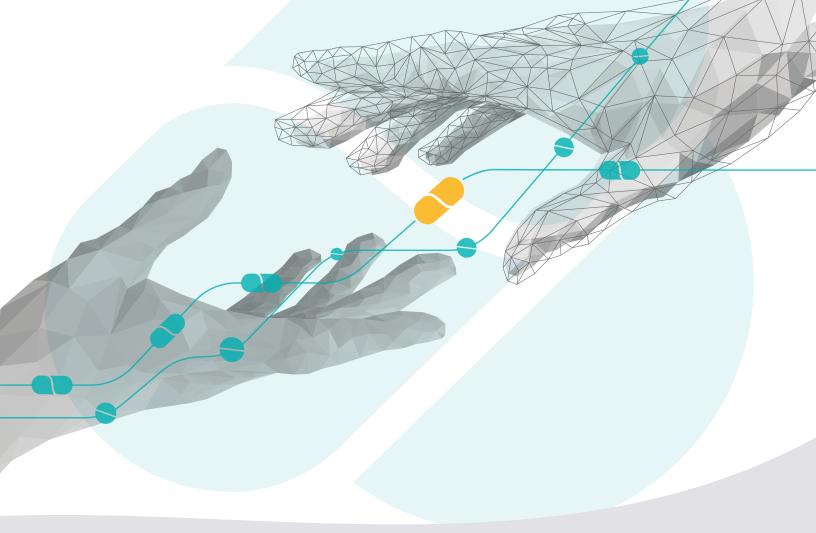


Integrated drug discovery services from Domainex

The research partner that creates innovative solutions







Introduction

Discovering the next generation of innovative drugs to improve the lives of patients worldwide is a truly global effort. At Domainex, we treat differences in geography not as a barrier but as an advantage. We celebrate internationalism and have written a blog about the multi-cultural nature of business including our employees, suppliers and clients. From our prime location in the thriving university city of Cambridge, UK, we are perfectly positioned to support clients in the USA. From communication to time management, we use our UK base to deliver our American clients a suite of advantages to make medicines research effective and efficient.



Shared common language

With no language barrier in UK-USA partnerships, communication is simple and efficient



Highly skilled scientists

Our team is built around experienced scientists from across multiple disciplines, who are capable of tackling the most difficult challenges, will propose innovative solutions, and will contribute valuable intellectual input to your project



Progress around the clock

Our overlapping working days combined with the commitment and flexibility of our team means our scientists will interact directly with you during your working day, and leave you safe in the knowledge that we are working hard on your behalf before your working day begins



Vibrant UK life sciences

By being a part of the UK's world-leading life sciences scene, you can have a high degree of certainty of continuity of service — a valued asset in times of heightened geopolitical uncertainty



Full capabilities

With our fully integrated services and state-of-the-art on-site facilities, outsourcing management is made simple. Our rapid design-make-test cycles enable project milestones to be achieved as quickly as possible.



Clear ownership of IP

Your IP is safe in our hands, guaranteeing you full ownership of, and exclusive rights to, your research results and products.

'Every compound counts'. It's not a slogan, it's the way we work. Whether that compound started its journey in the UK or the USA, our team at Domainex strive to make therapeutic breakthroughs on your behalf. With proprietary technologies and experience across therapeutic areas including cancer, immuno-oncology, inflammation, cardiovascular and respiratory diseases and diseases of the central nervous system (CNS), our integrated approach to medicines research combines the excellence of our scientists with your expertise, to ultimately deliver results.

Our integrated drug discovery approach

Domainex offers a range of drug discovery services, from protein production and assay development through to medicinal chemistry for lead optimization. Your dedicated project leader will listen to your needs and provide tailored and well-considered scientific solutions to support your project on a journey of discovery. Our integrated approach has a proven track record of success in delivering timely and cost-effective solutions, and generating new intellectual property for our clients.



Protein Production

Combinatorial
Domain Hunting

PoLiPa

X-ray
crystallography

PROTEIN CONSTRUCT

LeadBuilder
FragmentBuilder
High Throughput
Screening
Assay Biology



Hit-to-lead

Lead Optimisation



Protein production and assay development





If you need well-characterized, **high-quality protein** for your drug discovery programs or for other purposes, then Domainex's team of highly experienced protein scientists are at your disposal. We can offer you a complete solution, including construct design, cloning, protein expression (*E. coli*, insect cells, or mammalian cells), purification and characterization. We are able to produce and purify proteins in multi-milligram quantities, and our expertise in protein characterization means that you can have confidence that we will deliver you high quality protein that has passed our rigorous quality control process.



Our scientists will often use literature-informed or bioinformatics approaches for expression construct design, but our Combinatorial Domain Hunting (CDH) technology can be used for more challenging proteins. **CDH** is our patented technology for generating many thousands of variants of target proteins to identify suitable **novel constructs**, which can be used in assays or for structural biology.



Polymer Lipid Particle **(PoLiPa)** is our established generic platform for the efficient preparation of stable and highly purified **membrane proteins**, such as GPCRs, for use in drug discovery. Without the need for thermostabilizing mutations or detergents, our technology enables rapid access to pharmacologically intact membrane targets with versatile applications that are stable over several months.



Whether you are looking for high-quality assay biology solutions based on established techniques, or for more innovative assays, we will strive to deliver what you need. We offer a comprehensive suite of **assay development** services, including bespoke biophysical, biochemical and cell-based assays.

Hit identification

A critical factor for efficient prosecution of a successful drug discovery project is the quality of the chemical starting points. Domainex offers rapid and cost-effective compound screening services to match your needs.



LeadBuilder is Domainex's novel approach to **virtual screening**. At its heart is our NICE (Number of Interesting Chemical Entities) virtual database of 4.9 million compounds that has been assembled from known commercial vendor collections and filtered so that only compounds with lead-like properties remain. This collection can be screened in as little as 2 weeks to generate a virtual hit list of 500 – 1000 compounds.



FragmentBuilder is Domainex's **Fragment-Based Drug Design (FBDD)** platform that enables us to identify rapidly hits against your chosen target. Starting from a target gene, Domainex deploys its FBDD expertise in protein science, assay biology and medicinal chemistry to discover tractable, patentable leads cost-effectively.



At Domainex we offer high throughput screening (HTS) using a variety of assay formats (biochemical, cell-based) in 384- or 1536-well format. We can access commercially available diversity libraries of lead-like compounds or focused compounds directed towards target classes, modalities or therapeutic areas, on your behalf. Various library sizes are available, to suit your budget and project needs. Alternatively, if you would prefer to provide your own library, Domainex would be happy to screen it for you.



Our structural biologists can generate high-resolution **X-ray crystal structures** providing invaluable information for your program and enabling a structure guided hit-to-lead phase.

Hit-to-lead and lead optimization

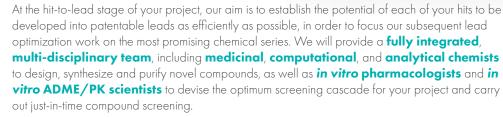












Our *in vitro* pharmacologists are fully enabled to run suites of biochemical, biophysical and cell-based assays in miniaturized (384-well typically) formats to minimize reagent consumption and maximize the flow of project-enabling data. They work closely with our medicinal chemists to minimize cycle times, thus facilitating real-time decision making on where to deploy chemistry resource to further enhance the pharmaceutical properties both within and across several series of compounds in parallel.

In addition, we have built a comprehensive platform of *in vitro* ADME/PK assays that generally utilize highly sensitive mass spectrometry read-outs. Our *in vitro* pharmacology and ADME/PK teams work very closely together to co-ordinate the running of multiple assays in parallel, providing information-rich data sets to our chemists.

Computational chemistry expertise can significantly improve the efficiency of your drug discovery project by helping to design compounds that have a high probability of binding to your target and the right molecular and physical properties to ultimately demonstrate good bioavailability. Our **Computer-Assisted Drug Design (CADD)** experts use state-of- the-art software and proactively develop new insights that are tailored to the particular characteristics of the binding pocket of your disease target.

Building partnerships

With an unrivalled track record of solving research challenges, our highly experienced scientific team delivers successful outcomes efficiently and quickly. We work closely and collaboratively with ambitious life science organizations globally. See what some of them have to say about working with Domainex:



Professor Helen Philippou – Professor of Translational Medicine, Lunac Therapeutics

"Domainex have provided us with integrated drug discovery services including medicinal chemistry, assay biology and ADME. The team were great to work with in a collaborative partnership. The scientists have contributed strategic insight and intellectual input across a range of disciplines. This support was invaluable in areas of expertise on the overall drug discovery process. I've really enjoyed working together with such a dedicated, highly skilled and professional team.

I would recommend Domainex to anyone looking for a provider of drug discovery services."



Professor Daniel Longley – Deputy Director of the Patrick G. Johnston Centre for Cancer Research, Queen's University Belfast

"It has been an absolute pleasure working with Dr Boffey and the Domainex team on this project. Without their first class medicinal chemistry expertise and the use of their Leadbuilder virtual screening platform at the outset of the program, we would not have been able to secure the support of The Wellcome Trust and advance the program to the stage it has now reached."



Marc Gleeson - CEO, Azura Ophthalmics

"Domainex with their deep industry experience, have been able to solve complex synthetic, medicinal, and analytical chemistry challenges to deliver project outcomes for Azura in a timely and efficient manner. They've been able to flexibly resource our program to deliver high quality results and support tight patent timelines."



Dr Jack Kennefick - Co-founder & Chief Executive Officer, Tagomics

"We have been really impressed with the service at Domainex, from initial discussion to delivery of the project. The Domainex team are attentive, well organized, and knowledgeable. They always take the time to understand our exact needs and consistently deliver high-quality results, in a time efficient manner."





Case Study 1

Domainex expertise

- Medicinal chemistry
- Computational chemistry
- ADMF
- Structure-based drug design
- Lead optimization

Disease area: Asthma

Der p 1 is a cysteine protease excreted by house dust mites (HDM) and is a major cause of allergic asthma. In collaboration with St George's University London and the University of Manchester, we set out to identify a candidate drug suitable for delivery by dry powder inhaler (DPI). The starting point was an irreversible peptide-based inhibitor deemed unsuitable for long-term administration in the allergy setting owing to concerns about its potential safety profile.

The Domainex team designed successfully a replacement for the irreversible pharmacophore by employing a reversible, covalent binding group that retained the benefits of a slow off-rate, but without the risk of adverse events. The computational chemistry team used structural information from published crystal structures of Der p 1 and related human cysteine peptidases to design-in exquisite

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Compound	Pl	P2	Р3	P4	P'	Der p 1 IC ₅₀ (nM)	Cat B IC ₅₀ (nM)	LogD _{7.4}
4	n-Bu	Me	benzyl	Ph	cyclohexyl	8	17	
5	i-Pr	Me	benzyl	Ph	cyclohexyl	18	52	
6	i-Pr	Me	benzyl	Ph	benzyl	12	50	
7	t-bu	Me	benzyl	Ph	cyclohexyl	9167	Not determined	
8	i-Pr	Me	t-butyl	Ph	cyclohexyl	14	378	3.9
9	i-Pr	Ме	C(Me) ₂ Ph	Ph	benzyl	42	446	
10	i-Pr	n-Pr	benzyl	Ph	cyclohexyl	164	67	
11	i-Pr	Me	t-butyl	7	benzyl	18	>2500	3.4
12	i-Pr	Me	t-butyl	7	Benzyl	6	274	-0.9
13	i-Pr	Me	t-butyl		cyclohexyl	13	231	2.8
14	i-Pr	Me	t-butyl	Ph	CH2Ph	9	512	3.2
15	i-Pr	Me	benzyl	Ph	×IQ	14	544	1.3
16	i-Pr	Me	t-butyl	Ph	×IQ	14	>2500	1.0
17	i-Pr	Me	benzyl	Ph	×IQ	9	88	1.7
18	i-Pr	Me	benzyl	Ph	× O-	17	>2500	-0.6
19	i-Pr	Me	benzyl		×IQ	20	540	2.0

Table 1: The impact of modifying P1, P2 and P3 on selectivity over Cathepsin B

Reference

Newton G et al. (2014), The discovery of potent, selective, and reversible inhibitors of the house dust mite peptidase allergen Der p 1: an innovative approach to the treatment of allergic asthma. J. Med. Chem., **57** (22), 9447-9462

selectivity, and improved stability to proteases in the lung. Physicochemical properties were fine-tuned to optimize lung retention, and this was confirmed by the long duration of action shown in allergy models where rodents were exposed to house dust mite pellets. Metabolic, plasma protein binding and oral absorption properties were also optimized to ensure low levels of systemic exposure, and hence reduced risks of adverse effects. The Domainex team demonstrated that compounds were compatible for use with dry powder inhalers by identifying compounds with stable crystalline forms that could be micronized to give particles of a size appropriate for inhaled delivery.

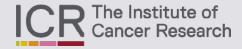
What was the successful outcome?

A candidate drug and a number of credible back-up compounds were identified from the primary series.

Work on the follow-up program led to a differentiated series with a non-covalent binding mode which demonstrated in vivo efficacy.







Case Study 2

Domainex expertise

- Virtual screening via LeadBuilder
- Hit identification
- Structure-based drug design
- ADME
- Medicinal chemistry
- Lead optimization

Disease area: Oncology

Tankyrase is a member of the PARP family which has been shown to play an important role in the Wnt signalling pathway. LeadBuilder was used to identify hit compounds that acted as tankyrase inhibitors. Previously published crystal structures showed tankyrase in a closed form, in which the active site was inaccessible to ligands. Domainex built a homology model of tankyrase using the closed conformation and a published crystal structure of PARP1 in an open conformation. This model was used to screen Domainex's NICE database of ~1.5 million commercially available compounds, from which ~1000 compounds were purchased. 59 hits were identified with IC $_{50}$ values between 100 nM and 10 μ M.

Supported by X-ray crystallography, a subsequent structure-based drug design program (incorporating integrated ADME to inform each iteration of medicinal chemistry, and early screening for other liabilities) generated several series of potent tankyrase inhibitors (< 20 nM tankyrase, < 100 nM in a Wnt reporter cell assay), with excellent selectivity over PARP1 (> 30 fold), and good ADME properties (e.g. oral bioavailability in rodents of > 50%). Lead compounds were shown to inhibit the growth of APC-null tumor xenografts.

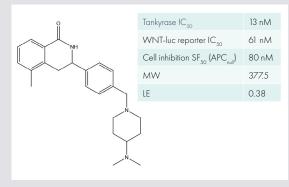


Figure 1: Example of an isoquinolone tankyrase inhibitor designed and synthesized at Domainex

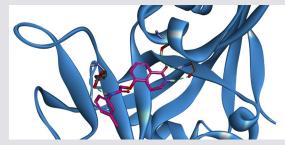


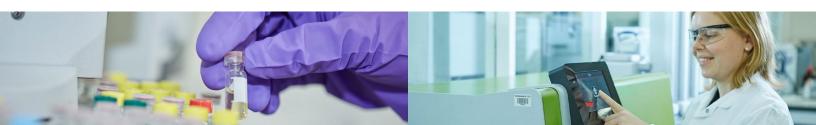
Figure 2: X-ray crystal structure of one of our compounds bound to tankyrase

Reference

Elliott R, Ashley J *et al.* (2015), Design and discovery of 3-aryl-5-substituted-isoquinolin-1-ones as potent tankyrase inhibitors. *Med. Chem. Commun.*, **6**, 1687-1692

What was the successful outcome?

The project team went from hit to candidate drug in less than 400 compounds. The project received further funding to generate a back-up candidate, and was subsequently out-licensed to a major pharmaceutical company to develop these compounds as anti-cancer drugs.



Imperial College London

Case Study 3

Domainex expertise

- Computational chemistry
- Medicinal chemistry
- Assay biology
- Lead optimization
- ADME
- X-ray crystallography

Disease area: Heart disease

Mitogen-Activated Protein Kinase Kinase Kinase Kinase 4 (MAP4K4), a serine-threonine kinase which activates the JNK signaling pathway, is activated in failing human hearts and relevant rodent models. Therefore, Professor Michael Schneider and his team at Imperial College London postulated that an inhibitor of MAP4K4 would be able to suppress human cardiac cell death, offering the possibility of cardioprotection following heart attacks.

An empirical screen was carried out against human MAP4K4 using about 1800 bioactive compounds, and the initial hit, DMX-4640 (Figure 1), was identified as a starting point for medicinal chemistry studies. We then undertook a program of rational drug design, including X-ray crystallography, to produce DMX-5804, a compound with a significantly greater water solubility and reduced metabolic clearance.

We demonstrated that MAP4K4 inhibition by DMX-5804 confers protection in hiPSC-CMs (Figure 2) and reduces ischemia-reperfusion injury in mice by >50% (Figure 3). The solubility and pharmacokinetic properties of DMX-5804 were insufficient for a human drug candidate in acute ischemic injury, where rapid intravenous infusion is desired. However, further structural modifications led to compounds with properties suitable for clinical candidates (Figure 1).

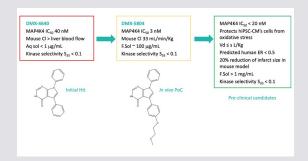


Figure 1: The medicinal chemistry progression from initial hit, DMX-4640, to the identification of pre-clinical candidates

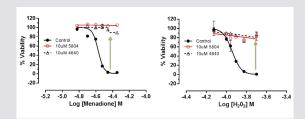


Figure 2: DMX-5804 confers 100% protection against two oxidative stress signals in cardiomyocyte stem cells

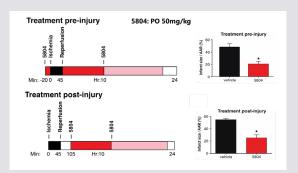


Figure 3: Initial proof of concept studies carried out in mice using 50mg/kg oral dose of DMX-5804

Reference

Schneider MD et al. (2019). MAP4K4 inhibition promotes survival of human stem cell-derived cardiomyocytes and reduces infarct size in vivo. Cell Stem Cell, 24 (4), 579-591

What was the successful outcome?

Using hiPSC-CMs as the most relevant platform for gene silencing and drug discovery, we designed small-molecule inhibitors of MAP4K4. DMX-58O4, a novel, potent, highly selective, small-molecule inhibitor of MAP4K4, was identified for proof of concept studies.

Additionally, we were able to further optimize the series and identified a number of compounds with properties suitable for a clinical candidate.



About Domainex

Domainex is a fully integrated drug discovery service company based at Cambridge, UK. We serve pharmaceutical, biotechnology, academic organizations and patient foundations globally. We have ambitious growth plans and are expecting to reach 110 biologists and chemists in the near future. We provide integrated services, from disease target selection to candidate drug nomination. We have a very strong reputation for contributing innovative ideas, undertaking high-quality experiments and for generating intellectual property on behalf of our clients. We strive to build strong, dynamic relationships. In 2021 we served over 60 clients from the United States, the UK, Europe, Japan and Australia and had a project renewal rate of over 80%.

How Can Domainex Help Your Drug Discovery Project?

Our highly experienced, multi-disciplined scientists – molecular biologists, protein biochemists, assay biologists, structural biologists, medicinal, computational, and bio/analytical chemists, in vitro pharmacologists and ADME scientists – will support you to advance your drug discovery projects towards drug development effectively and efficiently. We provide customized programs to address your specific needs at each stage of drug discovery. We draw from a wealth of expertise built up over the last 20 years across a wide range of drug targets and therapeutic areas. From our sites within Europe's leading bioscience hub at Cambridge, UK and with access to the very latest cutting-edge technologies, we are able to help you realize your goals and enrich your discovery pipeline.

Contacts

If you would like to know more about Domainex's discovery services, or speak to us regarding your own drug discovery needs, please contact us at: USA-enquries@domainex.co.uk

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