucleophilic aromatic substitution, Suzuki cross-coupling, and reduction of aldehyde to primary alcohol. The synthetic routes are illustrated in Scheme 0.0.1.

Scheme 0.0.1: Synthetic routes for the preparation of target compounds **6a-b** and **8a-b**.

This work has resulted in 14 new structures, of which two compounds were found to possess good EGFR-TK activity and may open up new areas of research. The phenyl group on the C-4 amine was found to have a significant impact on the EGFR-TK activity, as removal of this moiety proved detrimental. The chirality of the phenyl group was also found to influence the EGFR-TK activity, with significantly improved activity of the *(R)*-enantiomer relative to the racemate, and poor activity of the *(S)*-enantiomer. The findings of this thesis have demonstrated that it is possible to achieve high activity towards EGFR with a tertiary heterocyclic amine at C-4.

Sammendrag

Hensikten med denne masteroppgaven har vært å utføre en innledende struktur-aktivitet relasjonsstudie av tieno[2,3-*d*]pyrimidiner med rasemisk og enantioanriket 2-fenylpyrrolidine som substituent på C-4. Målet var å undersøke om det er nødvendig med et sekundært amin på C-4 for å oppnå EGFR-TK aktivitet, eller om det er mulig å oppnå god aktivitet med tertiære heterosykliske aminer. Denne studien har også inkludert strukturer med pyrrolidin som C-4 substituent, for å studere bidraget av fenylgruppen på aminet til EGFR-TK aktiviteten. Alle aminer og tieno[2,3-*d*]pyrimidiner har blitt syntetisert som en del av prosjektet, og et annet mål med denne masteroppgaven har vært å undersøke mulige synteseveier for å oppnå enantioanriket 2-fenylpyrrolidin med høy kjemisk og enantiomer renhet.

Enantioanriket 2-fenylpyrrolidin ble syntetisert i tre steg, med utgangspunkt i en Boc beskyttelse av pyrrolidin. Palladium-katalysert α-arylering gav Boc-beskyttede mellomprodukter, som ved avbeskyttelse gav enantioanriket amin. En alternativ syntesevei til enantioanriket amin via kiral oppløsning av rasemisk amin ble også utforsket, ved bruk av en kiral syre. Rasemisk 2-fenylpyrrolidin ble syntetisert i to steg, med utangspunkt i en to-steg, one-pot syntese fra pyrrolidin-2-on, etterfulgt av reduksjon av et konjugert dobbelbånd. Målstrukturene ble videre syntetisert gjennom to til tre syntetiske steg, herunder nukleofil aromatisk substitusjon, Suzuki kobling, og reduksjon av aldehyd til primær alkohol. Synteseveiene i denne masteroppgaven er illustrert i Skjema 0.0.2
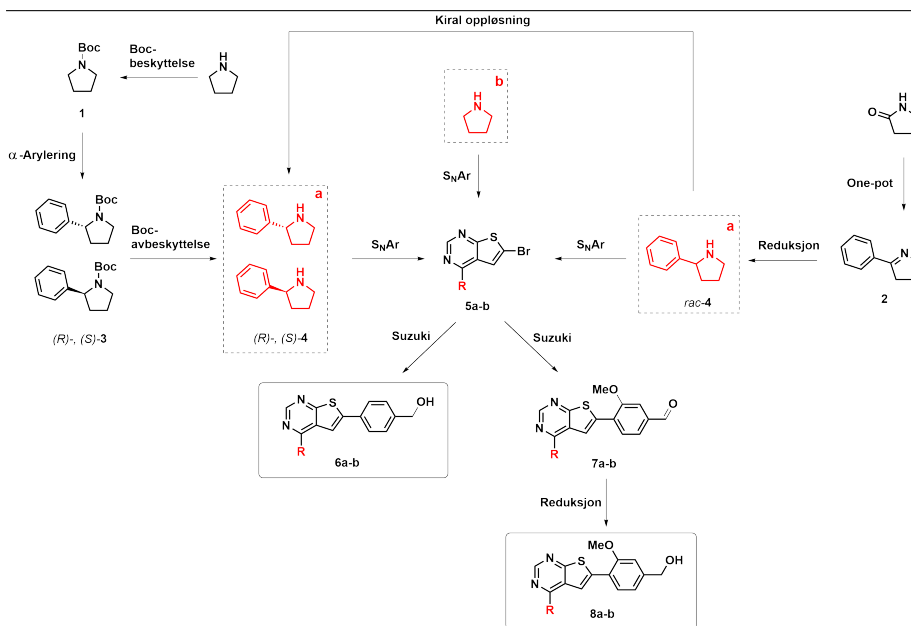
Scheme 0.0.2: Synteseveier til målstrukturene **6a-b**, og **8a-b**.

Denne oppgaven har resultert i 14 nye strukturer, hvorav to ble funnet å inneha god EGFR-TK-aktivitet og med potensial til å åpne nye forskningsområder. Fenylgruppen på C-4 aminet har vist seg å ha en signifikant påvirkning på EGFR-TK-aktiviteten, ettersom fjerning av denne gruppen hadde en betydelig negativ innvirkning. Også kiraliteten på fenylgruppen har vist seg å ha en påvikning på EGFR-TK-aktiviteten, med en markant økning i aktivitet for *(R)*-enantiomeren relativt til rasematet, og dårlig aktivitet for *(S)*-enantiomeren. Resultatene fra denne masteroppgaven har vist at det er mulig å oppnå god EGFR-TK aktivitet med et tertiært heterosyklisk amin på C-4.

Symbols and Abbreviations

∂	Chemical shift [ppm]
^{13}C NMR	Carbon Nuclear Magnetic Resonance
^{1}H NMR	Proton Nuclear Magnetic Resonance
APCI	Atmospheric Pressure Chemical Ionization
Ar	Aryl
ATP	Adenosine Triphosphate
bp	Boiling Point [°C
br	Broad signal
Conv	Conversion
COSY	Correlation Spectroscopy
CSF1R	Colony Stimulating Factor 1 Receptor
CSP	Chiral Stationary Phase
d	Doublet
Decomp	Decomponation
ee	Enantiomeric Excess
EGFR	Epidermal Growth Factor Receptor
eq	Equivalent(-s)
ESI	Electrospray Ionization
FGFR	Fibroblast Growth Factor Receptor
FRET	Fluorescence Resonance Energy Transfer
h	Hours

HMBC	Heteronuclear Multiple Bond Correlation
HPLC	High Performance Liquid Chromatography
HRMS	High Resolution Mass Spectrometry
HSQC	Heteronuclear Single Bond Correlation
IC_{50}	Half maximal inhibitory concentration
IR	Infrared
IRED	Imine Reductase
J	Coupling constant [Hz]
m	Multiplet
m/z	Mass per charge
min	Minutes
mp	Melting Point [°C
MS	Mass Spectroscopy
NSCLC	Non-Small Cell Lung Cancer
q	Quartet
R_f	Retention Factor
rt	Room Temperature
s	Singlet
t	Triplet
t_R	Retention Time [min]
TK	Tyrosine Kinase
TKI	Tyrosine Kinase Inhibitor
TKR	Tyrosine Kinase Receptor
TLC	Thin Layer Chromatography
UV	Ultraviolet
VGEFR	Vascular Endolethial Growth Factor Receptor

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Numbered Compounds

1 | Background

Cancer is one of the leading causes of death globally, as an affliction with high prevalence and mortality rates.[1] Three decades of research into the mechanisms of cancer pathogens have lead to the development of molecular targeted cancer therapy: the use of drugs or other substances to target specific biomacromolecules involved in the growth and spread of cancer.[2,3] Despite major advances in cancer therapy and drug development, targeted cancer therapy is inevitably challenged by drug resistance. As drug resistance is almost invariably succeeded by disease progression, the development of new anti-cancer agents remains an important objective in targeted cancer therapy.[4]

The growing arsenal of targeted anti-cancer agents can be categorized according to their respective effect on one or more hallmarks of cancer, as illustrated in Figure 1.1.[5] Sustained proliferative signalling is one such trait, allowing malignant cells to maintain constant growth by avoiding growth suppression.[6] Inhibition of sustained proliferative signalling may be done by use of tyrosine kinase inhibitors (TKIs), as for instance by targeting the epidermal growth factor receptor (EGFR). This treatment is primarily associated with lung-, breast-, and pancreatic cancer, with commercially available drugs like Erlotinib and Gefitinib.[7,8]
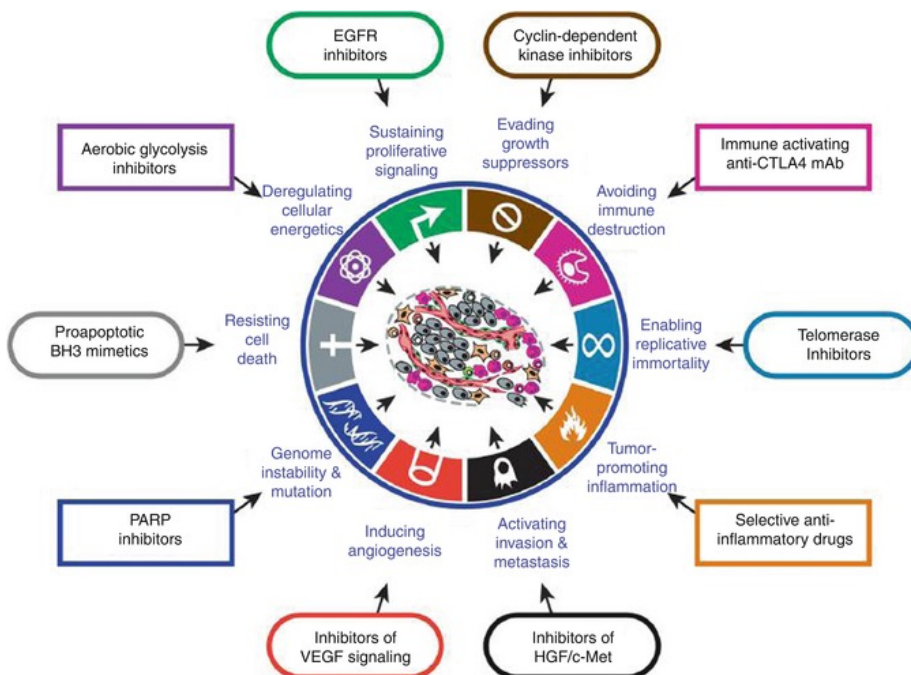
Figure 1.1: The hallmarks of cancer form the basis of contemporary drug development and targeted anti-cancer treatment.[5]

The Hoff/Sundby research group has focused on developing thieno-, pyrrolo-, and furo-pyrimidines as new EGFR-TKIs for use as anti-cancer agents. This work has resulted in a number of structure-activity studies, identifying several compounds with activity in the nanomolar range.[9–11] A biological assay of the purine TIA02-176 (Aarhus, Sundby, Hoff, work in progress), illustrated in Figure 1.2, found this compound to possess excellent EGFR-TK activity. Previous structure-reactivity studies on thieno[2,3-d]pyrimidines by Bugge, et al.,[11] have found that a secondary amine at C-4 is essential for the EGFR-TK activity, as illustrated in Figure 1.2. Preliminary results in the research group indicate that this is not necessarily the case.
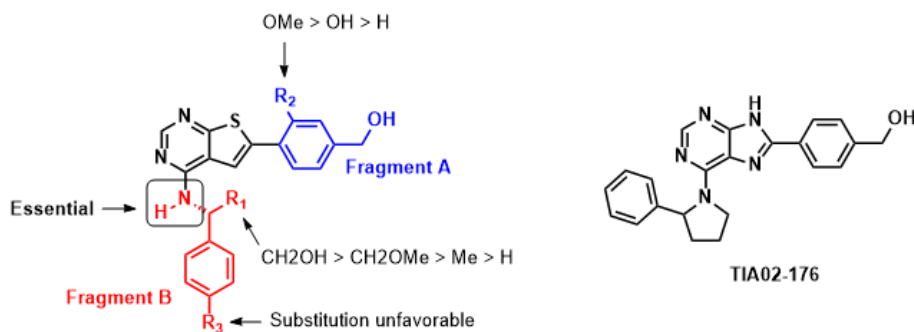
Figure 1.2: Structure-activity relationship model for thieno[2,3-*d*]pyrimidines.

The aim of this master's thesis was to perform a preliminary structure-activity relationship (SAR) study of thieno[2,3-*d*]pyrimidines with racemic and enantioenriched 2-phenyl-pyrrolidine as a substituent on C-4. An objective was investigate if secondary amines are essential substituents at C-4 by preparing thieno[2,3-*d*]pyrimidines with a tertiary heterocyclic amine. Compounds with pyrrolidine as a C-4 substituent were also investigated, to study the contribution of the phenyl moiety on the amine on the EGFR-TK activity. All amines and thieno[2,3-*d*]pyrimidines were synthesized as part of this project, and a secondary objective of this master's thesis was to investigate viable synthetic routes to obtain enantioenriched 2-phenylpyrrolidine with high chemical and enantiopurity.

2 | Introduction and Theory

2.1 EGFR

The epidermal growth factor receptor (EGFR, also known as ErbB-1/HER1) is a trans-membral protein belonging to the ErbB class of receptor tyrosine kinases (TKR).[12] The EGFR consists of an intracellular tyrosine kinase domain, a transmembrane lipo-philic domain, and an extracellular ligand-binding domain which will bind to receptor-specific ligands.[13] The EGFR, along with other tyrosine kinase receptors, including vascular endothelial growth factor receptors (VGEFR) and fibroblast growth factor receptors (FGFR), play a key role in regulating cellular processes such as cellular growth, differentiation and metabolism.[14]

Activation of the EGFR, illustrated in Figure 2.1, occur by binding of receptor-specific polypeptide ligands (growth factors) to the extracellular domain of the receptors. This induces receptor dimerization, forming homo- or heterodimeric receptor complexes, which leads to activation of the tyrosine kinase domain by phosphorylation of specific tyrosine residues.[12,14,15] These residues serve as docking sites for proteins containing specific domains, including the phosphotyrosine binding domain.[12] The introduction of these proteins leads to activation of the intracellular signalling pathways.
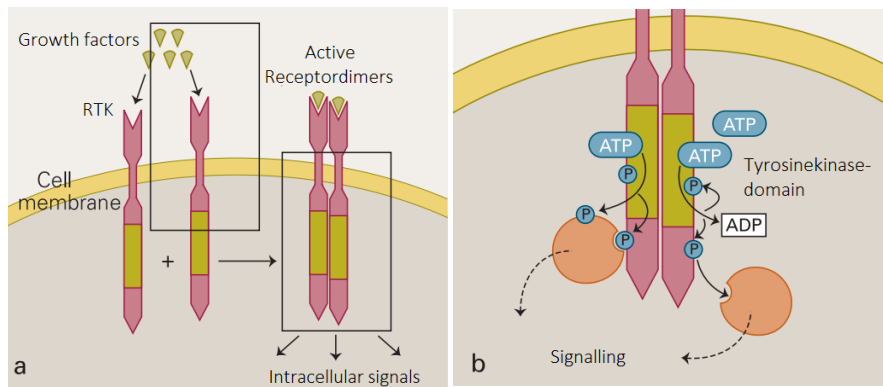
Figure 2.1: Activation of intracellular signalling of receptor tyrosine kinase (TKR). Binding of growth factors (a) induces receptor dimerization, which leads to activation of the tyrosine kinase domain by autophosphorylation (b).[16]

2.2   EGFR-TKI

Overexpression of the EGFR is associated with a variety of malignancies, including lung, breast and pancreatic cancer.[17] Aberrant activity of these receptors has been shown to play a key role in the development of tumor cells, making these receptors attractive targets in targeted cancer therapy.[18]

Several approaches have been developed for targeting, and inhibiting the EGFR signalling pathways, including the use of monoclonal anti-EGFR antibodies and recombinant proteins.[18] The use of tyrosine kinase inhibitors (TKI) as small, adenin triphosphate (ATP) analogues, has proven effective against human EGFR, and is now a standard clinical treatment for patients with advanced EGFR mutant non-small-cell lung cancer.[19] These molecules work by binding to the ATP binding sites on the tyrosine kinase domain, thus preventing autophosphorylation and activation of further downstream signalling pathways (as illustrated in Figure 2.2).[18]
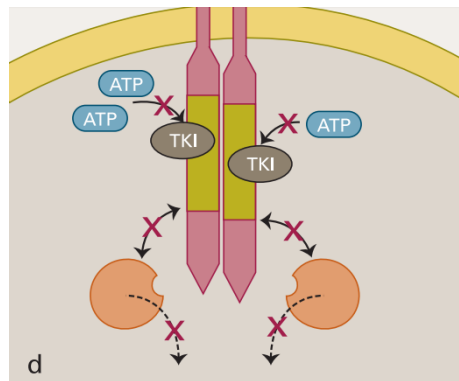
Figure 2.2: Blocking of the intracellular signalling of receptor tyrosine kinase (TKR) by use of a tyrosine kinase inhibitor (TKI).[16]

Several EGFR-TKIs are approved for therapeutic use worldwide, including Erlotinib, Gefitinib, and Afatinib for the treatment of non-small cell lung cancer (NSCLC) harbouring EGFR mutations (Figure 2.3).[20] Erlotinib and Gefitinib are first-generation EGFR-TKIs and act to block autophosphorylation and activation of downstream signalling. Unfortunately, most patients with EGFR-mutant NSCLC treated with EGFR-TKIs develop resistance within 9-14 months, and second-generation EGFR-TKIs, such as Afatinib, has therefore been developed to overcome the acquired resistance of the first-generation EGFR-TKIs.[21] Afatinib is an irreversible EGFR-TKI, unlike the reversible Erlotinib and Gefitinib, and acts to block the enzymatic activity of active ErbB receptor family members by forming irreversible covalent bonds. These bonds can inhibit the kinase activity until new receptors are formed, which may give longer action time of Afatinib compared to reversible EGFR-TKIs.[20] New generations of EGFR-TKIs continue to appear with the development of third-generation EGFR-TKIs (Osimertinib, Olmutinib)[22] to combat mutations altering the EGFR affinity for ATP (e.g. T790M mutation), and fourth-generation inhibitors (EAI045)[23] to battle mutations preventing irreversible EGFR-TKIs from forming covalent bonds (e.g. C797S mutation).[20]
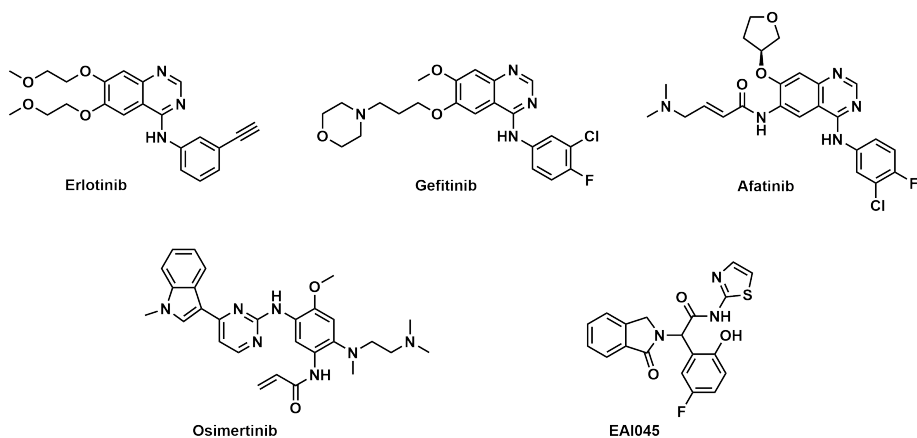
Figure 2.3: Structures of some different generation EGFR-TKIs: Erlotinib (first generation),
Gefitinib (first generation), Afatinib (second generation), Osimertinib (third gener-
ation), EAI045[24] (fourth generation).

2.3 Bioactive thienopyrimidines

Thienopyrimidines are heterocycles with a large diversity in bioactivity, and derivatives
of these compounds have been found to possess antimicrobial-,[25] antitubercular-,[26]
antiviral-,[27] anti-inflammatory-,[28] and anti-tumour[29] activity.

The use of thienopyrimidines as EGFR kinase inhibitors can be traced back to
the 1990s, as scaffold hopping of quinazolines and pyrrolopyrimidines found
thienopyrimidines to be potent EGFR-TKIs.[30,31] The recent years have seen a
growing interest in thienopyrimidines, as a number of potent thienopyrimidine-based
EGFR inhibitors have been published (Figure 2.4).[11,32,33] However, to the best of
the author's knowledge, as of June 2018, the only thienopyrimidine-based EGFR-
TKI to be approved for therapeutic use is Olmutinib (approved in South Korea in
2016[34]).[24,35]
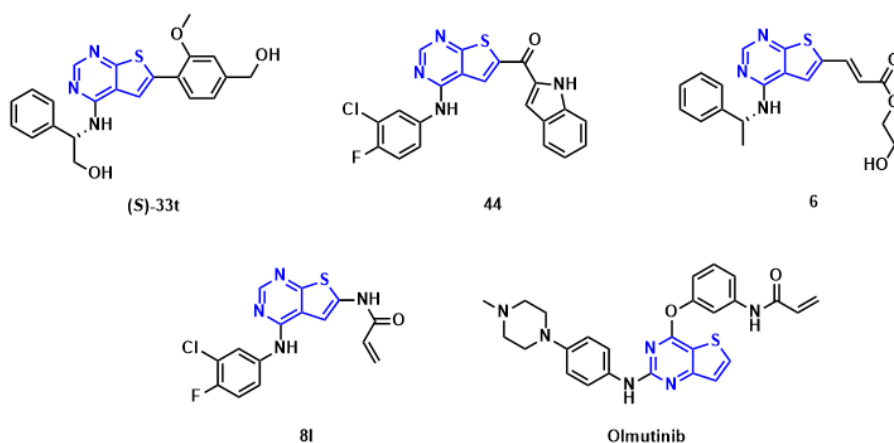
Figure 2.4: Structures of potent thienopyrimidine-based EGFR inhibitors, (S)-**33t**,[11] **44**,[32] **6**,[33] **8l**,[36] Olmutinib.[37]

2.4 Bioactive pyrrolidines

N-Heterocycles are important structural units in many biologically active compounds, and are attractive scaffolds in pharmaceutical development.[38] Substituted pyrrolidines are found in a large array of natural products, pharmaceauticals, and bioactive compounds (Figure 2.5). Some examples are Nicotine; alkaloid found in the nightshade family of plants (Solanaceae),[39] Kainic acid; naturally occuring neurotoxin isolated from seaweed (Digenea simplex),[40] Vildagliptin; an oral anti-diabetic drug,[41] Anisomycin; antibiotic inhibiting protein synthesis,[42] and Kaitocephalin; a glutamate receptor antagonist.[43]
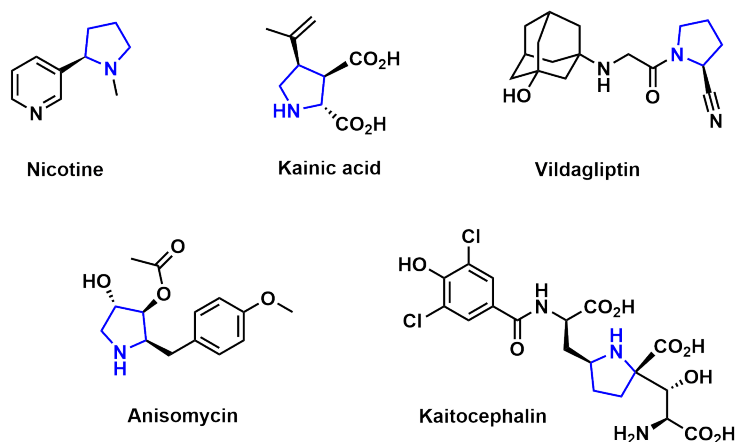
Figure 2.5: Structures of some natural products, pharmaceuticals, and bioactive compounds with substituted pyrrolidine as a structural unit.

2.5   Biological assay

Biological tests of the target compounds in this project have been done by Invitrogen, using the Z'-LYTE$^{©}$ biochemical assay.   This assay employs a fluorescence-based, coupled-enzyme format in which the peptide substrate is labeled with two fluorophores, one at each end.[44] As illustrated in Figure 2.6, this allows for in-tramolecular fluorescence resonance energy transfer (FRET) from the donor to the acceptor fluorophore, thus referred to as a FRET pair.[45] In this process, the excited donor fluorophore (e.g. Coumarin) transfers its excitation energy non-radiatively to the acceptor fluorophore (e.g. Fluorescein), thus causing the acceptor fluorophore to emit its characteristic fluorescence.[45]

During testing, the kinase transfers the gamma-phosphate of ATP to a single ty-rosine residue in a synthetic FRET-peptide.[44] In the case of a good kinase inhibitor the phosphate transfer is counteracted, yielding more non-phosphorylated peptide, as illustrated in Figure 2.6. In the following step, a site-specific protease recognizes and cleaves only the non-phosphorylated peptides, as phosphorylation suppresses cleavage by the Development Reagent.[44] The cleavage disrupts the FRET of the non-phosphorylated peptides, while FRET of the phosphorylated peptides is maintained.[44] This yields highly fluorescent cleaved peptide fragments and uncleaved FRET-peptides, both contributing to the fluorescence signals, as illustrated in Figure 2.6. Detection of the fluorescence signals gives the emission ratio as the ratio of donor emission to acceptor emission.[44] High emission ratio indicates larger amounts of non-phosphorylated FRET-peptide, hence better kinase inhibition.
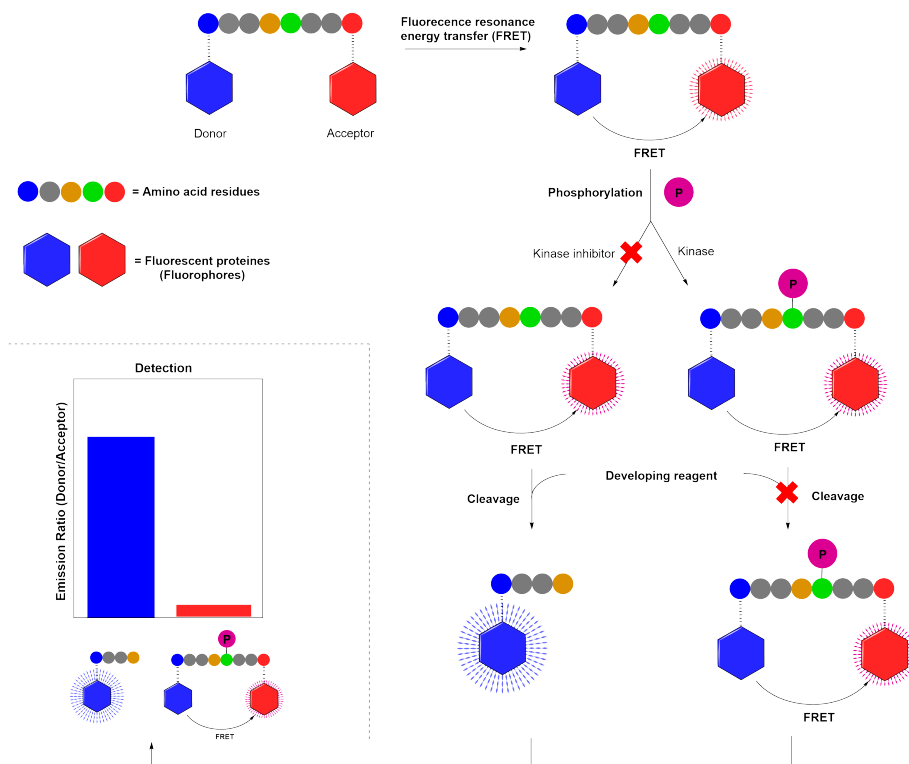
Figure 2.6: Biochemical assay of kinase inhibitors by use of the Z'-LYTE© biochemical assay.

2.6 Synthesis and choice of strategy

2.6.1 Asymmetric synthesis of 2-substituted pyrrolidines

The term *asymmetric synthesis* is used of "a chemical reaction (or reaction sequence) in which one or more new elements of chirality are formed in a substrate molecule and which produces the stereoisomeric (enantiomeric or diastereoisomeric) products in unequal amounts." (IUPAC def.[46]). As pyrrolidines are important precursors to biologically active compounds, asymmetric synthesis of these compounds is well documented, with possible synthetic routes including cycloaddition,[47,48] intramolecular addition of nucleophiles into imines,[49,50] and the reaction of organic azides with alkyldifluoroborane intermediates.[51]

The 2-substituted pyrrolidines *(R)*- and *(S)*-**4** have previously been prepared by asymmetric synthesis, employing a variety of synthetic routes.[52–55] Scheme 2.6.1 illustrates a selection of possible synthetic routes to *(R)*-**4**, of which the α-arylation and chiral resolution routes are employed in this master's thesis.
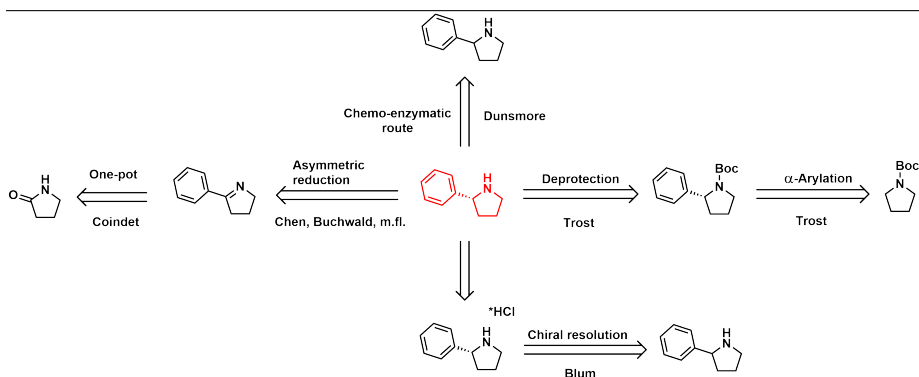
Scheme 2.6.1: Retrosynthetic analysis of *(R)*-2-phenylpyrrolidine (*(R)*-**4**). References: Trost *et al.*,[52] Dunsmore *et al.*,[53] Blum *et al.*,[54] Chen *et al.*,[55] Buchwald *et al.*,[56] Coindet *et al.*.[57]

Asymmetric hydrogenation

Asymmetric hydrogenation, or asymmetric reduction, is widely regarded as a highly efficient method for obtaining optically active amines.[55,58,59] The development of the technique originated in 1968,[60,61] and was pioneered by William S. Knowles[62] and Ryoji Noyori,[63] for which they were both rewarded with the Nobel Price in Chemistry 2001.

Enantioselective asymmetric hydrogenation allows for selective generation of chiral centres by use of specially designed catalysts, thus providing the possibility of synthesizing optically active 2-substituted pyrrolidines from cyclic imines. The hydrogenation of compound **2** to form *(R)*- or *(S)*-**3** or -**4** has already been investigated, and some known catalysts for these reactions includes Ru-,[55] Ti-,[56,64–66] and Ir-based catalysts.[67–69] Amine compounds are known to have a deleterious effect on the performance of both heterogeneous[70] and homogeneous[71] hydrogenation systems, and in-situ protection of the resulting 2-substituted pyrrolidine compound has therefore been found to be favourable to reduce the potential for product inhibition.[55,72,73] The synthesis of *(R)*-**3** from **2** was planned as a synthetic route in this master project.

Several mechanistic studies on asymmetric hydrogenation of cyclic amines have been conducted in an attempt to understand how the substrate coordinates to the metal center of the catalysts, and how hydrogen is transferred to the C=N.[56,65,74] A proposed catalytic cycle by Willoughby & Buchwald, using a chiral titanocene catalyst,[75] suggests a mechanism in which the imine reacts with a titanium hydride in a fast 1,2-insertion step, forming a titanium amide intermediate. This is followed by a slow reaction of the amide complex with hydrogen to produce the amine, which also regenerates the titanium hydride.[56] Later studies have found the formation of a dihydrido complex containing the imine, coordinated via the nitrogen atom, to play a key role in this catalytic cycle.[74] The proposed catalytic cycle for asymmetric

hydrogenation of cyclic amines is illustrated in Scheme 2.6.2. The mechanisms for reactions involving rhodium-, irridium-, and ruthenium-complexes are still under investigation.[55,76]
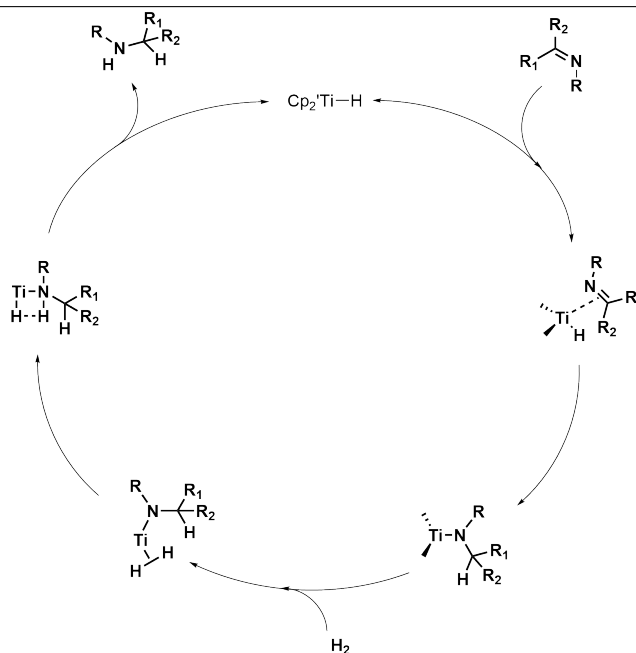
Scheme 2.6.2: Proposed catalytic cycle for the hydrogenation of cyclic amines, using a chiral titanocene catalyst.[56,65,74]

Despite the simplicity, good atom economy, and environmental friendliness of these reactions, enantioselective asymmetric hydrogenation reactions possess some practical limitations.[77] The specialized metal catalysts are often hard to obtain due to scarcity of the metals, patent-protected syntheses, and the high cost associated with them. The chosen ruthenium-based catalyst for the synthesis of *(R)*-**3**, [RuBArF(*p*-cymene)(*R,R*)-MsDPEN],[55] could be prepared from the precursor [RuCl(*p*-cymene)((*R,R*)-MSDPEN], by undergoing a simple ligand-exchange, as illustrated in Scheme 2.6.3.[78]
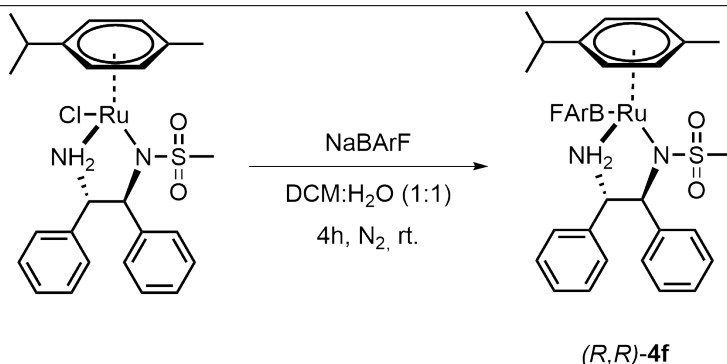
Scheme 2.6.3: Preparation of [RuBArF(*p*-cymene)(*R,R*)-MsDPEN] ((*R,R*)-**4f**)[55] from [RuCl(*p*-cymene)((*R,R*)-MSDPEN].[78]

The following synthesis of *(R)*-**3**, employing catalyst *(R,R)*-**4f**, utilizes pressurized H_2 at 10-80 atm. and 40 °C, as presented in Scheme 2.6.4.[55] Due to technical restrictions, H_2 at this pressure proved difficult to obtain. It may be possible to run the synthesis at lower hydrogen pressure, as the enantioselectivity is reportedly insensitive to changes in pressure.[55,56] However, along with difficulties obtaining the precursor for catalyst *R,R*-**4f** and time constraints, this reaction was ultimately not attempted.
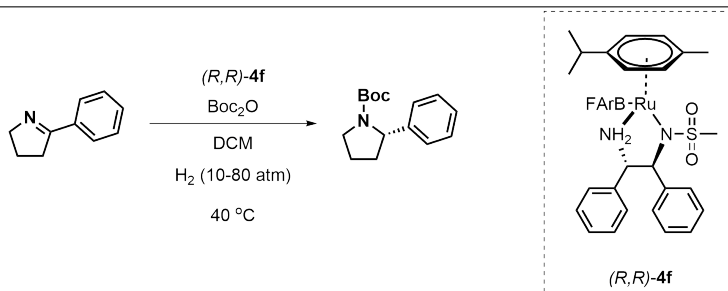
Scheme 2.6.4: Enantioselective asymmetric hydrogenation of **2** to *(R)*-**3**, using *(R,R)*-**4f** as the catalyst.[55]

Chiral resolution

Chiral resolution is a process in which a racemic compound is separated into its enantiomers, and is still an important tool for obtaining enantiopure products.[79,80] A downside of chiral resolution of racemates, compared to other asymmetric syntheses, is that a maximum yield of 50 % of the desired product can be obtained, with another 50 % being the undesired enantiomer.

Separation of enantiomers in a racemate may be achieved by reaction with enantio-pure chiral reagents, such as chiral acids or bases, to form a mixture of diastereomer intermediates.[54] Whereas the enantiomers are identical, the diastereomers may possess different chemical and physical properties and can be separated, as illustrated in Scheme 2.6.5. It is possible to also exploit the difference in reaction rates of enantiomers with chiral reagents in so-called kinetic resolution: one of the enantiomers reacts more rapidly, thus leaving an excess of the other enantiomers behind.[81] A widely utilized version of kinetic resolution is enzymatic kinetic resolution, employing biocatalysts, especially hydrolytic enzymes, to achieve high enantioselectivity and good yields.[82] A third method is the use of preparatory chromatography, such as chiral stationary phase HPLC (CSP-HPLC). For CSP-HPLC, the stationary phase is a chiral reagent that adsorbs one enantiomer more strongly than the other, allowing for direct separation of the enantiomers from the racemate.[83–85]
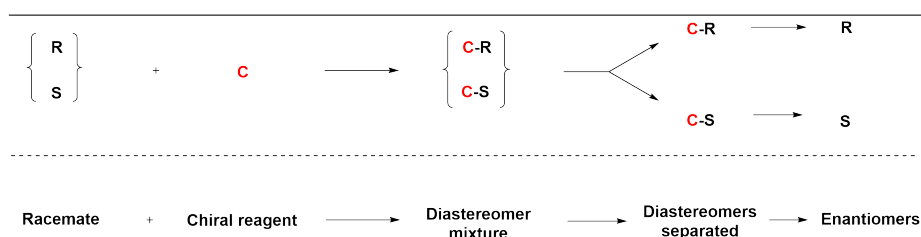
Scheme 2.6.5: Chiral resolution of a racemate by formation and separation of diastereomers.

Chiral resolution of *rac*-**4** to *(R)*-**4** has been reported by using dibenzoyl-*L*-tartaric acid as a chiral acid.[54] The *(R)*-enantiomer precipitates in the solution as a salt, allowing for recrystallization and removal of the salt for the formation of enantiopure *(R)*-**4**. To achieve high ee, the enantioenriched salt is submitted to sequential recrystallizations, with each cycle increasing the ee of the final product. Scheme 2.6.6 illustrates the chiral resolution of *rac*-**4** to *(R)*-**4**. Synthesis of the *(S)*-enantiomer is possible by substituting the chiral acid for dibenzoyl-*D*-tartaric acid.
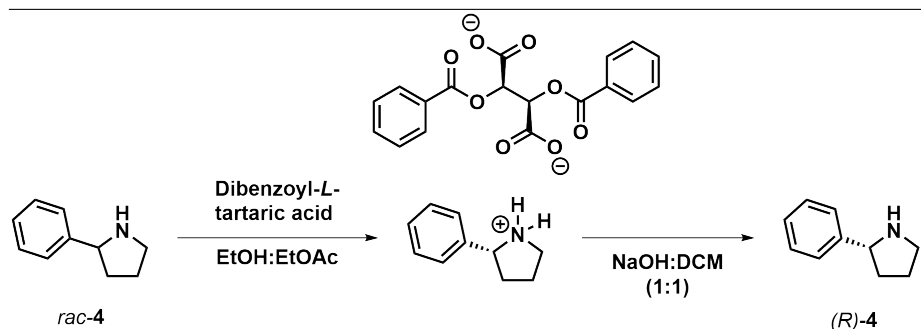
Scheme 2.6.6: Chiral resolution of *rac*-**4** to *(R)*-**4** by crystallization with dibenzoyl-L-tartaric acid.[54]

α-Arylation

An effective synthesis of 2-substituted pyrrolidines is the enantioselective, palladium-catalyzed α-arylation of *N*-Boc-pyrrolidine, illustrated in Scheme 2.6.7.[52,86] The reaction proceeds via an in-situ generated 2-pyrrolidinozinc reagent, allowing for a direct palladium-catalyzed arylation from **1** to *(R)*- or *(S)*-**3**. What makes this procedure favourable is that it provides a more direct route to enantioenriched 2-arylpyrrolidines, compared to longer synthetic routes,[87] while maintaining good yields and excellent enantiomeric excess.[86]
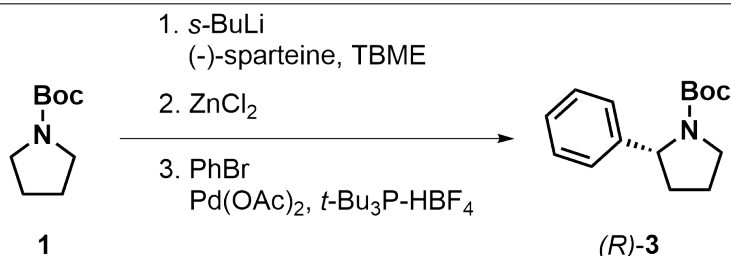
Scheme 2.6.7: Enantioselective, palladium-catalyzed α-arylation of *N*-boc-pyrrolidine to *(R)*-**3**.[52,86]

This methodology is based on Beak's sparteine-mediated, enantioselective deprotonation and lithiation of *N*-boc-pyrrolidine to form 2-pyrrolidinolithium compounds in high enantioselectivity.[88] The lithiation is followed by transmetallation with ZnCl$_2$ to generate a stereochemically rigid 2-pyrrolidinozinc reagent, which is coupled with aryl bromide in a Pd-catalyzed Negishi coupling.[86,89] Scheme 2.6.8 illustrates the enantioselective, sparteine mediated deprotonation and lithiation of *N*-Boc-pyrrolidine, followed by generation of the 2-pyrrolidinozinc reagent.
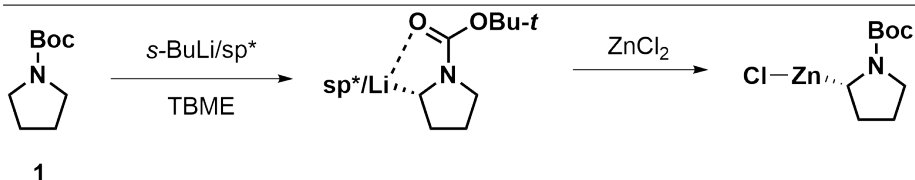
Scheme 2.6.8: Enantioselective, (-)-sparteine mediated deprotonation and lithiation of **1**. Transmetallation with ZnCl$_2$ generates the stereochemically rigid 2-pyrrolidinozinc.[86,89–91]

The lithiation is the enantiodiscriminating step of this synthesis, and thus the final enantiomeric excess is independent of the aryl bromide utilized in the Negishi-coupling.[52,89,91] Scheme 2.6.9 presents the catalytic cycle for the Negishi-coupling of this synthesis. The potential for β-H elimination in this catalytic Negishi cycle has been evaluated, and found products resulting from this pathway to be less than 8 %.[86]
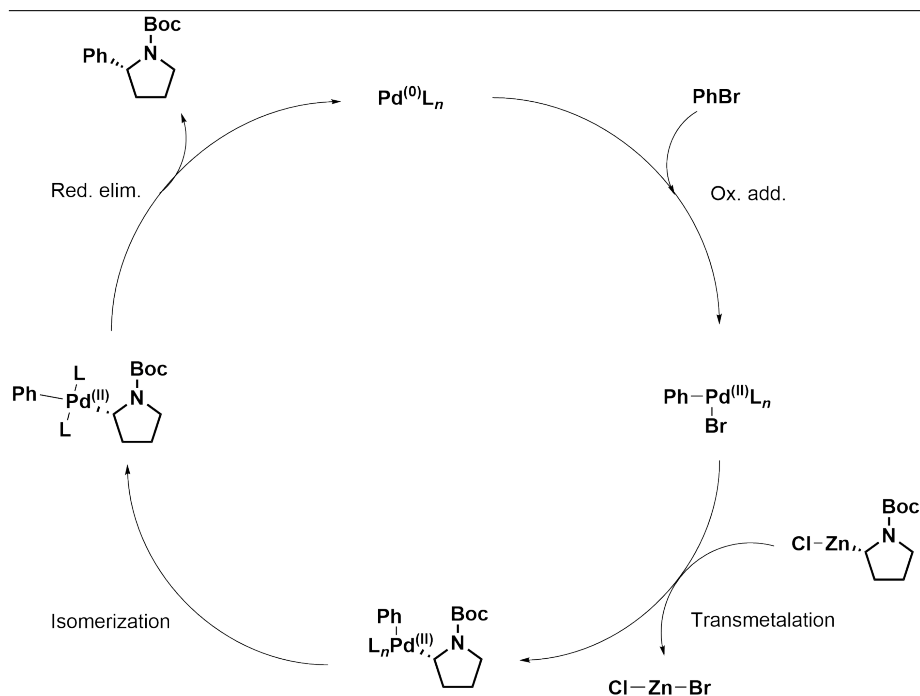
Scheme 2.6.9: Catalytic cycle for the transmetallation and Negishi-coupling in the one-pot α-arylation from **1** to *(R)*-**3**.[92]

Chemo-enzymatic route

The utilization of both chemical and biological approaches combined, have proven to be powerful tools for organic synthesis and has long been a topic of interest in asymmetric synthesis.[93,94] Such reactions combine the flexibility of chemical synthesis with the selectivity and efficiency of enzymatic methods, which also makes for "greener" synthesis by reducing downstream processing steps and allowing for potential in-situ regeneration of expensive co-factors.[93,95]

Several approaches have been suggested for chemo-enzymatic synthesis of cyclic amines, including asymmetric synthesis with transaminase,[96] deracemization with monoamine oxidase,[97] kinetic resolution with hydrolase,[98] and imine reductase.[99] Chemo-enzymatic deramemication of *rac*-**4** to *(R)*-**4** has been reported, using the specific monoamine oxidase Ile246Met/ Asn336Ser/Met348Lys/Thr384Asn/Asp385Ser (MAO-N-5) from *Aspergillus niger*. The process proceeds via a two-step, one-pot synthesis, in which the enzyme oxidizes only the *(S)*-enantiomer to the corresponding imine. The imine is reduced in-situ back to the racemic amine, with repeated cycles accumulating *(R)*-**4** in high yield and enantiomeric excess.[53] Scheme 2.6.10 presents this reaction.
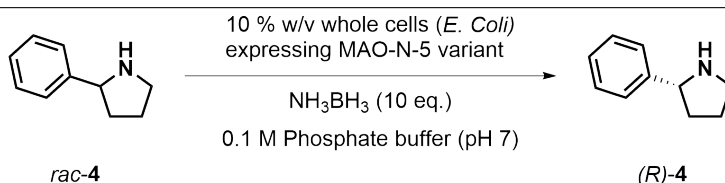
Scheme 2.6.10: Chemo-enzymatic deracemication of *rac*-**4** to *(R)*-**4**, by use of the enzyme Ile246Met/ Asn336Ser/Met348Lys/Thr384Asn/Asp385Ser (MAO-N-5) from *Aspergillus niger*.[53]

The direct reduction of prochiral imines is possible by NAD(P)H-dependent oxidoreductases, termed imine reductase (IRED). Reduction of **2** to *(S)*-**4** has been reported using *Ao*IRED from *Amycolatopsis orientalis* (Uniprot R4SNK4), with ee's up to 95 %.[100] The mechanism of IRED is not well understood, and it appears that the use of different oxidoreductases changes the mechanistic pathways as these employ different amino acid residues within the active site for substrate recognition.[100] Scheme 2.6.11 illustrates the imine reductase of **2** to *(S)*-**4** by *Ao*IRED.
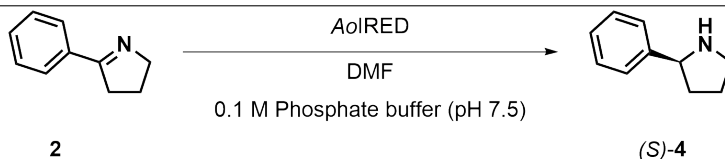
Scheme 2.6.11: Imine reductase (IRED) of **2** to *(S)*-**4**, using *Ao*IRED from *Amycolatopsis orientalis* (Uniprot R4SNK4).[100]

2.6.2   One-pot synthesis of 2-substituted pyrroline

Compound **2** can be synthesized from pyrrolidin-2-one in a two-step, one-pot reaction, starting with the protection of the amine by silylation, followed by a Grignard reaction.[57] Scheme 2.6.12 illustrates this reaction.
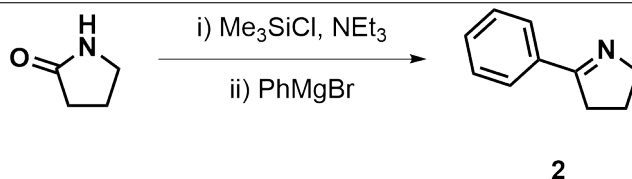
Scheme 2.6.12: Two-step, one-pot synthesis of **2** from pyrrolidin-2-one.[57]

Trimethylsilyl chloride may be used as the silylating agent, which acts as both a protecting group in the first reaction step, and a leaving group in the second step.[57] The

silyl group is predominantly located on the amine with some protection of the ketone group, as illustrated in Scheme 2.6.13.[57] Formation of **2** occur by a spontaneous deprotection, allowing for the formation of the double bond in the pyrroline moiety.
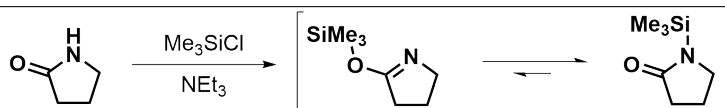
Scheme 2.6.13: Blocking of pyrrolidin-2-one with a trimethylsilyl group gives two products. The silyl group is predominantly located on the amine.[57]

2.6.3 Synthetic route for thieno[2,3-*d*]pyrimidines

Synthetic routes for preparing thieno[2,3-*d*]pyrimidines have been thoroughly investigated by Bugge.[101] A retrosynthetic analysis with 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine as the initial core, illustrated in Scheme 2.6.14, found nucleophilic aromatic substitution and Suzuki cross-coupling reactions to be viable options for the synthetic route of target structures **6-8**.
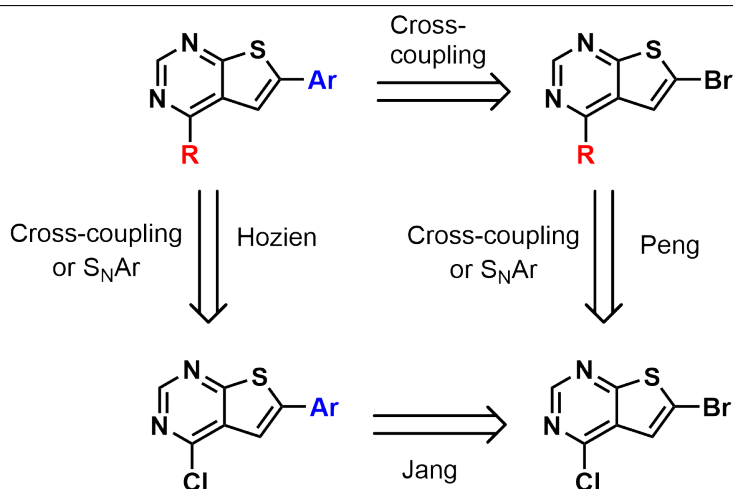
Scheme 2.6.14: Retrosynthetic analysis with 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine as the initial core. References: Hozien *et al.*,[102] Peng *et al.*[103]

Scheme 2.6.15 presents two synthetic routes, A and B, to the target structures. On similar structures, it has been found that nucleophilic aromatic substitution, followed by Suzuki cross-coupling (route B) was the most efficient synthetic route, due to low selectivity or incomplete conversion of route A. Route B was found to circumvent the selectivity issues, and also gave the highest overall yields.[101]
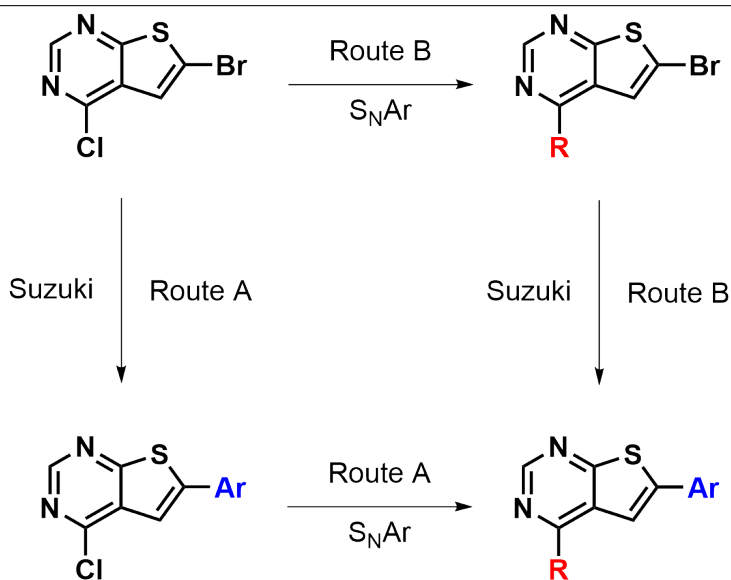
Scheme 2.6.15: Two possible routes, A and B, from 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine to the target structures.

The reduction of aldehydes **7a-b** to the corresponding primary alcohols **8a-b**, illustrated in Scheme 2.6.16, was done using sodium borohydride (NaBH$_4$) in THF:MeOH (3:1), as previously reported on similar structures.[9]
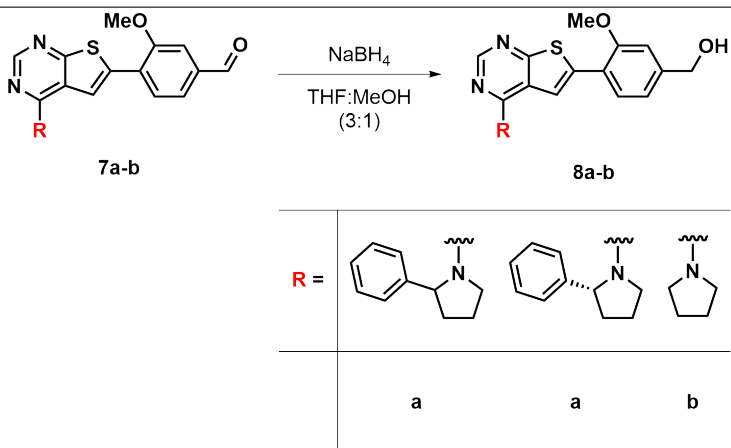
Scheme 2.6.16: Reduction of aldehydes **7a-b** to primary alcohols **8a-b**, using sodium borohydride (NaBH$_4$) as the reducing agent.[9]

2.6.4 Nucleophilic aromatic substitution

Thieno[2,3-*d*]pyrimidines consist of a fused pyrimidine- and thiophene moiety. While six-membered heterocycles such as pyrimidine are generally electron deficient, thiophene is an electron rich heterocycle, allowing for regioselective nucleophilic aromatic substitution on thieno[2,3-*d*]pyrimidines under the right conditions.[104]

Nucleophilic aromatic substitution may proceed through several mechanisms, including:

- Addition-elimination

- Elimination-addition

- Metal-catalyzed processes

Thieno[2,3-*d*]pyrimidines favour the addition-elimination mechanism, as the nitrogen atoms in the pyrimidine moiety acts to stabilize the reaction intermediate (Meisenheimer complex), thus activating the ring towards nucleophilic attack. The addition of the nucleophile forms a resonance-stabilized anion, followed by an elimination of the halogen leaving group, as illustrated in Scheme 2.6.17.
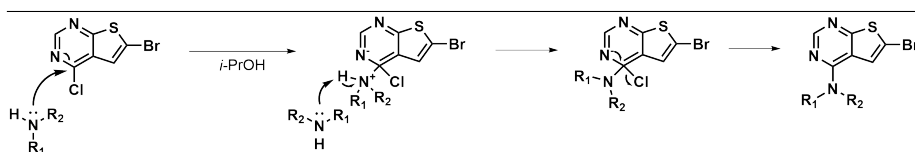
Scheme 2.6.17: Nucleophilic aromatic substitution via an addition-elimination mechanism.

The reaction rate of the nucleophilic aromatic substitution reaction is a product of several factors, including the chosen solvent, the leaving group, and the nucleophile. Considerably higher reaction rates has been found for S_NAr in dipolar aprotic solvents as opposed to polar solvents, due to increased stability of the Meisenheimer complex and increased basicity of the amine in aprotic solvents.[105,106] The addition of a hydrogen-bond acceptor co-solvent, such as a more basic non-nucleophilic amine, also has a favourable effect on the reaction rate. In addition to its inherent catalytic effect, the use of a co-solvent amine with higher pK_a than the nucleophilic amine gives an amine-amine aggregate which acts as a better nucleophile, compared to the amine without the co-solvent.[107]

The nature of the leaving group is a key factor for the reaction rate of nucleophilic aromatic substitution. Common leaving group are halides, although alkoxy, cyano, nitro and sulfonyl groups may also be displaced in highly activated systems. Because the addition of the nucleophile is the rate determining step of this reaction, the leaving group order of the halides is divergent from general S_N2 reactions, with F > Cl > Br > I. The rate determining step is also greatly facilitated by electron deficient aromatic reactants, hence perfect for regioselective substitution of thieno[2,3-*d*]pyrimidines.

2.6.5   Suzuki cross-coupling

The choice of method for the carbon-carbon coupling of thieno[2,3-*d*]pyrimidines **5a-b** with their respective aryl groups, was the Suzuki cross-coupling reaction. This palladium-catalyzed cross-coupling reaction has been a pioneer in cross-coupling since it was first published in 1981, as the large diversity of available boronic acids allows for a high degree of customization.[108] In combination with low cost and toxicity of substrates and catalysts, and mild reaction conditions, the Suzuki cross-coupling reaction has become the "golden standard" for preparation of biaryls and is ubiquitous in modern medicinal chemistry.[109]

The mechanism of Suzuki cross-coupling involves four steps:[110]

- Oxidative addition of aryl-halide to Pd(0) to form a Pd(II)-complex

- Exchange of the anion attached to Pd(II) for the anion of the base (Metathesis)

- Transfer of the organic group from the organo-metal to the Pd(II)-complex (Transmetallation)

- Reductive elimination of the substituted Pd(II)-complex to give carbon bond formation, and to re-form the Pd(0) catalyst.

The mechanism of the transmetallation step has long been a subject of debate, concerning the activation of the organoboron compound by the base.[111] Two prominent pathways have been considered (path A and B in Scheme 2.6.18), which differ in the way transmetallation is initiated by the hydroxide ion.[112]
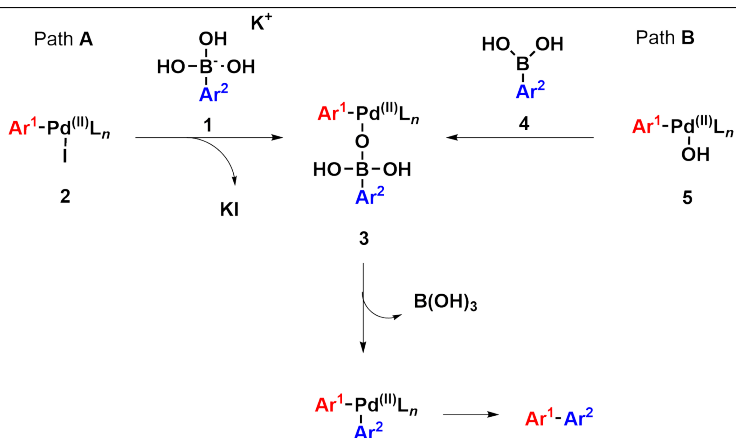
Scheme 2.6.18: Transmetallation pathways in Suzuki cross-coupling.[112,113]

Path A proceeds from a palladium halide complex (**2**), via a negatively charged aryl-trihydroxyboronate, while path B proceeds from a palladium hydroxide complex via a neutral arylboronic acid.[112] Both pathways converge at the same intermediate containing a Pd-O-B unit (**3**), whose structure was confirmed fairly recently.[113] Extensive

kinetic studies have established path B as the favoured route, and the generally accepted catalytic cycle for the Suzuki cross-coupling is presented in Scheme 2.6.19.[112,114–116]

Scheme 2.6.19: General catalytic cycle for the Suzuki cross-coupling reaction of arylhalides.[117]
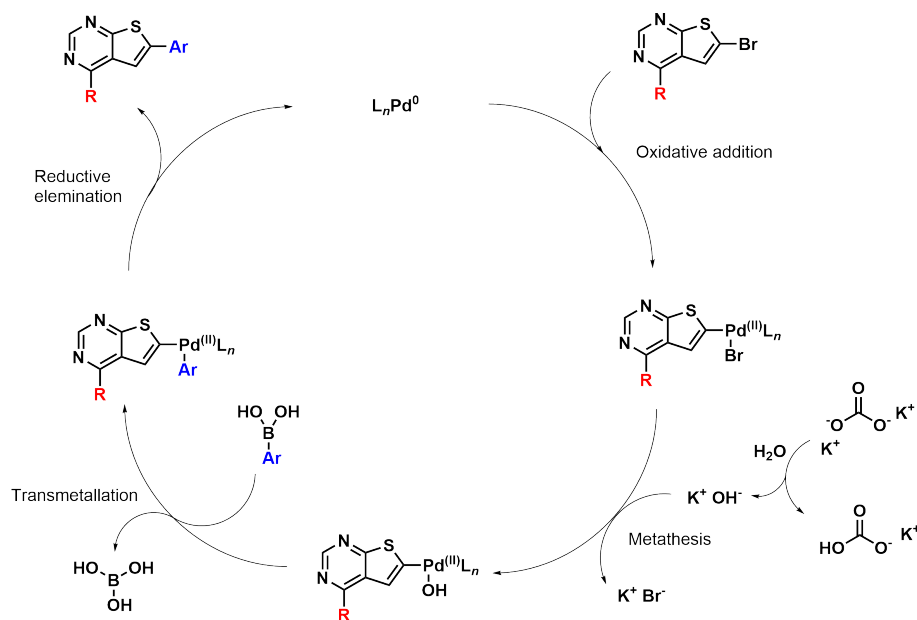
2.6.6 Reductions with NaBH$_4$

Reduction of aldehydes

The reduction of aldehydes to primary alcohols is done by treatment with either NaBH$_4$ or LiAlH$_4$. While both give the same overall result, the reaction conditions for using either one are rather different. Due to the high reactivity of LiAlH$_4$, this one reacts rapidly and violently with protic solvents, such as water or alcohols, and reactions employing this reducing agent must be protected from moisture. NaBH$_4$ is a milder reagent and allows for reductions to be carried out in water or methanol solution, which makes for an easier workup.[118,119] A general mechanism for the reduction of an aldehyde to a primary alcohol using NaBH$_4$ is presented in Scheme 2.6.20, illustrating the nucleophilic attack from the hydride ion on the carbonyl carbon, followed by protonation of the resulting alkoxide.
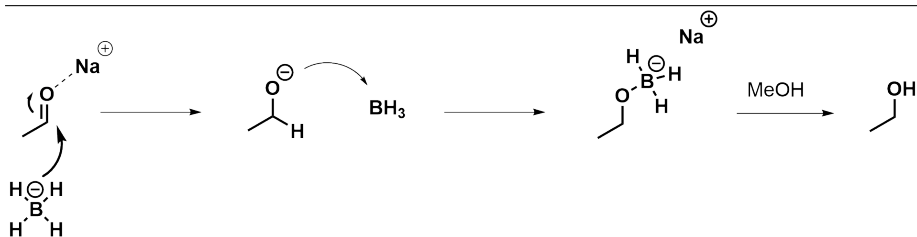
Scheme 2.6.20: Reduction of aldehyde with NaBH$_4$.

The mechanism of NaBH$_4$ reduction becomes more complex when taken into account that all four of the hydrides may eventually be transferred, meaning that several distinct reducing agents may be functioning throughout the course of the reaction.[120] Scheme 2.6.21 helps illustrate this complexity, as all species containing a B-H group may react either with the carbonyl or the solvent alcohol. As pointed out in this scheme, NaBH$_4$ may be used to reduce up to four equivalents of aldehyde, depending on the substrates and reaction conditions. This may be used along with methanol workup to form trimethyl borate from the boron complex, allowing for removal of the boron by drying to constant weight due to the low boiling point of trimethyl borate (bp. 69°C).[121–123]
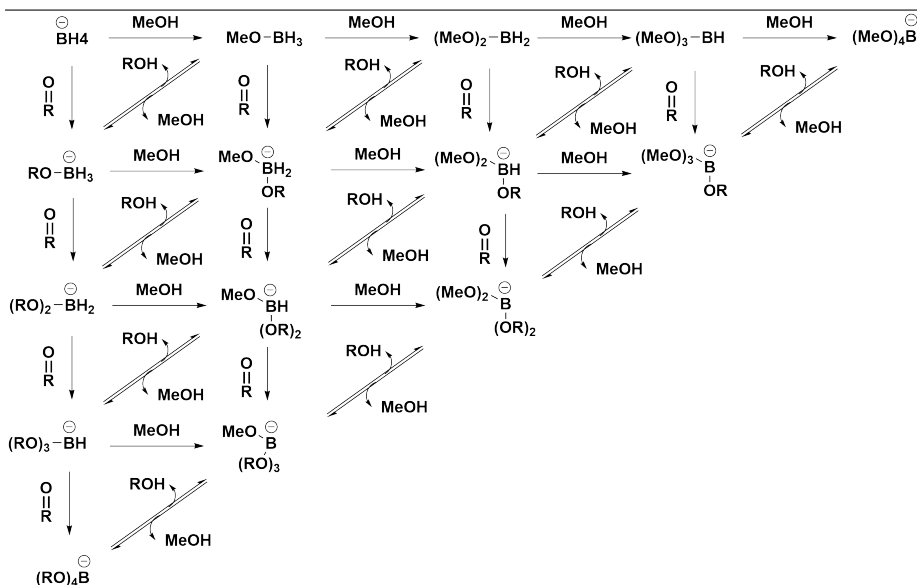
Scheme 2.6.21: Possible pathways in NaBH$_4$ carbonyl reduction. R=O in this case represents an aldehyde.

Reduction of imines

NaBH$_4$ has applications in a wide range of functional groups, and the use of this reducing agent for the reduction of imines conjugated with aromatic rings is well established.[119,124,125] As an inexpensive and mild reagent it was chosen for the reduction of compound **2** to *rac*-**4**, as illustrated in Scheme 2.6.22.[53] In this reaction, delocalization of the π-electrons occurs through conjugation of the carbon-nitrogen double bond with the phenyl group. This creates an electrophilic center, capable of being attacked by the nucleophilic borohydride anion.
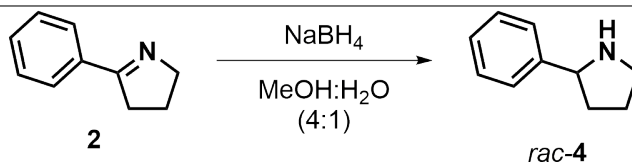
Scheme 2.6.22: Reduction of imine **2** with sodium borohydride (NaBH$_4$) to *rac*-**4**.

3 | Results and Discussion

The main focus of this master's thesis has been to carry out a preliminary SAR study of thieno[2,3d]pyrimidines with a tertiary heterocyclic amine as a substituent on C-4. A second objective has been to prepare these thieno[2,3-d]pyrimidine-based kinase inhibitors, using enanrioenriched and racemic 2-phenylpyrrolidine, and pyrrolidine as the substituent. The aim was to prepare new compounds with high chemical and enantiopurity, to study both the effect of chirality of the phenyl group of the C-4 amine, and the contribution it may provide to the EGFR-TK activity of these structures. This work has resulted in 14 new compounds, of which two show promising activity and may open new areas of research.
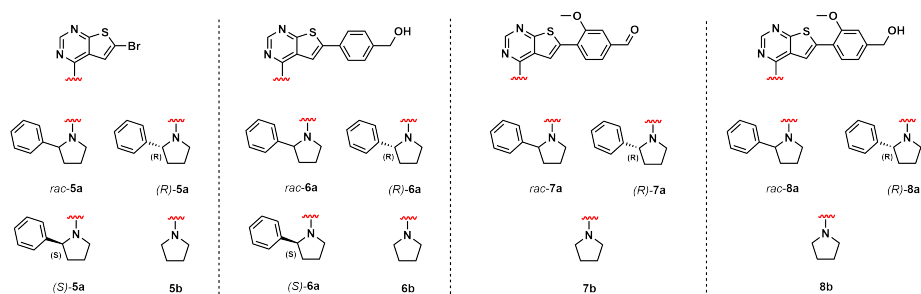
Figure 3.1: Structures of new thieno[2,3-d]pyrimidines **5-8**

This section will address the synthesis of precursor molecules, intermediates, and the synthesis and structure elucidation of structures **5-8**, followed by a review of the results from the bioassay of target structures **6a-b** and **8a-b**. The focus of the syntheses has been to achieve the target molecules with high purity, so the listed reaction parameters may not be optimized with respect to yield.

3.1   Boc protection
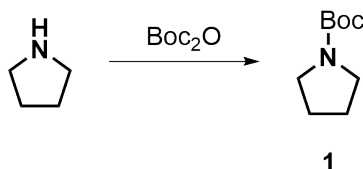
Scheme 3.1.1: Boc protection of pyrrolidine to compound **1**.

Compound **1** was synthesized as a precursor to *(R)-* and *(S)-***3**. The compound was synthesized via Boc protection of pyrrolidine, using Boc$_2$O.[126] The reaction time was 2 minutes, yielding 92-94 % of **1** as a yellowish liquid without the need for purification.

3.2   One-pot silylation and Grignard
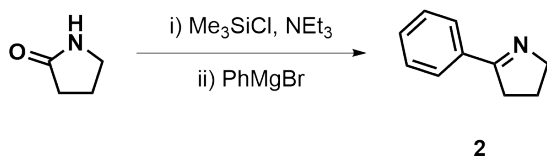
Scheme 3.2.1: Two-step, one-pot reaction of pyrrolidin-2-one to **2**.

Compound **2** was synthesized as a precursor to *rac*-**4**. The compound was synthesized in a two-step, one-pot reaction starting with a silylation, followed by addition of a Grignard reagent.[57] The obtained yields for this reaction were 12 - 36 %, with literature values ranging from 30 - 90 %.[53,57] The synthesis was performed several times under different conditions to tune conversion and yield.

The main issue of this reaction was to achieve full conversion of pyrrolidin-2-on during the initial silylation. The challenge encountered in this step was that the addition of Me$_3$SiCl to the reaction mixture caused a massive increase in viscosity of the solution. This made reflux and stirring of the reaction difficult, which in turn impacted the conversion of starting material. Small tweaks were done to improve this step, such as adding more solvent, changing reflux time, and substituting magnetic stirring for mechanical stirring. Table 3.1 presents an overview of these changes, and the overall effect on conversion and yield of the one-pot reaction.

Table 3.1: Scale, solvent volume, reflux time, stirring apparatus, conversion, and yield of the one-pot synthesis of compound **2**.

Scale [g]	Volume [mL][a]	Eq.[b]	Time [h][c]	Stirring	Conv. [%][d]	Yield [%][e]
1.03	15	1	24	Mag.	69	12
0.50	7.5	1	0.5	Mag.	95	14
0.71	10	2	4.5	Mag.	80	22
1.81	15	1.3	0.5	Mag.	89	35
5.07	75	1.1	2	Mech.	53	33
0.88	15	2.1	1	Mech.	92	35
0.52	10	2	0.5	Mech.	92	36

[a] Diethyl ether.
[b] Grignard reagent.
[c] Reflux time during silylation.
[d] Estimated from ^{1}H NMR.
[e] Isolated yield.

It is difficult to draw conclusions based on the results of Table 3.1, as there are considerable divergences in conversion and yield with the various parameters. The results obtained during these experiments were difficult to reproduce, something that has been previously noted with the use of a Grignard reagent in this transformation.[57] Some explanation for the erratic results may be the equilibrium that conforms during the silylation, where Me$_3$Si is attached to either the nitrogen or the oxygen of the lactam (Scheme 3.2.2). Provided only the N-protected isomer may react with the Grignard reagent to form **2**, this means that the yield is dependent on, inter alia, the amount of silyl-protected oxygen.
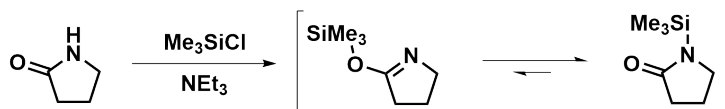
Scheme 3.2.2: The equilibrium between the O- or N-protected lactam of pyrrolidin-2-one after silylation. The equilibrium is shifted towards the protected nitrogen.

The formation of by-products could also be an explanation for the irregularities in the results. Some by-products are observed in the crude ^{1}H NMR of compound **2**, especially in the aromatic region, as presented in Figure 3.2.
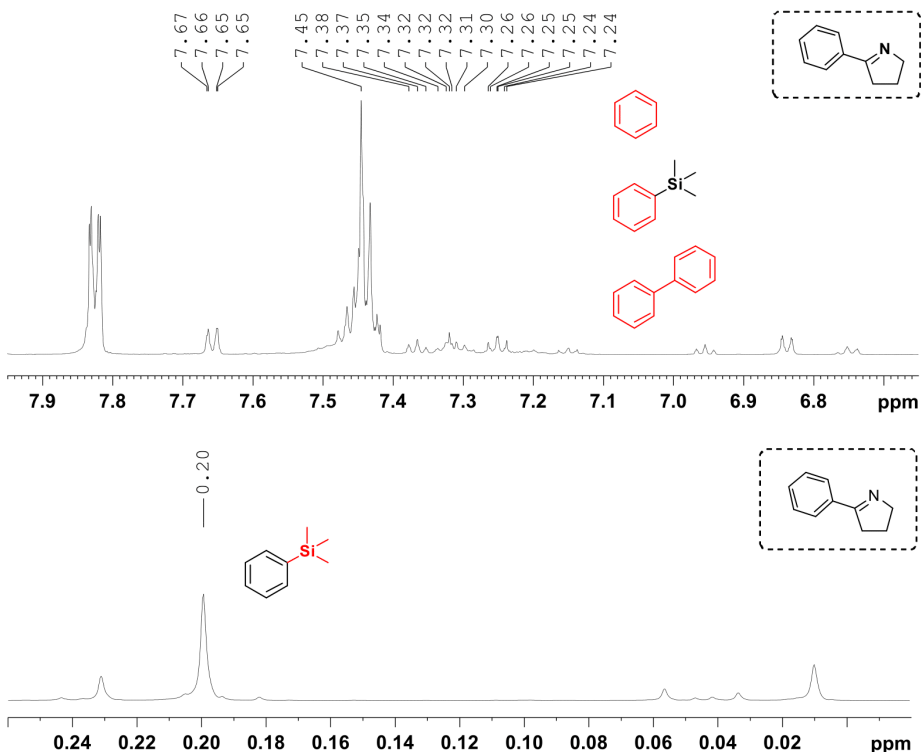
Figure 3.2: Presumed signals from trimethyl(phenyl)silane, benzene, and 1,1'-biphenyl in the ^{1}H NMR of crude material of compound **2**.

Some of these by-products arise from the reaction between the Grignard reagent and the silyllactam at silicon, as previously reported by Hua.[127] Scheme 3.2.3 illustrates this side reaction to form trimethyl(phenyl)silane. ^{1}H NMR spectrum of trimethyl(phenyl)silane in DMSO has not been found in the literature for reference; however, ^{1}H NMR modelling using integrated software on the computer-application "ChemBioDraw Ultra 14.0" suggest that this compound is present in the NMR of crude material of compound **2** as signals at 7.35-7.27 ppm and 0.20 ppm.[128] Figure 3.2 shows the presumed signals of trimethyl(phenyl)silane in the ^{1}H NMR spectrum of crude **2**.
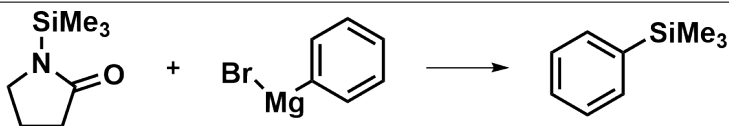
Scheme 3.2.3: Formation of the by-product trimethyl(phenyl)silane from the reaction between the Grignard reagent and the silyllactam at silicon.[127]

Addition of the Grignard reagent was done without purification of the silylated inter-mediate, which could lead to formation of by-products such as benzene due to insuffi-cient removal of ammonium salt from the silylation, as illustrated in Scheme 3.2.4.[129] Benzene may be present in the crude product of compound **2**, as a signal at 7.37 ppm in Figure 3.2.[130]
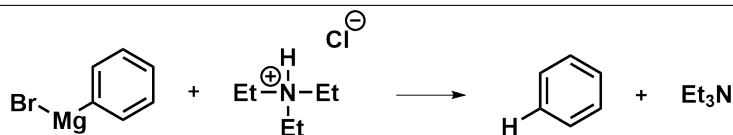
Scheme 3.2.4: Formation of the by-product benzene by reaction of the Grignard reagent with ammonium salt.

Another possibility for by-product is the formation of a 1,1'-biphenyl by homocou-pling of the Grignard reagent. This reaction could proceed via a radical type coupling or a Kumada cross-coupling, likely catalyzed by palladium-residues or other heavy-metal residues in the glassware or reagents, as illustrated in Scheme 3.2.5. Biphenyl may also be an impurity present in the Grignard reagent itself. ^{1}H NMR spectrum of 1,1'-biphenyl in DMSO has not been found in the literature for reference; however, ^{1}H NMR modelling using integrated software on the computer-application "ChemBio-Draw Ultra 14.0" suggest that this compound may be present in the crude material of **2** as signals at 7.66 ppm, 7.44 ppm, and 7.35 ppm.[131–133] Figure 3.2 shows the presumed signals of 1,1'-biphenyl in the ^{1}H NMR spectrum of crude **2**.
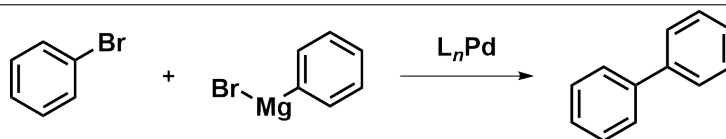
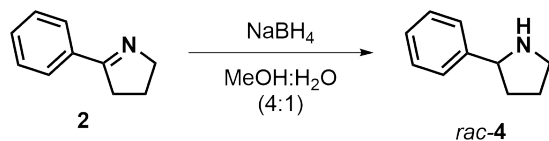
Scheme 3.2.5: Homocoupling of the Grignard reagent via a palladium-catalyzed Kumada cross-coupling.

Purification of the intermediate by distillation is a viable option to help avoid these side reactions; however, the gains of doing so have been reported as minuscule.[57]

3.3   Reduction of 2-phenylpyrroline

Scheme 3.3.1: Reduction of imine **2** to *(rac-***2**.

The reduction of imine **2** to *rac*-**4** was done using NaBH$_4$ as a reducing agent, and MeOH:H$_2$O (4:1) as solvent. The reaction was left for >12 hours, as previously reported by Dunsmore *et al.*.[53] Yields were in the range from 72-89 %, with a literature value of 97 % for this reaction.[53] Table 3.2 lists scale, reaction times, conversions, and yields for this reaction.

Table 3.2: Scale, reaction time, conversion, and yield for the reduction of compound **2** to *rac*-**4**.

Scale [g]	Time [h]	Conversion [%][a]	Yield [%][b]
3.16	17	>97	72
0.3	22	>97	77
1.36	22	>97	89

[a] Conversion determined by ^{1}H NMR.
[b] Isolated yield.

3.4   Palladium-catalyzed α-arylation
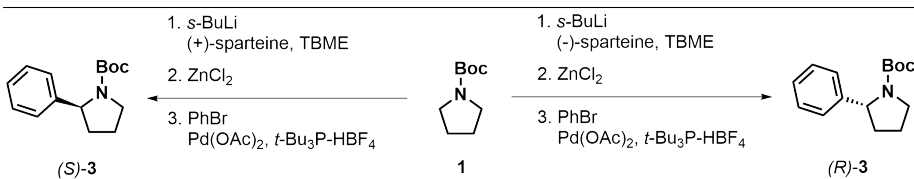
Scheme 3.4.1: Palladium-catalyzed α-arylation of compound **1** to *(R)*- and *(S)*-**3**.

The *(R)*- and *(S)*-enantiomers of **3** were synthesized from **1** in a palladium-catalyzed α-arylation, employing (-)-sparteine for the preparation of *(R)*-**3** and (+)-sparteine for *(S)*-**3**.[52,86] Both reactions were left for >12 hours, and achieved full conversion of starting material. Yields were 62 % for *(R)*-**3** and 70 % *(S)*-**3**. No other compounds were observed.

Enantiomeric analyses were done by analytical chiral stationary phase HPLC (CSP-HPLC), using a Chiralpak AD column. The analyses were done for these compounds, as opposed to the deprotected *(R)*- and *(S)*-**4**, due to more available experimental data and availability of HPLC-columns.[52][86] By tuning the available parameters, a near base line resolution was achieved with $R = 2.3$. Enantiomeric excess was found to be 86 % for *(R)*-**3**, and 96 % for *(S)*-**3** under these conditions. Parameters and corresponding retention times are listed in Table 3.3. Differences in ee are suspected due to divergences in the enantiopurity of (-)- and (+)-sparteine, as the experimental preparation of *(R)*- and *(S)*-**3** are close to identical.

| Flow [mL]	iPrOH/Hexane	$t_{R,1}$ [min.]	$t_{R,2}$ [min.]	α	R	N
0.8	1/99	8.5	9.0	1.05	1.79	10071
0.5	1/99	10.9	11.6	1.07	2.24	25688
0.3	1/99	24.9	26.7	1.07	2.3	23958
0.2	1/99	34.8		1	0	-
0.5	2/98	10.0		1	0	-
0.3	2/98	10.9	11.2	1.03	1.76	102949
0.2	2/98	24.0		1	0	-

Table 3.3: Optimization of base line separation for the enantiomers of *(R)*- and *(S)*-**3**.

^{1}H NMR specta of *(R)*- and *(S)*-**3** (Appendix C to D) show the presence of rotamers ($\sim$40:60) by splitting of the CH proton on the pyrrolidine moiety at 4.75 ppm, and the $(CH_3)_3$ protons of the Boc-group at 1.25 ppm. Figure 3.3 illustrates the rotamers of *(R)*-**3**.
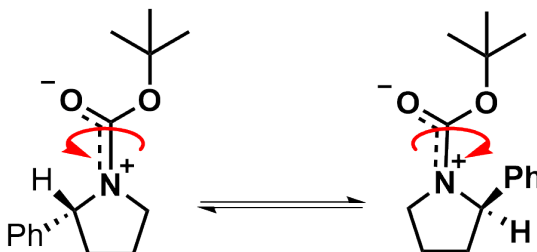
Figure 3.3: Illustration of the two rotamers of *(R)*-**3**.

To confirm that this splitting is indeed caused by rotamers, the compounds were analyzed by temperature gradient ^{1}H NMR. By increasing the temperature to 50 °C, the two CH peaks merge, while the $(CH_3)_3$ peaks only fully merge at around 80 °C. Figure 3.4 shows the effect of increased temperature on the ^{1}H NMR spectrum of *(R)*-**3**. The effect is the same for *(S)*-**3**, and the ^{1}H NMR spectra of the two compounds are close to identical.
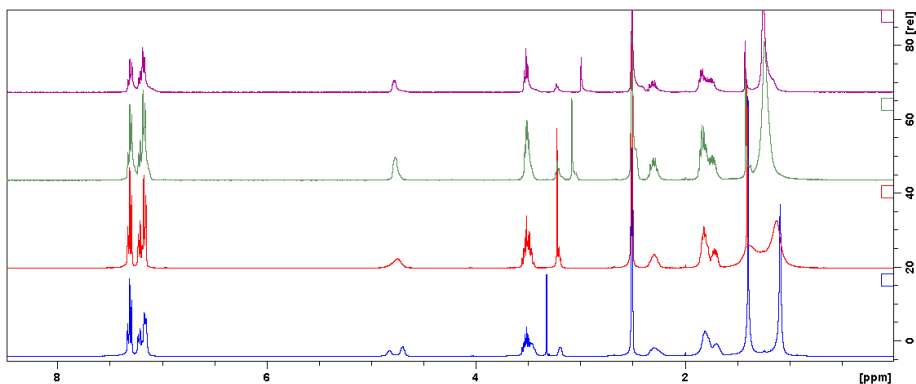
Figure 3.4: Temperature gradient ^{1}H NMR (400 MHz, DMSO-d_6) of *(R)*-**3**. From top to bottom: 100 °C, 80 °C. 50 °C, room temperature. Rotamers are visible at 4.75 ppm, and 1.25 ppm.

3.5 Boc deprotection
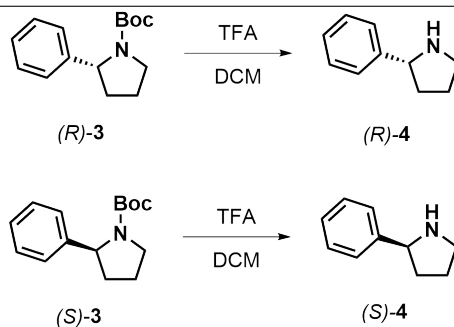
Scheme 3.5.1: Boc deprotection of compounds *(R)*- and *(S)*-**3** to *(R)*- and *(S)*-**4**.

(R)- and *(S)*-**4** were synthesized by Boc deprotection of *(R)*- and *(S)*-**3**. The deprotection was done using trifluoroacetic acid as described by Trost *et al.*,[52] with a reaction time of 4 hours. The reactions gave *(R)*- and *(S)*-**4** as mixtures with 0.61 eq. of TFA salt. The yields were 56 % and 49 % for *(R)*- and *(S)*-**4**, respectively.

Identification of the TFA salt was done by ^{19}F NMR, using hexafluorbenzene as an internal standard. The salt is visible as a signal at -73.5 ppm, with hexafluorbenzene at -162.6 ppm, as presented in Figure 3.5. The amount of TFA salt formed in the reactions was found by quantitative ^{19}F NMR, by referencing known amounts of *(S)*-**4** TFA salt with the internal standard. Calculations were done using the formula

$$\frac{M_P}{M_I} = \frac{I_P}{I_I} * \frac{N_I}{N_P},$$

where M_P/M_I is the molar ratio between product and internal standard, I_P/I_I is the integral ratio between product and internal standard, and N_I/N_P is the ratio between the number of ^{19}F atoms in the internal standard and the product.[134] As the experimental procedure for the preparation of *(R)*- and *(S)*-**4** are close to identical, it was estimated that both reactions yielded 0.61 equivalents of TFA salt. The yields were calculated based on the estimate of the formation of 39 % desired product.
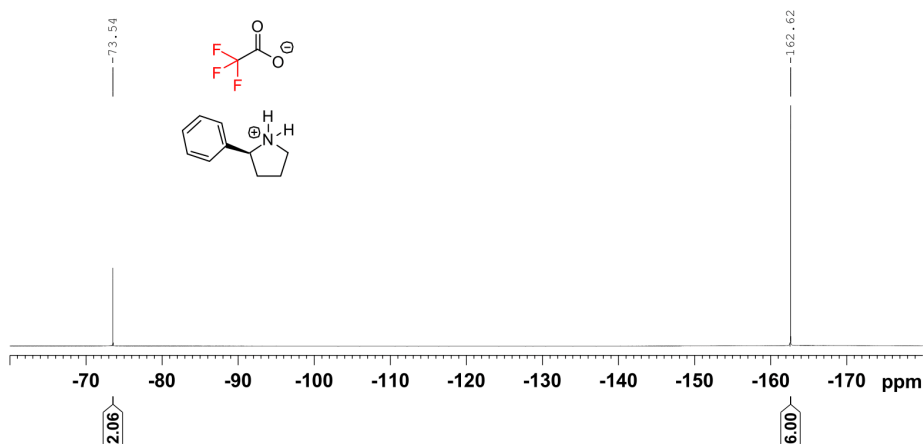
Figure 3.5: ^{19}F NMR of *(S)*-**4**, with hexafluorobenzene as the internal standard.

3.6   Chiral resolution of *rac*-2-phenylpyrrolidine
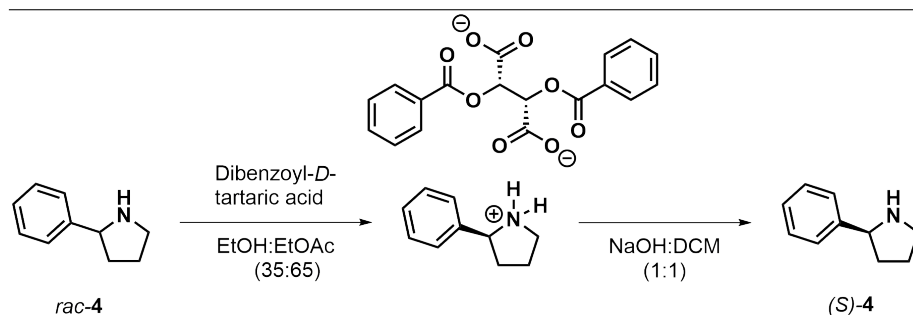

Scheme 3.6.1: Chiral resolution of compound *rac*-**4** to *(S)*-**4**.

Chiral resolution of racemic 2-phenylpyrrolidine was investigated as a synthetic route to enantioenriched 2-phenylpyrrolidines. Racemic **4** (*rac*-**4**) was mixed with dibenzoyl-*D*-tartaric acid and dissolved in a solution of 35 % ethanol in ethyl acetate. The solution was heated to boiling for 10 minutes and then slowly cooled to room temperature, to allow crystallization of the salt. The procedure yielded 31 % salt as

white crystals, which were then vigorously stirred in a mixture of NaOH:DCM (1:1) to remove the salt. *(S)*-2-phenylpyrrolidine was obtained in 49 % yield as an orange oil.

A sample of *(S)*-2-phenylpyrrolidine was Boc protected and analysed by CSP-HPLC to check the ee. Using the established eluent system for compounds *(R)*- and *(S)*-**3**, the ee was found to be 24 %, indicating a slight excess of the *(S)*-enantiomer. Subsequent recrystallization of the intermediary *(S)*-2-phenylpyrrolidine salt has been reported as a way to improve the enantioselectivity in this procedure;[54] however, due to time constraints this was not investigated in this project. Despite the low enantiopurity, this procedure succeeds in providing an alternate path to enantioenriched 2-phenylpyrrolidines from the racemic amine.

3.7 Amination of thienopyrimidines
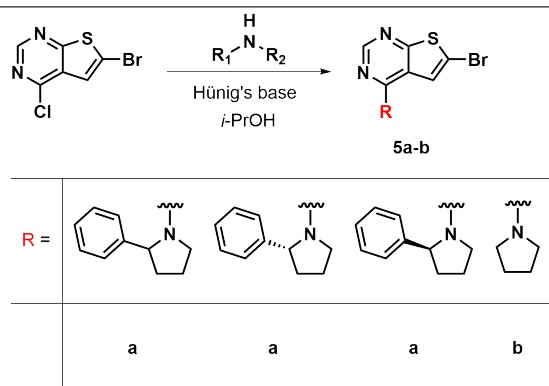
Scheme 3.7.1: Nucleophilic aromatic substitution of 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine to compounds **5a-b**.

Compounds **5a-b** were synthesized by nucleophilic aromatic substitution, using *i*-PrOH as solvent.[10] The reactions were stirred at 80 °C under an N_2 atmosphere, and all of the compounds were purified by silica gel column chromatography. Table 3.4 lists conversion, reaction times and yields for these reactions.

Table 3.4: Yields, conversion and reaction times for the synthesis of compounds **5a-b**

Reactant	Scale [mg]	Eq.[a]	Eq.[b]	Time [h]	Conv. [%][c]	Yield [%][d]	Product
rac-**4**	200	1.5	1.5	1.25	>97	80	*rac*-**5a**
(R)-**4**	302	14	1.4	24	82	74	*(R)*-**5a**
(S)-**4**	206	1.5	1.4	24	59	50	*(S)*-**5a**
Pyrrolidine	416	1.5	1.5	1.6	>97	91	**5b**

[a] Amine.
[b] Hünig's base.
[c] Conversion determined by ^{1}H NMR.
[d] Isolated yield.

Suboptimal conversion of starting material was observed in the synthesis of *(R)*- and *(S)*-**5a**, likely due to the presence of TFA salt in *(R)*- and *(S)*-**4**. With 0.61 equivalents of TFA salt, most of the amines would be unavailable as a nucleophile. It was thought that by adding a surplus of Hünig's base, this would act to deprotonate salt in addition to acting as co-base for the nucleophilic aromatic substitution. This was tested in the synthesis of *(R)*-**5a**, giving considerably better conversion compared to the synthesis of *(S)*-**5a**. To improve the conversion and yield of these two reactions, more co-base is needed to fully deprotonate the TFA salt. Alternatively, *(R)*- and *(S)*-**4** can be deprotonated prior to the nucleophilic aromatic substitution.

3.8 Suzuki cross-coupling
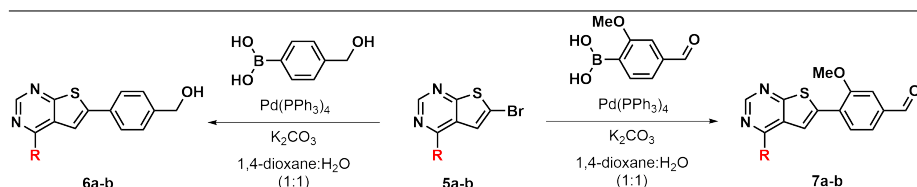
Scheme 3.8.1: Suzuki cross-coupling of **5a-b** to **6a-b** and **7a-b**.

Compounds **6a-b** were synthesized by Suzuki cross-coupling with the appropriate arylboronic acids. The reactions were stirred at 80 °C under an N$_2$ atmosphere, and all compounds were purified by silica gel column chromatography. Table 3.5 lists conversion, reaction times and yields for these reactions.

Table 3.5: Yields, scale, conversion, and reaction times for the synthesis of compounds **6a-b** and **7a-b**

Reagent	Scale [mg]	Time [min.]	Conv.[a] [%]	Yield[b] [%]	Product
rac-**5a**	111	25	>97	60	*rac*-**6a**
rac-**5a**	100	55	>97	93	*rac*-**6a**
rac-**5a**	131	30	>97	56	*rac*-**7a**
rac-**5a**	372	90	>97	99	*rac*-**7a**
(R)-**5a**	146	45	>97	92	*(R)*-**6a**
(R)-**6a**	172	45	>97	92	*(R)*-**7a**
(S)-**5a**	131	45	>97	92	*(S)*-**6a**
5b	103	55	>97	58	**6b**
5b	418	45	>97	94	**7b**

[a] Conversion determined by ^{1}H NMR.
[b] Isolated yield.

In the initial Suzuki cross-coupling reactions of *rac*-**5a** to *rac*-**6a** and *rac*-**7a**, ~24 % of by-products were formed, giving low yield of the desired products. The by-product from the synthesis of *rac*-**6a** was isolated and analysed, identifying this compound to be the product of dimerization of the arylboronic acid. The ^{1}H NMR spectrum of the dimer, 4,4'-bis(hydroxymethyl)biphenyl, is given in Appendix M, and NMR data are in accordance with previously reported values.[135] It is suspected that the dimerization was the result of insufficient degassing of the solvent, causing oxidation of the phosphine ligands of tetrakis(triphenylphosphine)palladium(0) and of Pd(0) to Pd(II), thus favouring formation of the dimer.[136,137] Evidence of this is that proper degassing resulted in minute amounts of by-product and increased the yield in later experiments.

CSP-HPLC analyses were done for compounds *(R)*- and *(S)*-**6a**, and *(R)*-**8a**, to cross-reference the ee of the final products to that of compounds *(R)*- and *(S)*-**3**. A chiral analysis of these compounds was developed, based on an analytic procedure on similar compounds by Bugge *et al.*,[11] using a Lux 5u Cellulose HPLC column. The ee of compounds *(R)*- and *(S)*-**6a** were both found to be 65 %, which marks a significant drop from the 86-96 % ee of *(R)*- and *(S)*-**3**, respectively. The ee of compound *(R)*-**8a** was found to be 84 %, which may be within standard deviation of the HPLC data. While no clear sources of racemization of compounds *(R)*- and *(S)*-**6a** have been found, a possibility may be palladium-catalyzed racemization. This has been reported for *(S)*-2-phenylpyrrolidine in a Buchwald-Hartwig N-Arylation,[138] and for amino acid derivatives in palladium-catalyzed cross-coupling reactions with the use of the Pd(PPh$_3$)$_4$ catalyst.[139] In these cases, racemization is possible due to β-hydrogen elimination or the presence of an acidic α-proton. Neither of these are likely sources for racemization in the Suzuki cross-coupling reactions in this project, as the amine is attached to the thieno[2,3*d*]pyrimidine core prior to the cross-coupling reactions. Racemization would most likely occur with the presence

of palladium-residues, either from glassware or reagents, during the S_NAr reactions; however, further studies are required to clarify these findings.

3.9   Reduction of aldehydes by sodium borohydride
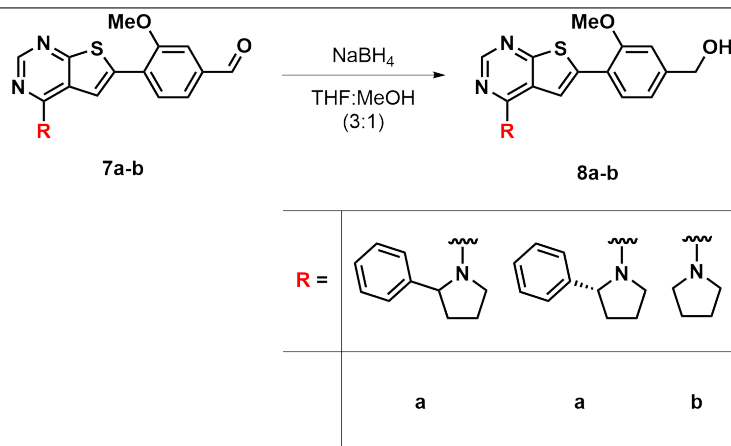
Scheme 3.9.1: Reduction of **7a-b** to **8a-b** with sodium borohydride (NaBH$_4$).

Compounds **8a-b** were synthesized by reduction of aldehyde to primary alcohol, using NaBH$_4$ as the reducing agent, and THF:MeOH as solvent. The reactions were stirred at room temperature under an N$_2$ atmosphere, and all compounds were purified by silica gel column chromatography. Table 3.6 lists scale, conversion, reaction times, and yields for these reactions.

Table 3.6: Yields, scale, conversion, reaction times, and yields for the synthesis of compounds **8a-b**

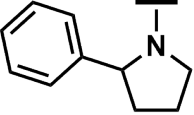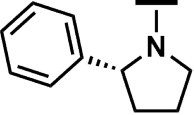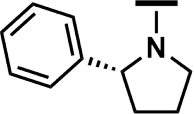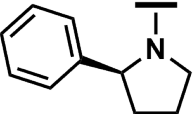| Reagent	Scale [mg]	Time [min.]	Conv. [%][a]	Yield [%][b]	Product
rac-**7a**	350	60	>97	48	*rac*-**8a**
(R)-**7a**	145	50	>97	>99	*(R)*-**8a**
7b	199	90	>97	67	**8b**
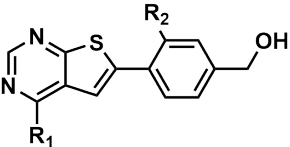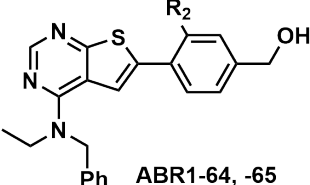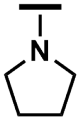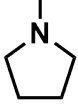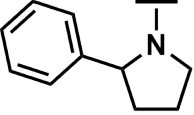
[a] Conversion determined by ^{1}H NMR.
[b] Isolated yield.

The low yield in the synthesis of *rac*-**8a** and **8b**, compared to *(R)*-**8a**, was due to poor solubility of these compounds, making the purification processes difficult.

## 3.10   *In vitro* testing of target compounds

The biological activity of the target compounds was tested towards EGFR-TK and CSF1R-TK in an enzymatic assay. Single-point analyses were performed at 100 nM and 500 nM concentration of the inhibitors for EGFR and CSF1R, respectively, and the results of these tests are given in Tables 3.7 and 3.9. Two previously synthesized thieno[2,3-*d*]pyrimidines, compounds **ABR1-65** and **ABR1-64**,[140] are also included in this biological evaluation. Table 3.8 lists the EGFR inhibitory potency of some reference compounds.

Table 3.7: EGFR (ErbB1) inhibitory potency of target compounds **6a-b** and **8a-b**, in addition to previously synthesized compounds **ABR1-65** and **ABR1-64**.[140]

Comp.	R_1	R_2	ee [%]	Inhibition[a] [%]	IC_{50} [nM]
6b		H	-	0	>1000
8b		OMe	-	-4	-
rac-**6a**		H	-	45	-
rac-**8a**		OMe	-	92	-
(R)-**6a**		H	65	73	30.6
(R)-**8a**		OMe	84	92	5.8
(S)-**6a**		H	65	5	811
ABR1-65[140]	-	H	-	8	860
ABR1-64[140]	-	OMe	-	49	115

[a] Mean value of two measurements at 100 nM concentration of the inhibitor.

Compounds *rac-* and *(R)-***8a** show promising results from the single-point analyses, with high activity towards EGFR. Considering the divergences in activity between *rac-*, *(R)-* and *(S)-***6a**, it appears that the chirality of the phenyl moiety has a significant effect on the EGFR-TK activity. As expected, the *(S)*-enantiomer has low activity, compared to the racemate, while the *(R)*-enantiomer appears to be more favourable. Figure 3.6 presents the IC_{50}-curves for compounds *(R)-* and *(S)-***6a**, and *(R)-***8a**, further illustrating the effect of the stereochemistry of the phenyl group on the activity of compounds *(R)-* and *(S)-***6a**.
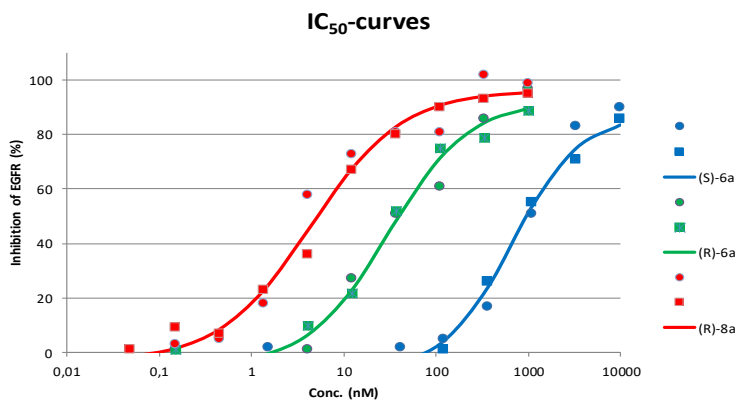
Figure 3.6: IC_{50}-curves for compounds *(R)-* and *(S)-***6a**, and *(R)-***8a**.

The positive effect of a correctly spaced phenyl moiety towards EGFR-TK is visible in the low activity of compounds **6b** and **8b**, compared to compounds *rac-***6a** and *rac-***8a**. The removal of the 2-substituted phenyl group proves to be detrimental for the EGFR-TK activity, which demonstrates the significance of this substituent to the activity of these compounds.

The effect of the methoxy moiety is visible by comparing *(R)-***6a** with compound *(R)-***8a**, and **ABR1-65** with **ABR1-64**. As illustrated by the IC_{50}-curves in Figure 3.6, the methoxy group appears to contribute favourably to the EGFR-TK activity, with a significant increase in activity from *(R)-***6a** to *(R)-***8a**. The contribution of the methoxy moiety has also been demonstrated in reference compounds *(R)-***5e** and **5t**.[10,11] (Table 3.8).

One objective of the thesis was to investigate if secondary amines are essential substituents at C-4. Based on the inhibition data for compounds *(R)-***5n** and **29n** (Table 3.8) and other previous work,[141] Bugge *et al.*[11] concluded that a secondary amine is essential for the EGFR-TK inhibitory activity; however, the findings of this master's thesis demonstrate that it is possible to achieve high activity towards EGFR with a tertiary heterocyclic amine. A hypothesis is that the EGFR-activity of thieno[2,3-*d*]pyrimidines is related to the conformation of the amino group, as opposed to the hydrogen donor ability. Further modelling and X-ray co-crystal structure determination are needed to fully answer these questions.

Table 3.8: EGFR inhibitory potency of reference compounds *(R)*-**1w**, -**5n**, -**29n**, -**5e**, and **5t**.[10,11]
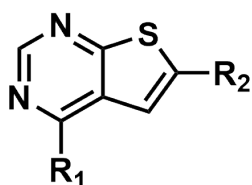
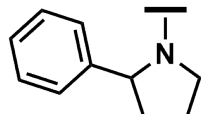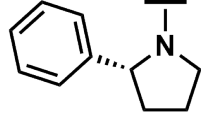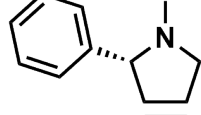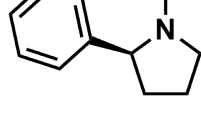| Comp.	R$_1$	R$_2$	Inhibition[a] [%]	IC$_{50}$[b] [nM]	SD[c]
(R)-**1w**[10]		**Br**	0	-	-
(R)-**5n**[11]			79	35	14
(R)-**29n**[11]			5	>1000	-
(R)-**5e**[11]			86	7	1
(R)-**5t**[11]			100	0.7[d]	0.2
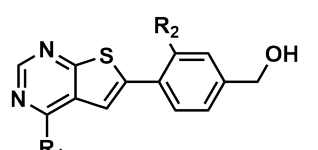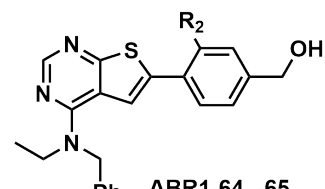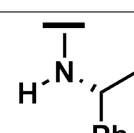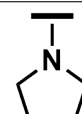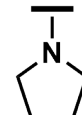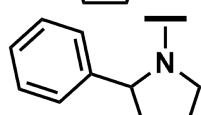
[a] Mean value of two measurements at 100 nM concentration of the inhibitor.
[b] Mean value of two titration curves (20 data points), unless otherwise stated.
[c] Standard deviation.
[d] Mean value of six titration curves (60 data points).

Table 3.9: CSF1R (FMS) inhibitory potency of reference compound *(R)*-**5t**,[11] target compounds **6a-b** and **8a-b**, and previously synthesized compounds **ABR1-65** and **ABR1-64**.[140]

(R)-**5t, 6a-b, 8a-b**   **ABR1-64, -65**

Comp.	R_1	R_2	ee [%]	Inhibition[a] [%]
(R)-**5t**[11]	(H–N, Ph)	OMe	ND	2
6b	(pyrrolidine N)	H	-	55
8b	(pyrrolidine N)	OMe	-	68
rac-**6a**	(2-phenylpyrrolidine N)	H	-	41
rac-**8a**	(2-phenylpyrrolidine N)	OMe	-	71
(R)-**6a**	(2-phenylpyrrolidine N)	H	65	65
(R)-**8a**	(2-phenylpyrrolidine N)	OMe	84	86
(S)-**6a**	(2-phenylpyrrolidine N)	H	65	20
ABR1-65[140]	-	H	3	
ABR1-64[140]	-	OMe	11	

[a] Mean value of two measurements at 500 nM concentration of the inhibitor.

Compound *(R)*-**8a** seems to possess good CSF1R activity, and is considerably more active than the racemic *rac*-**8a**. Looking at structures *(R)*-, *(S)*- and *rac*-**6a**, the conformation of the phenyl moiety on the C-4 substituent appears to have an impact on the CSF1R activity. However, compounds **6b** and **8b** without the phenyl group have activity almost on a par with, or higher than compounds *rac*-**6a** and -**8a**. It may be that the conformation of the pyrrolidine moiety is of importance, and that the addition of a 2-substituted phenyl group contributes either positively or negatively to the CFS1R activity, depending on the chirality of this substituent.

3.11 Compound characterization

Compounds **5-8**, excluding *rac*-**5a** and *rac*-**6a**, were previously unknown, and their structures are therefore elucidated in this section. Compounds *rac*-**5a** and *rac*-**6a** were synthesized as part of the specialization project of autumn 2017.[140] However, due to contamination of the final products, these structures have been resynthesized in this master project and are therefore included in this structure elucidation. ^{1}H NMR, ^{13}C NMR, IR and MS have been used for elucidation and confirmation of the target structures, and the spectra from these analyses can be found in the appendices.

General remarks

IR spectroscopy

IR spectroscopy of the elucidated compounds show generally similar results, and so the spectra are not discussed individually. Instead, some general remarks are discussed in this section.

Aromatic C-H and C=C stretches are visible in all spectra at 3000 cm^{-1}, along with aromatic C-H bending at 860-680 cm^{-1}. For compounds *(R)*- and *rac*-**7a**, a characteristic C=O stretch for aldehyde is visible at $\sim$ 1690 cm^{-1}. For compounds **6a-b** and **8a-b** a broad O-H stretch at 3307-3314 cm^{-1}, along with the strong C-O stretches at 1050-1150 cm^{-1}, suggest the presence of an alcohol group.[142]

NMR

A general procedure for the structure elucidation of the target compounds from ^{1}H, ^{13}C, and 2D NMR, is elaborated in this section. Numbering of positions in the thieno[2,3-*d*]pyrimidine and pyrrolidine moiety are given in Figure 3.7
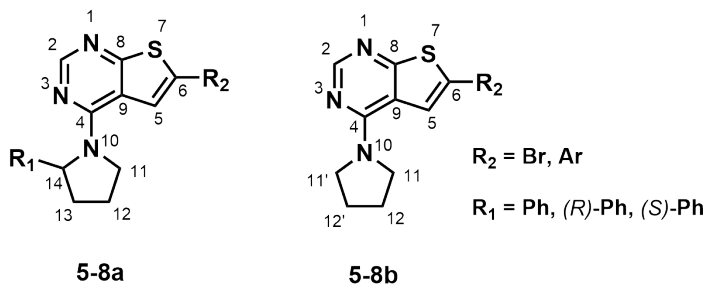
Figure 3.7: General numbering of positions in compounds **5-8**

The C-2 proton of the thieno[2,3-*d*]pyrimidine moiety is easily spotted in the ^{1}H spectra as a singlet at 8.20-8.30 ppm, and is used as a landmark in the structure elucidations. From this proton, the C-4, C-8, and C-9 carbons may be assigned from long-range ^{1}H-^{13}C coupling (HMBC), as illustrated in Figure 3.8. Comparing this signal in the spectra of compounds **5-8a** and **5-8b** show a significant broadening in the signals of the former. This broadening is also applied to the proton on C-5, as it is easily spotted as a singlet at 7.74-8.17 ppm for compounds **5-8b** while appearing as a very broad signal in the aromatic region of compounds **5-8a**. These differences clearly show the effect of dynamic processes in the structures of compounds **5-8a** as a result of the 2-substituted phenyl group.

The broad signal of the C-5 proton in compounds **5-8a** makes it difficult to assign in these structures; however, the HMBC couplings to this proton are clearly visible in the spectra of compounds **5-8b**, which also allow for the identification of carbon C-6 by cross-referencing with the long-range coupling of the C-2 proton to C-8 and C-9 (see Figure 3.8). While these HMBC couplings are not always visible, especially in the spectra of compounds **5-8a**, the shifts of the thieno[2,3-*d*]pyrimidine moiety remain fairly consistent in all of the structures.
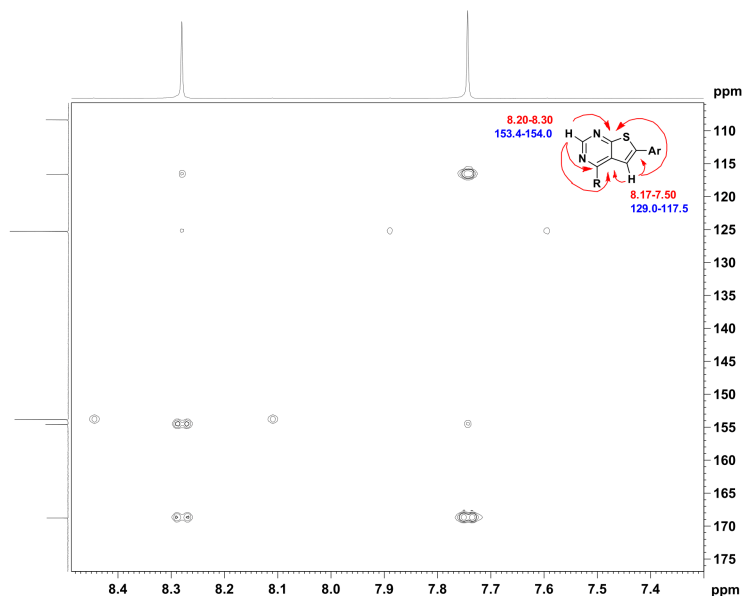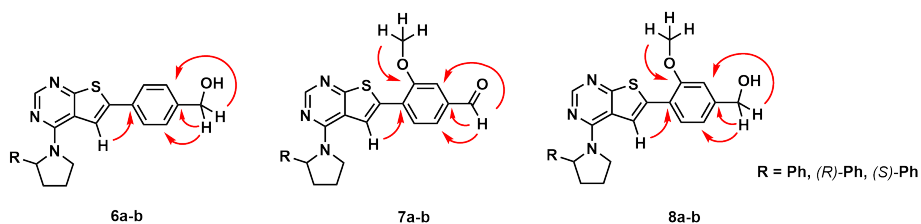
Figure 3.8: HMBC coupling of the proton on C-2 (8.29 ppm) to carbons C-4 (155.5 ppm), C-8 (167.3 ppm), and C-9 (117.4 ppm), and of the proton on C-5 (7.90 ppm) to C-6 (136.6 ppm), C-8 (167.3 ppm), and C-9 (117.4 ppm) in compound **6b**.

The protons on C-12 and C-13 for compounds **5-8a** are visible as broad signals in ^{1}H NMR. The associated carbons of these signals are poorly visible in all the ^{13}C spectra, and the C-H couplings are not visible in the HSQC spectra. These shifts are assigned from HMBC by coupling of the C-14 proton to C-12 and C-13, as illustrated in Figure 3.9. Although this coupling is only observed in the HMBC spectrum of *rac*-**4**, these shifts should be applicable to the other compounds as the introduction of the aryl-group at C-6 does not appear to have any significant impact on the chemical shifts of the C-4 substituent in Tables 3.10-3.17.
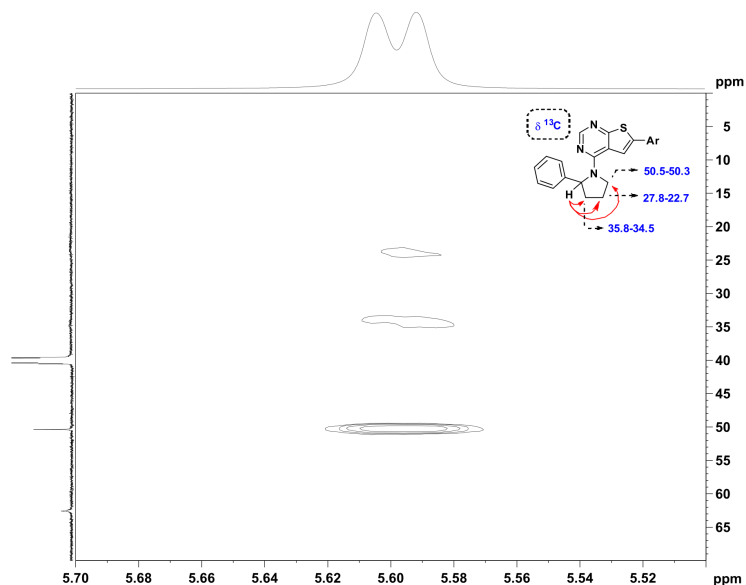
Figure 3.9: HMBC coupling of the C-14 proton to C-11-C-13 in compound *rac*-**4**

The aromatic shifts of compounds **6-8** may be difficult to elucidate, especially when attempting to distinguish these from the aromatic group on the C-4 substituent of compounds **6-8a**. Some key signals to help assign these shifts by HMBC are the CH$_2$OH protons for compounds **6a-b** and **8a-b**, the aldehyde proton for compounds **7a-b**, and the methoxy protons for compounds **7a-b** and **8a-b**. Figure 3.10 illustrates the HMBC couplings of these protons to the C-6 aromatic carbons. The proton on C-5 also helps identify the shifts on the C-6 aryl group by HMBC coupling to the nearest carbon on the aromatic ring, as illustrated in Figure 3.10.

Figure 3.10: HMBC coupling of the CH$_2$OH-, aldehyde-, and methoxy protons to the aromatic ring on C-6 in compounds **6-8**.

COSY spectra are primarily used to confirm the proton shifts in the pyrrolidine moiety on C-4, and to identify the neighbouring CH$_2$ protons to the alcohol group, as illustrated in Figure 3.11.

6a-b, 8a-b
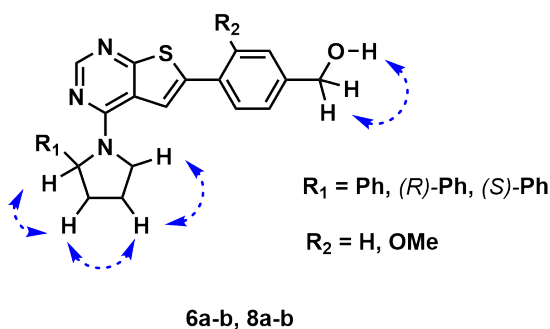
Figure 3.11: COSY coupling of the CH_2 and OH protons, and of the protons on C-11-C-13 in compounds **6a-b** and **8a-b**.

3.11.1 Compound *rac*-5a

Compounds *rac*-, *(R)*-, and *(S)*-**5a** have near identical ^{1}H and ^{13}C NMR shifts, and only the structure elucidation of *rac*-**5a** is therefore presented in this section. The structure of *rac*-**5a** with numbering positions is illustrated in Figure 3.12.

NMR

The results from ^{1}H and ^{13}C NMR of *rac*-**5a** are presented in Table 3.10. Long range ^{1}H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compounds *rac*-, *(R)*-, and *(S)*-**5a** are given in Appendix H to J.
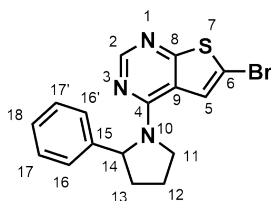
Figure 3.12: Numbering of positions in compound *rac*-**5a**

Table 3.10: Assigned ^{1}H NMR (400 MHz) and ^{13}C NMR (150 MHz) shifts for compound *rac*-**5a** in DMSO-d_6.

Position	[13]C [ppm]	[1]H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.7	8.24 (s, 1H)		
4	154.7			
5	128.9	7.78 (br s, 1H)		
6	108.7			
8	169.1			
9	116.6			
11	50.4	4.20 (br s, 1H) 3.92 (br s, 1H)	11, 12	
12	24.0[a]	2.00 (br s, 1H) 1.89 (br s, 1H)		
13	34.4[b]	2.40 (br s, 1H) 1.89 (br s, 1H)		
14	62.6	5.60 (d, J = 7.7, 1H)	13,16/16'	4,11-16
15	143.6			
16/16'	125.9	7.23-7.21 (m, 2H)	14,17,18	14-18
17/17'	128.9	7.31-7.30 (m, 2H)	16,17	15-18
18	127.1	7.23-7.21 (m, 1H)	17	16,17

[a] Poorly visible in ^{13}C NMR. Shift confirmed by HMBC.
[b] Poorly visible in ^{13}C NMR. Shift confirmed by HMBC.

3.11.2 Compound 5b

The structure of **5b** with numbering positions is illustrated in Figure 3.13.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.11. Long range [1]H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compound **6b** are given in Appendix K.
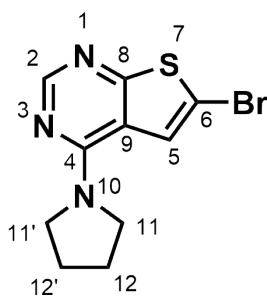
Figure 3.13: Numbering of positions in compound **5b**

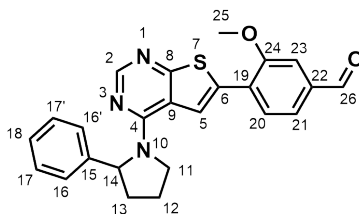**Table 3.11:** Assigned ^{1}H NMR (600 MHz) and ^{13}C NMR (150 MHz) shifts for compound **5b** in DMSO-d_6.

Position	^{13}C [ppm]	^{1}H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.8	8.28 (s, 1H)		4,8,9
4	154.6			
5	125.3	7.74 (s, 1H)		8,9
6	108.3			
8	168.8			
9	116.6			
11/11'	49.1	3.71 (br s, 4H)	12/12'	
12/12'	25.5	1.96 (br s, 4H)	11/11'	

3.11.3 Compound *rac*-6a

Compounds *rac*-, *(R)*-, and *(S)*-**6a** have near identical ^{1}H and ^{13}C NMR shifts, and only the structure elucidation of *rac*-**6a** is therefore presented in this section. The structure of *rac*-**6a** with numbering positions is illustrated in Figure 3.14.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.14. Long range ^{1}H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compounds *rac*-, *(R)*-, and *(S)*-**6a** are given in Appendix L to O.
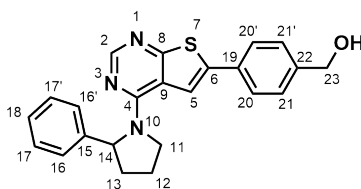
Figure 3.14: Numbering of positions in compound *rac*-**6a**

Table 3.12: Assigned ^{1}H NMR (600 MHz) and ^{13}C NMR (150 MHz) shifts for compound *rac*-**6a** in DMSO-d_6.

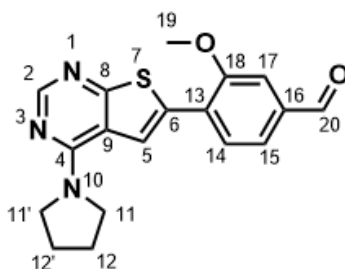| Position	^{13}C [ppm]	^{1}H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.5	8.27 (br s, 1H)		
4	155.6			
5	128.9	7.56-7.53 (m, 1H)		19
6	136.7			
8	167.6			
9	117.9			
11	50.5	4.29 (br s, 1H) 4.0 (br s, 1H)		
12	24.0[a]	1.93 (br s, 1H) 2.04 (br s, 1H)	13	
13	34.4[b]	2.43 (br s, 1H) 1.93 (br s, 1H)	12	
14	62.9	5.70 (br s, 1H)		
15	144.0			
16/16'	126.0	7.28 (br s, 2H)		
17/17'	129.0	7.31-7.30 (m, 2H)	15	
18	127.1	7.21-7.19 (m, 1H)		16/16'
19	132.5			
20/20'	132.0	7.64-7.61 (m, 2H)		
21/21'	127.6	7.39-7.38 (m, 2H)	23	19,23
22	143.5			
23	62.9	4.54 (d, J = 5.4 Hz, 2H)	21,OH	21,22
OH		5.27 (t, J = 5.7 Hz, 1H)	23	22,23

[a] Poorly visible in ^{13}C NMR. Shift confirmed by HMBC.
[b] Poorly visible in ^{13}C NMR. Shift confirmed by HMBC.

3.11.4   Compound 6b

The structure of **6b** with numbering positions is illustrated in Figure 3.15.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.13. Long range ^{1}H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compound **6b** are given in Appendix P.
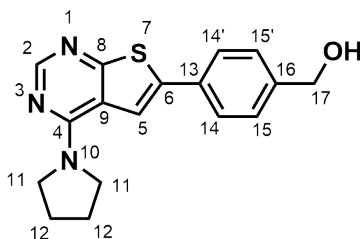
Figure 3.15: Numbering of positions in compound **6b**

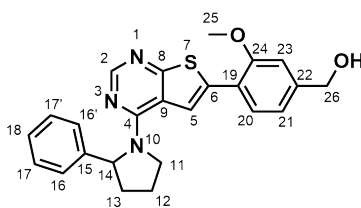**Table 3.13:** Assigned ^{1}H NMR (600 MHz) and ^{13}C NMR (150 MHz) shifts for compound **6b** in DMSO-d_6.

Position	^{13}C [ppm]	^{1}H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.6	8.29 (s, 1H)		4,8,9
4	155.5			
5	117.5	7.90 (s, 1H)		6,8,9,13,
6	136.6			
8	167.3			
9	117.4			
11/11'	49.2	3.82 (br s, 4H)	12/12'	
12/12'	25.5	1.99 (br s, 4H)	11/11'	
13	132.1			
14/14'	126.1	7.75-7.33 (m, 2H)	15/15'	6,14/14',16
15/15'	127.5	7.40-7.39 (m, 2H)	14/14',17	13,15,17
16	143.4			
17	63.0	4.54 (d, J = 5.7 Hz, 2H)	15/15',OH	15/15',16
OH		5.26 (t, J = 5.7 Hz, 1H)	17	16,17

3.11.5 Compound *rac*-7a

Compounds *rac*-, and *(R)*-**7a** have near identical ^{1}H and ^{13}C NMR shifts, and only the structure elucidation of *rac*-**7a** is therefore presented in this section. The structure of *rac*-**7a** with numbering positions is illustrated in Figure 3.16.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.16. Long range ^{1}H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compounds *rac-*, and *(R)*-**7a** are given in Appendix Q to R.

Figure 3.16: Numbering of positions in compound *rac-***7a**

Table 3.14: Assigned ^{1}H NMR (600 MHz) and ^{13}C NMR (150 MHz) shifts for compound *rac-***7a** in DMSO-d_6.

Position	[13]C [ppm]	[1]H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.9	8.31 (1H)		
4	155.7			
5	122.3	7.80-7.55 (m, 1H)		
6	131.1			
8	168.8			
9	116.0			
11	50.4	4.26 (br s, 1H) 4.0 (br s, 1H)		
12	27.8 [a]	1.95 (br s, 1H) 2.04 (br s, 1H)	13	
13	35.7 [b]	2.44 (br s, 1H) 1.95 (br s, 1H)	12	
14	62.7	5.72 (br s, 1H)		
15	143.8			
16/16'	126.0	7.33-7.29 (m, 2H)		
17/17'	129.0	7.33-7.29 (m, 2H)		
18	127.2	7.23-7.20 (m, 1H)		16/16'
19	127.9			
20	129.0	7.80-7.55 (m, 1H)		
21	123.3	7.80.7.55 (m, 1H)		
22	137.0			
23	112.3	7.80-7.55 (m, 1H)		
24	156.2			
25	56.6	4.0 (br s, 3H)		
26	192.7	10.0 (s, 1H)		21,22,23

[a] Poorly visible in ^{13}C NMR. Shift confirmed by HSQC.
[b] Poorly visible in ^{13}C NMR. Shift confirmed by HSQC.

3.11.6 Compound 7b

The structure of **7b** with numbering positions is illustrated in Figure 3.17.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.15. Long range [1]H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compound **7b** are given in Appendix S.

Figure 3.17: Numbering of positions in compound **7b**

Table 3.15: Assigned ^{1}H NMR (600 MHz) and ^{13}C NMR (150 MHz) shifts for compound **7b** in DMSO-d_6.

Position	^{13}C [ppm]	^{1}H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	154.0	8.33 (s, 1H)		4,8,9
4	155.6			
5	121.9	8.17 (s, 1H)		6,9,13
6	130.9			
8	168.6			
9	116.0			
11/11'	49.2	3.85 (br s, 4H)		
12/12'	25.5	2.01 (br s, 4H)		
13	128.1			
14	129.1	8.14-8.13 (m, 1H)	15	6,16,18
15	123.3	7.63-7.60 (m, 1H)	14	13,17,18,20
16	136.9			
17	112.2	7.63-7.60 (m, 1H)		13,15,18,20
18	156.1			
19	56.7	4.04 (s, 3H)		17,18
20	192.7	10.02 (s, 1H)		

3.11.7   Compound *rac*-8a

Compounds *rac*-, and *(R)*-**8a** have near identical ^{1}H and ^{13}C NMR shifts, and only the structure elucidation of *rac*-**8a** is therefore presented in this section. The structure of *rac*-**8a** with numbering positions is illustrated in Figure 3.18.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.18. Long range ^{1}H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compounds *rac*-, and *(R)*-**8a** are given in Appendix T to U.

Figure 3.18: Numbering of positions in compound *rac*-**8a**

Table 3.16: Assigned ¹H NMR (600 MHz) and ¹³C NMR (150 MHz) shifts for compound *rac*-**8a** in DMSO-d_6.

Position	¹³C [ppm]	¹H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.3	8.28 (s, 1H)		
4	155.5			
5	120.4	7.86-7.53 (m, 1H)		
6	132.0			
8	167.9			
9	116.1			
11	50.3	4.22 (br s, 1H) 3.98 (br s, 1H)	12	
12	22.7 ª	1.92 (br s, 1H) 2.01 (br s, 1H)	11	
13	35.5 ᵇ	2.42 (br s, 1H) 1.92 (br s, 1H)	14	
14	62.6	5.69 (s, 1H)	13	
15	144.0			
16/16'	126.0	7.32-7.31 (m, 2H)		14
17/17'	129.0	7.32-7.31 (m, 2H)		
18	127.1	7.22-7.20 (m, 1H)		
19	128.3			
20	129.0	7.86-7.53 (m, 1H)		
21	119.3	6.98-6.97 (m, 1H)		23,26
22	145.1			
23	110.5	7.10 (s, 1H)		21,26
24	155.9			
25	56.2	3.87 (s, 3H)		
26	63.0	4.53 (d, J = 5.0 Hz, 2H)	OH	21,22,23
OH		5.29 (t, J = 5.5 Hz, 1H)	26	22,26

ª Poorly visible in ¹³C NMR. Shift confirmed by HSQC.
ᵇ Poorly visible in ¹³C NMR. Shift confirmed by HSQC.

3.11.8   Compound 8b

The structure of **8b** with numbering positions is illustrated in Figure 3.19.

NMR

The results from ^{1}H and ^{13}C NMR are presented in Table 3.17. Long range [1]H-^{13}C coupling was used to assign quaternary carbons. ^{1}H NMR, ^{13}C NMR, IR and MS spectra for compound **8b** are given in Appendix V.
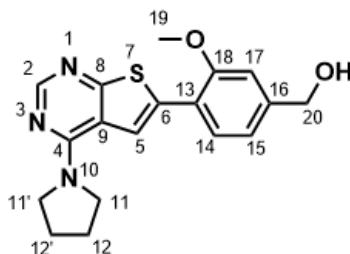
Figure 3.19: Numbering of positions in compound **8b**

Table 3.17: Assigned ^{1}H NMR (600 MHz) and ^{13}C NMR (150 MHz) shifts for compound **8b** in DMSO-d_6.

Position	[13]C [ppm]	[1]H [ppm] (mult., J [Hz], int.)	COSY	HMBC
2	153.4	8.29 (s, 1H)		4,8
4	155.9			
5	118.9	7.95 (s, 1H)		4,6,8,9,13
6	132.7			
8	167.7			
9	116.1			
11/11'	49.1	3.81 (br s, 4H)		
12/12'	25.5	2.0 (br s, 4H)		
13	120.6			
14	128.4	7.78 (s, 1H)	15	6,16,17,18
15	119.3	7.01 (s, 1H)	14	13,17,20
16	145.0			
17	110.5	7.13 (s, 1H)		13,16,18,20
18	155.9			
19	56.2	3.95 (s, 3H)		
20	63.1	4.55 (d, J = 5.7 Hz, 1H)	OH	15,16,17
OH		5.29 (t, J = 5.8 Hz, 1H)	20	16,20

4 | Concluding Remarks and Further Work

4.1 Summary

Investigation of several potential synthetic routes to *(R)*- and *(S)*-2-phenylpyrrolidine (*(R)*- and *(S)*-**4**) found palladium-catalyzed α-arylation to be a viable and simple method for obtaining these compounds. The compounds were synthesized from pyrrolidine, starting with Boc protection to form compound **1** in 92-94 % yield. Palladium-catalyzed α-Arylation of **1** gave the Boc-protected 2-phenylpyrrolidines *(R)*- and *(S)*-**3** in 62 % and 70 % yield, and with 81 % and 96 % ee, respectively. Differences in ee of *(R)*- and *(S)*-**3** is suspected due to divergences in the enantiopurity of (-)- and (+)-sparteine. Boc deprotection of compounds *(R)*- and *(S)*-**3** with TFA gave *(R)*- and *(S)*-**4** in 56 % and 49 % yield, respectively.

Racemic 2-phenylpyrrolidine (*rac*-**4**) was prepared from pyrrolidin-2-one in a two-step, one-pot procedure involving a silylation and Grignard reaction, followed by an imine reduction of the intermediary 2-phenylpyrroline (**2**). Achieving full conversion of starting material during the initial silylation proved challenging, and some erratic results were observed with the use of the Grignard reagent. Suspected causes for these issues are the attachment of Me_3Si to both the nitrogen and the oxygen of pyrrolidin-2-one, and possible interactions between the Grignard reagent and the silyllactam at silicon. Attempted optimization of the one-pot reaction gave compound **2** in 12-36 % yield. Reduction of **2** with sodium borohydride gave *rac*-**4** in 72-89 % yield.

Nucleophilic aromatic substitution of 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine with amines *rac*-, *(R)*-, and *(S)*-**4**, in addition to pyrrolidine, gave compounds **5a-b** in 50-91 % yield. Suboptimal conversion of the thieno[2,3-*d*]pyrimidine was observed in the synthesis of *(R)*- and *(S)*-**5a**, likely due to the presence of TFA salt in these amines. A surplus of Hünig's base was added in the reaction of *(R)*-**5a** to help deprotonate the salt, which gave considerably higher conversion and yield compared to the reaction of *(S)*-**5a**.

Suzuki cross-coupling of thieno[2,3-*d*]pyrimidines **5a-b** was done using catalyst

Pd(PPh$_3$)$_4$, and yielded compounds **6-7** in 60-99 % yield. Dimerization of the boronic acid was observed in the reactions of *rac*-**6a** and -**7a**, likely due to oxidation of the palladium catalyst because of insufficient degassing of the solvents. Proper degassing resulted in minute amounts of by-product and increased the yield of later reactions.

Reduction of thieno[2,3-*d*]pyrimidine aldehydes **7a-b** to primary alcohols **8a-b** was done using sodium borohydride, and gave full conversion and 48-99 % yields. Loss of yield from workup and purification was due to poor solubility of these compounds.

Target compounds *(R)*- and *(S)*-**6a**, and *(R)*-**8a** were analysed by CSP-HPLC to check the ee of the final products. The ee was found to be 65 % for compounds *(R)*- and *(S)*-**6a** and 84 % for *(R)*-**8a**, marking a significant decrease in enantiopurity from compounds *(R)*- and *(S)*-**3**. A suspected cause is racemization in the S$_N$Ar reactions.

This masters's thesis has presented preliminary results in the structure-activity relationship study of thieno[2,3-*d*]pyrimidines with heterocyclic tertiary amines as a substituent on C-4. Of the 14 new thieno[2,3-*d*]pyrimidines, seven new thienopyrimidine-based inhibitor candidates were prepared. Single point analysis identified two of these compounds, *rac*- and *(R)*-**8a**, as potent EGFR-TK inhibitors with IC_{50} values of 30.6 and 5.8 nM, respectively. This demonstrates that it is possible to obtain high EGFR-TK activity with tertiary amines at C-4. The single-point analyses further indicate that the 2-substituted phenyl moiety on the amines may contribute favourably to the EGFR-TK activity, depending on the stereochemistry. The *(R)*-enantiomers were found to exhibit better inhibition compared to the racemates, while the *(S)*-enantiomer exhibited poor activity. Removal of the phenyl group proved detrimental for the EGFR-TK activity, which supports the proposed theory that the EGFR-TK activity of thieno[2,3-*d*]pyrimidines is related to the conformation of the C-4 amino group, as opposed to hydrogen donor ability.

Single point analyses on CSF1R activity were also performed, and found compound *(R)*-**8a** to possess good activity with 86 % inhibition. Analyses found compounds without the 2-substituted phenyl group on the amine (compounds **6b** and **8b**) to possess activity on a par with, or higher than the racemates. This may suggest that the conformation of the pyrroline moiety is of importance to the CSF1R activity, and that the 2-substituted phenyl group interacts favourably or unfavourably, depending on the chirality. The activity of compounds *(S)*-**6a** (5 %) and *(R)*-**6a** (73 %), with opposite stereochemistry of the phenyl group, supports this statement.

4.2   Further work

The discovery of compounds *rac*- and *(R)*-**8a** with high EGFR-TK facilitates further investigation of thieno[2,3-*d*]-, pyrrolo-, and furopyrimidines with a tertiary amine as the substituent on C-4. These structures may open new areas of research with the preparation of different heterocyclic or non-heterocyclic tertiary amines as the

substituent. As an extension of this master's thesis, further analysis, modelling and X-ray co-crystal structure determination of the compounds *rac-* and *(R)-***8a** should be performed for future SAR studies.

As the chirality of the 2-substituted phenyl moiety on the C-4 substituent has been found to influence the EGFR-TK activity, investigation of the possible racemization in the reactions from compounds *(R)-* and *(S)-***3** to target compounds *(R)-* and *(S)-***6a** and *(R)-***8a** is imperative for future studies. A viable method for this study would be to perform chiral analyses after each reaction step, to check for any changes in ee. Methods for preparing the tertiary amines with higher ee could also be further investigated. Some alternate synthetic procedures with high enantioselectivity have been presented in this master's thesis, including asymmetric hydrogenation and chemoenzymatic routes. A method for improving upon the enantiopurity of the already synthesized compounds by recrystallization has also been suggested, which may present compounds with improved EGFR-TK activity.

5 | Experimental

5.1 General

6-bromo-4-chlorothieno[2,3-*d*]pyrimidine had been synthesized and purified as part of previous studies in our research group, and the available compound was used in the reactions mentioned below.[143] Unless otherwise stated, all other reagents in this project were of synthetical of analytic grade and were supplied by Sigma-Aldrich.

5.1.1 Separation techniques

Thin-layer chromatography (TLC) was used to follow the reaction progress, and to check for conversion of starting materials (Silica gel on aluminium sheets, F_{254}). Silica gel column chromatography was performed using Silica gel 60A from EMD Millipore, pore size 40-63 μm.

5.1.2 Spectroscopic analysis

NMR analyses was performed using Bruker 400 MHz Avance III HD equipped with a 5-mm SmartProbe z-gradient probe and SampleCase, and Bruker 600 MHz Avance III HD equipped with a 5-mm cryogenic CP-TCI z-gradient probe and SampleCase. DMSO-d_6 was used as solvent for the analyses, and the chemical shifts, reported in δ (ppm), are relative to the solvent peak of DMSO-d_6 (2.50 ppm in ^{1}H NMR and 39.52 ppm in ^{13}C NMR[130]). Traces of water are usually present when using DMSO-d_6, and can be detected at 3.33 ppm (s) in ^{1}H NMR.[130] Traces of solvents may also be present in the spectra: EtOAc (1.99 ppm (s), 4.03 ppm (q), 1.17 ppm (t)), n-pentane (0.86 ppm (t), 1.27 ppm (m)), dichloromethane (5.76 ppm (s)), 1,4-dioxane (3.57 ppm (s)).[130] Apiezon brand H grease (0.82-0.88 ppm (m), 1.24 ppm (br s)) may also be present in some of the spectra.[130] The coupling constants (J) are given in Hz, and the signals are defined according to their multiplicity: s (singlet), d (doublet), t (triplet), q (quartet). Multiplets (m) are defined as intervals.

HPLC was performed using an Agilent 1100 Series instrument with a 1260 infinity degasser, 1100 Series quaternary pump, 1100 Series autosampler, and a 1200 Series diode array detector. The software used was Agilent ChemStation. Spectra were recorded for wavelengths 230 nm, 254 nm, and 280 nm. The column was an Agilent Eclipse XDB-C18 (5 μm, 4.6 x 150 mm) with a flow of 0.8 mL/min. The

method started with isocratic eluation with water:CH$_3$CN (90:10) for 5 min followed
by a linear gradient ending at water:CH$_3$CN (0:100) over 35 min. Chiral HPLC
analyses were conducted through two methods. Method A: Chiralpak AD 0.46 x 25
cm chiral column (part no. 00CE-FE167), eluting with hexane:i-PrOH, 99:1, flow
rate: 0.3 mL/min, detection at 254 nm. Method B: Lux 5u Cellulose-1 4.6 x 250
mm chiral column (part no. 00G-4459-E0), eluting with hexane (cont. 0.2% diethyl
amine):i-PrOH, 85:15, flow rate: 1.5 mL/min, detection at 254 nm.

Accurate mass determination in positive and negative mode was performed on a
"Synapt G2-S" Q-TOF instrument from Water^{TM}. Samples were ionized by the use
of ASAP probe (APCI) or ESI probe. Calculated exact mass and spectra processing
was done by Waters^{TM} Software Masslynx V4.1 SCN871.

The IR analyses was performed using Bruker Alpha ECO-ATR FTIR-spectrometer,
and processing software OPUS.

5.1.3   Melting point

The melting point analyses was performed using a Stuart automatic melting point
SMP40 instrument (1 °C/min).

5.1.4   Optical rotation

Due to technical issues with the polarimeter, optical rotation measurements were not
available for this master's thesis.

5.2   General procedure for α-arylations

N-Boc pyrrolidine (0.37 mL, 2.13 mmol) was dissolved in $tert$-butyl methyl ether (4.5
mL) and added (+)- or (-)-sparteine (500 mg, 2.1 mmol), depending on the desired
enantiomer. The solution was cooled to -78 °C using dry ice, before adding s-BuLi
(1.56 mL of a 1.4 M solution in cyclohexane, 2.18 mmol) dropwise slowly over 13
minutes. The solution was stirred at -78 °C for three hours. ZnCl$_2$ (1.3 mL of a 1 M so-
lution in diethyl ether, 1.3 mmol) was added dropwise slowly over 11 minutes, and the
solution was stirred at -78 °C for an 30 minutes. The mixture was allowed to warm to
room temperature and stir for an additional 30 minutes. Bromobenzene (0.20 mL, 1.91
mmol) was added, followed by Pd(OAc)$_2$ (20.5 mg, 0.10 mmol) and t-Bu$_3$P−HBF$_4$
(32.0 mg, 0.11 mmol), and the solution was stirred for 13 hours. Aqueous ammonia
(2.0 mL) was added, and the solution was stired for one hour. The mixture was filtered
through Celite and washed with $tert$-butyl methyl ether. The filtrate was washed with
1 N HCl and water, dried over MgSO$_4$, filtered, and concentrated in vacuo. The crude
product was purified by silica gel column chromatography.

5.3 General procedure for Boc-deprotection

Compound *(R)*- or *(S)*-**3** (227-293 mg, 0.92-1.19 mmol) was dissolved in CH_2Cl_2 (10 mL), and cooled to 0 °C. TFA (19.6-24.8 mmol) was added, and the solution was stirred at room temperature for 4 hours. The solvent was removed in a rotary evaporator, before adding EtOAc (15 mL), 2M NaOH (10 mL), and NEt_3 (2.5 mL). The phases were separated, and the aquatic phase was extracted with more EtOAc.

5.4 General procedure for nucleophilic aromatic substitution

6-Bromo-4-chlorothieno[2,3-*d*]pyrimidine (0.3-2.5 g) was mixed with the selected amine (~1.5 eq.), Hünig's base (1.5-14 eq.), and *i*-PrOH (2-8 mL). The solution was stirred at 80 °C under a N_2 atmosphere until full conversion. The mixture was then cooled to room temperature and diluted with water (40 mL) and diethyl ether (40 mL). After phase separation, the water phase was extracted with more diethyl ether (2 x 40 mL). The combined organic phase was washed with saturated aq NaCl solution (15-20 mL), filtered and concentrated in vacuo. The crude product was purified by drying under reduced pressure to constant weight, and by silica gel column chromatography.

5.5 General procedure for Suzuki cross-coupling

Compound **5a-b** (100-500 mg) was mixed with the selected arylboronic acid (1.2 eq.), fine powdered K_2CO_3 (3 eq.), $Pd(PPh_3)_4$ (0.01 eq.) and 1,4-dioxane:water (1:1 by vol.%, 2-12 mL). The 1,4-dioxane:water mixture was degassed using nitrogen. The reaction was stirred at 80 °C for 4.5-20 hours under a N_2 atmosphere. The solvent was removed and the product was diluted with water (50-100 mL), and diethyl ether or dichloromethane (50-100 mL). After phase separation the water phase was extracted with more diethyl ether or dichloromethane (2 x 25 mL). The combined organic phase was washed with saturated aq NaCl solution (25-50 mL), filtered and concentrated in vacuo. The crude product was purified by drying under reduced pressure to constant weight and by silica gel column chromatography.

5.6 General procedure for reduction using NaBH$_4$

Compound **7a-b** was dissolved in THF:MeOH (3:1, 20 mL) and mixed with $NaBH_4$ (1 eq.). The reaction was stirred for 30 minutes at room temperature before additional $NaBH_4$ (1 eq.) was added. The mixture was stirred for another 30 minutes, and concentrated to about 5 mL. The residue was diluted with EtOAc (50 mL) and water (50 mL), and the water phase was extracted with more EtOAc (2 x 25 mL). The combined organic phase was washed with saturated aq NaCl solution (25 mL), filtered and concentrated in vacuo. The product was purified by drying under reduced pressure to constant weight.

5.6.1 *N*-Boc-pyrrolidine (1)[126]

Pyrrolidine (1.0 mL, 12 mmol) was added to di-*tert*-butyl
dicarbonate (2.62 g, 12.0 mmol), and the mixture was
stirred at room temperature for 2 minutes. The solu-
tion was diluted with water, and extracted with EtOAc.
The organic phase was washed with saturated aq. NaCl
solution, filtered and concentrated in vacuo to yield
1.89 g (11.0 mmol, 92 %) of compound **1** as a yellow
oil.
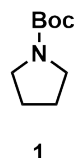
Spectroscopic data (Appendix A): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 3.19-
3.18 (m, 4H), 1.77-1-76 (m, 4H), 1.39 (s, 9H). ^{1}H NMR data was in accordance with
previously reported values.[126]

5.6.2 *(R)*-*N*-Boc-2-phenylpyrrolidine (*(R)*-3)[52][86]

Compound *(R)*-**3** was prepared as described in section 5.2, using
(-)-sparteine (500 mg, 2.13 mmol). The reaction was left for
13 hours, and purification by silica gel column chromatography
(*n*-pentane:EtOAc, 6:1, R_f = 0.48) yielded 329 mg (1.33 mmol, 62
%) of *(R)*-**3** as a white, crystalline solid; mp = 62.6-63.5 °C (lit.[86]
61.9-62.7 °C); Chiral HPLC (Method A): t_R = 24.9 (major), t_R =
26.7 (minor).
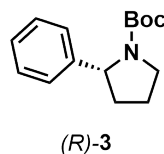
Spectroscopic data (Appendix C): ^{1}H NMR (400 MHz, DMSO-d_6) (40:60 mix-
ture of rotamers) δ: 7.33-7.29 (m, 2H), 7.23-7.15 (m, 3H), 4.82 (br s, 0.4H), 4.70
(br s, 0.6H), 3.56-3.46 (m, 2H), 2.30 (br s, 1H), 1.81-1.78 (m, 2H), 1.70 (br s, 1H),
1.40 (s, 4H), 1.09 (s, 5H). ^{1}H NMR data was in accordance with previously reported
values.[86][52][89]

5.6.3 *(R)*-2-Phenylpyrrolidine (*(R)*-4)[52]

Compound *(R)*-**4**) was prepared as described in
Section 5.3, starting with *(R)*-**3** (293 mg, 1.19
mmol). The reaction time was 4 hours, and pu-
rification by silica gel column chromatography (*n*-
pentane:EtOAc, 9:1, R_f = 0.11) yielded 248 mg (0.66
mmol, 56 %) of *(R)*-**4**. The product was an orange
oil.
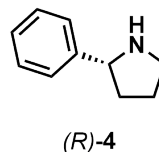
Spectroscopic data (Appendix F): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 7.42-7.40
(m, 2H), 7.38-7.34 (m, 2H), 7.31-7.27 (m, 1H), 4.26 (t, J = 8.1, 1H), 3.20-3.14 (m,
1H), 3.13 (br s, 1H) (missing), 3.10-3.04 (m, 1H), 2.27-2.19 (m, 1H), 1.98-1.81 (m,
2H), 1.77-1.67 (m, 1H). ^{1}H NMR data was in accordance with previously reported
values.[144]

5.6.4   *(S)-N*-Boc-2-phenylpyrrolidine (*(S)*-3)[52][86]

Compound *(S)*-**3** was prepared as described in section 5.2, using (+)-sparteine (500 mg, 2.13 mmol). The reaction was for 17 hours, and purification by silica gel column chromatography (*n*-pentane:EtOAc, 6:1, R_f = 0.67) yielded 374 mg (70 %, 1.51 mmol) of *(S)*-**3** as a white, crystalline solid; mp = 62.6-63.5 °C (lit.[86] 61.9-62.7 °C); Chiral HPLC (Method A): t_R = 24.9 (minor), t_R = 26.7 (major).
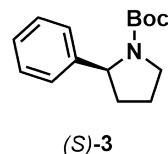
Spectroscopic data (Appendix D): ^{1}H NMR (400 MHz, DMSO-d_6) (40:60 mixture of rotamers) δ 7.33-7.29 (m, 2H), 7.23-7.15 (m, 3H), 4.82 (br s, 0.4H), 4.70 (br s, 0.6H), 3.56-3.46 (m, 2H), 2.30 (br s, 1H), 1.80-1.78 (m, 2H), 1.70 (br s, 1H), 1.40 (s, 4H), 1.09 (s, 5H). ^{1}H NMR data was in accordance with previously reported values.[86][52][89]

5.6.5   *(S)*-2-Phenylpyrrolidine (*(S)*-4)

The synthesis of compound *(S)*-**4** was carried out through two different synthetic routes:

Boc deprotection[52]

Compound *(S)*-**4** was prepared as described in Section 5.3, starting with *(S)*-**3** (227 mg, 0.92 mmol). The reaction time was 4 hours, and purification by silica gel column chromatography (*n*-pentane:EtOAc, 9:1, R_f = 0.11) yielded 168 mg (0.45 mmol, 49 %) of *(S)*-**4**. The product was an orange oil.
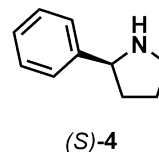
Chiral resolution[54]

Compounds *(rac)*-**4** was dissolved in EtOH:EtOAC (35 %, 6 mL) and added Dibenzoyl-*D*-tartaric acid (1.24 g, 3.45 mmol). The solution was heated to reflux for 10 minutes, and then allowed to cool to room temperature for 24 hours. The solution was refrigerated for an additional 24 hours, and the crystals were then collected and rinced with cold EtOAc. The crystals were dissolved in 2 M NaOH:CH$_2$Cl$_2$ (50 mL) and stirred vigorously for 15 minutes. Phase separation and extraction with CH$_2$Cl$_2$ (10 mL), followed by concentration in vacuo, yielded 500 mg (3.40 mmol, 49 %) of *(S)*-**4** as an orange oil; Chiral HPLC (Method A): t_R = 22.3 (minor), t_R = 23.5 (major).

Spectroscopic data (Appendix G): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 7.43-7.42 (m, 2H), 7.39-7.36 (m, 2H), 7.32-7.29 (m, 1H), 4.33-4.28 (m, 1H), 3.23-3.17 (m, 1H), 3.14-3.07 (m, 1H), 3.13 (br s, 1H) (missing), 2.28-2.21 (m, 1H), 1.99-1.85 (m, 2H), 1.81-1.73 (m, 1H). ^{1}H NMR data was in accordance with previously reported values.[144]

5.6.6   2-Phenylpyrroline (2)[57]

Pyrrolidin-2-one (524 mg, 6.16 mmol) was dissolved in dry ethyl
ether (14 mL) and added NEt$_3$ (0.90 mL, 6.5 mmol). The solution
was cooled to 0 °C in an ice bath. Me$_3$SiCl (0.82 mL, 6.5 mmol)
was added dropwise over 20 minutes using a syringe pump, and
the reaction mixture was stirred vigorously using a mechanical
stirrer. The reaction was heated to reflux, and stirred under reflux
for 1 hour. After allowing to cool to room temperature, the
reaction was filtered and washed with EtOAc. The filtrate was evaporated in vacuo,
and dissolved in dry ethyl ether (10 mL). PhMgBr (4.20 mL of a 3M solution in diethyl
ether, 12.6 mmol) was added to the solution under N$_2$ atmosphere. The solution was
heated to reflux, and stirred at this temperature for 3 hours. The solution was allowed
to cool to room temperature, before it was quenched with 1M HCl (6.3 mL). The
layers were separated, and the aqueous phase was made basic with 2M NaOH (6.3
mL). The aqueous phase was extracted with EtOAc (3*15 mL), and the combined
organic phases were washed with brine (15 mL), filtered and concentrated in vacuo.
The product was purified by silica-gel column chromatography (EtOAc:n-pentane,
2:1, R_f = 0.43), yielding 322 mg (2.22 mmol, 36 %) of **2** as an orange oil.
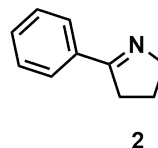
Spectroscopic data (Appendix B): ^{1}H NMR (400 MHz, DMSO-d_6) δ 7.85-7.83
(m, 2H), 7.46-7.45 (m, 3H), 3.97-3.93 (m, 2H), 2.94-2.90 (m, 2H), 1.99-1.91 (m, 2H).
^{1}H NMR data was in accordance with previously reported values.[57]

5.6.7   2-Phenylpyrrolidine (*rac*-4)

Compound **2** (1.36 g, 9.34 mmol) was dissolved in MeOH:H$_2$O
(4:1, 25 mL) and added NaBH$_4$ (405 mg, 10.7 mmol). The
solution was stirred for 22 hours at room temperature. Then 2M
HCl (15 mL) was added to bring the pH of the solution to 0-2, and
the solution was stirred for an additional 30 minutes. 2M NaOH
(15 mL) was added to bring the pH of the solution to 13-14, and
the organic phase was extracted with dichloromethane (3 x 25
mL). The product was purified by column chromatography (EtOAc:n-pentane, 1:1,
R_f = 0.44), yielding 1.22 g (8.29 mmol, 89 %) of *rac*-**4** as a yellowish oil.
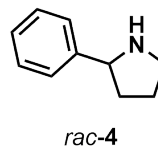
Spectroscopic data (Appendix E): (400 MHz, DMSO-d_6) δ 7.37-7.35 (m, 2H),
7.30-7.26 (m, 2H), 7.20-7.17 (m, 1H), 4.01 (t, J = 7.62 Hz, 1H), 3.13 (br s, 1H),
3.05-3.00 (m, 1H), 2.91-2.85 (m, 1H), 2.14-2.06 (m, 1H), 1.83-1.67 (m, 2H), 1.51-1.42
(m, 1H). ^{1}H NMR data was in accordance with previously reported values.[53]

5.6.8   6-Bromo-4-(2-phenylpyrrolidin-1-yl)thieno[2,3-*d*]--pyrimidine (*rac*-5a)

Compound *rac*-**5a** was prepared as described in Section 5.4, using 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine (230 mg, 0.92 mmol), *rac*-**4** (200 mg, 1.36 mmol) and Hünig's base (0.24 mL, 1.36 mmol). The reaction time was 1.25 hours, and purification by silica gel column chromatography (*n*-pentane:EtOAc, 6:1, R_f = 0.24) yielded 265 mg (0.74 mmol, 80 %) of *rac*-**5a** as a white, crystalline solid; mp = 145.1-146.3 °C; HPLC purity: > 99%, t_R=26.8 min.
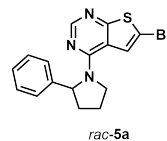
Spectroscopic data (Appendix H): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 8.24 (s, 1H), 7.78 (br s, 1H), 7.31-7.30 (m, 2H), 7.23-7.21 (m, 3H), 5.60 (d, *J* = 7.8 Hz, 1H), 4.20 (br s, 1H), 3.92 (br s, 1H), 2.40 (br s, 1H), 2.01 (br s, 1H), 1.89 (br s, 2H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 169.1, 154.7, 153.7, 143.6, 128.9 (2C), 127.1, 125.9 (2C), 125.4, 116.6, 108.7, 62.6, 50.4, 35.0, 23.8; IR (neat, cm^{-1}): 2971, 2873, 1541, 1469, 1304, 1061, 840, 770, 698; HRMS(ESI/APCI, *m/z*): 360.0165 (calcd. C$_{16}$H$_{14}$^{79}BrN$_3$S, 360.0170, [M+H]$^+$).

5.6.9   *(R)*-6-Bromo-4-(2-phenylpyrrolidin-1-yl)thieno[2,3-*d*]--pyrimidine (*(R)*-5a)

Compound *(R)*-**5a** was prepared as described in Section 5.4, using 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine (302 mg, 1.21 mmol), *(R)*-**4** (248 mg, 1.69 mmol), Hünig's base (3.0 mL, 17 mmol) and *i*-PrOH (8 mL). The reaction time was 24 hours, and purification by silica gel column chromatography (*n*-pentane:EtOAc, 6:1, R_f = 0.38) yielded 320 mg (0.89 mmol, 74 %) of *(R)*-**5a** as a yellow oil.
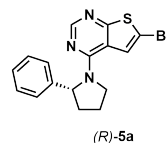
Spectroscopic data (Appendix I): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 8.24 (br s, 1H), 7.78 (br s, 1H), 7.32-7.29 (m, 2H), 7.23-7.21 (m, 3H), 5.60 (d, J = 7.6 Hz, 1H), 4.19 (br s, 1H), 3.91 (br s, 1H), 2.38 (br s, 1H), 1.99 (br s, 1H), 1.90-1.88 (m, 2H); ^{13}C NMR (100 MHz, DMSO-d_6) δ: 169.1, 154.6, 153.7, 143.6, 128.9 (2C), 127.1, 125.9 (2C), 125.4, 116.6, 108.7, 62.6, 50.4, 34.4, 23.6; IR (neat, cm^{-1}): ; HRMS(ESI/APCI, *m/z*) 360.0169 (calcd. C$_{16}$H$_{14}$^{79}BrN$_3$S, 360.0170, [M+H]$^+$)

5.6.10   *(S)*-6-Bromo-4-(2-phenylpyrrolidin-1-yl)thieno[2,3-*d*]-pyrimidine (*(S)*-5a)

Compound *(S)*-**5a** was prepared as described in Section 5.4, using 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine (203 mg, 0.81 mmol), *(S)*-**4** (206 mg, 0.81 mmol), Hünig's base (0.19 mL, 1.1 mmol) and *i*-PrOH (8 mL). The reaction time was 24 hours, and purification by silica gel column chromatography (*n*-pentane:EtOAc, 6:1, R_f
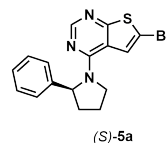
= 0.38) yielded 147 mg (0.41 mmol, 50 %) of *(S)*-**5a** as a yellow oil.

Spectroscopic data (Appendix J): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 8.24 (s, 1H), 7.78 (br s, 1H), 7.32-7.29 (m, 2H), 7.23-7.21 (m, 3H), 5.60 (d, J = 7.7 Hz, 1H), 4.19 (br s, 1H), 3.92 (m, 1H), 2.38 (br s, 1H), 2.0 (br s, 1H), 1.90-1.88 (m, 2H); ^{13}C NMR (100 MHz, DMSO-d_6) δ: 169.1, 154.6, 153.7, 143.6, 128.9 (2C), 127.1, 125.9 (2C), 125.4, 116.6, 108.7, 62.6, 50.4, 34.6, 24.7; IR (neat, cm^{-1}) ; HRMS(ESI/APCI, *m/z*) 360.0166 (calcd. $C_{16}H_{14}{}^{79}BrN_3S$, 360.0170, [M+H]$^+$)

5.6.11   6-Bromo-4-(pyrrolidin-1-yl)thieno[2,3-*d*]--pyrimidine (5b)

Compound **5b** was prepared as described in Section 5.4, using 6-bromo-4-chlorothieno[2,3-*d*]pyrimidine (416 mg, 1.67 mmol), pyrrolidine (0.21 mL, 2.56 mmol) and Hünig's base (0.42 mL, 2.41 mmol). The reaction time was 1.5 hours, and drying yielded 432 mg (1.52 mmol, 91 %) of **5b** as a brownish solid; mp = 131.2-132.2 °C.

5b
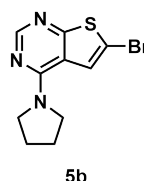
Spectroscopic data (Appendix K): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.28 (s, 1H), 7.74 (s, 1H), 3.71 (br s, 4H), 1.96 (br s, 4H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 168.8, 154.6, 153.8, 125.3, 116.6, 108.3, 49.1 (2C), 25.5 (2C); IR (neat, cm^{-1}): 2971, 2871, 1534, 1495, 1341, 1218, 1026, 849, 770; HRMS (ESI/APCI, *m/z*): 283.9861 (calcd., $C_{10}H_{11}N_3S^{79}Br$, 283.9857, [M+H]$^+$).

5.6.12   (4-(4-(2-Phenylpyrrolidin-1-yl)thieno[2,3-*d*]--pyrimidin-6-yl)phenyl)methanol (*rac*-6a)

Compound *rac*-**6a** was prepared as described in Section 5.5, starting with compound *rac*-**5a** (100 mg, 0.28 mmol) and 4-(hydroxymethyl)phenyl boronic acid (51.0 mg, 0.35 mmol). The reaction time was 55 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f = 0.12) yielded 100 mg (0.26 mmol, 93 %) of *rac*-**6a** as a yellowish solid; mp = 171.8-173.0 °C; HPLC purity: 97%, t_R = 22.6 min.

rac-**6a**
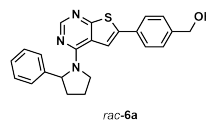
Spectroscopic data (Appendix L): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 8.27 (br s, 1H), 7.64-7.61 (m, 2H), 7.56-7.53 (m, 1H), 7.39-7.38 (m, 2H), 7.31-7.30 (m, 2H), 7.28 (br s, 2H), 7.21-7.19 (m, 1H), 5.70 (s, 1H), 5.27 (t, J = 5.7 Hz, 1H), 4.54 (d, J = 5.4 Hz, 2H), 4.29 (br s, 1H), 4.00 (br s, 1H), 2.43 (br s, 1H), 2.04 (br s, 1H), 1.93 (m, 2H); ^{13}C NMR (100 MHz, DMSO-d_6) δ: 167.6, 155.6, 153.5, 144.0, 143.5, 136.7, 132.5, 132.0 (2C), 128.9 (3C), 127.6 (2C), 127.1, 126.0 (2C), 117.9, 62.9, 62.6, 50.5, 34.4, 24.0; IR (neat, cm^{-1}): 3309, 2969, 1552, 1451, 1316, 1062, 1021, 773, 701; HRMS(ESI/APCI, *m/z*): 388.1479 (calcd. $C_{23}H_{23}N_3O_2S$, 388.1484, [M+H]$^+$).

5.6.13 (R)-(4-(4-(2-Phenylpyrrolidin-1-yl)thieno[2,3-d]--pyrimidin-6-yl)phenyl)methanol ((R)-6a)

Compound *(R)*-**6a** was prepared as described in Section 5.5, starting with *(R)*-**5a** (146 mg, 0.405 mmol). The reaction time was 45 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f = 0.25) yielded 145 mg (0.374 mmol, 92 %) of *(R)*-**6a** as a brown solid, mp = 200.1-202.4 °C; HPLC purity: 99%, t_R = 22.6 min; Chiral HPLC (Method B): t_R = 12.2 (minor), t_R = 14.8 (major).

(R)-**6a**
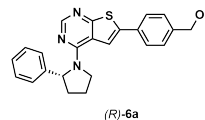
Spectroscopic data (Appendix N): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.26 (br s, 1H), 7.64-7.61 (m, 2H), 7.57-7.54 (m, 1H), 7.40-7.39 (m, 2H), 7.31-7.30 (m, 2H), 7.28 (br s, 2H), 7.22-7.20 (m, 1H), 5.71 (br s, 1H), 5.26 (t, J = 5.7 Hz, 1H), 4.54 (d, J = 5.5, 2H), 4.30 (br s, 1H), 4.0 (br s, 1H), 2.43 (br s, 1H), 2.04 (br s, 1H), 1.93 (br s, 2H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 167.6, 155.6, 153.5, 144.0, 143.5, 136.7, 132.5, 131.5 (2C), 129.0 (3C), 127.6 (2C), 127.1, 126.0 (2C), 117.9, 62.9, 62.6, 50.5, 34.5, 24.3; IR (neat, cm^{-1}): 3314, 3026, 2973, 2875, 1552, 1506, 1345, 1062, 773, 701; HRMS(ESI/APCI, *m/z*): 388.1487 (calcd. C$_{23}$H$_{21}$N$_3$OS, 388.1484, [M+H]$^+$).

5.6.14 (S)-(4-(4-(2-Phenylpyrrolidin-1-yl)thieno[2,3-d]--pyrimidin-6-yl)phenyl)methanol ((S)-6a)

Compound *(S)*-**6a** was prepared as described in Section 5.5, starting with *(S)*-**5a** (131 mg, 0.362 mmol). The reaction time was 45 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f = 0.25) yielded 129 mg (0.33 mmol, 92 %) of *(S)*-**6a** as a brown solid; mp = 200.1-202.4 °C; HPLC purity: 98%, t_R = 22.6 min; Chiral HPLC (Method B): t_R = 12.3 (major), t_R = 14.8 (minor).

(S)-**6a**
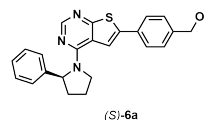
Spectroscopic data (Appendix O): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.27 (br s, 1H), 7.64-7.61 (m, 2H), 7.57-7.54 (m, 1H), 7.40-7.39 (m, 2H), 7.32-7.31 (m, 2H), 7.28 (br s, 2H), 7.22-7.20 (m, 1H), 5.71 (br s, 1H), 5.26 (t, J = 5.7 Hz, 1H), 4.54 (d, J = 5.5, 2H), 4.30 (br s, 1H), 4.04-3.98 (m, 1H), 2.43 (br s, 1H), 2.04 (br s, 1H), 1.93 (br s, 2H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 167.6, 155.6, 153.5, 144.0, 143.5, 136.7, 132.5, 132.0 (2C), 129.0 (3C), 127.6 (2C), 127.1, 126.0 (2C), 117.9, 62.9, 62.6, 50.5, 34.5, 24.4; IR (neat, cm^{-1}): 3311, 3026, 2973, 2873, 1552, 1495, 1345, 1180, 1021, 772, 701; HRMS (ESI/APCI, *m/z*): 388.1482 (calcd., C$_{23}$H$_{22}$N$_3$OS, 388.1484, [M+H]$^+$).

5.6.15 (4-(4-(Pyrrolidin-1-yl)thieno[2,3-*d*]pyrimidin-6-yl)--phenyl)methanol (6b)

Compound **6b** was prepared as described in Section 5.5, starting with **5b** (103 mg, 0.36 mmol). The reaction time was 55 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 9:1, R_f= 0.20) yielded 64.0 mg (0.21 mmol, 58 %) of **6b** as a white solid; mp = 216.5-221.3 °C; HPLC purity: 99%, t_R = 18.1 min.

6b
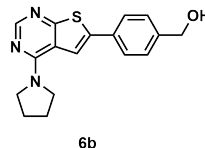
Spectroscopic data (Appendix P): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.29 (s, 1H), 7.90 (s, 1H), 7.75-7.33 (m, 2H), 7.40-7.39 (m, 2H), 5.26 (t, J = 5.7 Hz, 1H), 4.54 (d, J = 5.7 Hz, 2H), 3.82 (br s, 4H), 1.99 (br s, 4H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 167.3, 155.5, 153.6, 143.4, 136.6, 132.1, 127.5 (2C), 126.1 (2C), 117.5, 117.4, 63.0, 49.2 (2C), 25.5 (2C); IR (neat, cm^{-1}): 3274, 2971, 2870, 1558, 1500, 1331, 1178, 811, 723, 696; HRMS (ESI/APCI, *m/z*): 312.1175 (calcd., $C_{17}H_{18}N_3OS$, 312.1171, [M+H]+).

5.6.16 3-Methoxy-4-(4-(2-phenylpyrrolidin-1-yl)--thieno[2,3-*d*]pyrimidin-6-yl)benzaldehyde (*rac*-7a)

Compound *rac*-**7a** was prepared as described in Section 5.5, starting with *rac*-**5a** (372 mg, 1.03 mmol). The reaction time was 1.5 hours, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f= 0.16) yielded 422 mg (1.02 mmol, 99 %) of *rac*-**7a** as a yellow solid; mp = 205.5-207.0 °C.

rac-7a
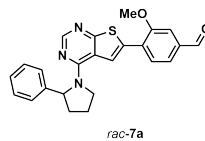
Spectroscopic data (Appendix Q): ^{1}H NMR (400 MHz, DMSO-d_6) δ: 10.0 (s, 1H), 8.31 (m, 2H), 7.80-7.55 (m, 4H), 7.33-7.29 (m, 4H), 7.23-7.20 (m, 1H), 5.72 (br s, 1H), 4.26 (br s, 1H), 4.0 (br s, 3H), 2.44 (br s, 1H), 2.04 (br s, 1H), 1.95 (br s, 2H); ^{13}C NMR (100 MHz MHz, DMSO-d_6) δ: 192.7, 168.8, 156.2, 155.7, 153.9, 143.8, 137.0, 131.1, 129.0 (3C), 127.9, 127.2, 126.0 (2C), 123.3, 122.3, 116.0, 112.3, 62.7, 56.6, 50.4, 35.7, 27.8; IR (neat, cm^{-1}): 2971, 2876, 2731, 1690, 1546, 1474, 1302, 1121, 1030, 777, 702; HRMS (ESI/APCI, *m/z*): 416.1427 (calcd., $C_{24}H_{22}N_3O_2S$, 416.1433, [M+H]$^+$).

5.6.17 (3-Methoxy-4-(4-(2-phenylpyrrolidin-1-yl)-thieno[2,3-*d*]pyrimidin-6-yl)phenyl)methanol (*rac*-8a)

Compound rac-**8a** was synthesized as described in   Section 5.6, starting with *rac*-**7a** (350 mg, 0.84 mmol). The reaction time was 1 hours, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f = 0.28) yielded 167 mg (0.40 mmol, 48 %) of *rac*-**8a** as a yellowish solid; mp = 177-179.2 °C; HPLC purity: 96%, t_R = 22.5 min.

rac-**8a**
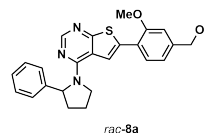
Spectroscopic data (Appendix R): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.28 (s, 1H), 7.86-7.53 (m, 2H), 7.32-7.31 (m, 4H), 7.22-7.20 (m, 1H), 7.1 (s, 1H), 6.98-6.97 (m, 1H), 5.69 (s, 1H), 5.29 (t, J = 5.5 Hz, 1H), 4.53 (d, J = 5.0 Hz, 2H), 4.22 (br s, 1H), 3.98 (br s, 1H), 3.87 (s, 3H), 2.42 (br s, 1H), 2.01 (br s, 1H), 1.92 (br s, 2H), ^{13}C NMR (150 MHz, DMSO-d_6) δ: 167.9, 155.9, 155.5, 153.3, 145.1, 144.0, 132.0, 129.0 (3C), 128.3, 127.1, 126.0 (2C), 120.4, 119.3, 116.1, 110.5, 63.0, 62.6, 56.2, 50.3, 35.5, 22.7; IR (neat, cm^{-1}): 3297, 3206, 2938, 2873, 1550, 1451, 1388, 1215, 1160, 1063, 775, 701; HRMS (ESI/APCI, *m/z*): 418.1586 (calcd., C$_{24}$H$_{24}$N$_3$O$_2$S, 418.1589, [M+H]$^+$).

5.6.18 *(R)*-3-Methoxy-4-(4-(2-phenylpyrrolidin-1-yl)-thieno[2,3-*d*]pyrimidin-6-yl)benzaldehyde (*(R)*-7a)

Compound *(R)*-**7a** was prepared as described in Section 5.5, starting with *(R)*-**5a** (172 mg, 0,48 mmol). The reaction time was 45 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f = 0.19) yielded 183 mg (0.44 mmol, 92 %) of *(R)*-**7a** as a yellowish solid; mp = 156.6-158.2 °C (decomp.).

(R)-**7a**
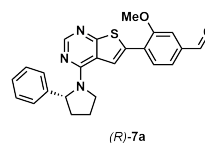
Spectroscopic data (Appendix S): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 10.0 (s, 1H), 8.30 (s, 1H), 7.88-7.61 (m, 4H), 7.33-7.29 (m, 4H), 7.22-7.21 (m, 1H), 5.71 (br s, 1H), 4.30 (br s, 1H), 4.0 (br s, 4H), 2.44 (br s, 1H), 2.04 (br s, 1H), 1.94 (br s, 2H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 192.7, 168.8, 156.2, 155.7, 153.9, 143.8, 137.0, 131.1, 129.0 (3C), 127.9, 172.2, 126.0 (2C), 123.3, 122.3, 116.0, 112.3, 62.7, 56.6, 50.4, 35.7, 27.3; IR (neat, cm^{-1}) 2969, 2877, 1690, 1570, 1474, 1345, 1214, 1122, 1030, 816, 702; HRMS (ESI/APCI, *m/z*): 416.1439 (calcd., C$_{24}$H$_{22}$N$_3$O$_2$S, 416.1433, [M+H]$^+$).

5.6.19 *(R)*-(3-Methoxy-4-(4-(2-phenylpyrrolidin-1-yl)--thieno[2,3-*d*]pyrimidin-6-yl)phenyl)methanol (*(R)*-8a)

Compound *(R)*-**8a** was prepared as described in Section 5.6, starting with *(R)*-**7a** (145 mg, 0.35 mmol). The reaction time was 50 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 1:1, R_f = 0.12) yielded 146 mg (0.35 mmol, > 99%) of *(R)*-**8a** as a white solid; mp = 162.0-163.5 °C; HPLC purity >99%, t_R = 22.5 min; Chiral HPLC (Method B): t_R = 14.6 (minor), t_R = 17.7 (major).

(R)-8a
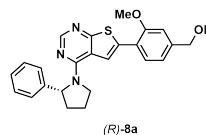
Spectroscopic data (Appendix T): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.27 (br s, 1H), 7.59 (m, 2H), 7.34-7.27 (m, 4H), 7.23-7.19 (m, 1H), 7.10 (s, 1H), 6.98-6.97 (m, 1H), 5.69 (br s, 1H), 5.28 (t, J = 5.7 Hz, 1H), 4.54 (d, J = 5.7 Hz, 2H), 4.24 (br s, 1H), 4.0 (br s, 1H), 3.87 (s, 3H), 2.43 (br s, 1H), 2.02 (br s, 1H), 1.92 (br s, 2H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 167.9, 155.9, 155.5, 153.1, 145.1, 144.0, 132.8, 129.0 (3C), 128.3, 127.1, 126.0 (2C), 120.4, 119.3, 116.1, 110.5, 63.0, 62.6, 56.2, 50.3, 35.8, 22.7; IR (neat, cm^{-1}): 3301, 3027, 2972, 2876, 1551, 1474, 1389, 1216, 1063, 849, 701; HRMS (ESI/APCI, *m/z*): 418.1591 (calcd., $C_{24}H_{24}N_3O_2S$, 418.1589, [M+H]$^+$).

5.6.20 3-Methoxy-4-(4-(pyrrolidin-1-yl)thieno[2,3-*d*]--pyrimidin-6-yl)benzaldehyde (7b)

Compound **7b** was synthesized as described in Section 3.5, starting with **5b** (418 mg, 1.47 mmol). The reaction time was 45 minutes, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 9:1, R_f = 0.32) yielded 468 mg (1.38 mmol, 94 %) of **7b** as a yellow solid; mp = 185.6-186.5 °C.

7b
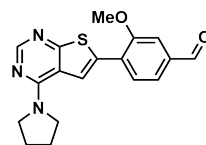
Spectroscopic data (Appendix U): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 10.02 (s, 1H), 8.33 (s, 1H), 8.17 (s, 1H), 8.14-8.13 (m, 1H), 7.63-7.61 (m, 2H), 4.04 (s, 3H), 3.85 (br s, 4H), 2.01 (br s, 4H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 192.7, 168.6, 156.1, 155.6, 154.0, 136.9, 130.9, 129.1, 128.1, 123.3, 121.9, 116.0, 112.2, 56.7, 49.2 (2C), 25.5 (2C); IR (neat, cm^{-1}): 2971, 2872, 2729, 1688, 1552, 1483, 1386, 1122, 1028, 815, 775, 736; HRMS (ESI/APCI, *m/z*): 340.1124 (calcd., $C_{18}H_{18}N_3O_2S$, 340.1120, [M+H]$^+$).

5.6.21 (3-Methoxy-4-(4-(pyrrolidin-1-yl)thieno[2,3-*d*]--pyrimidin-6-yl)phenyl)methanol (8b)

Compound **8b** was synthesized as described in Section 5.6, starting with **7b** (199 mg, 0.59 mmol). The reaction time was 1.5 hours, and purification by silica gel column chromatography (EtOAc:*n*-pentane, 9:1, R_f = 0.11) yielded 134 mg (0.39 mmol, 67 %) of **8b** as a yellowish solid; mp = 199.0-201.1 °C (decomp.); HPLC purity: 98%, t_R = 18.5 min.

8b
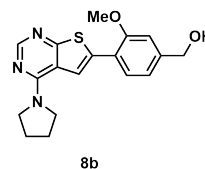
Spectroscopic data (Appendix V): ^{1}H NMR (600 MHz, DMSO-d_6) δ: 8.29 (s, 1H), 7.94 (s, 1H), 7.79-7.78 (m, 1H), 7.12 (s, 1H), 7.01-7.0 (m, 1H), 5.29 (t, 5.8 Hz, 1H), 4.55 (d, J = 5.7 Hz, 2H), 3.92 (s, 3H), 3.81 (br s, 4H), 2.0 (br s, 4H); ^{13}C NMR (150 MHz, DMSO-d_6) δ: 167.7, 155.9, 155.4, 153.4, 145.0, 132.7, 128.4, 120.6, 119.3, 118.9, 116.1, 110.5, 63.1, 56.2, 49.1 (2C), 25.5 (2C); IR (neat, cm^{-1}): 3256, 2969, 2871, 1578, 1500, 1330, 1125, 1028, 773; HRMS (ESI/APCI, *m/z*): 342.1282 (calcd., C$_{18}$H$_{20}$N$_3$O$_2$S, 342.1276, [M+H]$^+$).

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Appendix

A | Spectroscopic Data - Compound I

3.5 3.0 2.5 2.0 1.5 ppm

4.01
4.08
9.15

10 9 8 7 6 5 4 3 2 1 ppm

3.31
3.19
3.18
2.50
1.77
1.76
1.39

N–Boc
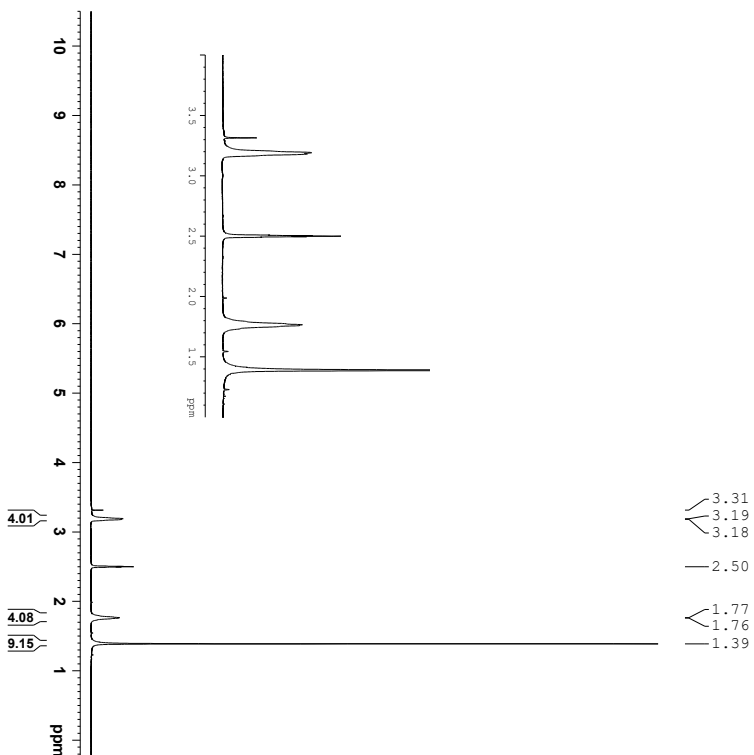
Current Data Parameters
NAME   ABR1-02-Crude
EXPNO   2
PROCNO   1

F2 - Acquisition Parameters
Date_   20180205
Time   11.30
INSTRUM   spect
PROBHD   5 mm PABBO BB/
PULPROG   zg30
TD   65536
SOLVENT   DMSO
NS   8
DS   2
SWH   8012.820 Hz
FIDRES   0.122266 Hz
AQ   4.0894465 sec
RG   128.065
DW   62.400 usec
DE   6.50 usec
TE   298.0 K
D1   1.00000000 sec
TD0   1

======== CHANNEL f1 ========
SFO1   400.1324710 MHz
NUC1   1H
P1   9.50 usec
PLW1   17.00000000 W

F2 - Processing parameters
SI   65536
SF   400.1300033 MHz
WDW   EM
SSB   0
LB   0.30 Hz
GB   0
PC   1.00
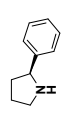
^{1}H NMR (400 MHz, DMSO-d_6) spectrum of compound **1**.
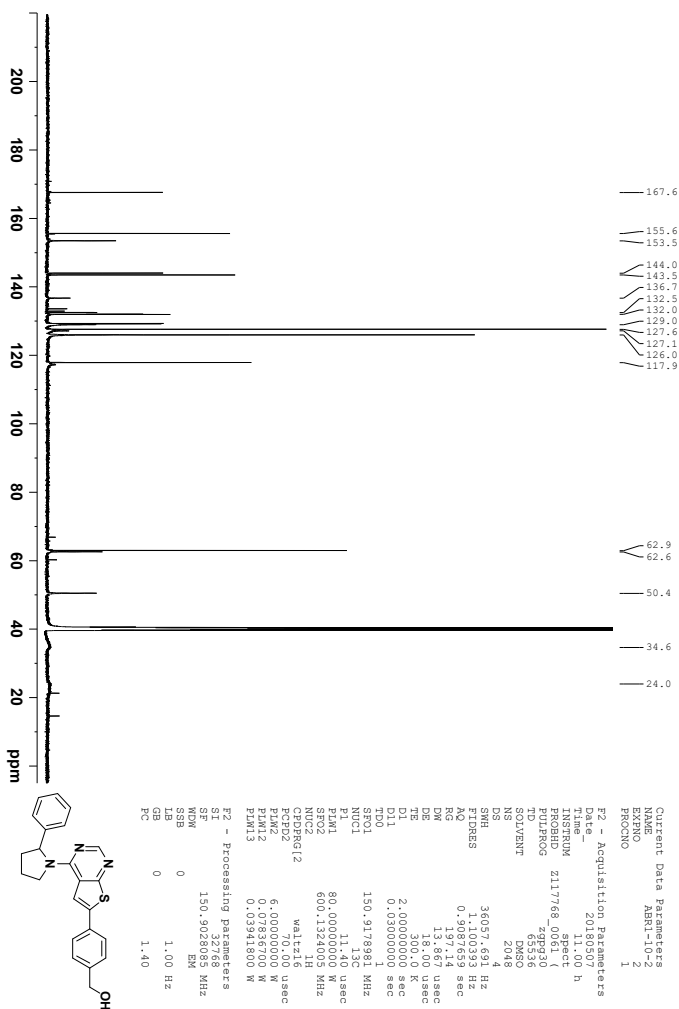
B | Spectroscopic Data - Compound 2
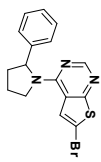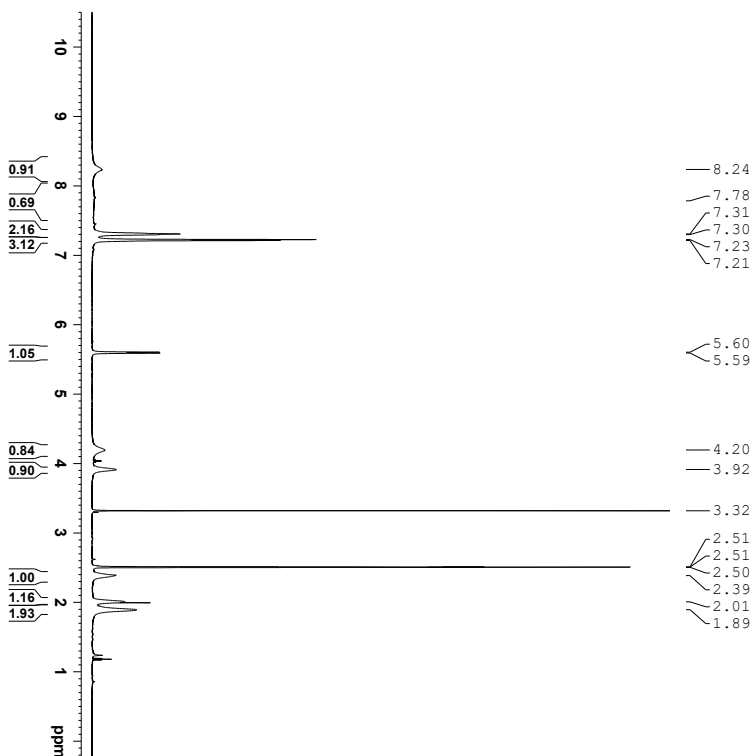

Current Data Parameters
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EXPNO   1
PROCNO   1

F2 - Acquisition Parameters
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Time   11.10
INSTRUM   spect
PROBHD   5 mm PABBO BB/
PULPROG   zg30
TD   65536
SOLVENT   DMSO
NS   8
DS   2
SWH   8012.820 Hz
FIDRES   0.122266 Hz
AQ   4.0894465 sec
RG   82.93
DW   62.400 usec
DE   6.50 usec
TE   297.9 K
D1   1.00000000 sec
TD0   1

======== CHANNEL f1 ========
SFO1   400.1324710 MHz
NUC1   1H
P1   9.50 usec
PLW1   17.00000000 W

F2 - Processing parameters
SI   65536
SF   400.1300000 MHz
WDW   EM
SSB   0
LB   0.30 Hz
GB   0
PC   1.00

7.85
7.84
7.83
7.46
7.45

3.97
3.96
3.95
3.93
3.34
2.94
2.92
2.90
2.90
2.51
1.99
1.97
1.95
1.93
1.91

2.01
3.06
2.04
2.14
2.07

^{1}H NMR (400 MHz, DMSO-d_6) spectrum of compound **2**.

C | Spectroscopic Data - Compound (R)-3

^{1}H NMR (400 MHz, DMSO-d_6) spectrum of compound *(R)*-**3**.

```
Print of window 38: Current Chromatogram(s)
Data File  : C:\CHEM32\...INASE\THOMAS A KINASE_RENHETSANALYSE 2018-03-20 12-09-31\034-0201.D
Sample Name : ABR1-03-Rflow0.3
=================================================================
Acq. Operator   : Thomas A                        Seq. Line :    2
Acq. Instrument : HPLC                            Location : Vial 34
Injection Date  : 3/20/2018 1:06:48 PM              Inj :    1
                                                Inj Volume : 10.000 µl
Acq. Method     : C:\CHEM32\1\DATA\KINASE\THOMAS A KINASE_RENHETSANALYSE 2018-03-20 12-09-
                  31\RENHETSANALYSE FELLES_NEUTRAL.M
Last changed    : 3/20/2018 12:09:31 PM by Thomas A
Analysis Method : C:\CHEM32\1\METHODS\FELLES\RENHETSANALYSE FELLES_NEUTRAL.M
Last changed    : 6/13/2018 6:16:27 PM by AC
                  (modified after loading)
```


Current Chromatogram(s)

DAD1 A, Sig=254,4 Ref=off (KINASE\THOMAS A KINASE_RENHETSANALYSE 2018-03-20 12-09-31\034-0201.D)

HPLC 6/13/2018 6:22:41 PM AC   Page   1 of 1

CSP-HPLC (Chiralpak AD 0.46 x 25 cm, hexane:*i*-PrOH, 99:1, flow 0.3 mL/min, detection at 254 nm) of compound *(R)*-**3**

D | Spectroscopic Data - Compound (S)-3

The image is the NMR spectrum

^{1}H NMR (400 MHz, DMSO-d_6) spectrum of compound *(S)*-**3**.

E ǀ Spectroscopic Data - *rac*-4

^{1}H NMR (400 MHz, DMSO-d_6) spectrum of compound (*rac*)-**4**.

^{1}H NMR (400 MHz, DMSO-d_6) spectrum of compound (R)-4.

7.41
7.39
7.37
7.35
7.32
7.31
7.29
4.33
4.32
4.31
4.30
4.29
4.28
3.21
3.21
3.20
3.19
3.17
3.17
3.12
3.12
3.11
3.10
3.10
3.10
3.09
2.50
2.27
2.26
2.25
2.24
2.23
1.99
1.95
1.94
1.92
1.90
1.88
1.79
1.78
1.77
1.76
1.75
1.74

1.88
2.05
1.09
0.97
0.93
1.01
1.00
2.01
1.08

¹H NMR (400 MHz, DMSO-d_6) spectrum of compound (S)-4.

¹H NMR (600 MHz, DMSO-*d₆*) spectrum of compound *(rac)*-**5a**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *(rac)*-**5a**.

COSY spectrum of compound *rac*-**5a**.
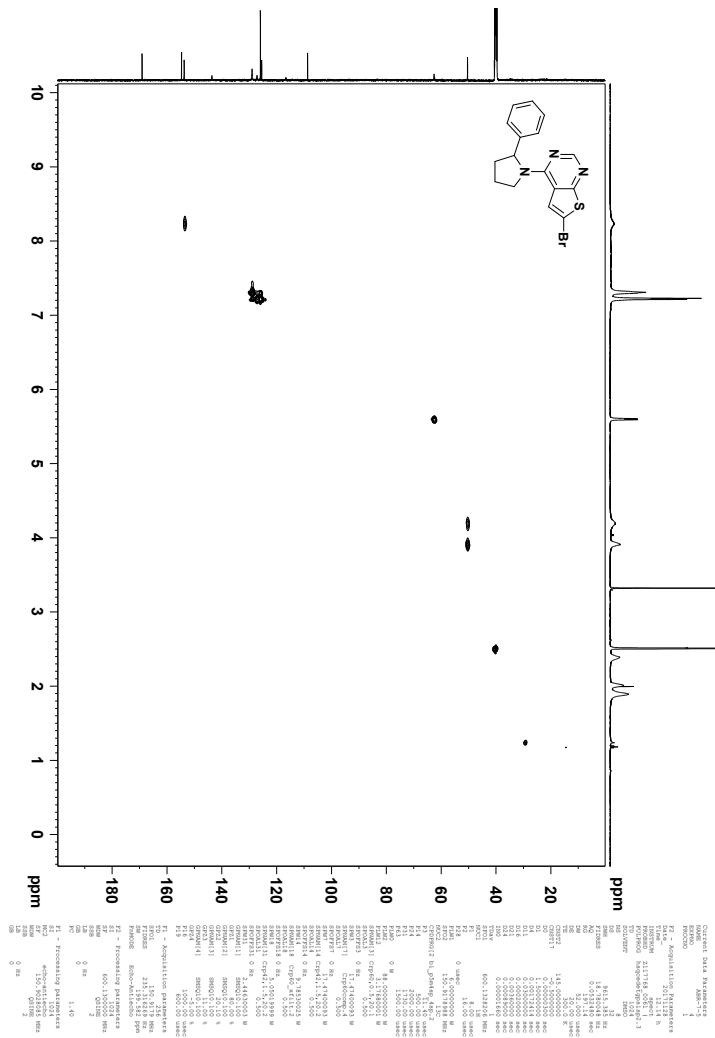
HSQC spectrum of compound *rac*-**5a**.
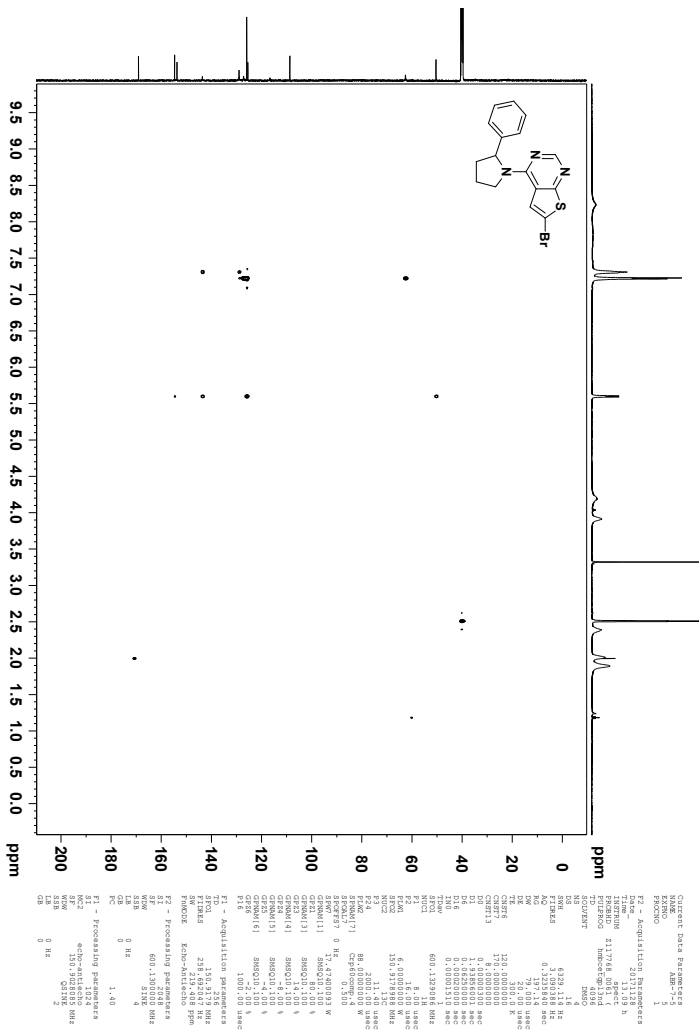
HMBC spectrum of compound *rac*-**5a**.
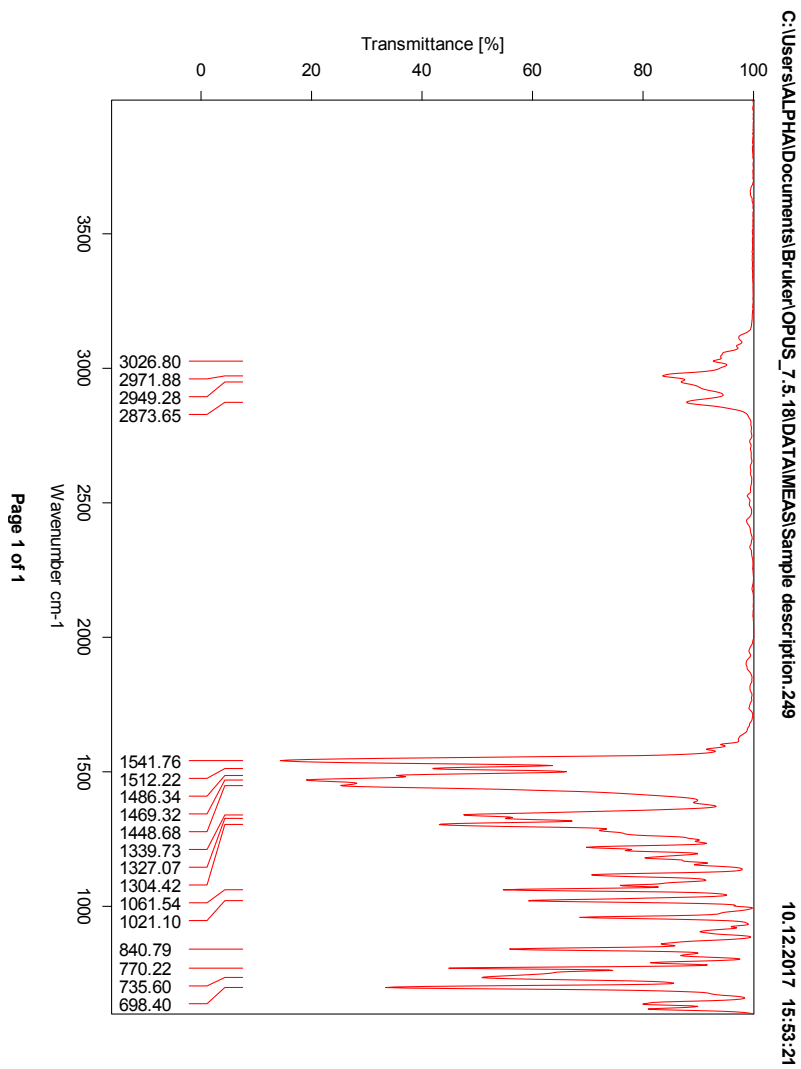
IR spectrum of compound *rac*-**5a**.

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
2848 formula(e) evaluated with 4 results within limits (up to 50 closest results for each mass)
Elements Used:
C: 1-500   H: 0-1000   N: 0-20   O: 0-100   S: 0-1   Br: 0-3

2017-439 43 (0.863) AM2 (Ar,35000.0,0.00,0.00); Cm (41:45)
1: TOF MS ASAP+
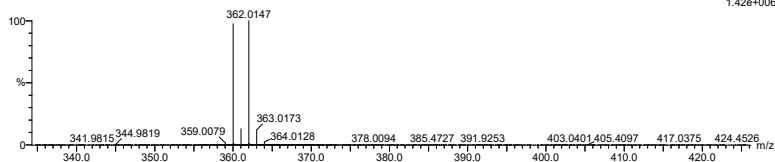
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
360.0165	360.0170	-0.5	-1.4	10.5	1220.7	9.472	0.01	C16 H15 N3 S Br
	360.0164	0.1	0.3	18.5	1224.5	13.291	0.00	C13 H2 N11 O S
	360.0168	-0.3	-0.8	7.5	1211.2	0.000	99.99	C8 H11 N9 O3 Br
	360.0163	0.2	0.6	4.5	1234.7	23.528	0.00	C7 H10 N3 O14

Minimum: -1.5
Maximum: 5.0   2.0   50.0

Mass spectrum of compound *rac*-**5a**.

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound (R)-5a.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *(R)*-**5a**.

COSY spectrum of compound *(R)*-**5a**.
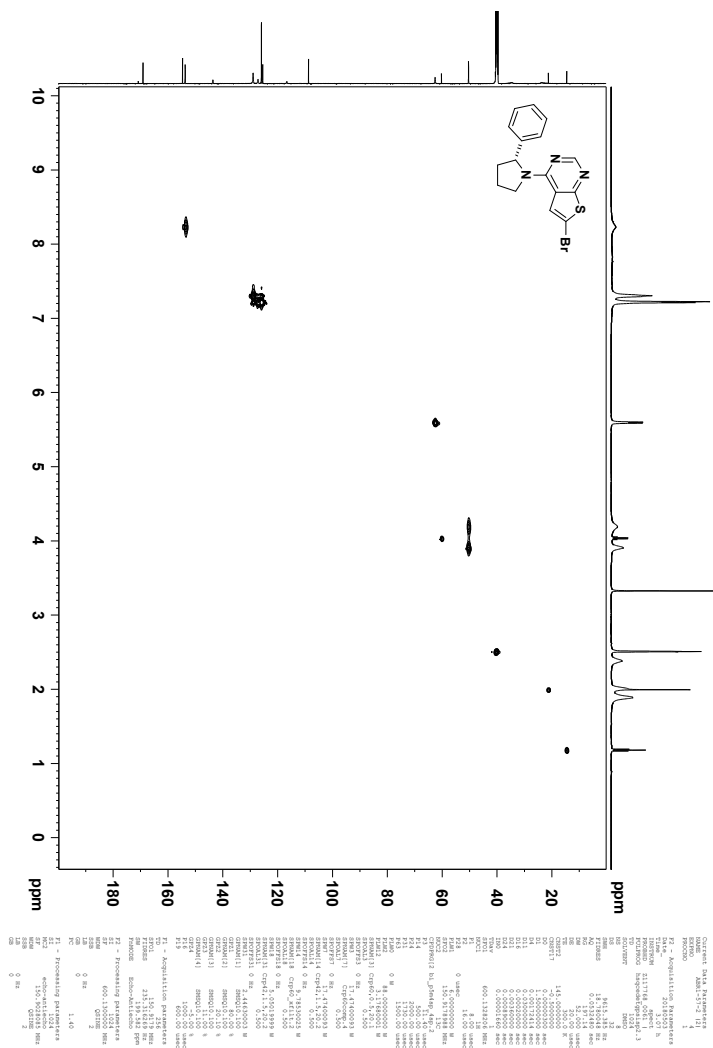
HSQC spectrum of compound *(R)*-**5a**.
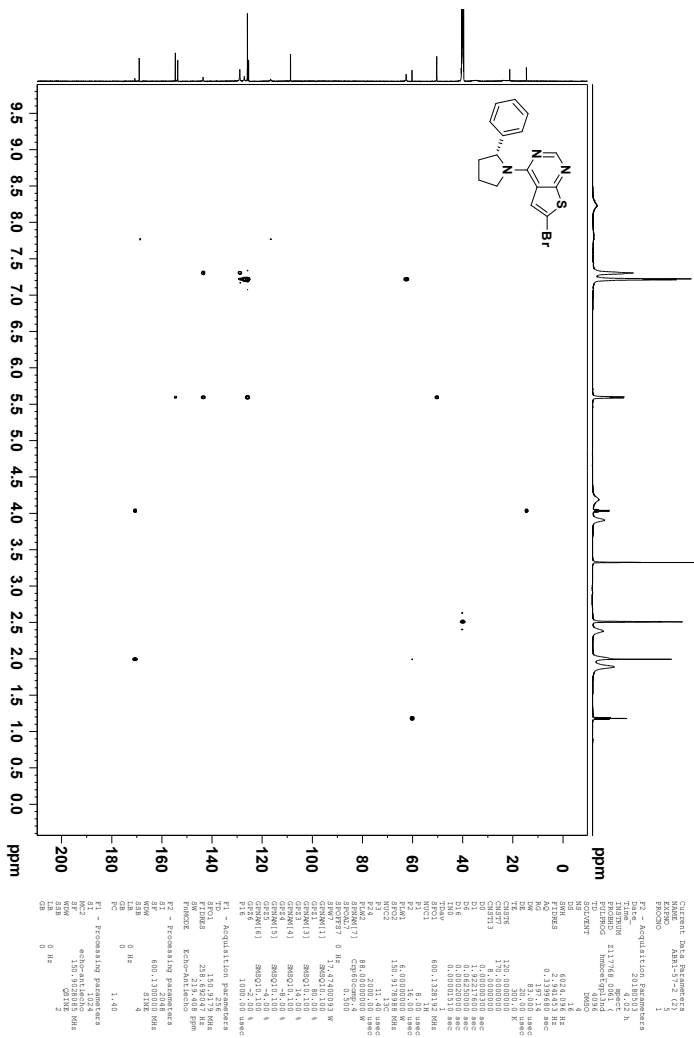
HMBC spectrum of compound *(R)*-**5a**.
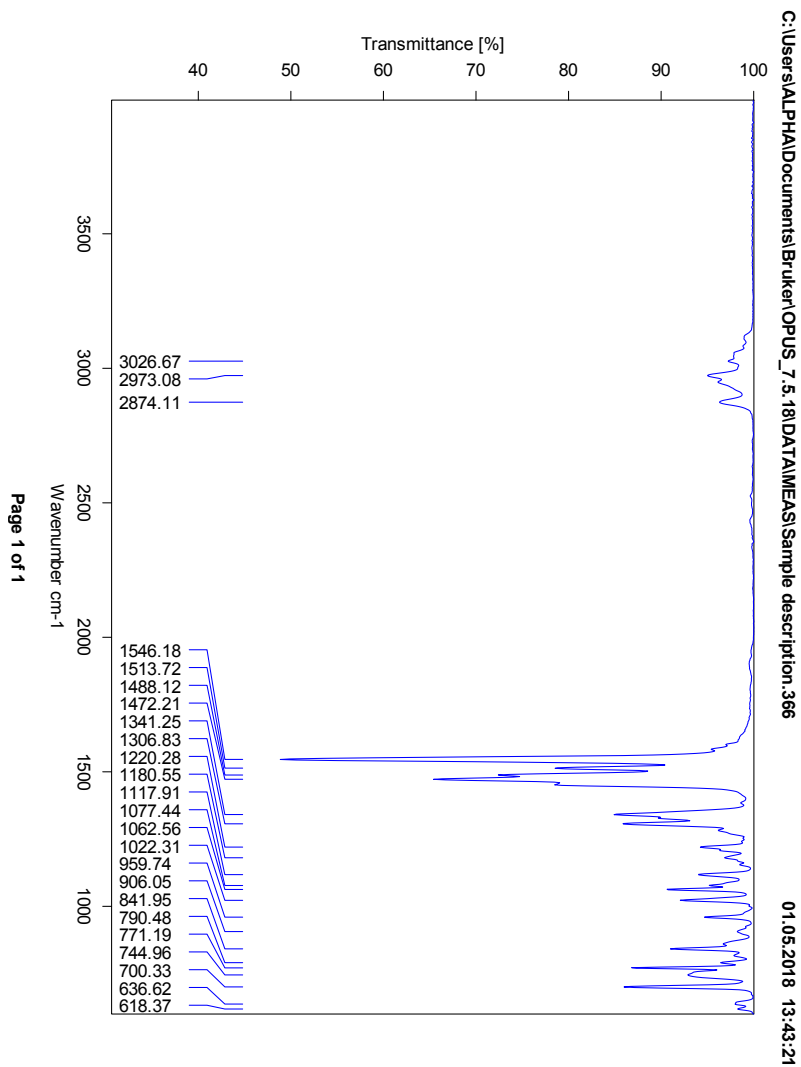
IR spectrum of compound *(R)*-**5a**.

Elemental Composition Report   **Page 1**

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
4829 formula(e) evaluated with 7 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-10   Na: 0-1   S: 0-2   Br: 0-4

2018-174 21 (0.431) AM2 (Ar,35000.0,0.00,0.00); Cm (19:22)
1: TOF MS ASAP+
3.67e+006
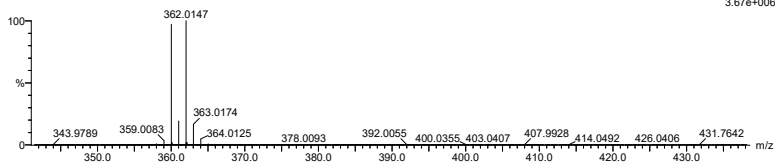
Minimum:			-1.5			
Maximum:		5000.0	2.0	50.0		

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
360.0166	360.0170	-0.4	-1.1	10.5	1462.3	0.001	99.95	C16 H15 N3 S Br
	360.0168	-0.2	-0.6	7.5	1470.2	7.823	0.04	C8 H11 N9 O3 Br
	360.0164	0.2	0.6	1.5	1471.4	9.041	0.01	C8 H19 N5 O2 S2 Br
	360.0171	-0.5	-1.4	3.5	1474.6	12.261	0.00	C10 H16 N3 O5 Na Br
	360.0171	-0.5	-1.4	3.5	1480.7	18.358	0.00	C8 H14 N3 O9 S2
	360.0167	-0.1	-0.3	14.5	1480.8	18.472	0.00	C15 H7 N5 O3 Na S
	360.0161	0.5	1.4	5.5	1481.0	18.646	0.00	C7 H11 N7 O5 Na S2

Mass spectrum of compound *(R)*-**5a**.

¹H NMR (600 MHz, DMSO-*d*₆) spectrum of compound *(S)*-**5a**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *(S)*-**5a**.

COSY spectrum of compound *(S)*-**5a**.
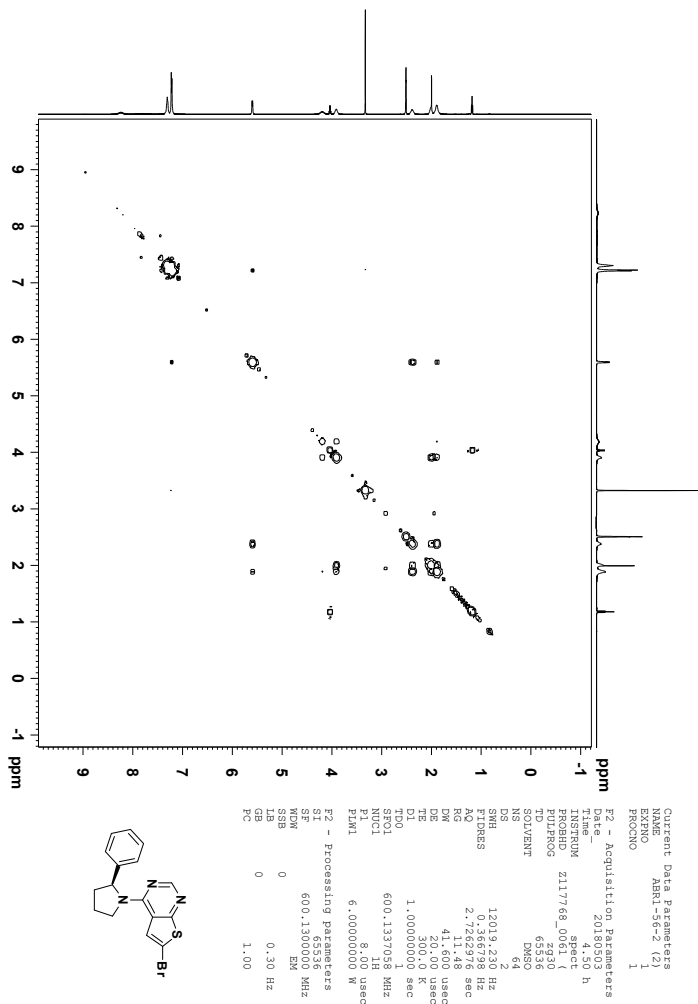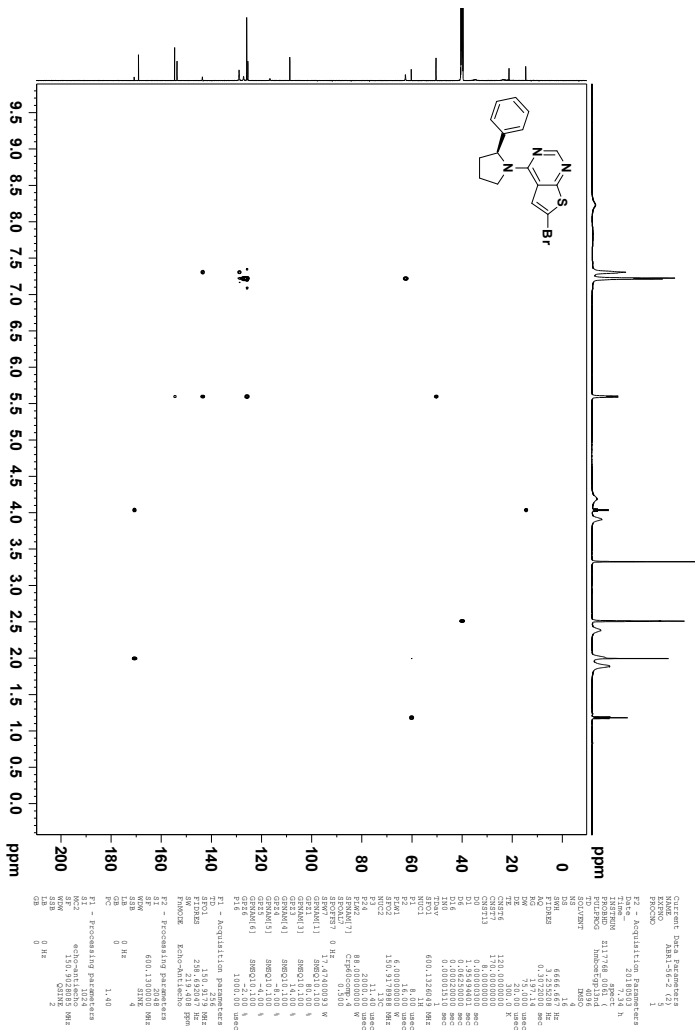
HSQC spectrum of compound *(S)*-**5a**.

HMBC spectrum of compound *(S)*-**5a**.
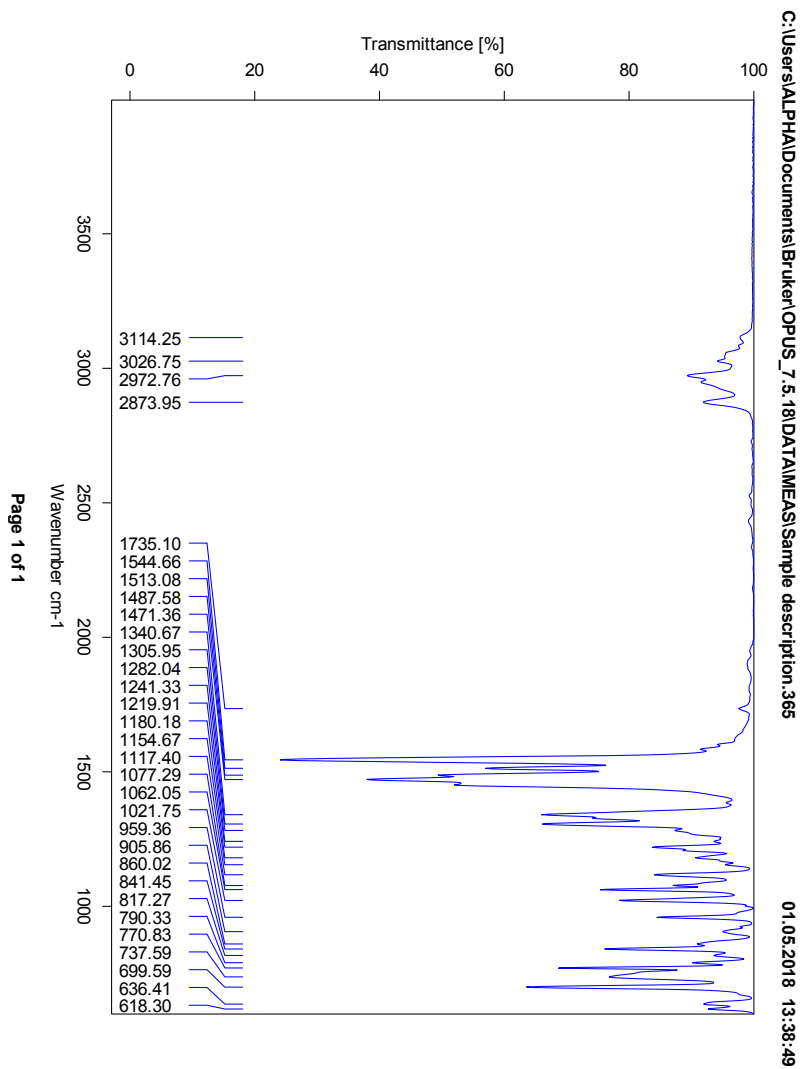
C:\Users\ALPHA\Documents\Bruker\OPUS_7.5.18\DATA\MEAS\Sample description.365

Transmittance [%]

Wavenumber cm-1

Page 1 of 1

3114.25
3026.75
2972.76
2873.95

1735.10
1544.66
1513.08
1487.58
1471.36
1340.67
1305.95
1282.04
1241.33
1219.91
1180.18
1154.67
1117.40
1077.29
1062.05
1021.75
959.36
905.86
860.02
841.45
817.27
790.33
770.83
737.59
699.59
636.41
618.30

01.05.2018 13:38:49

IR spectrum of compound *(S)*-**5a**.

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
4829 formula(e) evaluated with 9 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-10   Na: 0-1   S: 0-2   Br: 0-4

2018-175 21 (0.431) AM2 (Ar,35000.0,0.00,0.00); Cm (17:21)
1: TOF MS ASAP+

3.96e+006
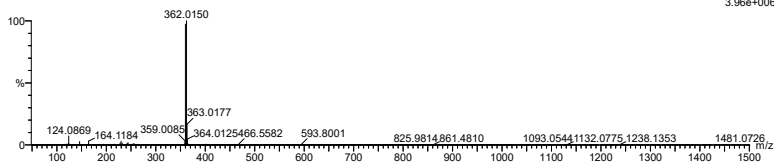
Minimum:			-1.5			
Maximum:		5000.0	2.0	50.0		

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
360.0169	360.0170	-0.1	-0.3	10.5	1423.9	0.000	99.99	C16 H15 N3 S Br
	360.0168	0.1	0.3	7.5	1433.4	9.516	0.01	C8 H11 N9 O3 Br
	360.0164	0.5	1.4	1.5	1435.5	11.682	0.00	C8 H19 N5 O2 S2 Br
	360.0171	-0.2	-0.6	3.5	1437.7	13.838	0.00	C10 H16 N3 O5 Na Br
	360.0174	-0.5	-1.4	-0.5	1443.7	19.816	0.00	C11 H24 N O2 Br2
	360.0167	0.2	0.6	14.5	1443.9	20.028	0.00	C15 H7 N5 O3 Na S
	360.0171	-0.2	-0.6	3.5	1443.9	20.031	0.00	C8 H14 N3 O9 S2
	360.0176	-0.7	-1.9	9.5	1451.3	27.434	0.00	C8 H6 N7 O10
	360.0174	-0.5	-1.4	23.5	1452.1	28.282	0.00	C23 H3 N3 O Na

Mass spectrum of compound *(S)*-**5a**.

8.28
7.74
3.71
3.31
2.50
1.96

1.00
1.15
4.49
4.47

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound **5b**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound **5b**.

COSY spectrum of compound **5b**.

HSQC spectrum of compound **5b**.

HMBC spectrum of compound **5b**.

C:\Users\ALPHA\Documents\Bruker\OPUS_7.5.18\DATA\MEAS\Sample description.359

Transmittance [%]

40   50   60   70   80   90   100

3500

3000

2971.20
2871.16

2500

Page 1 of 1

Wavenumber cm-1

2000

1553.78
1512.09
1495.17
1481.66
1459.59
1341.55
1327.19
1308.48
1218.08
1116.61
1067.73
1026.46
958.86
849.14
813.30
769.56

1000

01.05.2018 11:34:39
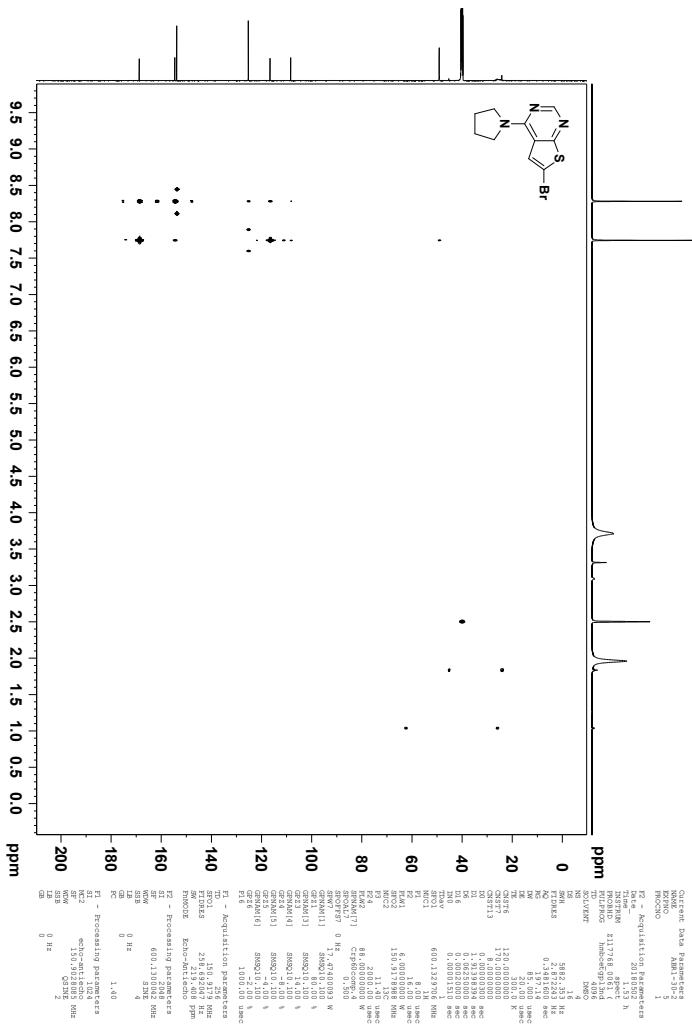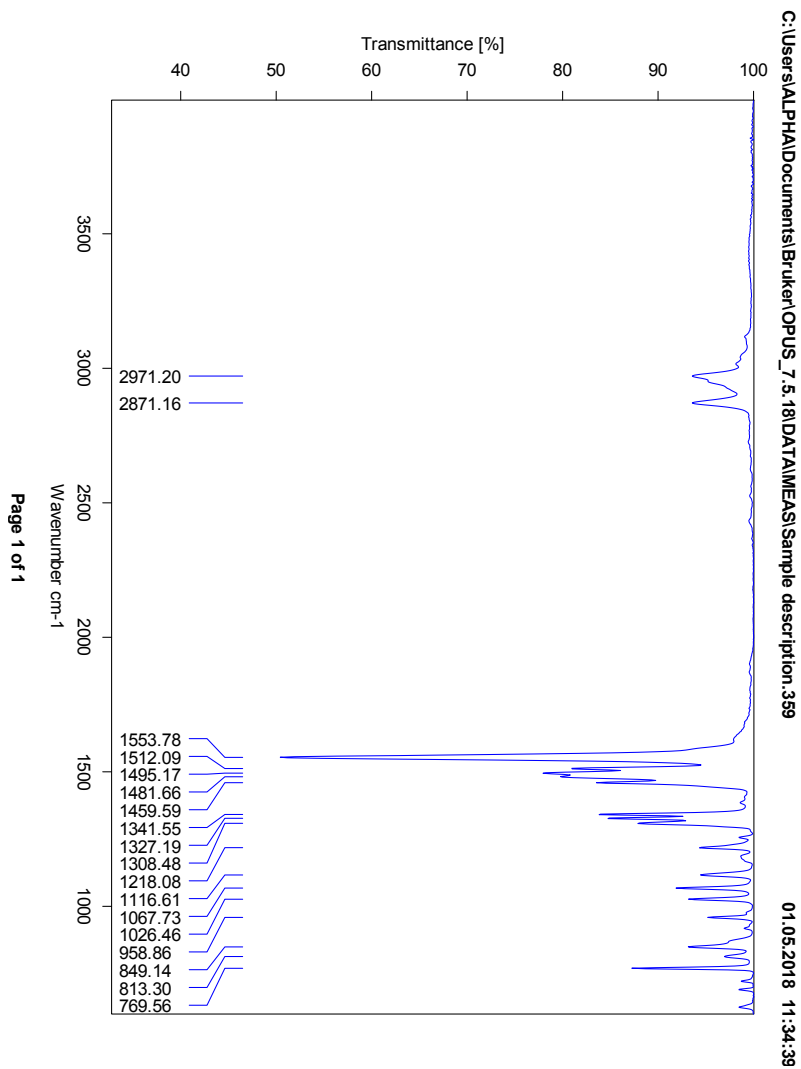
IR spectrum of compound **5b**.

Elemental Composition Report   **Page 1**

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
1804 formula(e) evaluated with 4 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   S: 0-3   Br: 0-2

2018-248B 91 (1.793) AM2 (Ar,35000.0,0.00,0.00); Cm (91:92)
1: TOF MS ASAP+
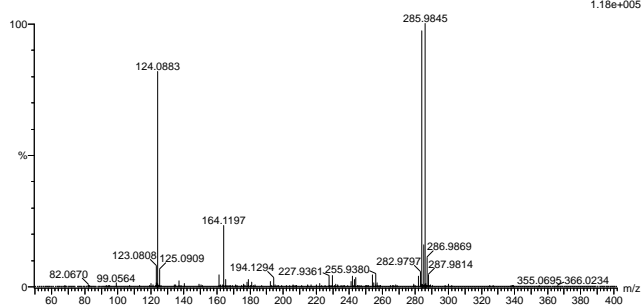
Minimum:				-1.5			
Maximum:		5.0	2.0	50.0			
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%) Formula
283.9861	283.9863	-0.2	-0.7	5.5	1022.0	12.438	0.00 C2 H2 N7 O10
	283.9858	0.3	1.1	-0.5	1018.1	8.476	0.02 C2 H10 N3 O9 S2
	283.9865	-0.4	-1.4	8.5	1020.1	10.477	0.00 C10 H6 N O7 S
	283.9857	0.4	1.4	6.5	1009.6	0.000	99.98 C10 H11 N3 S Br

Mass spectrum of compound **5b**.

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound *rac*-**6a**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *rac*-**6a**.
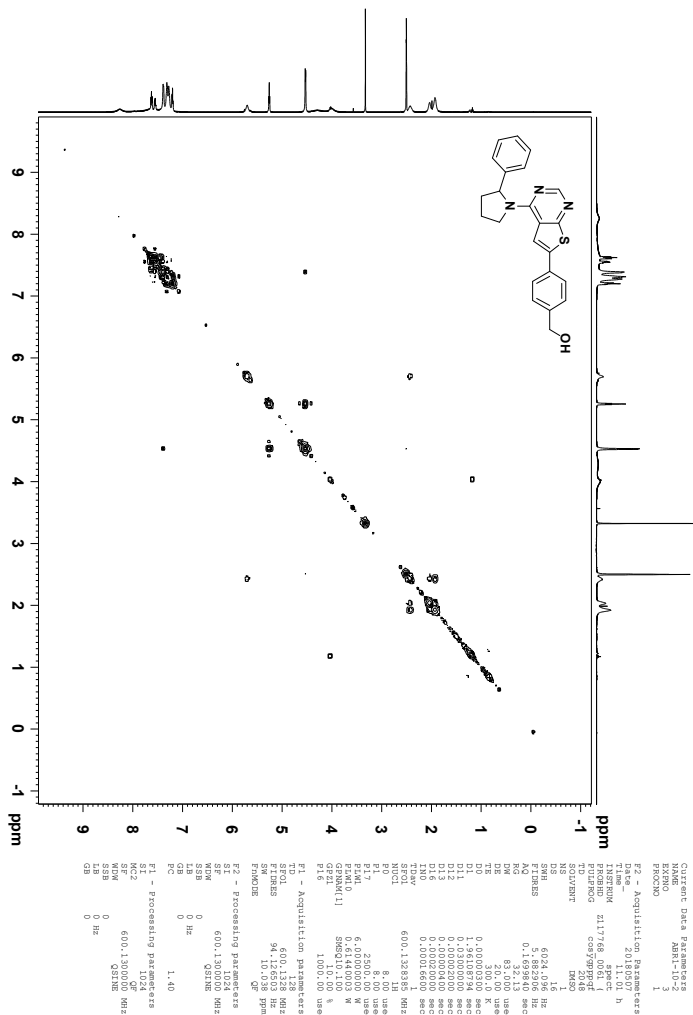
COSY spectrum of compound *rac*-**6a**.
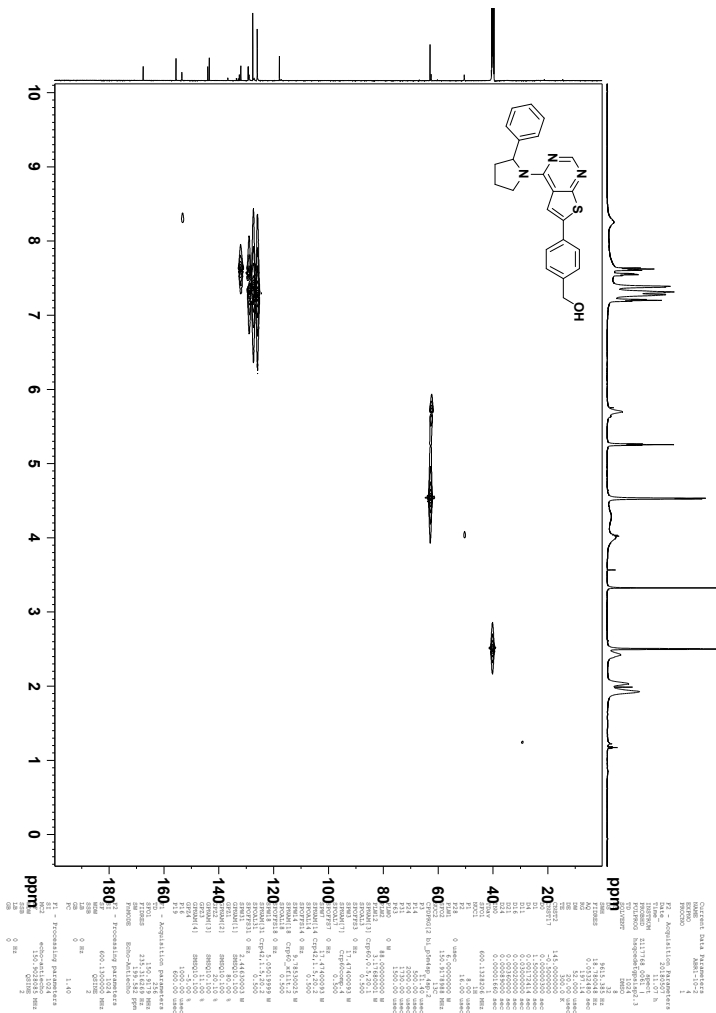
HSQC spectrum of compound *rac*-**6a**.
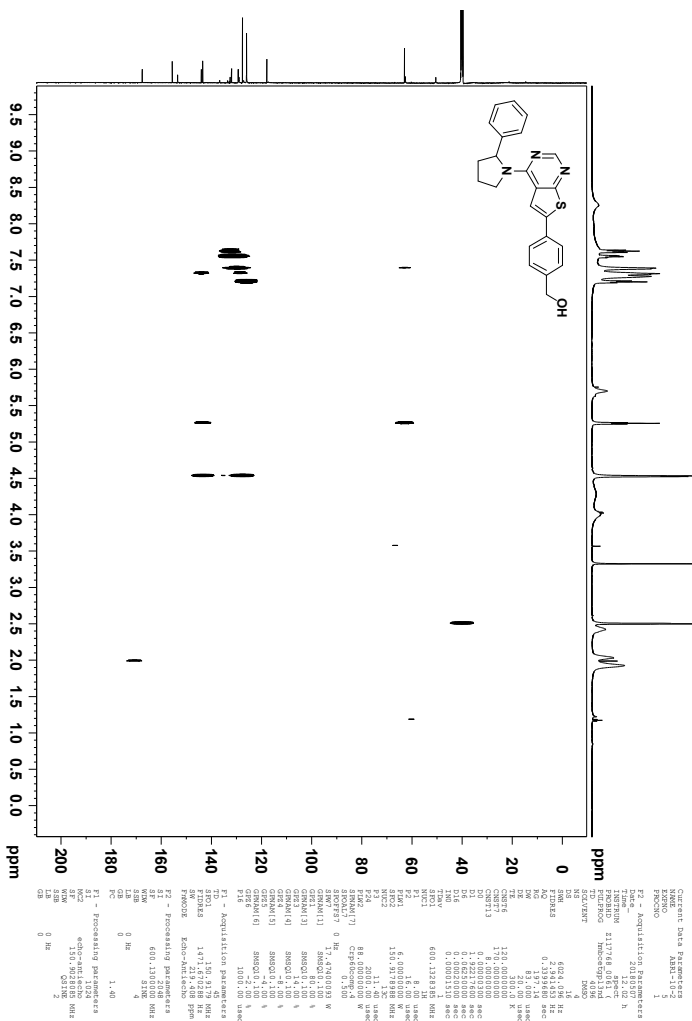
HMBC spectrum of compound *rac*-**6a**.
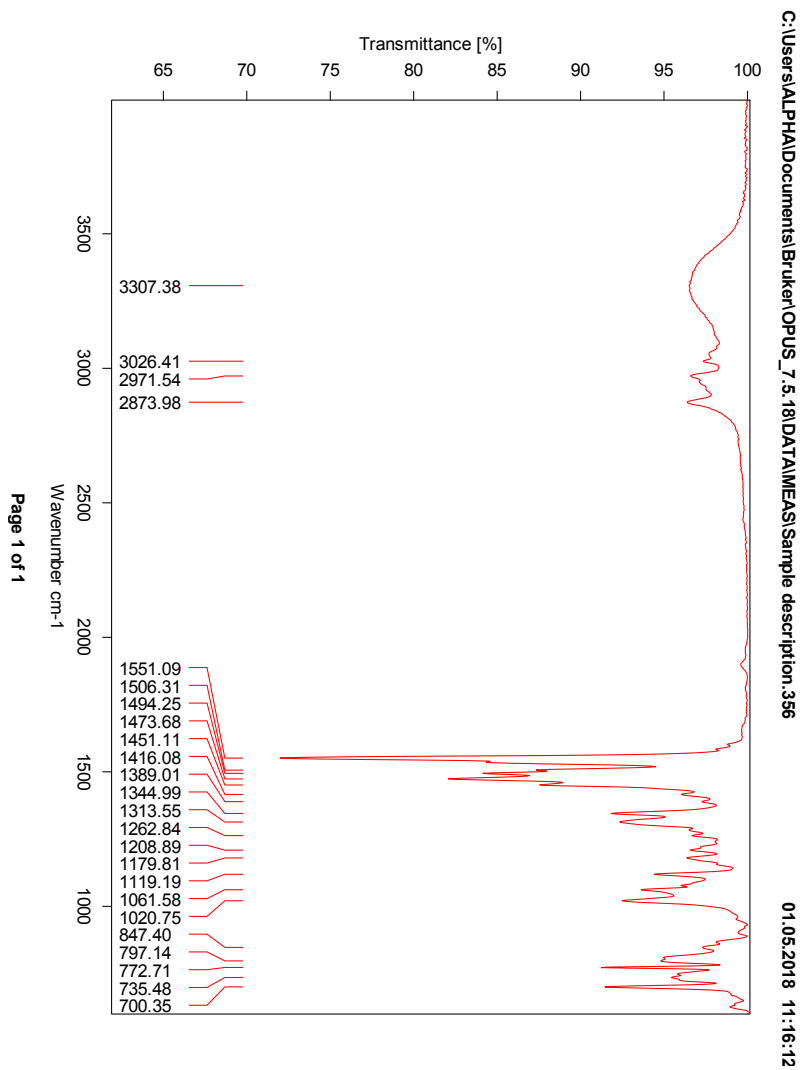
IR spectrum of compound *rac*-**6a**.

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
4739 formula(e) evaluated with 4 results within limits (up to 50 closest results for each mass)
Elements Used:
C: 1-500   H: 0-1000   N: 0-20   O: 0-100   S: 0-2   Br: 0-3

2017-440 118 (2.309) AM2 (Ar,35000.0,0.00,0.00); Cm (113:118)
1: TOF MS ASAP+

6.38e+005
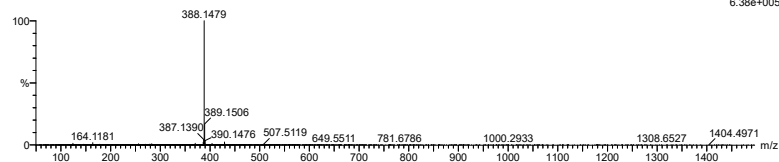
Minimum:			-1.5				
Maximum:	5.0	2.0	50.0				

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
388.1479	388.1484	-0.5	-1.3	14.5	921.1	0.451	63.69	C23 H22 N3 O S
	388.1482	-0.3	-0.8	11.5	922.4	1.728	17.77	C15 H18 N9 O4
	388.1475	0.4	1.0	2.5	922.5	1.756	17.28	C7 H22 N11 O6 S
	388.1477	0.2	0.5	5.5	925.1	4.371	1.26	C15 H26 N5 O3 S2

Mass spectrum of compound *rac*-**6a**.

Spectroscopic Data - Dimer by-product
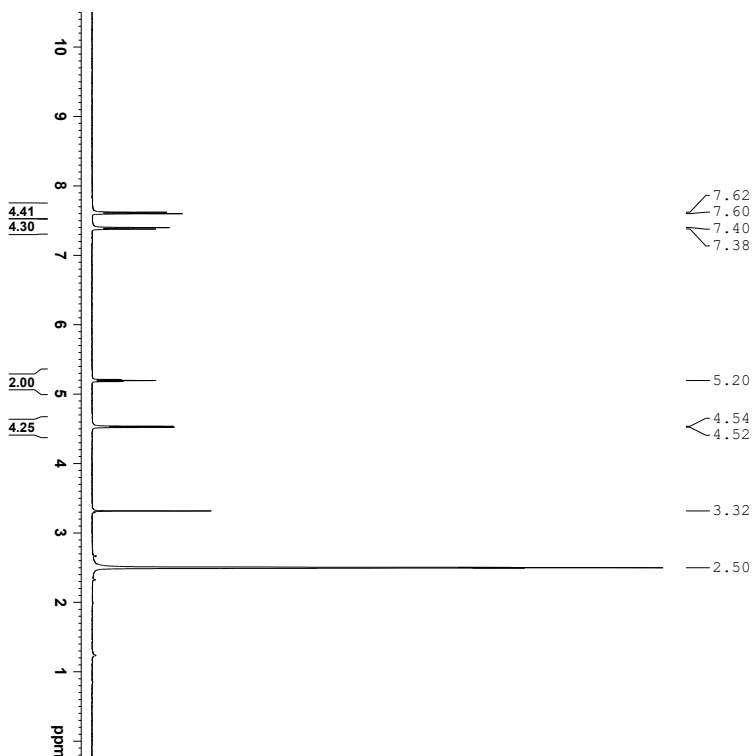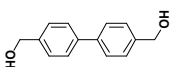
^{1}H NMR (400 MHz, DMSO-d_6) spectrum of [1,1'-biphenyl]-4,4'-diyldimethanol, as a byproduct in the synthesis of *rac*-**6a**.

Spectroscopic Data - Compound *(R)*-6a

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound *(R)*-**6a**.

^{13}C NMR (150 MHz, DMSO-*d*$_6$) spectrum of compound *(R)*-**6a**.

COSY spectrum of compound *(R)*-**6a**.

HSQC spectrum of compound *(R)*-**6a**.
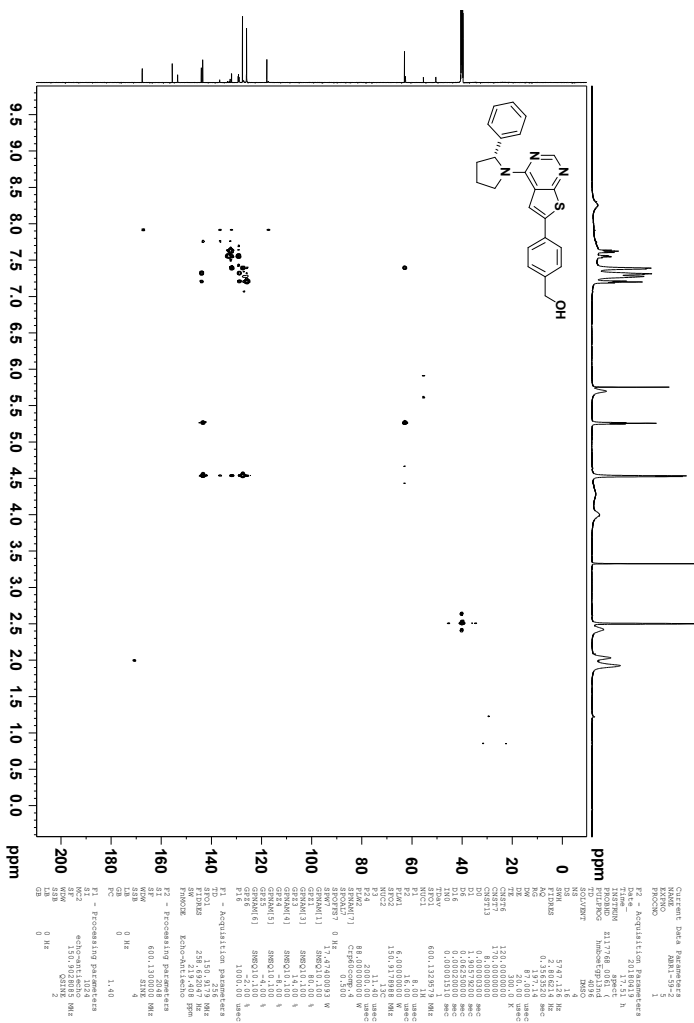
HMBC spectrum of compound *(R)*-**6a**.
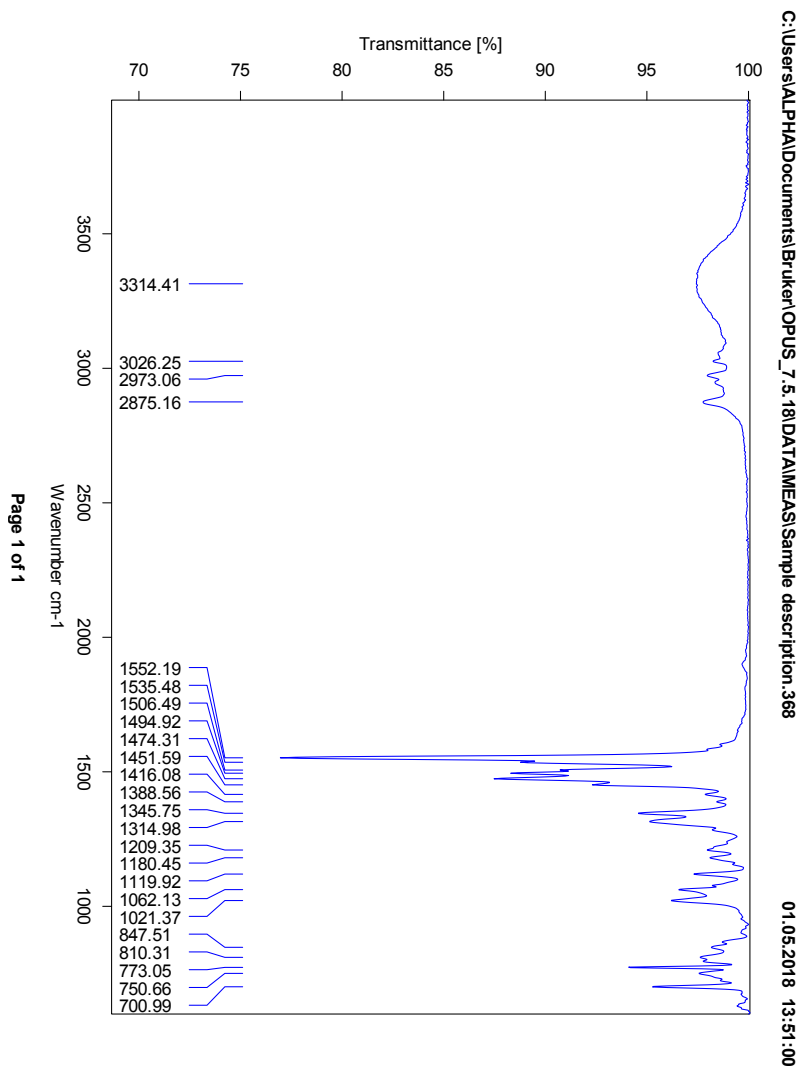
IR spectrum of compound *(R)*-**6a**.

Single Mass Analysis
Tolerance = 2.0 PPM / DBE: min = -100.0, max = 100.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
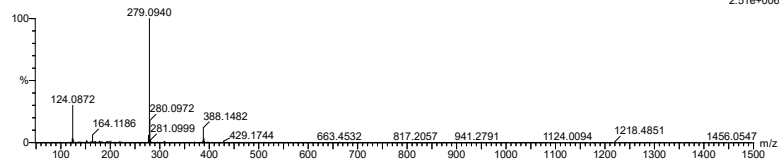7239 formula(e) evaluated with 8 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   S: 0-2
2018-211 133 (2.602) AM2 (Ar,35000.0,0.00,0.00); Cm (128:133)
1: TOF MS ASAP+

2.70e+007
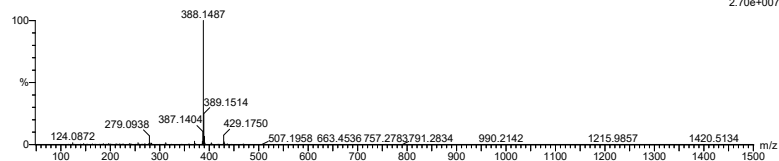
```
Minimum:                        -100.0
Maximum:            5000.0 2.0    100.0

Mass       Calc. Mass  mDa    PPM    DBE    i-FIT   Norm    Conf(%) Formula
388.1487   388.1484    0.3    0.8    14.5   1614.0  0.008   99.19   C23 H22 N3 O S
           388.1489   -0.2   -0.5   -3.5   1618.8  4.819   0.81    C10 H30 N O12 S
           388.1482    0.5    1.3    11.5   1625.6  11.654  0.00    C15 H18 N9 O4
           388.1490   -0.3   -0.8   -70.5  1626.0  12.049  0.00    C4 H155 N4 O4 S2
           388.1482    0.5    1.3    -12.5  1626.2  12.220  0.00    C2 H34 N3 O14 S2
           388.1481    0.6    1.5    -69.5  1628.8  14.772  0.00    C3 H151 N4 O9
           388.1494   -0.7   -1.8   -64.5  1628.8  14.857  0.00    C4 H147 N8 O5
           388.1487    0.0    0.0    -6.5   1629.6  15.573  0.00    C2 H26 N7 O15
```
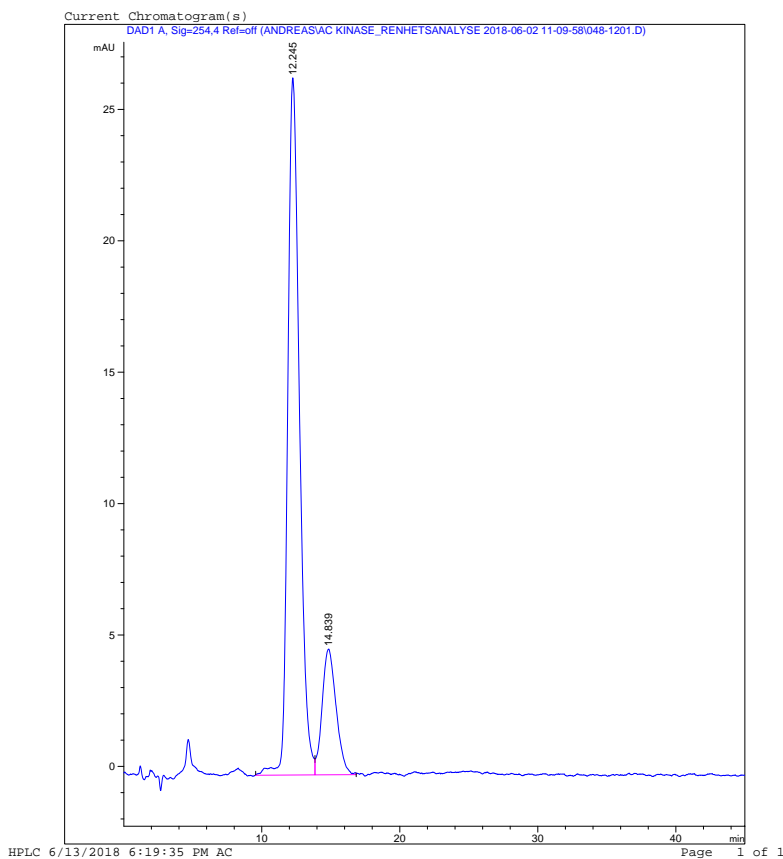
Mass spectrum of compound *(R)*-**6a**.

```
Print of window 38: Current Chromatogram(s)
Data File  : C:\CHEM32\1\DATA\ANDREAS\AC KINASE_RENHETSANALYSE 2018-06-02 11-09-58\049-1001.D
Sample Name : ABR1-59-2 85/15,DEA, FLOW 1.5
=====================================================================
Acq. Operator   : AC                          Seq. Line :  10
Acq. Instrument : HPLC                          Location : Vial 49
Injection Date  : 6/2/2018 7:35:19 PM              Inj :   1
                                            Inj Volume : 5.000 µl
Acq. Method       : C:\CHEM32\1\DATA\ANDREAS\AC KINASE_RENHETSANALYSE 2018-06-02 11-09-58\
                    ANDREAS CHIRAL 85 FLOW 1,5.M
Last changed     : 5/31/2018 10:18:31 AM by AC
Analysis Method : C:\CHEM32\1\METHODS\FELLES\RENHETSANALYSE FELLES_NEUTRAL.M
Last changed     : 6/13/2018 6:16:27 PM by AC
                    (modified after loading)
Additional Info : Peak(s) manually integrated
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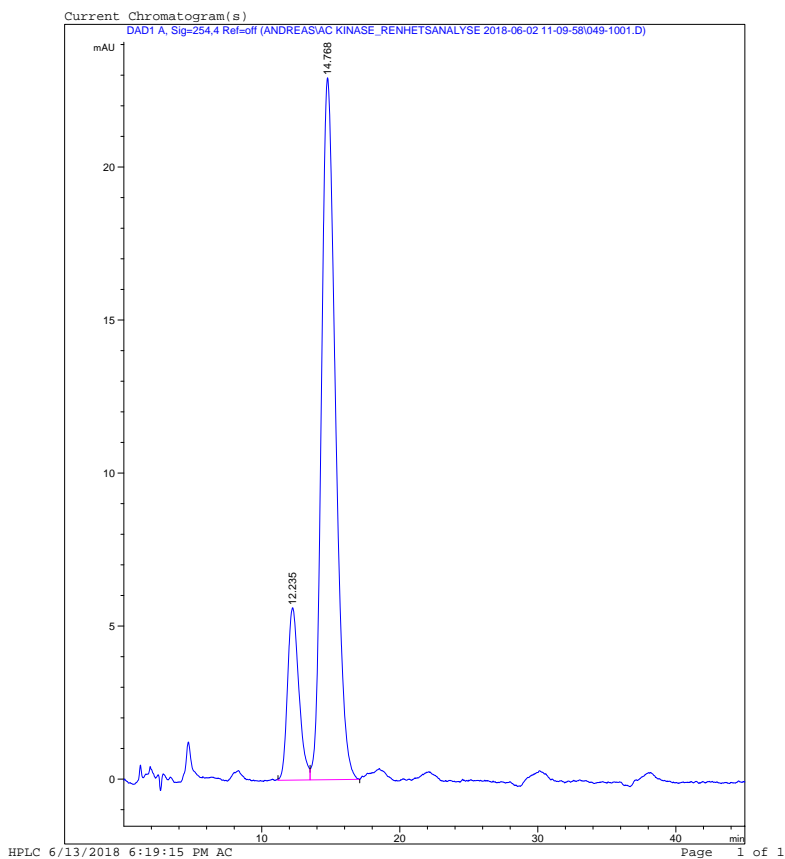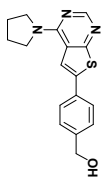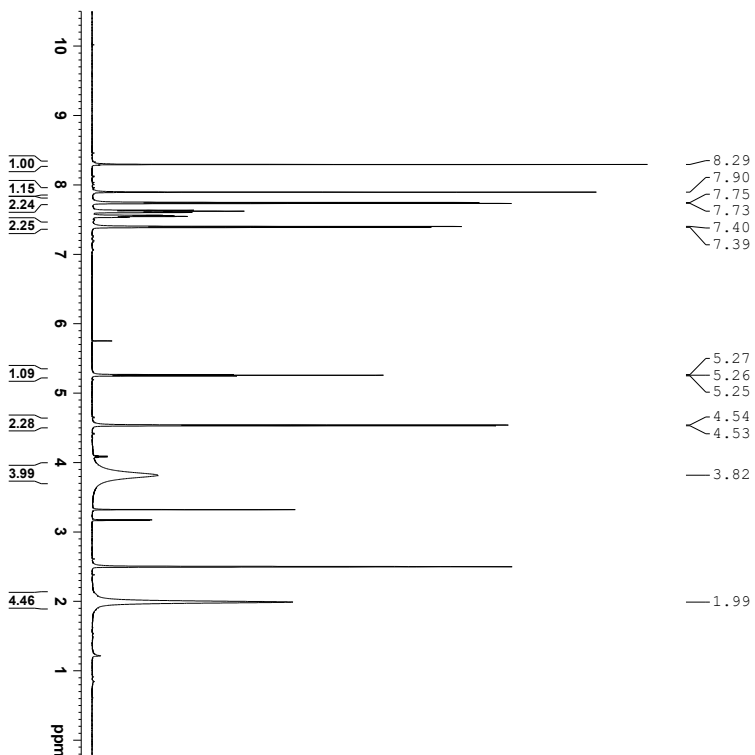

CSP-HPLC (Lux 5u Cellulose-1 4.6 x 250 mm, hexane(cont. 0.2% diethyl amine):*i*-PrOH, flow 1.5 mL/min, detection at 254 nm) of compound *(R)*-**6a**
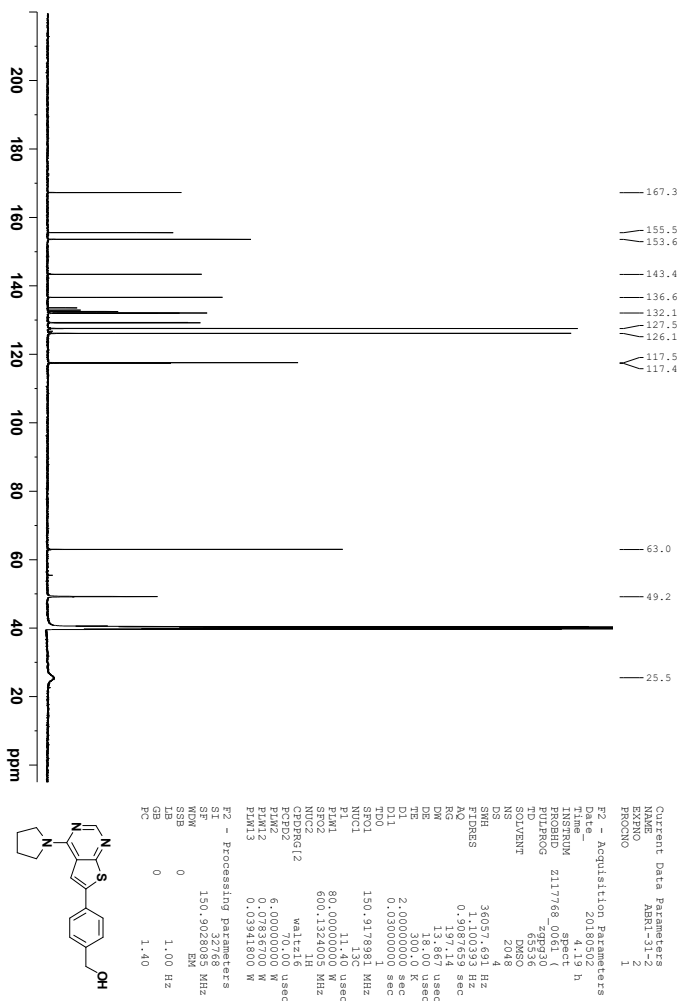
Spectroscopic Data - Compound *(S)*-6a
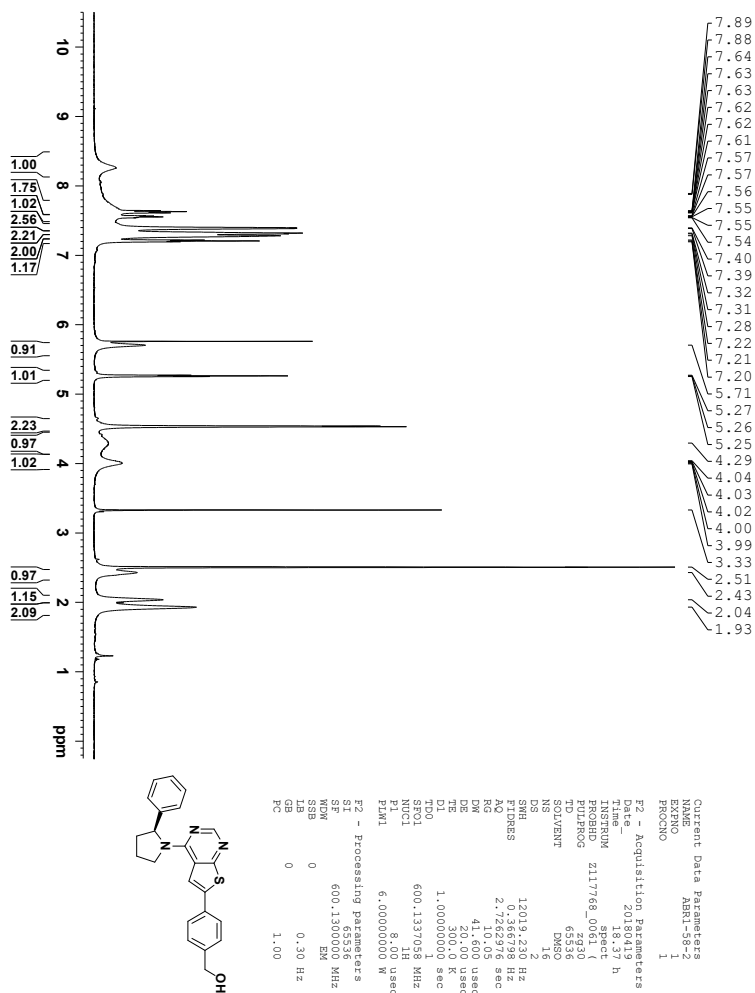
^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound *(S)*-**6a**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *(S)*-**6a**.

COSY spectrum of compound *(S)*-**6a**.

HSQC spectrum of compound *(S)*-**6a**.
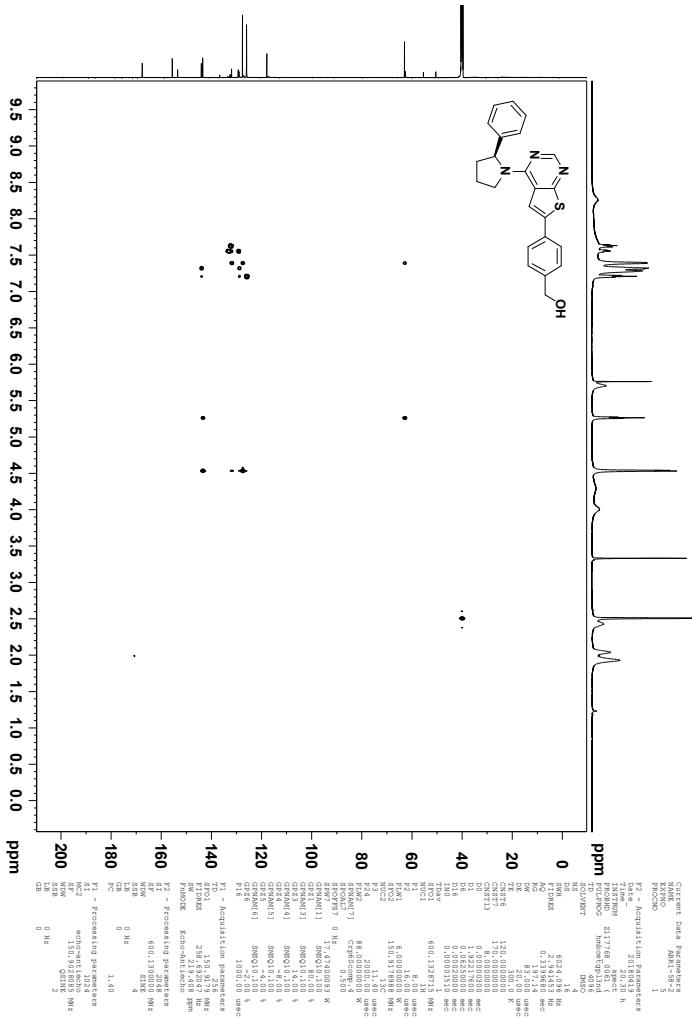
HMBC spectrum of compound *(S)*-**6a**.
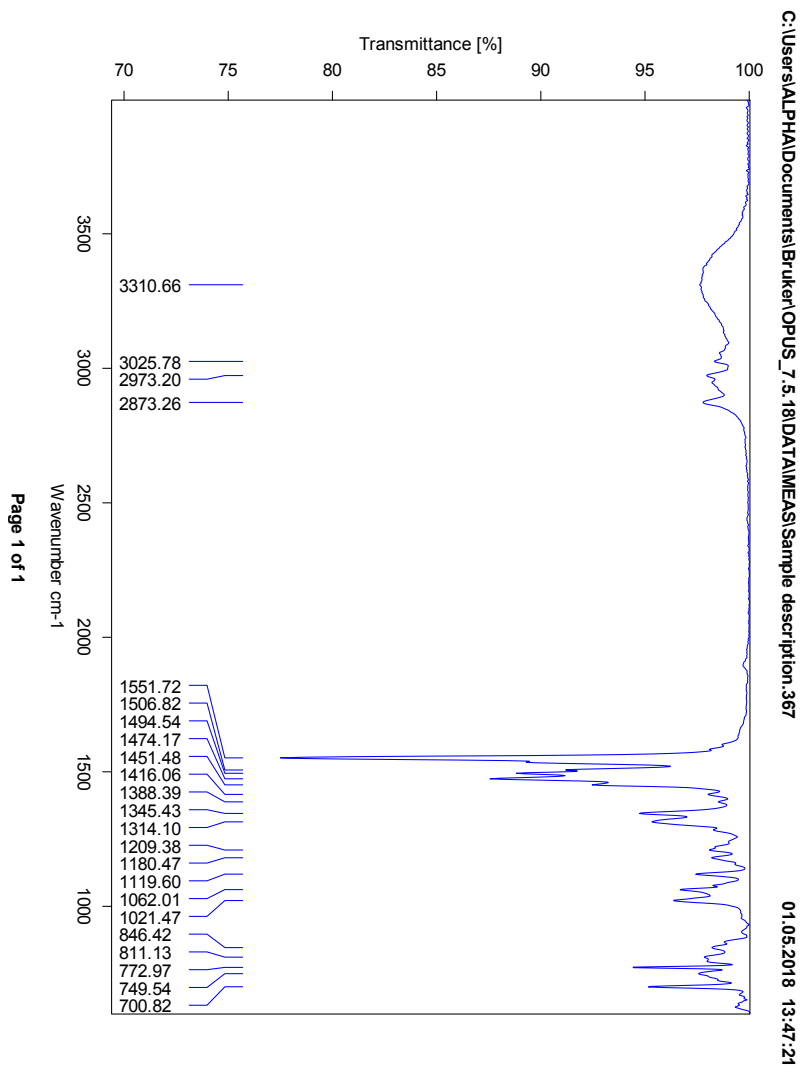
Transmittance [%]

C:\Users\ALPHA\Documents\Bruker\OPUS_7.5.18\DATA\MEAS\Sample description.367

Wavenumber cm-1

Page 1 of 1

3310.66

3025.78
2973.20
2873.26

1551.72
1506.82
1494.54
1474.17
1451.48
1416.06
1388.39
1345.43
1314.10
1209.38
1180.47
1119.60
1062.01
1021.47
846.42
811.13
772.97
749.54
700.82

01.05.2018 13:47:21

IR spectrum of compound *(S)*-**6a**.

Single Mass Analysis
Tolerance = 5.0 PPM   /   DBE: min = -1.5, max = 100.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
3485 formula(e) evaluated with 10 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   Na: 0-1   S: 0-2
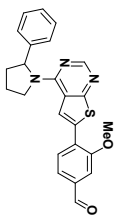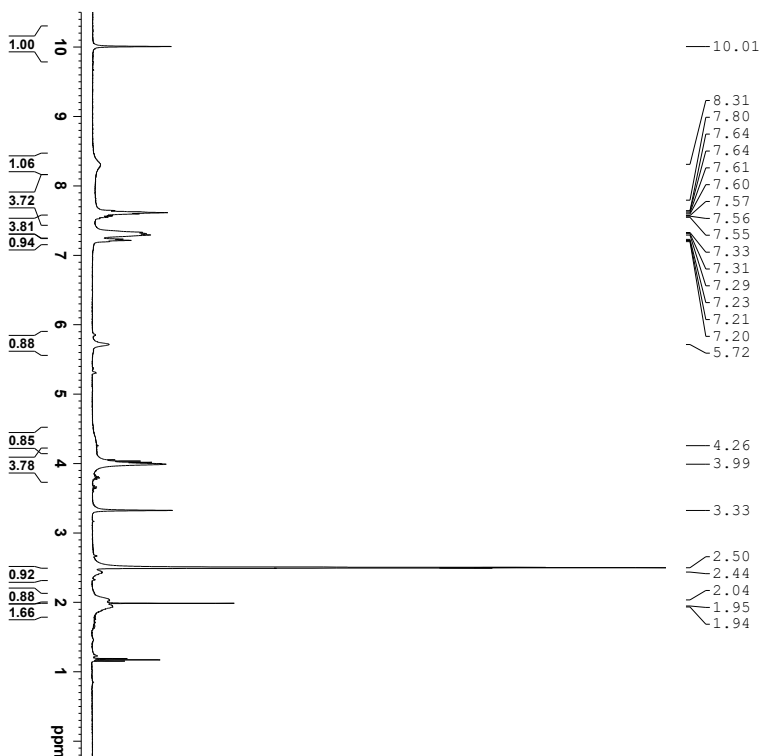2018-214 68 (1.345) AM2 (Ar,35000.0,0.00,0.00); Cm (68:73)
1: TOF MS ASAP+

2.51e+006

100% ┤
                    279.0940

%─ ┤
        124.0872
                    280.0972
            164.1186   388.1482
        281.0999
0 ┼─────────────────────────────────────── m/z
   100   200   300   400   500   600   700   800   900   1000 1100 1200 1300 1400 1500
                              429.1744   663.4532   817.2057   941.2791   1124.0094   1218.4851   1456.0547

Minimum: -1.5
Maximum: 5000.0 5.0   100.0

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
388.1482	388.1484	-0.2	-0.5	14.5	868.2	0.000	100.00	C23 H22 N3 O S
	388.1493	-1.1	-2.8	6.5	881.4	13.241	0.00	C18 H27 N3 O Na S2
	388.1491	-0.9	-2.3	3.5	881.5	13.298	0.00	C10 H23 N9 O4 Na S
	388.1478	0.4	1.0	-1.5	881.6	13.391	0.00	C9 H27 N5 O8 Na S
	388.1477	0.5	1.3	5.5	882.4	14.164	0.00	C15 H26 N5 O3 S2
	388.1464	1.8	4.6	0.5	883.5	15.291	0.00	C14 H30 N O7 S2
	388.1482	0.0	0.0	11.5	884.6	16.419	0.00	C15 H18 N9 O4
	388.1498	-1.6	-4.1	12.5	884.8	16.596	0.00	C18 H19 N7 O2 Na
	388.1485	-0.3	-0.8	7.5	884.8	16.607	0.00	C17 H23 N3 O6 Na
	388.1468	1.4	3.6	6.5	884.9	16.727	0.00	C14 H22 N5 O8

Mass spectrum of compound *(S)*-**6a**.

```
Print of window 38: Current Chromatogram(s)
Data File  : C:\CHEM32\1\DATA\ANDREAS\AC KINASE_RENHETSANALYSE 2018-06-02 11-09-58\048-1201.D
Sample Name : ABR1-58-52 85/15,DEA, FLOW 1.5
==================================================================
Acq. Operator  : AC                        Seq. Line :  12
Acq. Instrument : HPLC                      Location : Vial 48
Injection Date  : 6/2/2018 9:27:24 PM          Inj :   1
                                           Inj Volume : 5.000 µl
Acq. Method    : C:\CHEM32\1\DATA\ANDREAS\AC KINASE_RENHETSANALYSE 2018-06-02 11-09-58\
                 ANDREAS CHIRAL 85 FLOW 1,5.M
Last changed   : 5/31/2018 10:18:31 AM by AC
Analysis Method : C:\CHEM32\1\METHODS\FELLES\RENHETSANALYSE FELLES_NEUTRAL.M
Last changed   : 6/13/2018 6:16:27 PM by AC
                 (modified after loading)
Additional Info : Peak(s) manually integrated
```

CSP-HPLC (Lux 5u Cellulose-1 4.6 x 250 mm, hexane(cont. 0.2% diethyl amine):*i*-PrOH, flow 1.5 mL/min, detection at 254 nm) of compound *(S)*-**6a**

Spectroscopic Data - Compound 6b

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound **6b**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound **6b**.

COSY spectrum of compound **6b**.
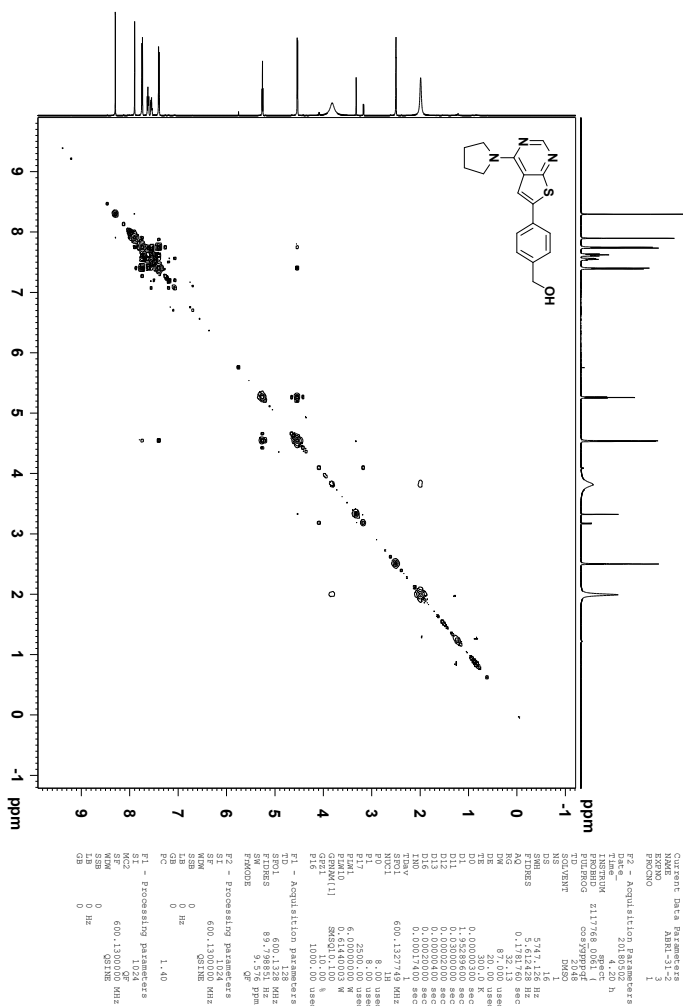
HSQC spectrum of compound **6b**.
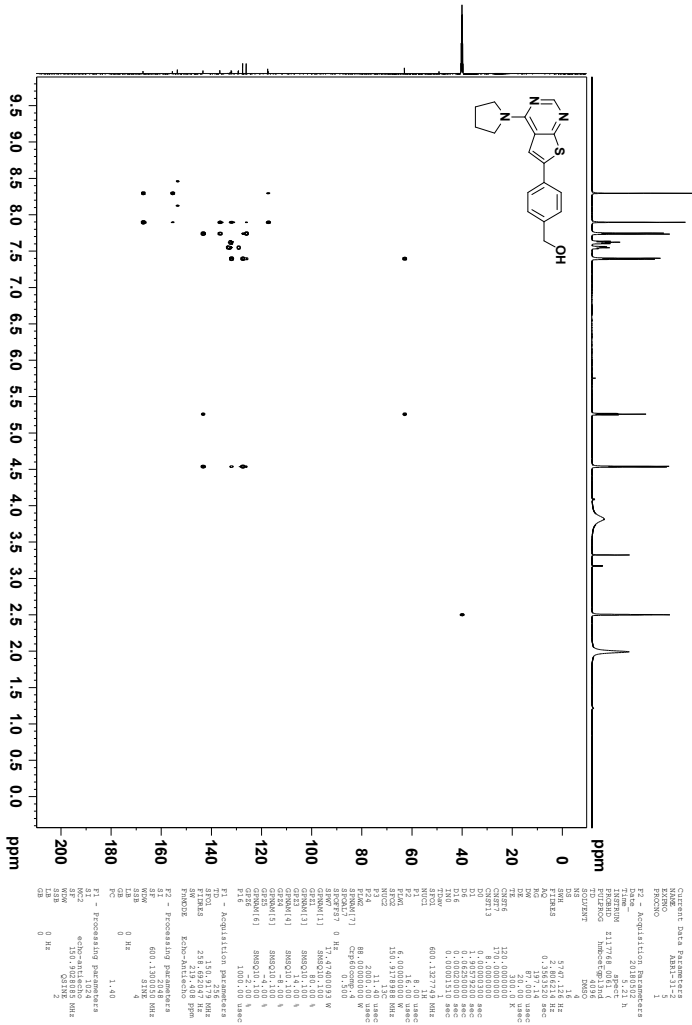
HMBC spectrum of compound **6b**.
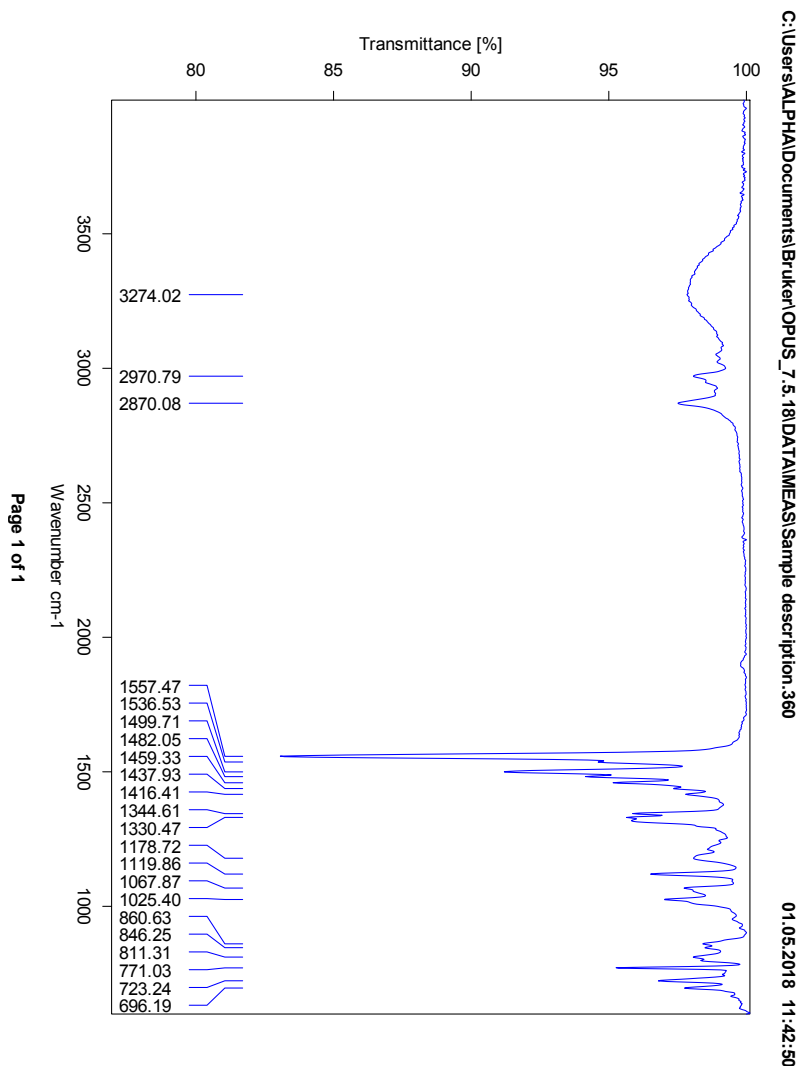
IR spectrum of compound **6b**.

Elemental Composition Report　　　　　　　　　　　　　　　　**Page 1**

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
1411 formula(e) evaluated with 1 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   S: 0-3
2018-247 91 (1.793) AM2 (Ar,35000.0,0.00,0.00); Cm (89:91)
1: TOF MS ASAP+
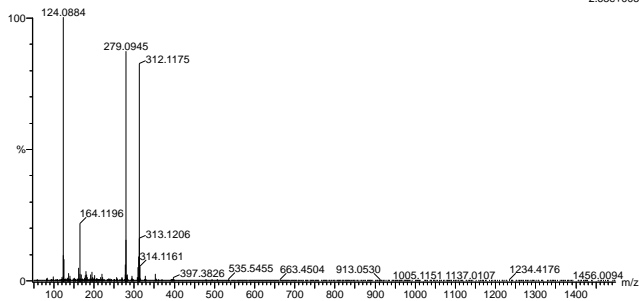
Minimum:			-1.5				
Maximum:		5.0	2.0	50.0			
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%) Formula
312.1175	312.1171	0.4	1.3	10.5	913.7	n/a	n/a C17 H18 N3 O S

Mass spectrum of compound **6b**.

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound *rac*-**7a**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *rac*-**7a**.

COSY spectrum of compound *rac*-**7a**.
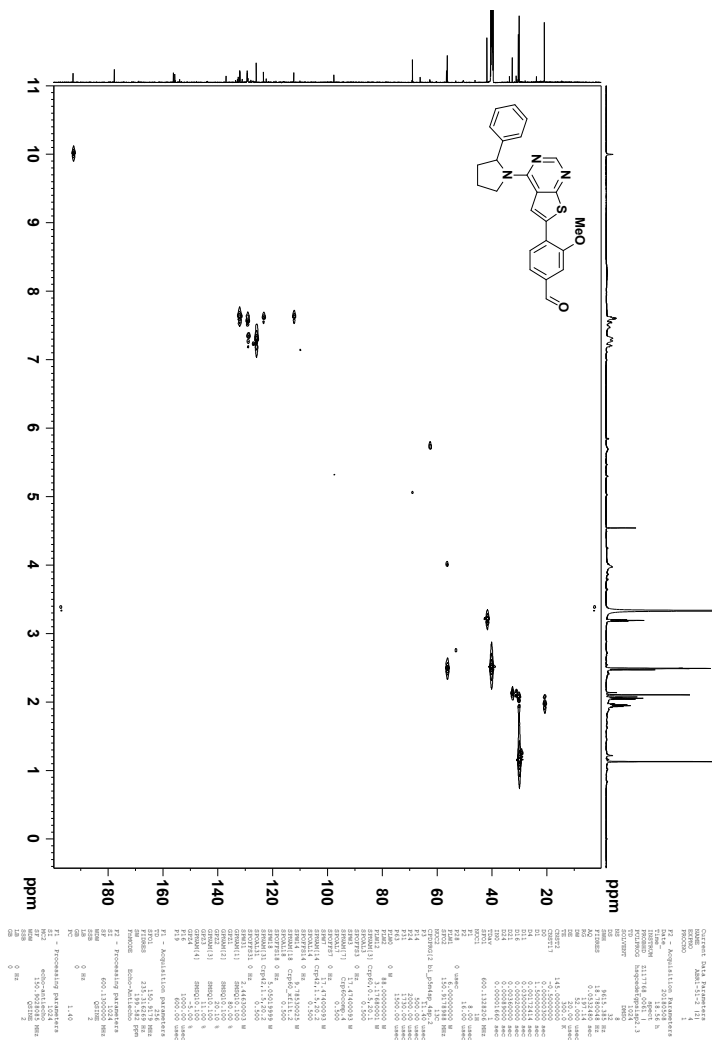
HSQC spectrum of compound *rac*-**7a**.
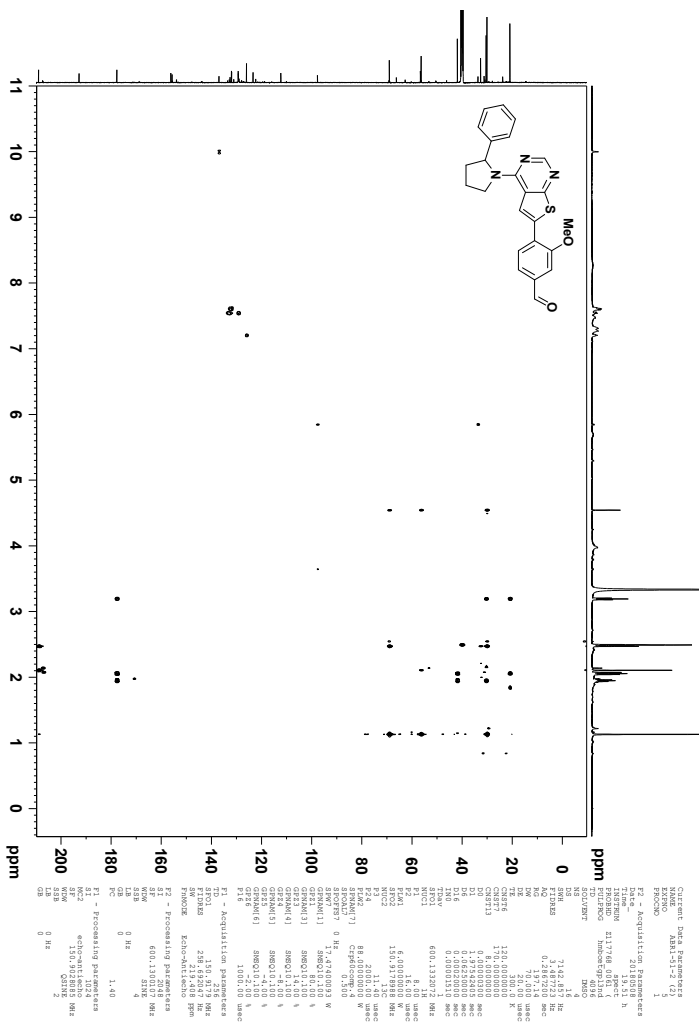
HMBC spectrum of compound *rac*-**7a**.
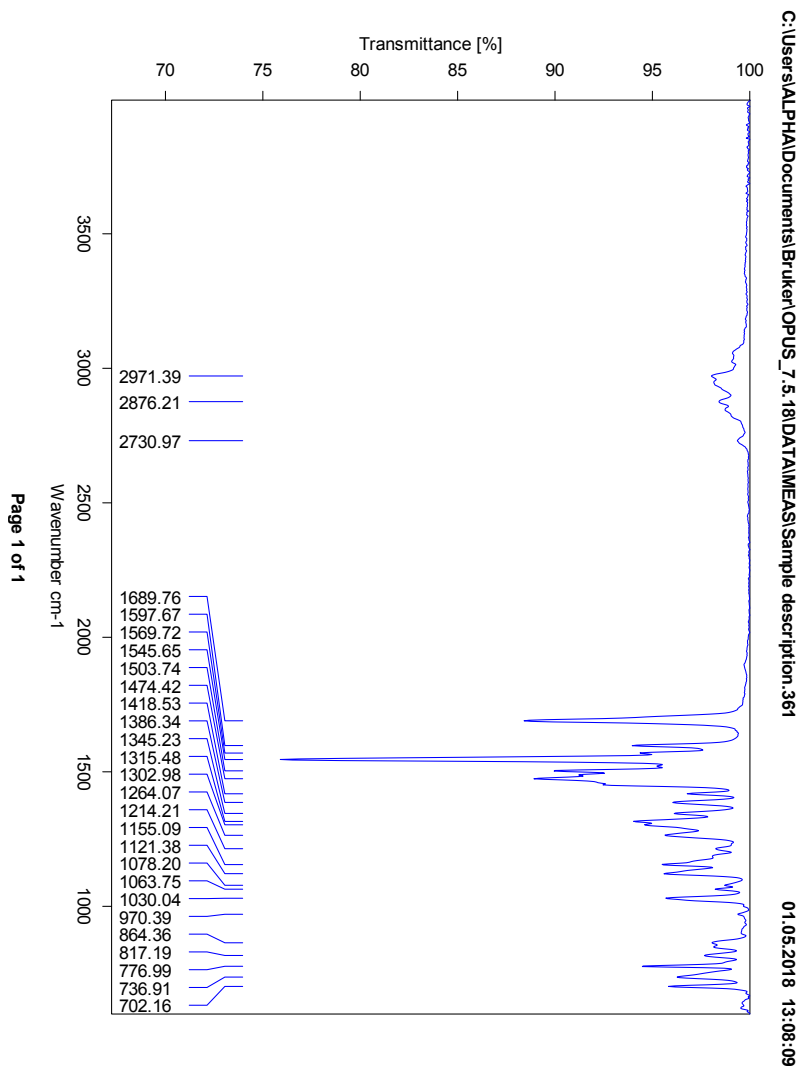
IR spectrum of compound *rac*-**7a**.

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
6815 formula(e) evaluated with 6 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-10   Na: 0-1   S: 0-2   Br: 0-4

2018-176 62 (1.224) AM2 (Ar,35000.0,0.00,0.00); Cm (61:62)
1: TOF MS ASAP+
                                                                                                                                              2.39e+004
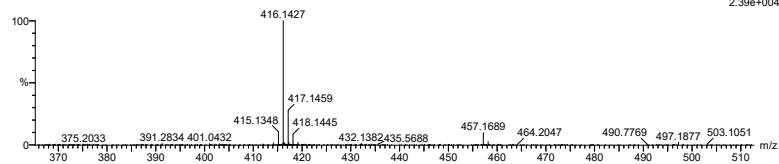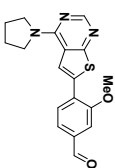


```
Minimum:                              -1.5
Maximum:                5000.0  2.0   50.0

Mass        Calc. Mass  mDa   PPM   DBE   i-FIT  Norm   Conf(%)  Formula
416.1427    416.1433    -0.6  -1.4  15.5  431.2  0.000  100.00   C24 H22 N3 O2 S
            416.1426     0.1   0.2   6.5  446.1  14.847   0.00   C16 H26 N5 O4 S2
            416.1427     0.0   0.0  -0.5  446.1  14.881   0.00   C10 H27 N5 O9 Na S
            416.1431    -0.4  -1.0  12.5  446.4  15.128   0.00   C16 H18 N9 O5
            416.1434    -0.7  -1.7   8.5  446.8  15.565   0.00   C18 H23 N3 O7 Na
            416.1426     0.1   0.2   6.5  453.6  22.369   0.00   C18 H28 N5 Na Br
```

Mass spectrum of compound *rac*-**7a**.

^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound _(R)_-**7a**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *(R)*-**7a**.
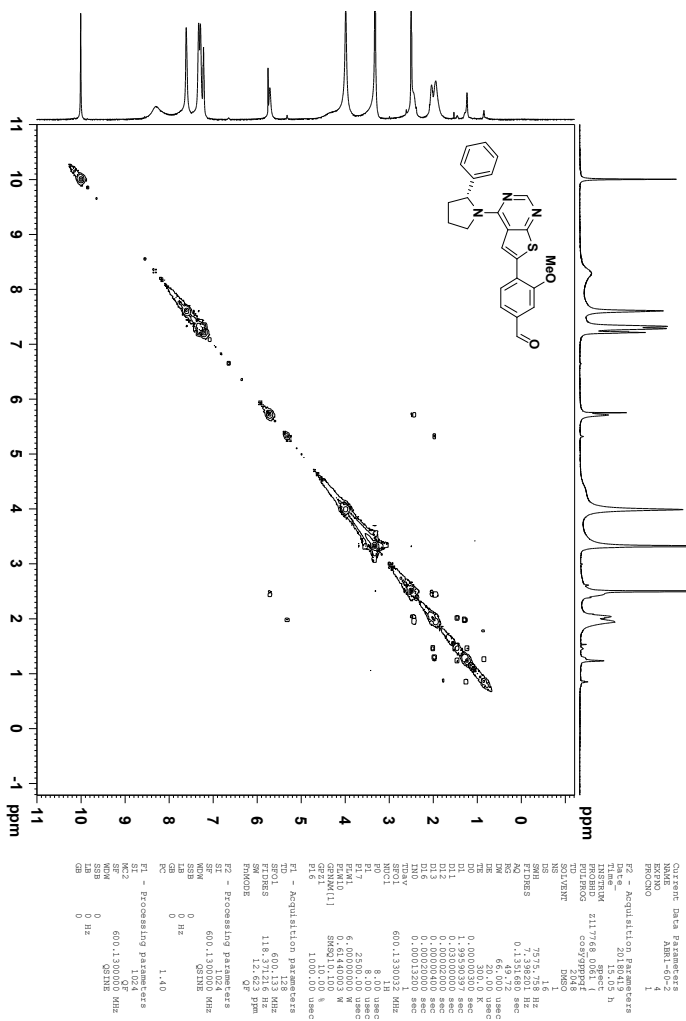
COSY spectrum of compound *(R)*-**7a**.
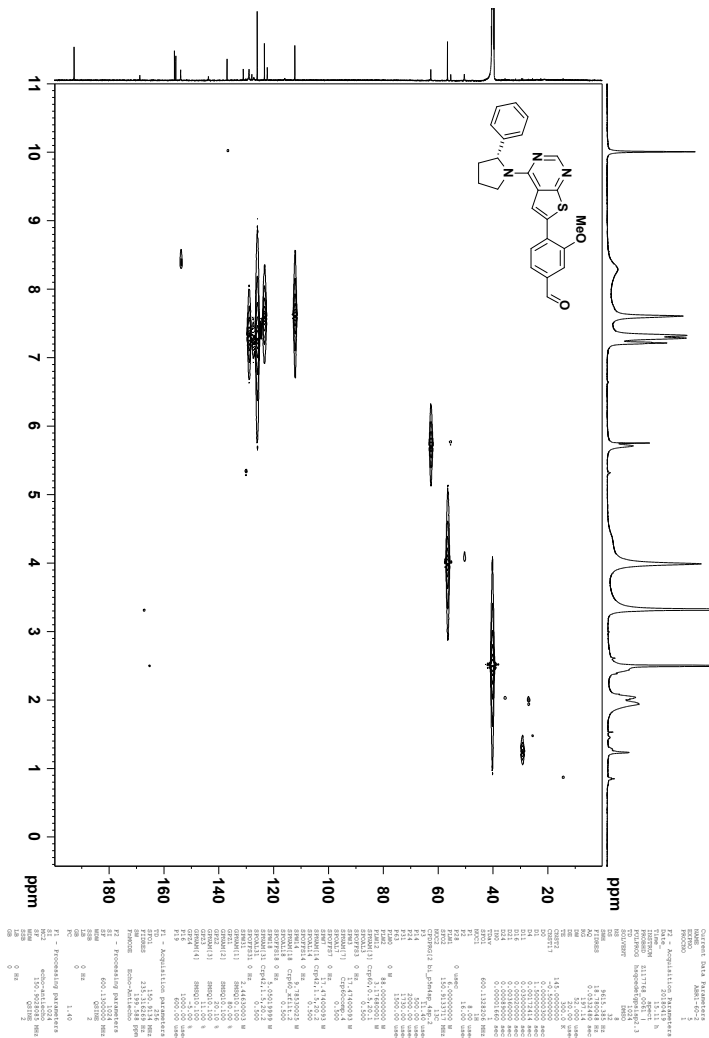
HSQC spectrum of compound *(R)*-**7a**.
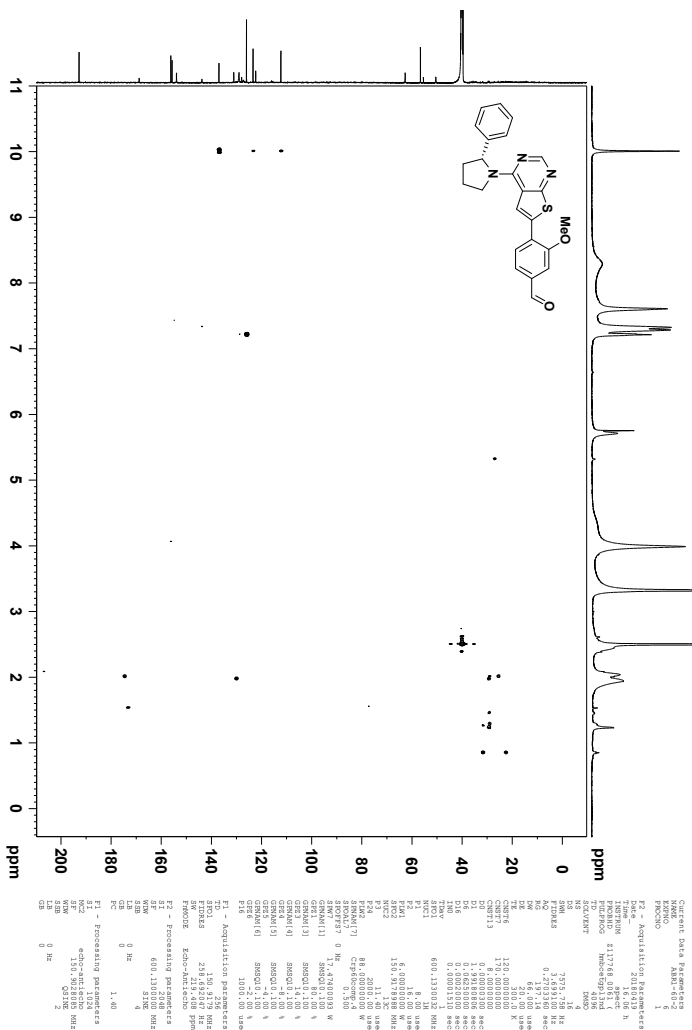
HMBC spectrum of compound *(R)*-**7a**.
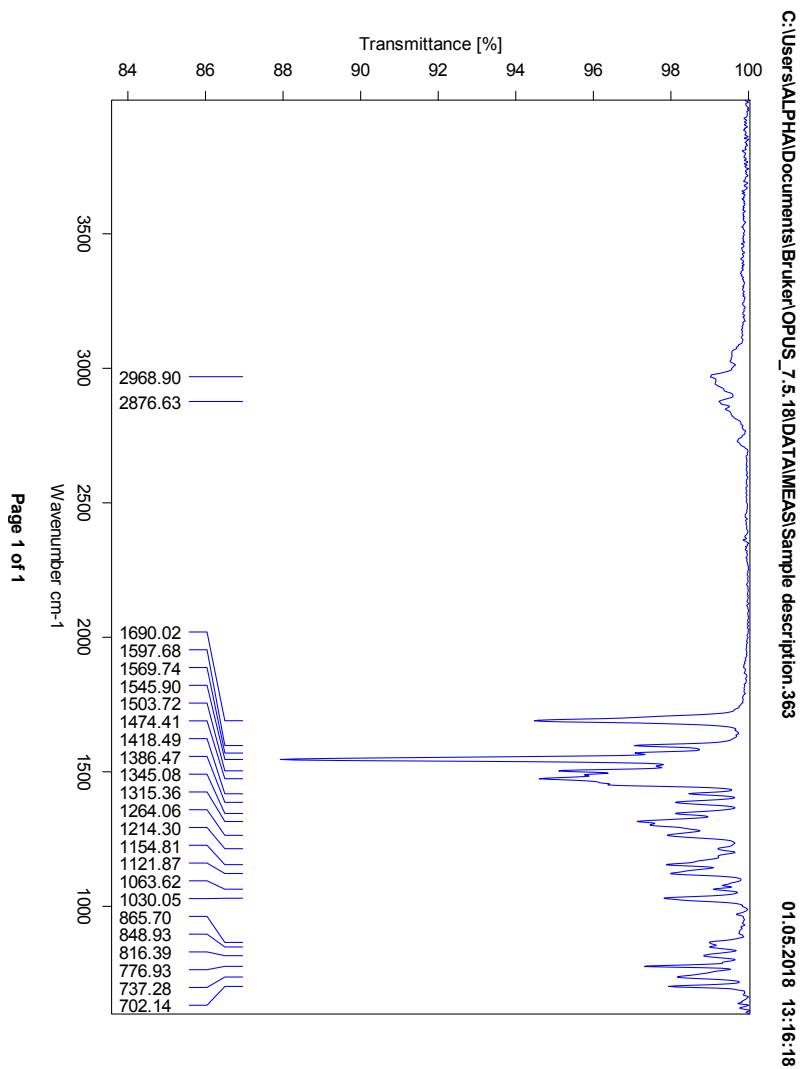
IR spectrum of compound *(R)*-**7a**.

Elemental Composition Report   **Page 1**

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
2625 formula(e) evaluated with 4 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   S: 0-3
2018-246 135 (2.636) AM2 (Ar,35000.0,0.00,0.00); Cm (131:136)
1: TOF MS ASAP+
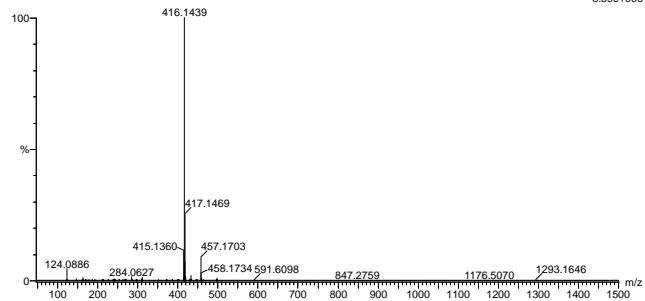
Minimum:				-1.5			
Maximum:		5.0	2.0	50.0			
Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%) Formula
416.1439	416.1439	0.0	0.0	24.5	1251.9	11.790	0.00 C32 H18 N
	416.1440	-0.1	-0.2	11.5	1248.4	8.262	0.03 C17 H22 N9 S2
	416.1433	0.6	1.4	15.5	1240.1	0.000	99.97 C24 H22 N3 O2 S
	416.1431	0.8	1.9	12.5	1250.1	10.008	0.00 C16 H18 N9 O5

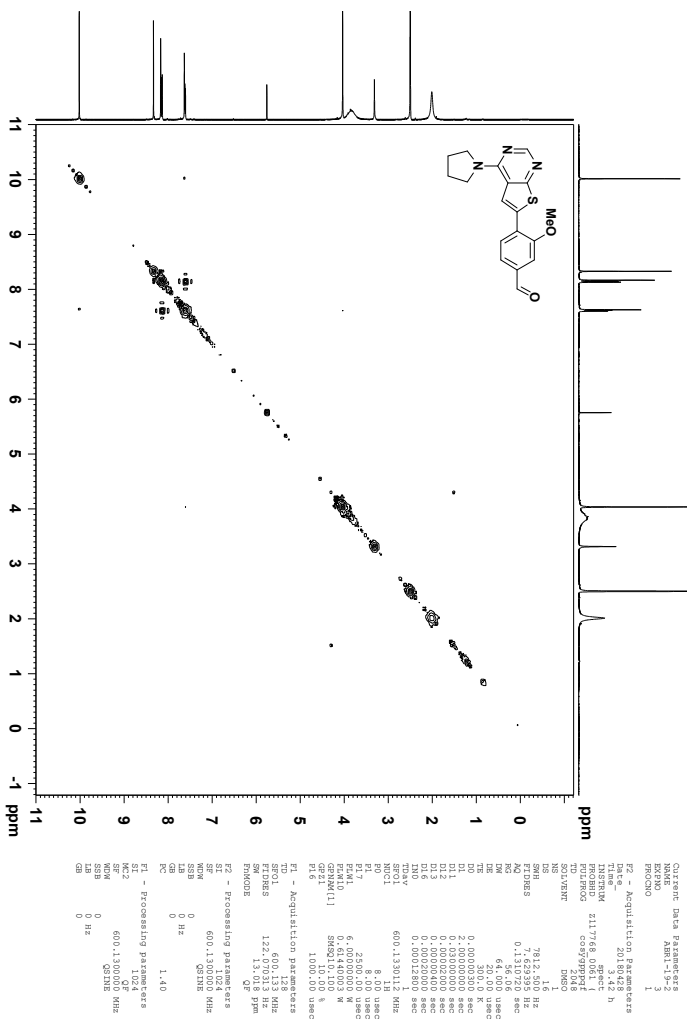Mass spectrum of compound *(R)*-**7a**.

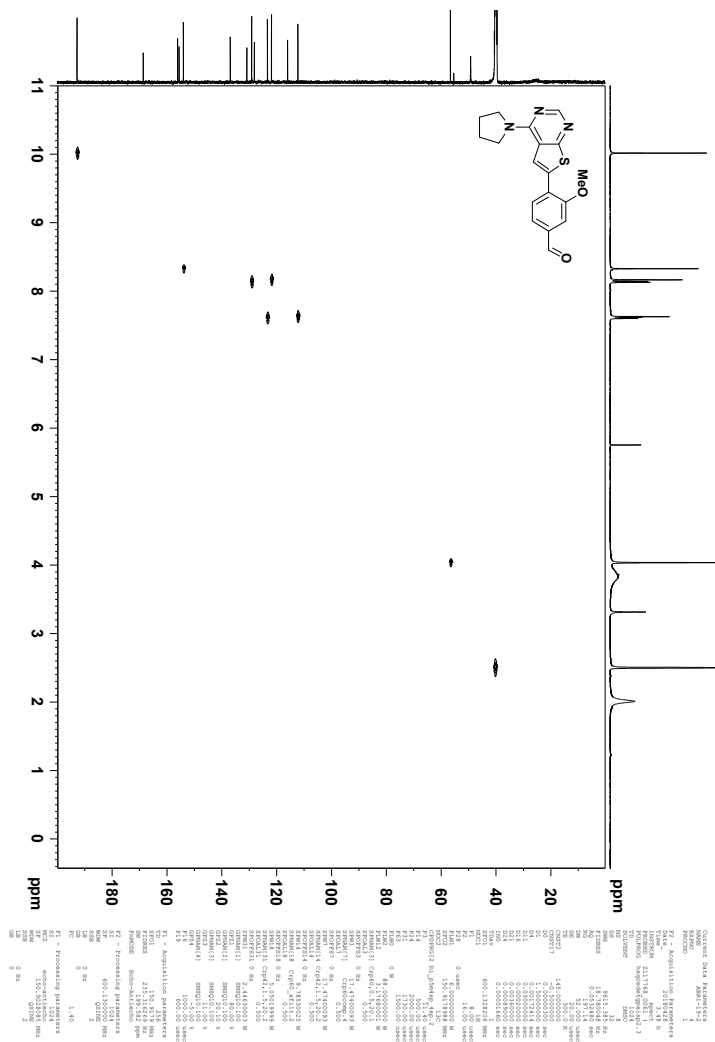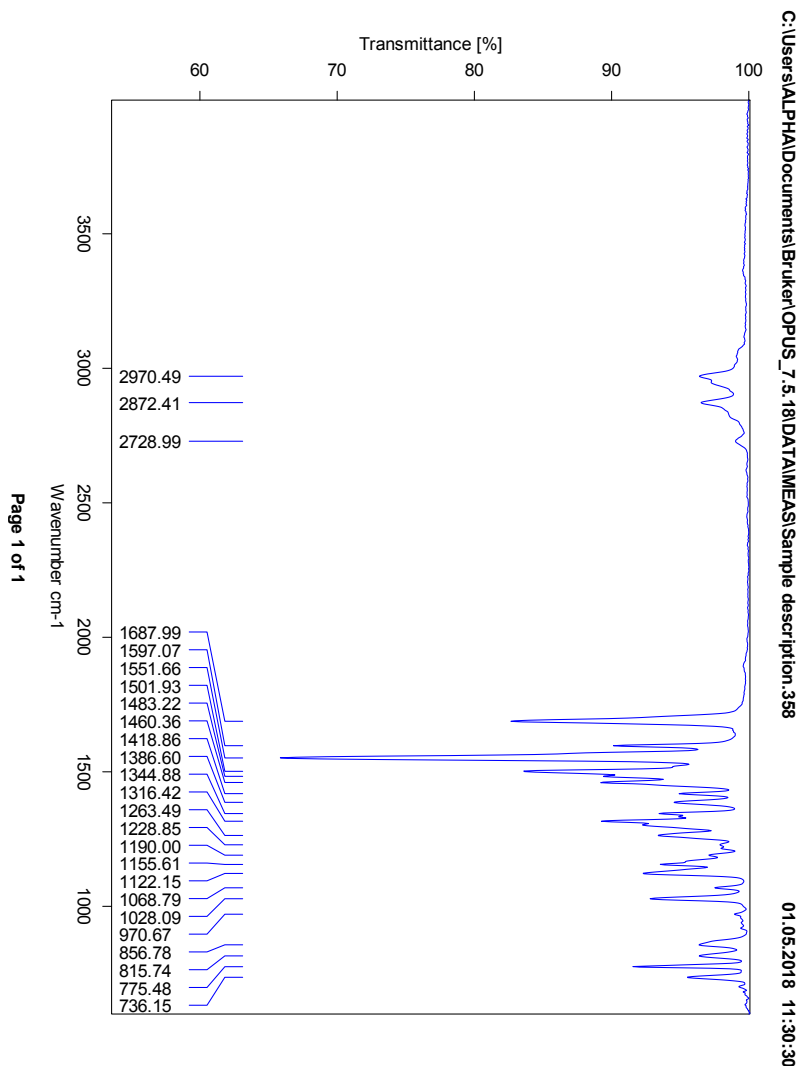^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound **7b**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound **7b**.

COSY spectrum of compound **7b**.

HSQC spectrum of compound **7b**.

HMBC spectrum of compound **7b**.

IR spectrum of compound **7b**.

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
3762 formula(e) evaluated with 9 results within limits (all results (up to 1000) for each mass)
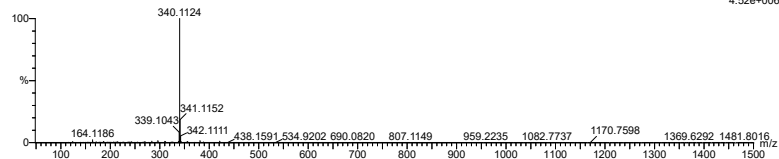Elements Used:
C: 0-500   H: 0-1000   B: 0-1   N: 0-50   O: 0-50   S: 0-2
2018-84 140 (2.740) AM2 (Ar,35000.0,0.00,0.00); Cm (138:142)
1: TOF MS ASAP+

4.52e+006

```
Minimum:                        -1.5
Maximum:              5.0   2.0  50.0

Mass       Calc. Mass  mDa   PPM   DBE   i-FIT   Norm  Conf(%) Formula
340.1124   340.1120    0.4   1.2   11.5  1547.5  0.001 99.88   C18 H18 N3 O2 S
           340.1127   -0.3  -0.9   7.5   1554.4  6.866  0.10   C11 H18 N9 S2
           340.1125   -0.1  -0.3   4.5   1556.3  8.841  0.01   C3 H14 N15 O3 S
           340.1118    0.6   1.8   8.5   1557.8  10.331 0.00   C10 H14 N9 O5
           340.1126   -0.2  -0.6   20.5  1560.3  12.831 0.00   C26 H14 N
           340.1118    0.6   1.8   16.5  1564.1  16.635 0.00   C17 H11 B N7 O
           340.1123    0.1   0.3   -1.5  1569.2  21.669 0.00   C4 H19 B N5 O12
           340.1118    0.6   1.8   3.5   1573.3  25.825 0.00   C2 H15 B N15 O S2
           340.1123    0.1   0.3   9.5   1574.4  26.867 0.00   C2 H7 B N19 O2
```
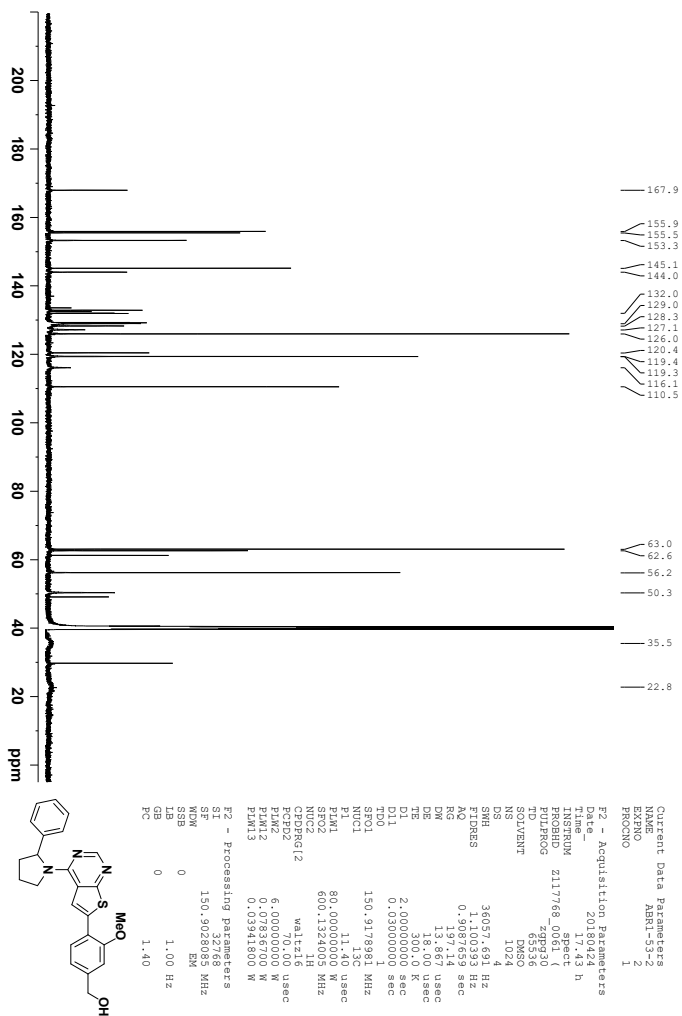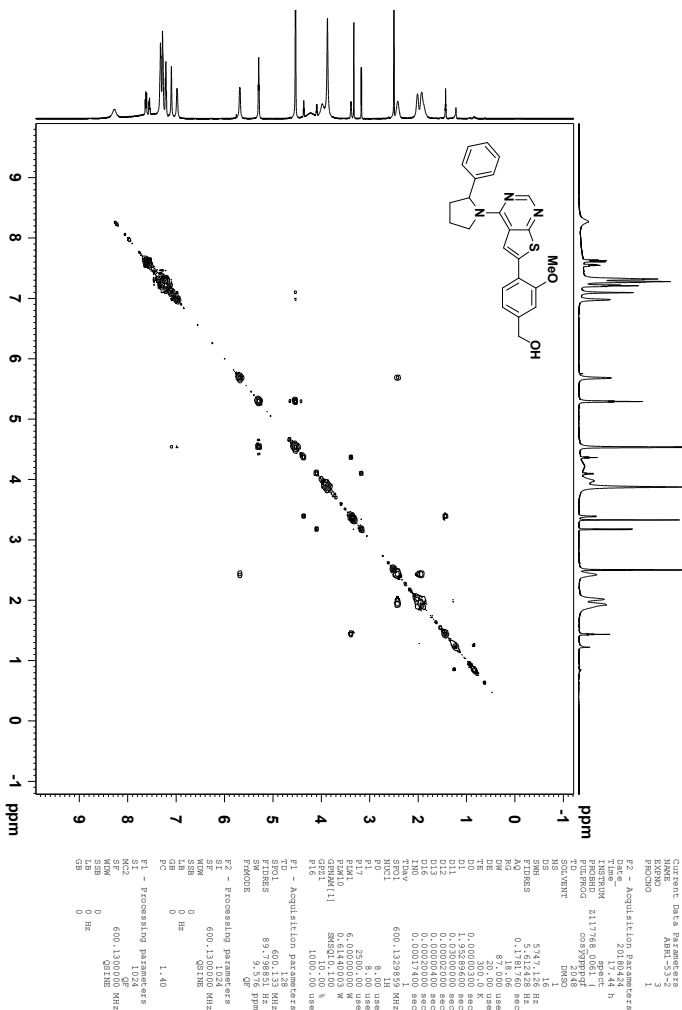
Mass spectrum of compound **7b**.

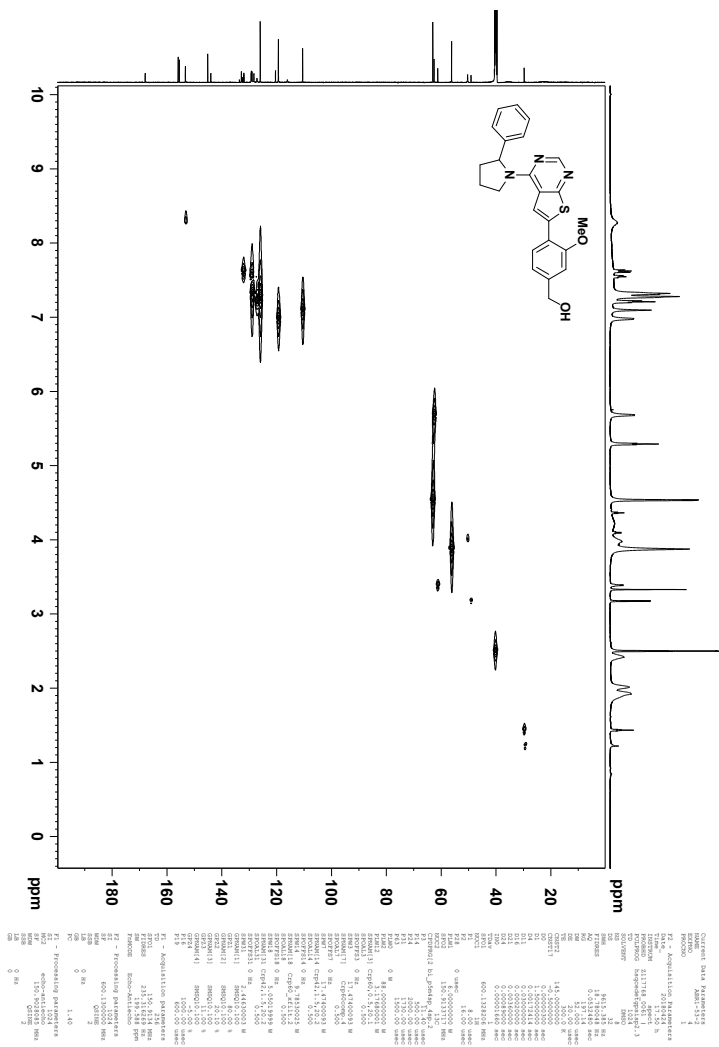# Spectroscopic Data - Compound *rac*-8a
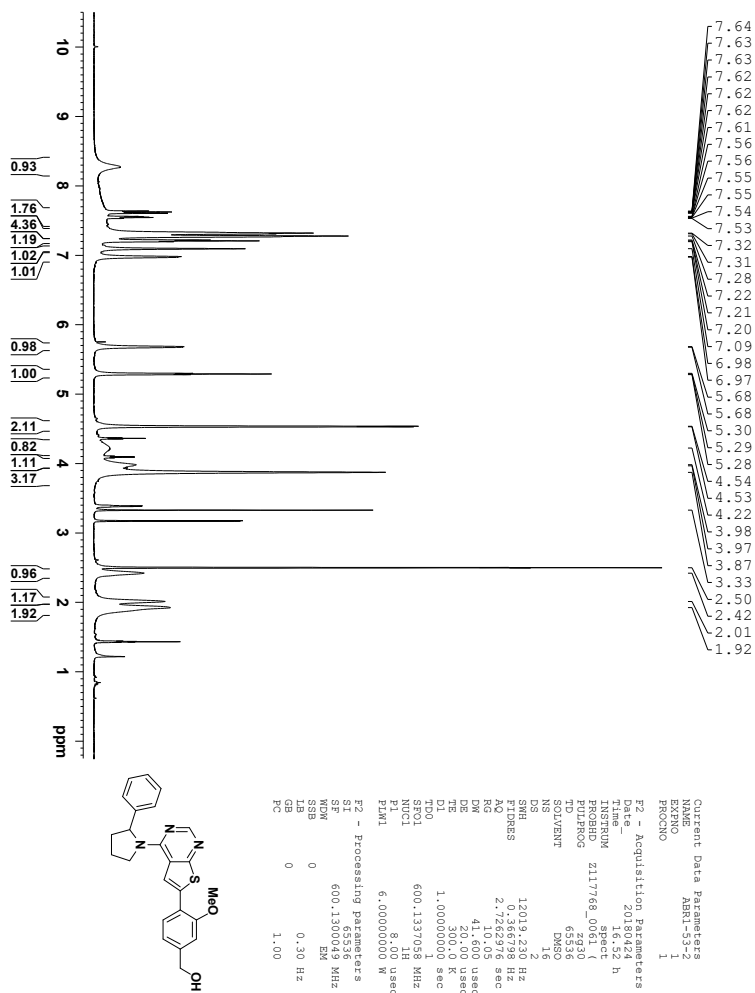
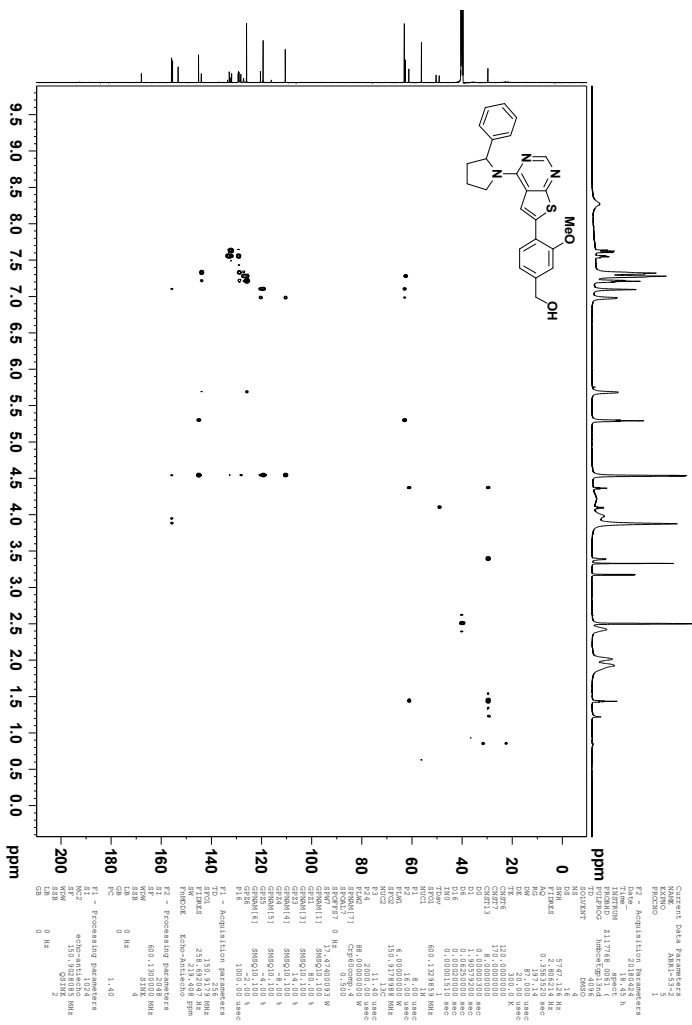^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound *rac*-**8a**.

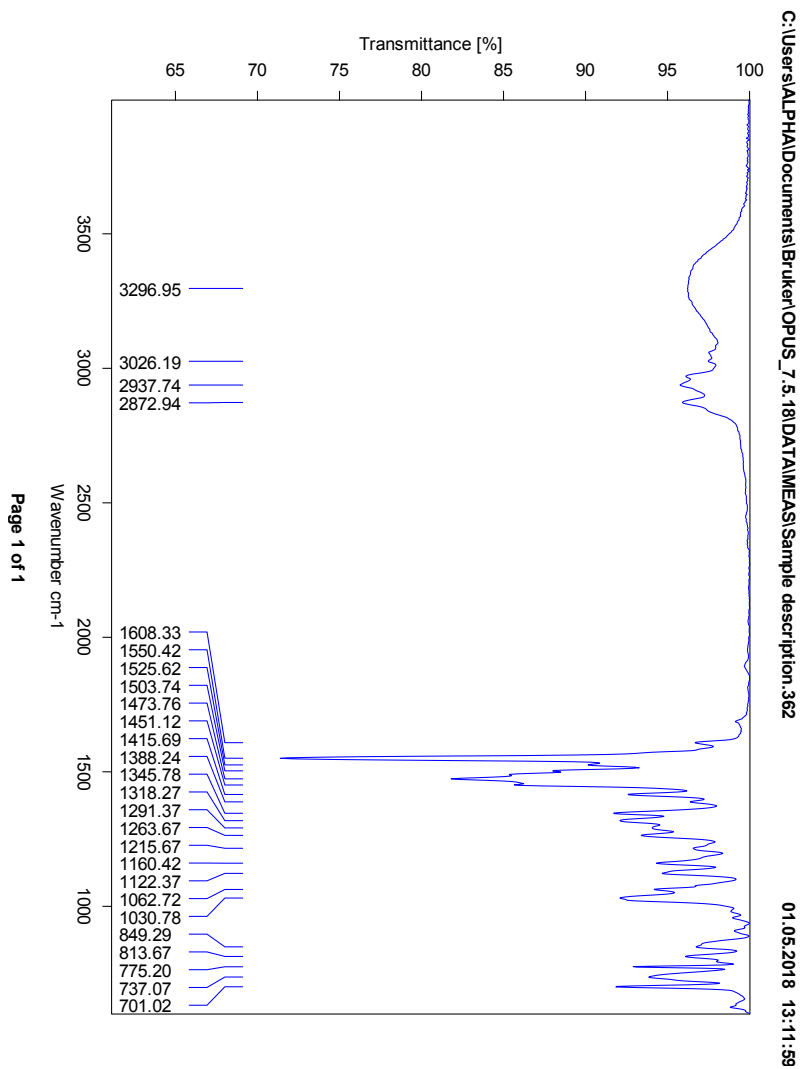^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *rac*-**8a**.

COSY spectrum of compound *rac*-**8a**.

HSQC spectrum of compound *rac*-**8a**.

HMBC spectrum of compound *rac*-**8a**.

IR spectrum of compound *rac*-**8a**.

Single Mass Analysis
Tolerance = 2.0 PPM / DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
6889 formula(e) evaluated with 7 results within limits (all results (up to 1000) for each mass)
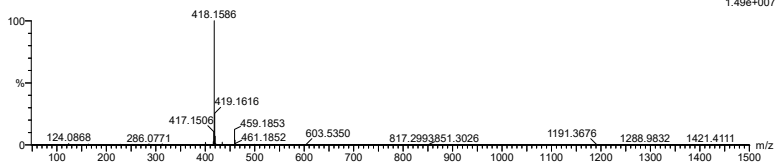Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-10   Na: 0-1   S: 0-2   Br: 0-4

2018-177 149 (2.912) AM2 (Ar,35000.0,0.00,0.00); Cm (144:149)
1: TOF MS ASAP+


Minimum:			-1.5			
Maximum:		5000.0	2.0	50.0		

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
418.1586	418.1589	-0.3	-0.7	14.5	1400.4	0.001	99.90	C24 H24 N3 O2 S
	418.1584	0.2	0.5	-1.5	1407.5	7.057	0.09	C10 H29 N5 O9 Na S
	418.1583	0.3	0.7	5.5	1409.6	9.111	0.01	C16 H28 N5 O4 S2
	418.1587	-0.1	-0.2	11.5	1411.0	10.555	0.00	C16 H20 N9 O5
	418.1590	-0.4	-1.0	7.5	1411.1	10.681	0.00	C18 H25 N3 O7 Na
	418.1593	-0.7	-1.7	3.5	1416.3	15.887	0.00	C19 H33 N O4 Br
	418.1582	0.4	1.0	5.5	1416.6	16.174	0.00	C18 H30 N5 Na Br

Mass spectrum of compound *rac*-**8a**.
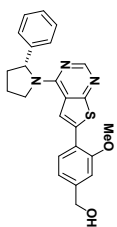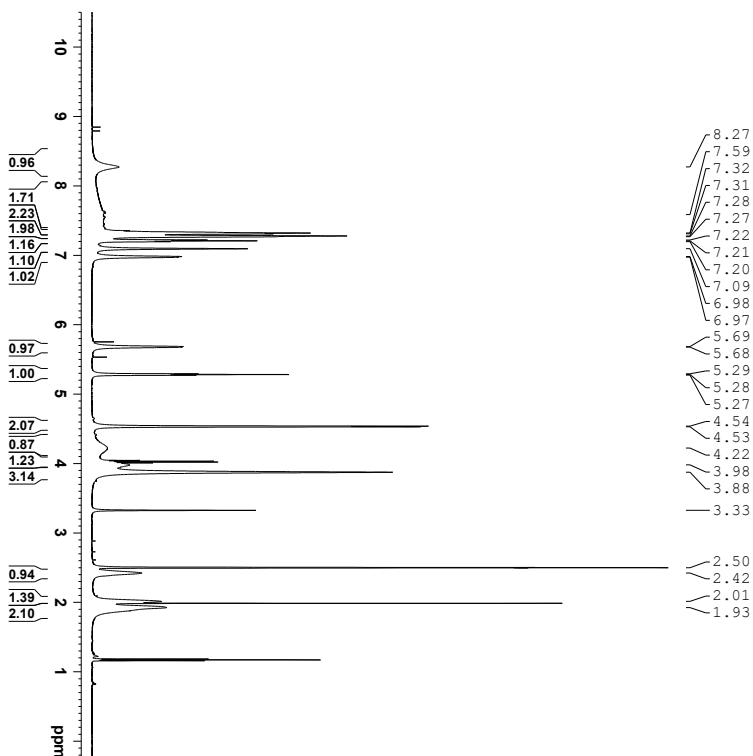
^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound *(R)*-**8a**.
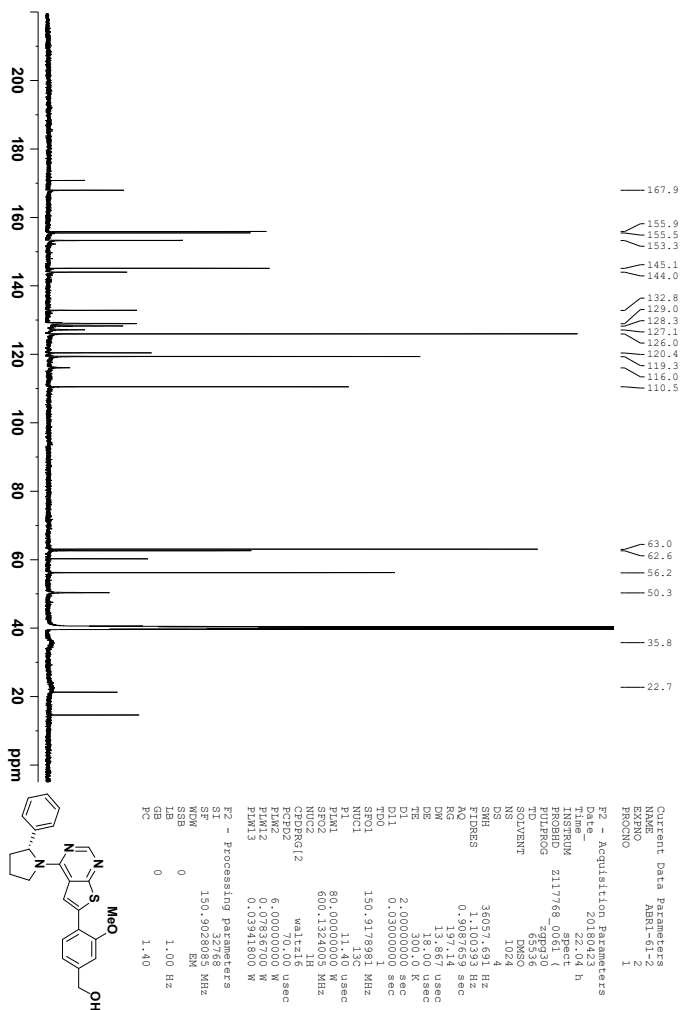
^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound *(R)*-**8a**.
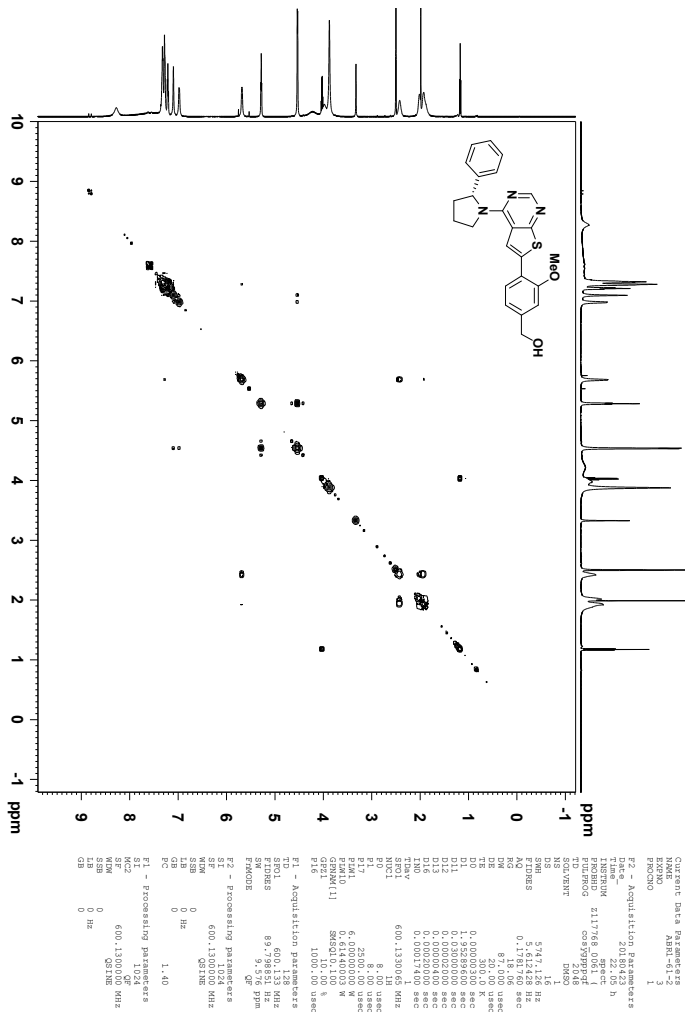
COSY spectrum of compound *(R)*-**8a**.
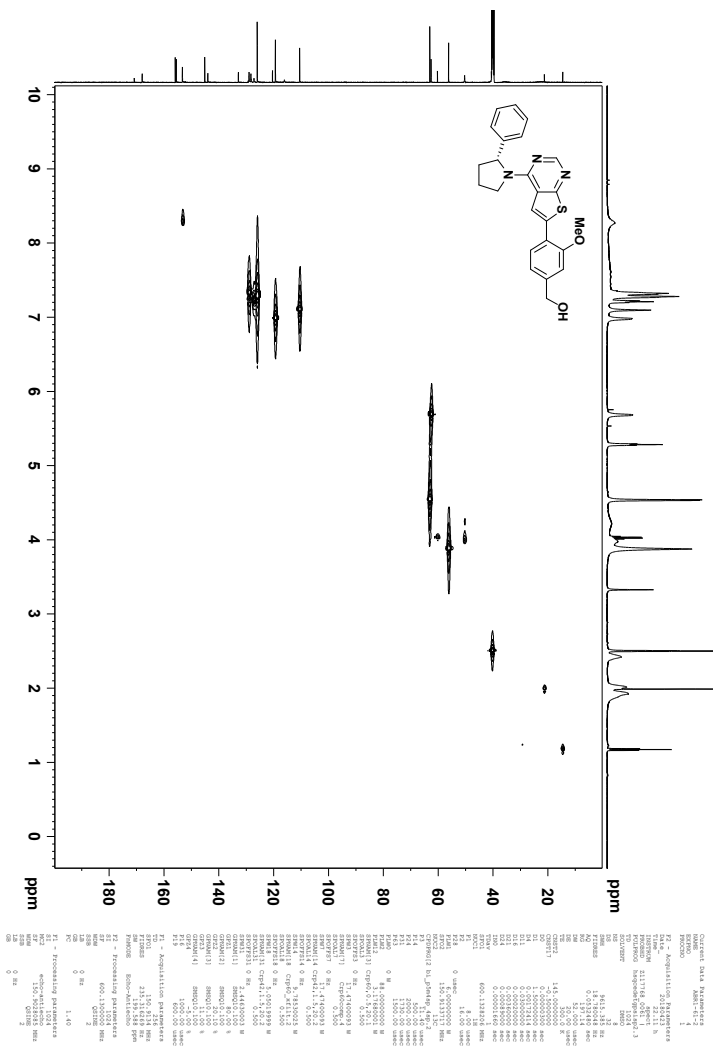
HSQC spectrum of compound *(R)*-**8a**.
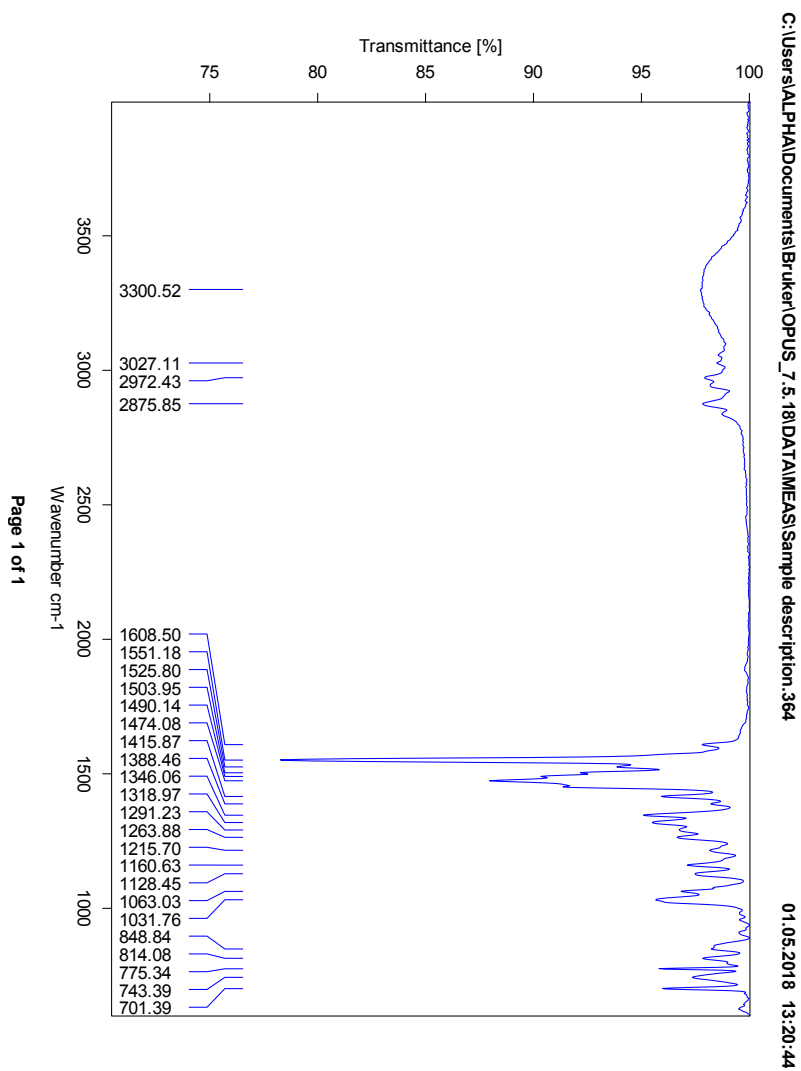
HMBC spectrum of compound *(R)*-**8a**.

Transmittance [%]

C:\Users\ALPHA\Documents\Bruker\OPUS_7.5.18\DATA\MEAS\Sample description.364

01.05.2018 13:20:44

Wavenumber cm-1

Page 1 of 1

3300.52

3027.11
2972.43
2875.85

1608.50
1551.18
1525.80
1503.95
1490.14
1474.08
1415.87
1388.46
1346.06
1318.97
1291.23
1263.88
1215.70
1160.63
1128.45
1063.03
1031.76
848.84
814.08
775.34
743.39
701.39

IR spectrum of compound *(R)*-**8a**.

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 100.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions
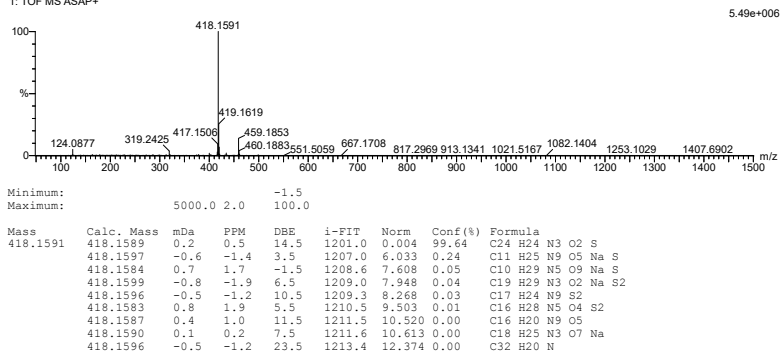4042 formula(e) evaluated with 9 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   Na: 0-1   S: 0-2
2018-225 144 (2.808) AM2 (Ar,35000.0,0.00,0.00); Cm (137:144)
1: TOF MS ASAP+

5.49e+006


```
Minimum:                     -1.5
Maximum:        5000.0 2.0   100.0

Mass        Calc. Mass  mDa   PPM   DBE   i-FIT   Norm    Conf(%)  Formula
418.1591    418.1589    0.2   0.5   14.5  1201.0  0.004   99.64    C24 H24 N3 O2 S
            418.1597   -0.6  -1.4   3.5   1207.0  6.033   0.24     C11 H25 N9 O5 Na S
            418.1584    0.7   1.7  -1.5   1208.6  7.608   0.05     C10 H29 N5 O9 Na S
            418.1599   -0.8  -1.9   6.5   1209.0  7.948   0.04     C19 H29 N3 O2 Na S2
            418.1596   -0.5  -1.2   10.5  1209.3  8.268   0.03     C17 H24 N9 S2
            418.1583    0.8   1.9   5.5   1210.5  9.503   0.01     C16 H28 N5 O4 S2
            418.1587    0.4   1.0   11.5  1211.5  10.520  0.00     C16 H20 N9 O5
            418.1590    0.1   0.2   7.5   1211.6  10.613  0.00     C18 H25 N3 O7 Na
            418.1596   -0.5  -1.2   23.5  1213.4  12.374  0.00     C32 H20 N
```

Mass spectrum of compound *(R)*-**8a**.

```
Print of window 38: Current Chromatogram(s)
Data File  : C:\CHEM32\1\DATA\ANDREAS\AC KINASE_RENHETSANALYSE 2018-06-02 11-09-58\050-0801.D
Sample Name : ABR1-61-2 85/15,DEA, FLOW 1.5
==================================================================
Acq. Operator   : AC                          Seq. Line :   8
Acq. Instrument : HPLC                         Location : Vial 50
Injection Date  : 6/2/2018 5:43:15 PM          Inj :   1
                                               Inj Volume : 5.000 µl
Acq. Method     : C:\CHEM32\1\DATA\ANDREAS\AC KINASE_RENHETSANALYSE 2018-06-02 11-09-58\
                  ANDREAS CHIRAL 85 FLOW 1,5.M
Last changed    : 5/31/2018 10:18:31 AM by AC
Analysis Method : C:\CHEM32\1\METHODS\FELLES\RENHETSANALYSE FELLES_NEUTRAL.M
Last changed    : 6/13/2018 6:16:27 PM by AC
                  (modified after loading)
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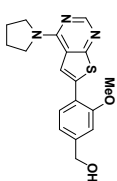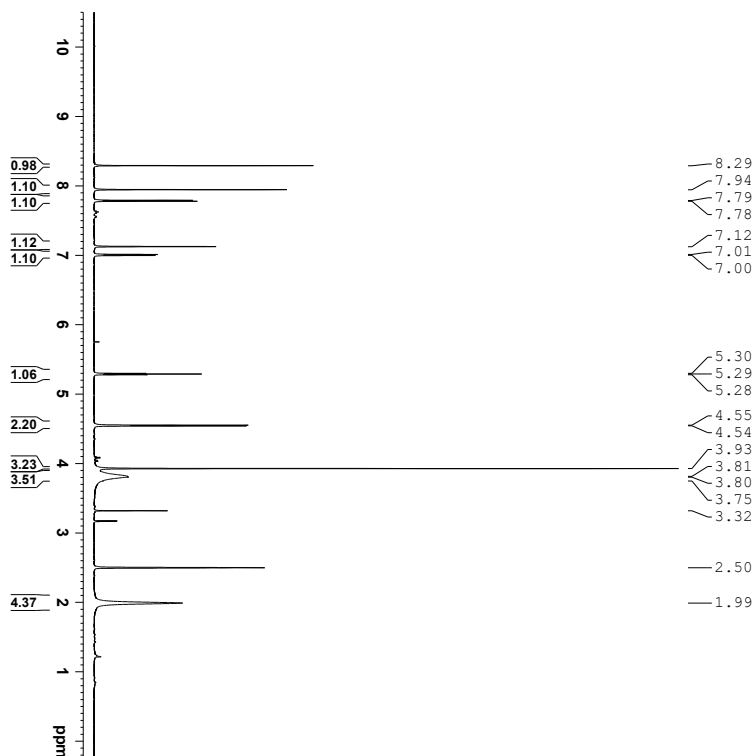
Current Chromatogram(s)

HPLC 6/13/2018 6:18:48 PM AC   Page   1 of 1
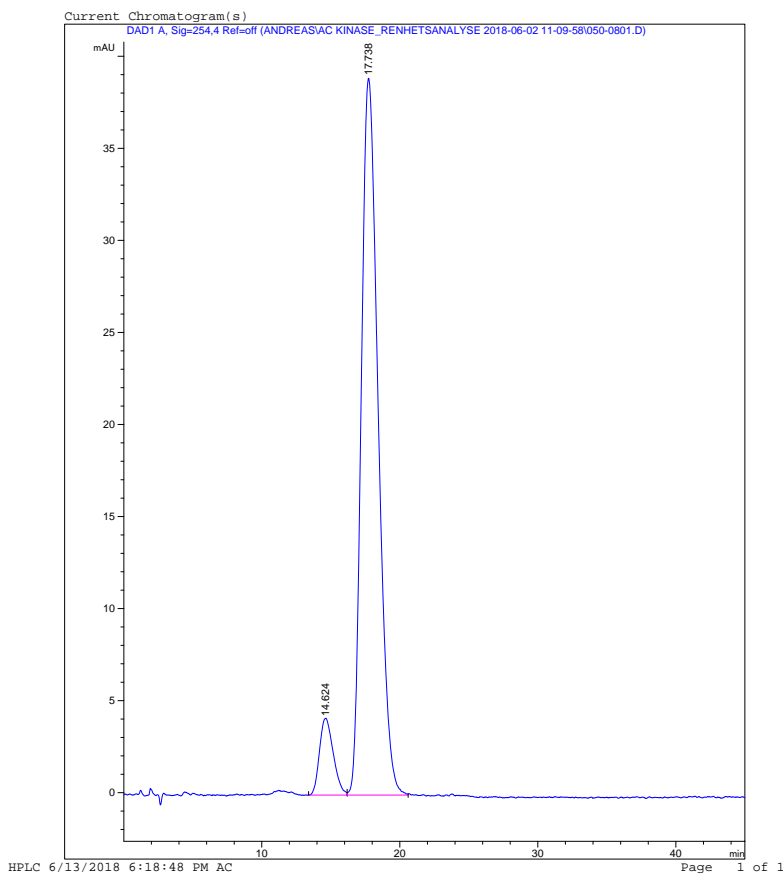
CSP-HPLC (Lux 5u Cellulose-1 4.6 x 250 mm, hexane(cont. 0.2% diethyl amine):*i*-PrOH, flow 1.5 mL/min, detection at 254 nm) of compound *(R)*-**8a**
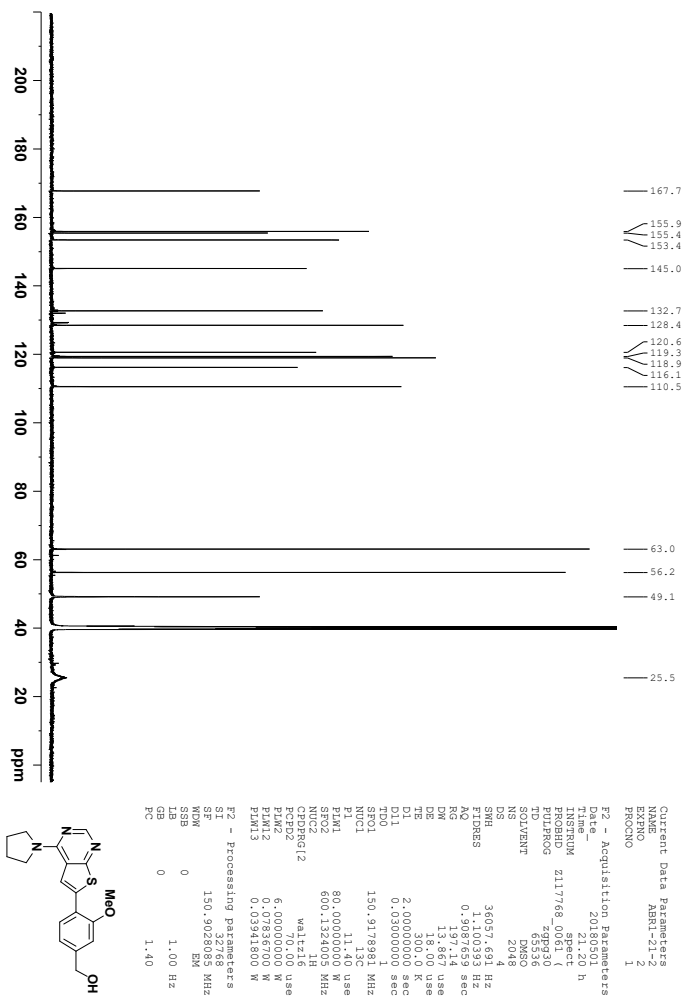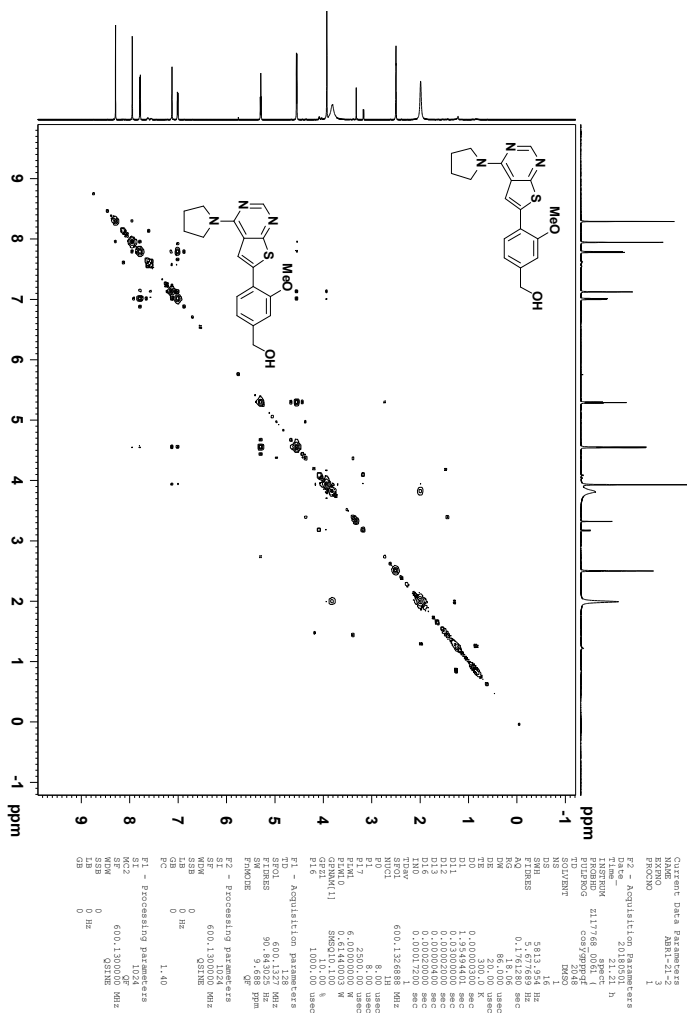
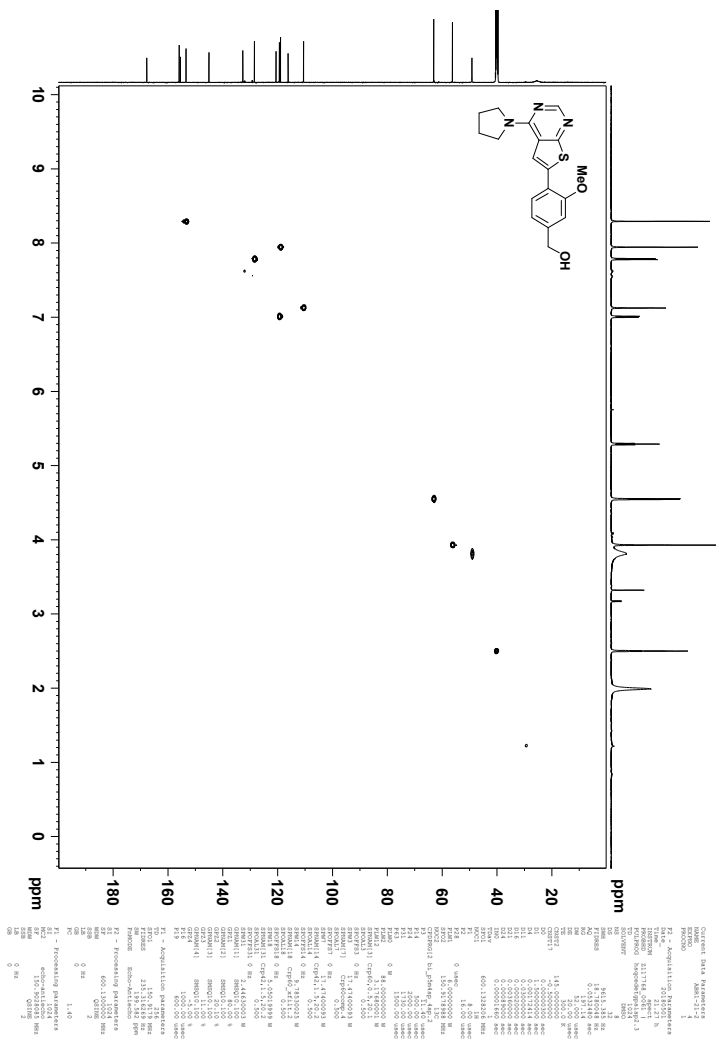# V | Spectroscopic Data - Compound 8b

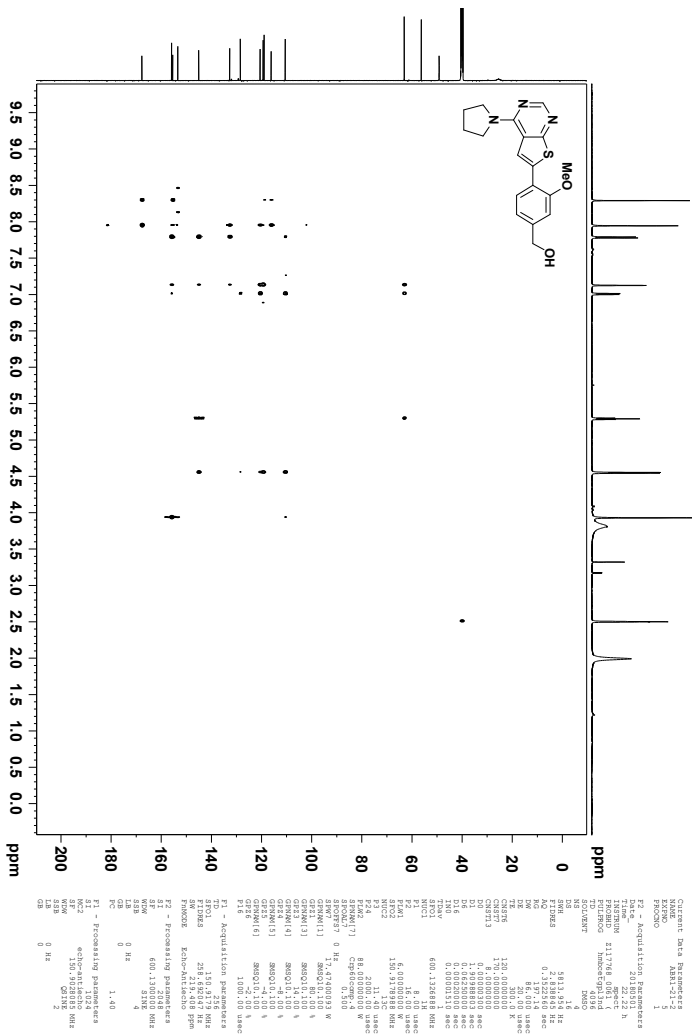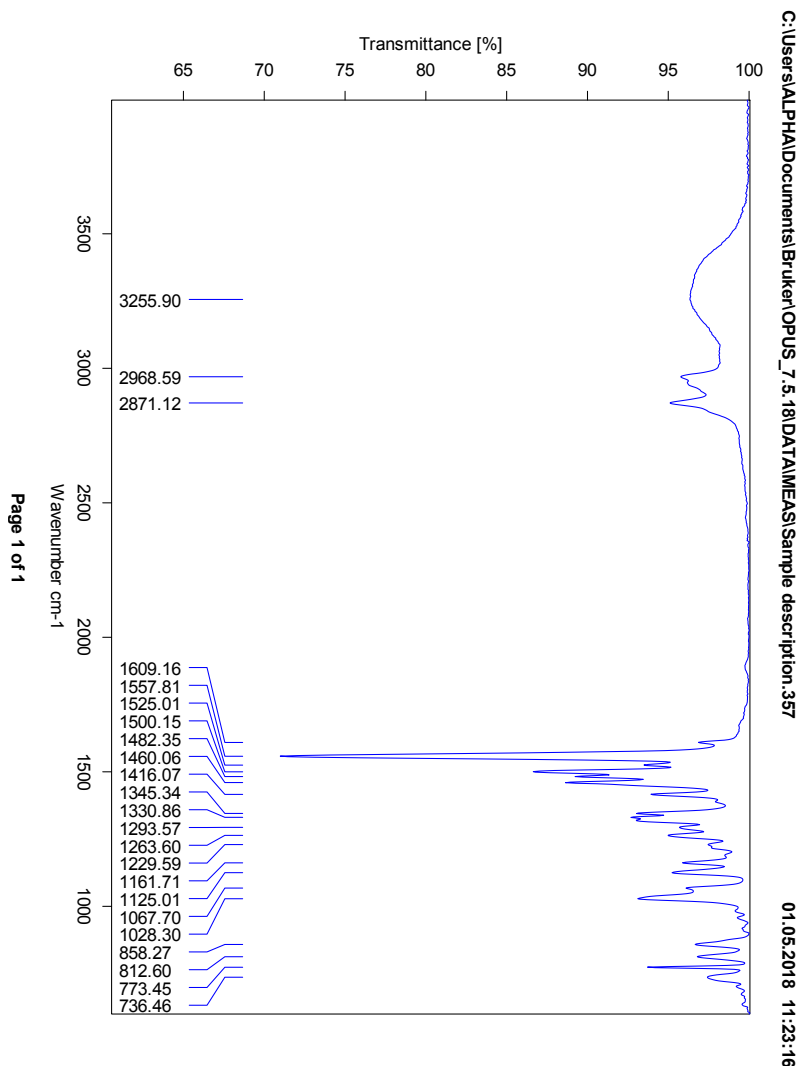^{1}H NMR (600 MHz, DMSO-d_6) spectrum of compound **8b**.

^{13}C NMR (150 MHz, DMSO-d_6) spectrum of compound **8b**.

COSY spectrum of compound **8b**.

HSQC spectrum of compound **8b**.

HMBC spectrum of compound **8b**.

IR spectrum of compound **8b**.

Elemental Composition Report   **Page 1**

Single Mass Analysis
Tolerance = 2.0 PPM   /   DBE: min = -1.5, max = 50.0
Element prediction: Off
Number of isotope peaks used for i-FIT = 3

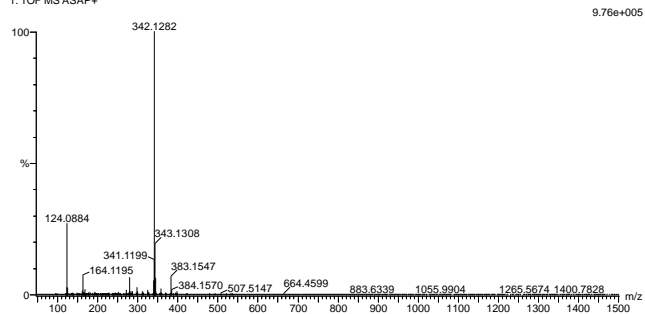Monoisotopic Mass, Even Electron Ions
3002 formula(e) evaluated with 4 results within limits (all results (up to 1000) for each mass)
Elements Used:
C: 0-500   H: 0-1000   N: 0-10   O: 0-20   S: 0-3   Br: 0-2

2018-249 94 (1.843) AM2 (Ar,35000.0,0.00,0.00); Cm (91:94)
1: TOF MS ASAP+
9.76e+005

Minimum:			-1.5				
Maximum:		5.0	2.0	50.0			

Mass	Calc. Mass	mDa	PPM	DBE	i-FIT	Norm	Conf(%)	Formula
342.1282	342.1276	0.6	1.8	10.5	1229.2	0.000	99.99	C18 H20 N3 O2 S
	342.1283	-0.1	-0.3	6.5	1238.1	8.924	0.01	C11 H20 N9 S2
	342.1283	-0.1	-0.3	19.5	1244.1	14.883	0.00	C26 H16 N
	342.1280	0.2	0.6	-0.5	1244.1	14.929	0.00	C13 H29 N O4 Br

Mass spectrum of compound **8b**.