



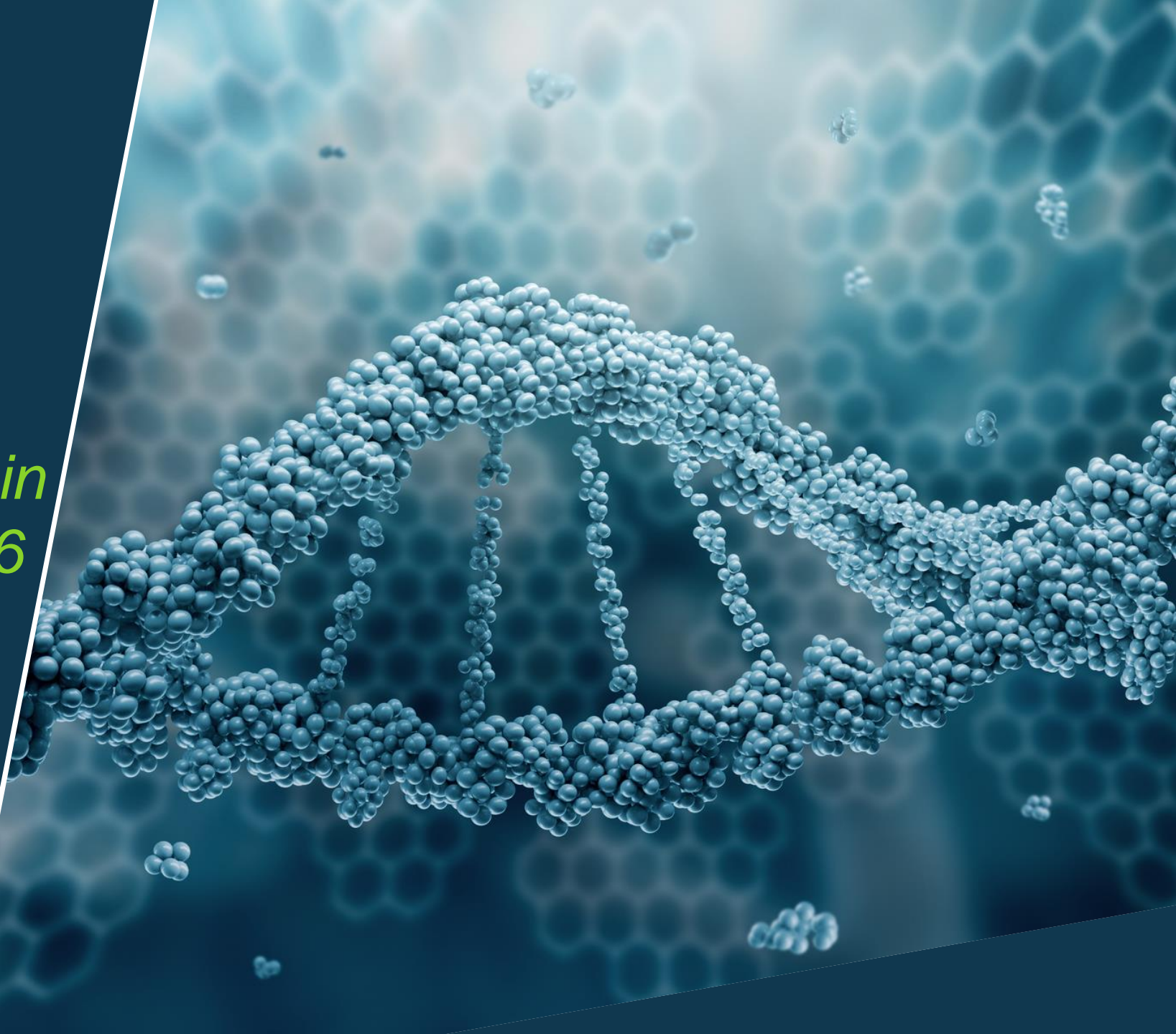
Donated Chemical Probe

*Adrenergic α_{2B}
Antagonist – in vitro & in
vivo iv Probe BAY-6096*



December 6th, 2023

**Presenter:
Daniel Meibom
On behalf of the team**

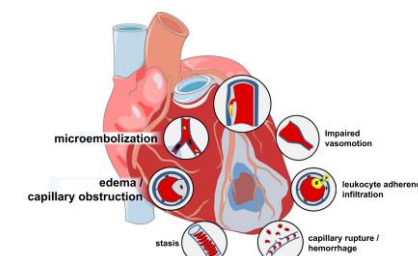




Adrenergic α_{2B} Antagonist iv Probe BAY-6096

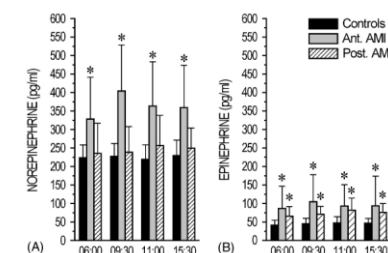
Rationale for inhibition of the adrenergic α_{2B} receptor (GPCR) in heart diseases

- Myocardial infarction (MI) is typically treated by reopening of the occluded coronary artery with a catheter placed at the site of stenosis
- Paradoxically, reperfusion not only terminates myocardial ischemia but inflicts injury by itself, the so-called reperfusion damage



Basic Res. Cardiol. 2019, 45

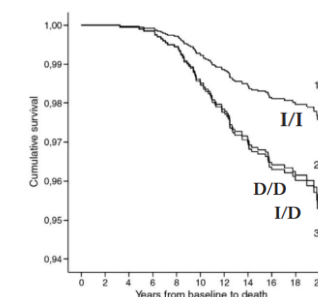
- Adrenaline and noradrenaline are endogeneous ligands of the α_{2B} receptor, which is expressed in small coronary arteries
- Catecholamine concentrations are elevated during MI which should lead to enhanced vasoconstriction and increased infarct sizes
- A genetic variant of the α_{2B} receptor with prolonged receptor activation is associated with an increased risk for MI and sudden cardiac death



Clin. Cardiol. 2007, 326

	D/D	I/I
All subjects (n = 912)		
Men at risk	192	256
Acute coronary events*	15 (7.8)	10 (3.9)
FINMONICA classification†	10; 5; 0	5; 1; 4
Ischemic findings in exercise test*	57 (30)	52 (20)

J. Am. Coll. Cardiol. 2001, 1516



Am. Heart J. 2009, 615

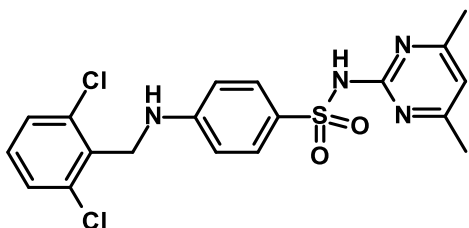
An α_{2B} antagonist for iv administration during and after reperfusion might lead to reduced infarct sizes and improved outcomes



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

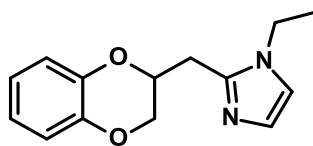
Reference compounds

Example C in WO2003/008387A1, Preclinical, Juvantia Pharma



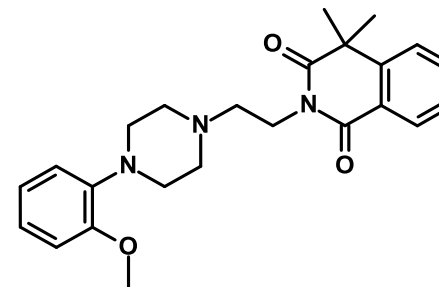
α_{2B} -IC ₅₀	24 nM
α_{1A}	>20000x
α_{2A}	36x
α_{2C}	>249x
Sol _{6.5}	6 mg/l

Imiloxan, Preclinical, Syntex Laboratories



α_{2B} -IC ₅₀	655 nM
α_{1A}	>15x
α_{2A}	4x
α_{2C}	2x
Sol _{6.5}	483 mg/l

ARC-239, Preclinical, Thomae GmbH



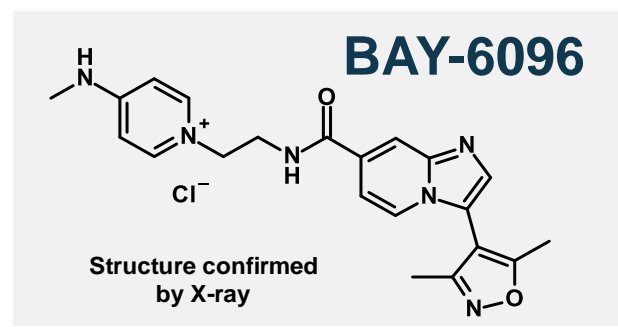
α_{2B} -IC ₅₀	99 nM
α_{1A}	0.1x
α_{2A}	74x
α_{2C}	2x
Sol _{6.5}	80 mg/l

- Published reference compounds are less selective and lack a broader selectivity data set (inside & outside of target family)
- There is currently no adrenergic α_{2B} antagonist available which is soluble enough to investigate α_{2B} mediated pharmacology after iv administration



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Technical in vitro profile



Potency vs α_{2B} [nM]	
Cell-based hIC ₅₀	14
Cell-based hIC ₅₀ (mutant*)	25
Binding hK _i	21
Cell-based rIC ₅₀	13
Cell-based dIC ₅₀	25

Physchem	
LogD @ pH 7.5	1.2
BEI / LLE (based on hIC ₅₀)	20 / 6.7
Sol @ pH 7.4 [g/L], cryst. material	>90
MW / MW _{cation} / TPSA [g/mol / Å ²]	427 / 392 / 88
Stability (pH 7.4, 13 weeks, RT)	yes

in vitro DMPK Properties						
Caco2 permeability	P _{app} (A-B) [nm/s]		P _{app} (B-A) [nm/s]		efflux ratio	
	3		9		3	
Metabolic stability			CL [L/h/kg]		F _{max} [%]	
	rat hepatocytes		10 ⁻⁴		100%	
	human hepatocytes		10 ⁻⁴		100%	
CYP inhibition IC ₅₀ [μM]	1A2	2C8	2C9	2D6	3A4	3A4 preinc.
	>20	>20	>30	>20	>20	>20 (no TDI)
CYP1A2 & 3A4 induction [μM]	> 211					

Selectivity	
In-house kinase panel (22 kinase assays)	hDDR2: 1.4 μM Rest >20 μM
Panlabs @ 10 μM	See next slide

Safety	
Ames	negative
hERG, hNa _v 1.5, hCa _v 1.2, hK _{ir} 2.1 IC ₅₀ [μM]	>10

h = human, r = rat, d = dog

* J. Am. Coll. Cardiol. 2001, 1516; Am. Heart J. 2009, 615

- BAY-6096 is a potent and selective α_{2B} antagonist
- BAY-6096 shows very high aqueous solubility at pH 7.4 from crystalline material



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Selectivity profile in more detail (off-targets & target family)

Assay Name	Conc.	% Inh.	Assay Name	Conc.	% Inh.	Assay Name	Conc.	% Inh.
Aldehyde Reductase	10 μ M	-2	Androgen (Testosterone)	10 μ M	-3	Motilin	10 μ M	-10
ATPase, Na ⁺ /K ⁺ , Heart, Pig	10 μ M	-2	Angiotensin AT ₁	10 μ M	5	Muscarinic M ₁	10 μ M	-4
Carbonic Anhydrase II	10 μ M	3	Angiotensin AT ₂	10 μ M	4	Muscarinic M ₂	10 μ M	6
Cholinesterase, Acetyl, ACES	10 μ M	15	Bradykinin B ₁	10 μ M	-1	Muscarinic M ₃	10 μ M	0
Cyclooxygenase COX-1	10 μ M	-6	Bradykinin B ₂	10 μ M	-2	Muscarinic M ₄	10 μ M	4
Cyclooxygenase COX-2	10 μ M	-9	Cannabinoid CB ₁	10 μ M	-12	Nicotinic Acetylcholine	10 μ M	9
HMG-CoA Reductase	10 μ M	-15	Cannabinoid CB ₂	10 μ M	-4	Opiate δ_1 (OP1, DOP)	10 μ M	-5
Leukotriene LTC ₄ Synthase	10 μ M	6	Dopamine D ₁	10 μ M	-7	Opiate κ (OP2, KOP)	10 μ M	1
Lipoxygenase 15-LO	10 μ M	0	Dopamine D _{2L}	10 μ M	-2	Opiate μ (OP3, MOP)	10 μ M	0
Monoamine Oxidase MAO-A	10 μ M	-14	Dopamine D _{2S}	10 μ M	-5	Progesterone PR-B	10 μ M	12
Monoamine Oxidase MAO-B	10 μ M	6	Dopamine D ₃	10 μ M	-5	Purinergic P2X	10 μ M	-7
Nitric Oxide Synthase, Neuronal (nNOS)	10 μ M	-15	Endothelin ET _A	10 μ M	-2	Purinergic P2Y	10 μ M	11
Nitric Oxide Synthetase, Inducible (iNOS)	10 μ M	6	Endothelin ET _B	10 μ M	-2	Serotonin (5-Hydroxytryptamine) 5-HT _{1A}	10 μ M	25
Peptidase, Angiotensin Converting Enzyme	10 μ M	1	Estrogen ER α	10 μ M	6	Serotonin (5-Hydroxytryptamine) 5-HT _{2A}	10 μ M	1
Phosphodiesterase PDE3	10 μ M	3	GABA _A , Chloride Channel, TBOB	10 μ M	-4	Serotonin (5-Hydroxytryptamine) 5-HT _{2B}	10 μ M	0
Phosphodiesterase PDE4D2	10 μ M	-1	GABA _A , Flunitrazepam, Central	10 μ M	-2	Serotonin (5-Hydroxytryptamine) 5-HT _{2C}	10 μ M	0
Phosphodiesterase PDE5	10 μ M	4	GABA _B , Non-Selective	10 μ M	2	Transporter, Adenosine	10 μ M	-2
Thromboxane Synthase	10 μ M	-8	Glucocorticoid	10 μ M	1	Transporter, Dopamine (DAT)	10 μ M	20
Adenosine A ₁	10 μ M	1	Glutamate, AMPA	10 μ M	-1	Transporter, GABA	10 μ M	0
Adenosine A _{2A}	10 μ M	-8	Glutamate, Kainate	10 μ M	0	Transporter, Norepinephrine (NET)	10 μ M	3
Adenosine A ₃	10 μ M	1	Glutamate, NMDA, Agonism	10 μ M	-16	Transporter, Serotonin (5-Hydroxytryptamine) (SERT)	10 μ M	0
			Glutamate, NMDA, Glycine	10 μ M	4	Vasopressin V _{1A}	10 μ M	-2
			Growth Hormone Secretagogue (GHS, Ghrelin)	10 μ M	2			
			Histamine H ₁	10 μ M	1			
			Histamine H ₂	10 μ M	0			
			Histamine H ₃	10 μ M	2			
			Insulin	10 μ M	-3			

Adrenergic receptor	IC ₅₀ [nM]	Selectivity factor vs h α_{2B} IC ₅₀
h α_{2B}	14	1x
h α_{1A}	5516	394x
r α_{1B}	>10000*	>714x
h α_{1D}	>10000*	>714x
h α_{2A}	10000	725x
h α_{2C}	>11830	>845x
h β_1	>10000*	>714x
h β_2	>10000*	>714x
h β_3	>10000*	>714x

h = human, r = rat

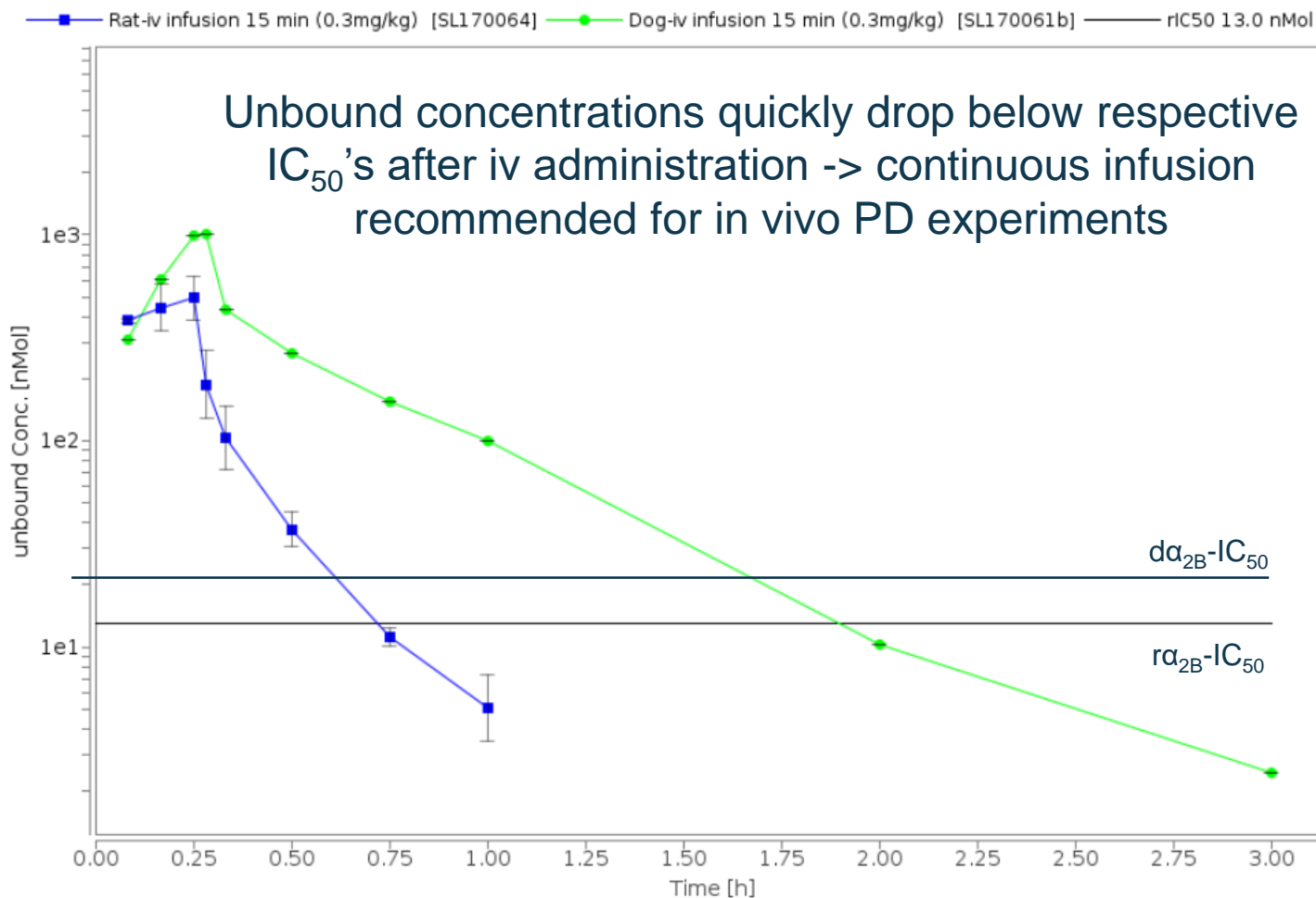
* no significant effect at 10 μ M, an IC₅₀ >10 μ M is assumed

**BAY-6096 shows good selectivity vs 70 off-targets
Meets criteria >30x in target family**



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

In vivo PK data



Rat PK data:

CL_p 4.3 l/h/kg
 V_{ss} 0.5 l/kg
 MRT_{iv} 0.1 h
 f_u 79%

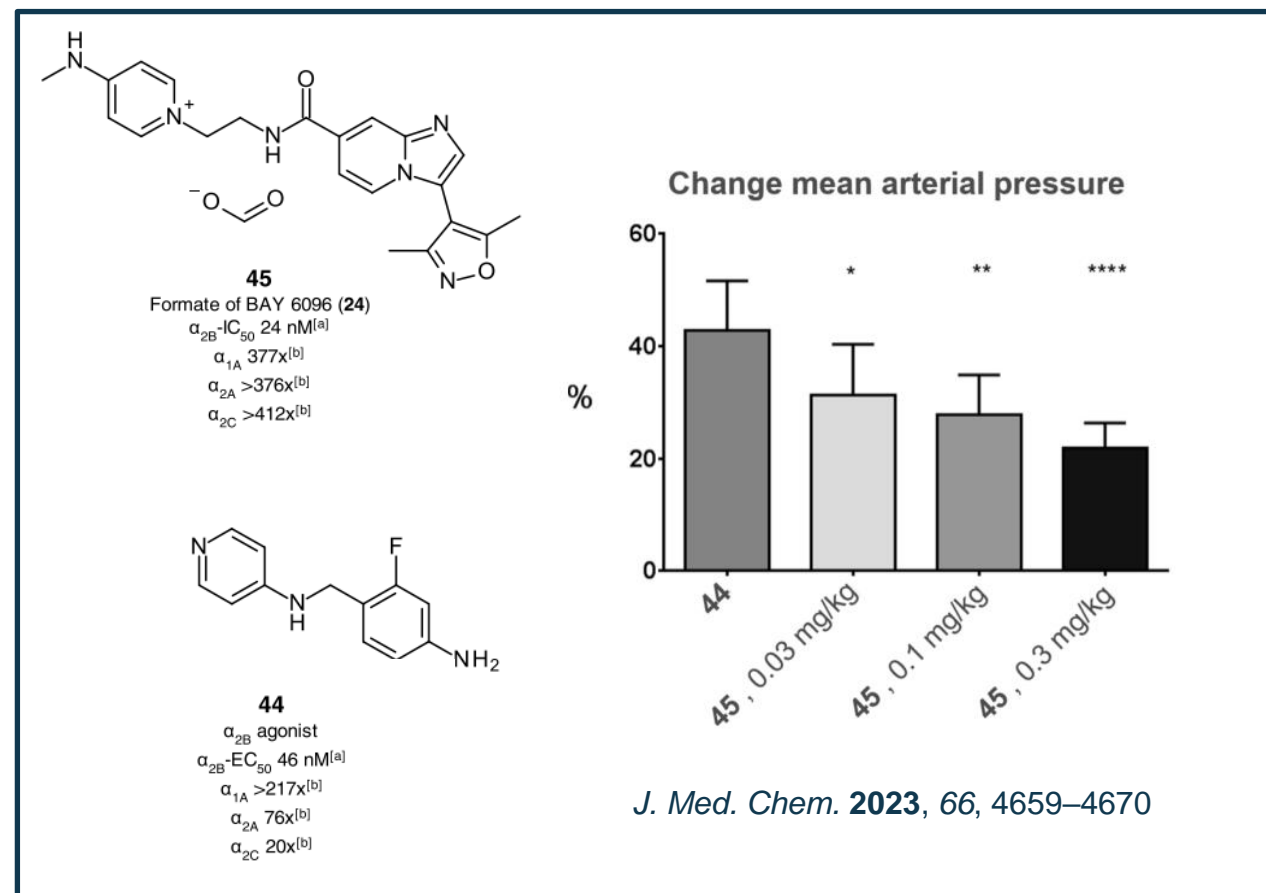
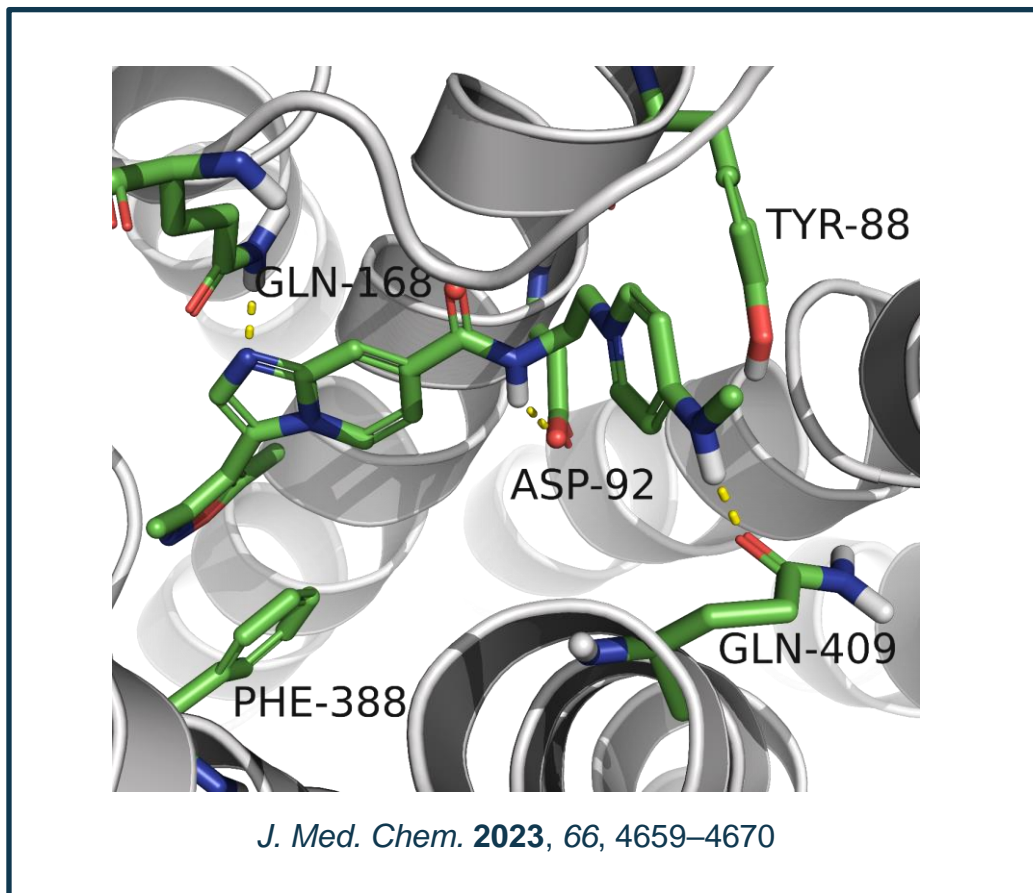
Dog PK data:

CL_p 1.6 l/h/kg
 V_{ss} 0.6 l/kg
 MRT_{iv} 0.4 h
 f_u 82%

BAY-6096 shows high clearance in rats and dogs (continuous infusion recommended)

Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Hints for target engagement

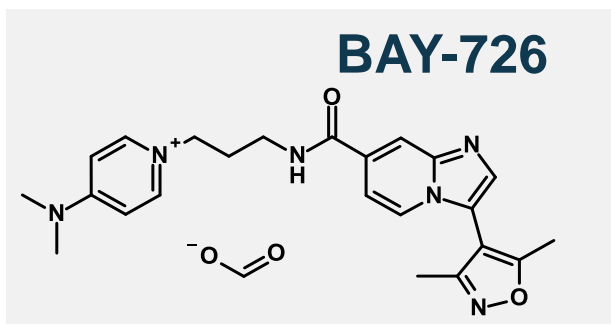


- Docking of BAY-6096 suggests key interactions with receptor supported by SAR (see publication for details)
- The formate of BAY-6096 (**45**) dose dependently decreases the blood pressure increase induced by α_{2B} agonist **44**, thereby suggesting a role for α_{2B} receptors in vascular constriction in rats (see publication for details)



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

In vitro profile of negative control BAY-726



Potency vs α_{2B} [nM]	
Cell-based hIC ₅₀	7100
Cell-based hIC ₅₀ (mutant)	n.d.
Binding hK _i	n.d.
Cell-based rIC ₅₀	n.d.
Cell-based dIC ₅₀	10000

Physchem	
LogD @ pH 7.5	1.3
BEI / LLE (based on hIC ₅₀)	12 / 4.0
Sol@pH 7.4 [mg/L], cryst. material	>500
MW / MW _{cation} / TPSA [g/mol / Å ²]	465 / 420 / 80
Stability (pH 1 / 7 / 10, 24 h, 37 °C)	yes

in vitro DMPK Properties

Caco2 permeability	P _{app} (A-B) [nm/s]		P _{app} (B-A) [nm/s]		efflux ratio	
		2		4		2
Metabolic stability			CL [L/h/kg]		F _{max} [%]	
	rat hepatocytes		10 ⁻⁴		100%	
	human hepatocytes		0.3		75	
CYP inhibition IC ₅₀ [μM]	1A2	2C8	2C9	2D6	3A4	3A4 preinc.
	>20	>20	>20	>20	>20	>20
CYP1A2 & 3A4 induction [μM]	> 65					

Selectivity

In-house kinase panel (22 kinase assays)	hDDR2: >20 μM Rest >20 μM
IC ₅₀ vs h α_{2A} , h α_{2C} , h α_{1A} [nM]	>10000

Safety

Ames	n.d.
hERG, hNa _v 1.5, hCa _v 1.2, hK _{ir} 2.1 IC ₅₀ [μM]	>10

n.d. = not determined
h = human, r = rat, d = dog

Negative control BAY-726 is >500x less active on the human α_{2B} receptor than the probe BAY-6096
Negative control BAY-726 shows no activity on the human α_{2A} , α_{2C} and α_{1A} receptor



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Summary / Conclusion

Probe criteria	
Inhibitor potency: goal is < 100 nM (IC ₅₀)	Meets criteria Cell-based h α_{2B} -IC ₅₀ : 14 nM (binding hK _i : 21 nM)
Selectivity within target family: goal is > 30-fold	Meets criteria All selectivity factors within adrenergic receptor family at least >393x
Selectivity outside target family: describe the off-targets (which may include both binding and functional data)	Meets criteria Clean in a panel of 70 off-targets at 10 μ M and 21 kinases at 20 μ M* Hits from inhouse 3NN target prediction# outside target family all devalidated
On target cell activity for cell-based targets: goal is < 1 μ M IC ₅₀ /EC ₅₀	Meets criteria The α_{2B} receptor is located on the cellular surface Cell-based h α_{2B} -IC ₅₀ : 14 nM; Recommended concentration for use: 100 nM
On target cell activity for secreted targets: appropriate alternative such as mouse model or other mechanistic biological assay, e.g., explant culture	Meets criteria Docking of BAY-6096 suggests key interactions with receptor supported by SAR Decrease of blood pressure increase induced by an α_{2B} agonist in vivo
Neg ctrl: <i>in vitro</i> potency – > 100 times less; Cell activity – >100 times less potent than the probe	Meets criteria Negative control >500 times less active than probe (cell-based h α_{2B} assay)

We ask for acceptance of α_{2B} antagonist BAY-6096 as chemical probe, accompanied by BAY-726 as negative control

nearest neighbour search in internal and external bioactivity databases

* see slide 4 & 5 for details

h = human



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Project team / Acknowledgement

Journal of
**Medicinal
Chemistry**



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Article

BAY-6096: A Potent, Selective, and Highly Water-Soluble Adrenergic α_{2B} Antagonist

Daniel Meibom,* Jutta Meyer, Clemens-Jeremias von Buehler, Karl D. Collins, Stefanie Maassen, Kersten Matthias Gericke, Jörg Hüser, Joachim Mittendorf, Nuria Ortega Hernandez, Jens Schamberger, Jan Stampfuss, Alexander Straub, Afra Torge, Norbert Witowski, and Frank Wunder



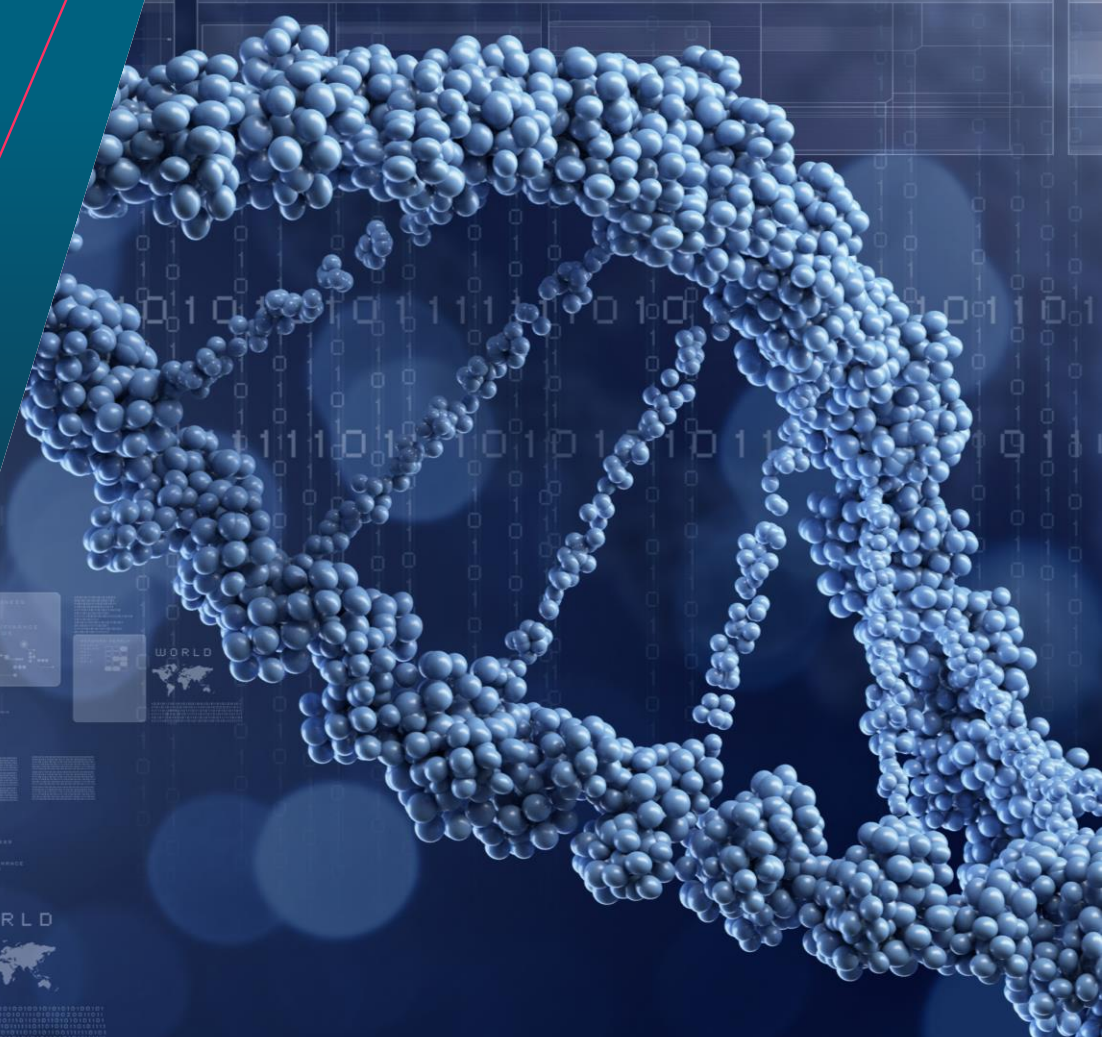
Cite This: *J. Med. Chem.* 2023, 66, 4659–4670



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Thank You





Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Assays in more detail

Characterization on adrenoceptor reporter cells: Adrenoceptor α_{1A} antagonism was tested on a recombinant human α_{1A} receptor CHO cell line, also expressing recombinant mtAeq (mitochondrial aequorin). Adrenoceptor α_{2A} antagonism was tested on a recombinant human α_{2A} -G α_{16} receptor fusion protein CHO cell line (PerkinElmer Life Sciences), also expressing recombinant mtAeq. Adrenoceptor α_{2B} antagonism was tested on a recombinant human α_{2B} receptor CHO cell line (PerkinElmer Life Sciences), also expressing recombinant mtAeq. Adrenoceptor α_{2C} antagonism was tested on a recombinant human α_{2C} receptor HEK cell line, also expressing a chimeric G protein (G α_{q3}) and Clytin. Adrenoceptor α_{2B} antagonism was also tested in a CHO cell line expressing the human α_{2B} receptor deletion variant (del Glu301-Glu303) and recombinant mtAeq.

Cells were cultured at 37°C and 5% CO₂ in Dulbecco's modified Eagle's medium/NUT mix F12 with L-glutamine, supplemented with 10% (v/v) inactivated fetal calf serum, 1 mM sodium pyruvate, 0.9 mM sodium bicarbonate, 50 U/ml penicillin, 50 µg/ml streptomycin, 2.5 µg/ml amphotericin B and 1 mg/ml geneticin. Cells were passaged using enzyme-free/Hank's-based cell dissociation buffer. All cell culture reagents were obtained from Invitrogen (Carlsbad, USA).

Luminescence measurements were performed on opaque 384-well microtiter plates. 2000 cells/well were plated in a volume of 25 µl and were cultured for 1 day at 30°C and 5% CO₂ in cell culture medium containing coelenterazine (α_{2A} and α_{2B} : 5 µg/ml; α_{1A} and α_{2C} : 2.5 µg/ml). Serial dilutions of the test compounds (10 µl) in Tyrode (130 mM NaCl, 5 mM KCl, 20 mM HEPES, 2 mM CaCl₂, 1 mM MgCl₂, 4.8 mM NaHCO₃ at pH 7.4) were applied to the cells. After 5 minutes norepinephrine was added to the cells (35 µl, final concentration: EC₅₀ - EC₈₀) and the emitted light was measured for 50 seconds using a charge-coupled device (CCD) camera (Hamamatsu Corporation, Shizuoka, Japan) in a light tight box. Curve fitting and calculation of IC₅₀/EC₅₀ values was performed using GraphPad Prism Software (version 8.0, GraphPad Software Inc., San Diego, CA, USA).

Reversibility of receptor binding was tested in washout experiments. After antagonist treatment of the recombinant α_{2B} receptor reporter cells for 5 min, the supernatant was removed. Cells were washed twice with Tyrode (35 µl). 5 min later cells were stimulated with norepinephrine.

Adrenergic receptor agonists were characterized using the reporter cell lines described above. Serial dilutions of the test compounds (10 µl) in Tyrode were applied to the cells and measurements were performed using the FLIPR® Tetra system (Molecular Devices, Sunnyvale, CA, USA).

The α_{2B} receptor agonist **44** was identified by uHTS. **44** stimulates reporter cell lines with EC₅₀ values of 3,500 nM (α_{2A}), 46 nM (α_{2B}), 900 nM (α_{2C}) and >10,000 nM (α_{1A}), respectively.

alpha2B Human Adrenoceptor GPCR Binding Antagonist Radioligand LeadHunter Assay - TW

Item: 203710

Assay Information

Assay Type: Biochemical

Assay Sub Type: Binding

Detection Method: Radiometric

Measured Response: Scintillation

Testing Information

Procedure Summary:

This assay measures binding of [³H]Rauwolscine to human adrenergic α_{2B} receptor CHO-K1 cells stably transfected with a plasmid encoding the human adrenergic α_{2B} receptor are used in modified Tris-HCl pH 7.4 buffer using standard techniques. A 10 µg± aliquot of membrane is incubated with 2.5 nM [³H]Rauwolscine for 60 minutes at 25°C. Non-specific binding is estimated in the presence of 10 µM prazosin. Membranes are filtered and washed 3 times and filters are counted to determine [³H]Rauwolscine specifically bound. Compounds are screened at 10 µM.
Note: ‡Membrane protein may change from lot to lot, the concentration used will be adjusted if necessary.

Ligand: [³H] Rauwolscine

Ligand Kd (nM): 2.1

Ligand Concentration: 2.5 nM

Non Specific: 10 µM Prazosin

Incubation: 60 min at 25°C

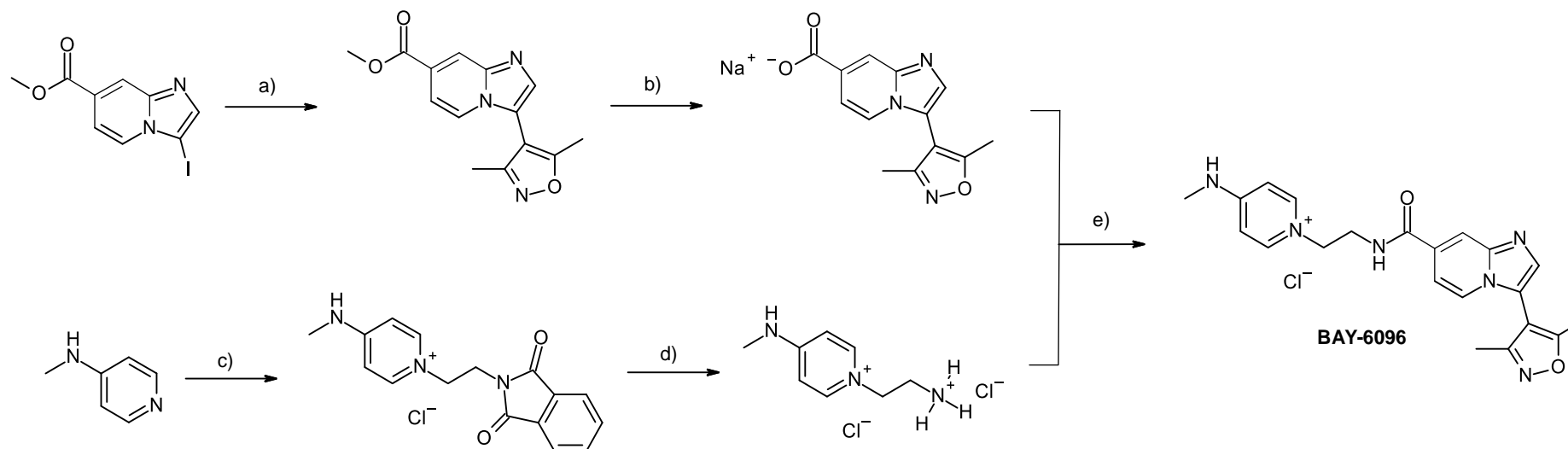
Control Inhibitor: Yohimbine

<https://www.eurofindiscovery.com/catalog/alpha2b-human-adrenoceptor-gpcr-binding-antagonist-radioligand-leadhunter-assay-tw/203710>



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Synthesis of BAY-6096



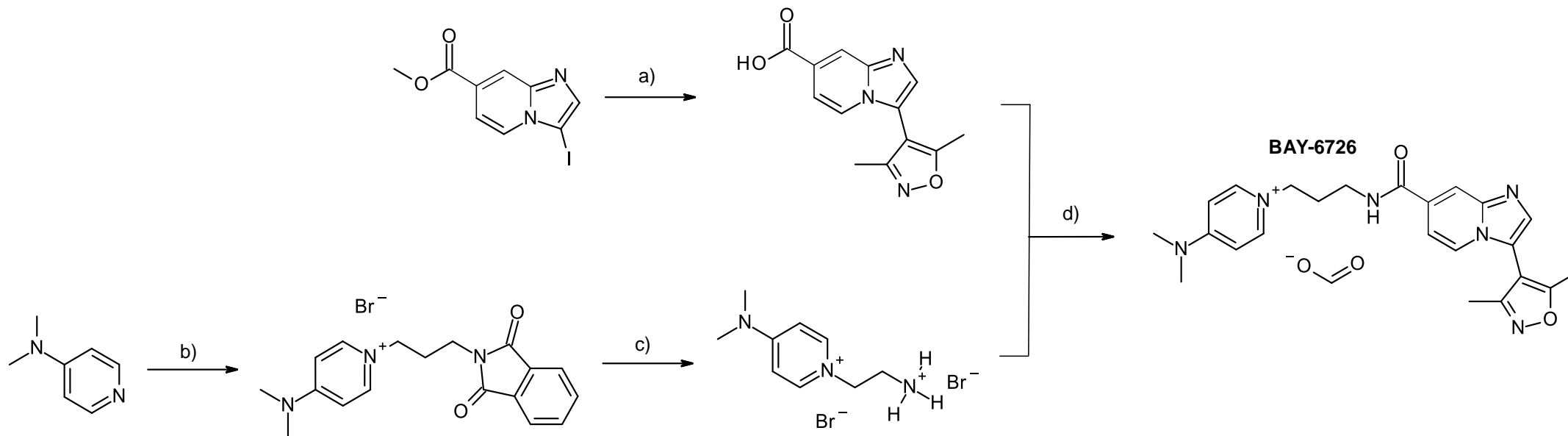
Synthesis of BAY-6096. Reagents and conditions: a) (3,5-dimethyl-1,2-oxazol-4-yl)boronic acid, [1,1-bis(diphenylphosphino)ferrocene]dichloropalladium(II), CsF, DMF, 90°C, overnight, 54%; b) NaOH aq. 1 M, THF, MeOH, r.t., 30 min, 95%; c) N-(2-chloroethyl)-phthalimide, DMF, 110°C, overnight, 63%; d) HCl aq. conc., 100°C, 3 d, 95%; e) EDC·HCl, DMAP, DCM, r.t., overnight, 63%. aq. = aqueous; conc. = concentrated; DCM = dichloromethane; DMF = N,N-dimethylformamide; EDC·HCl = N-(3-dimethylaminopropyl)-N'-ethylcarbodiimide hydrochloride; MeOH = methanol; r.t. = room temperature; THF = tetrahydrofuran.

BAY-6096 was synthesized in a convergent sequence of overall 5 steps



Adrenergic α_{2B} Antagonist iv Probe BAY-6096

Synthesis of BAY-6726



Synthesis of BAY-6726. Reagents and conditions: a) (3,5-dimethyl-1,2-oxazol-4-yl)boronic acid, tetrakis(triphenylphosphine) palladium (0), K_2CO_3 , DME/ H_2O , $75^\circ C$, 48h, 46%; b) 2-(3-bromopropyl)-1H-isoindole-1,3(2H)-dione, DMF, $110^\circ C$, overnight, 79%; c) HBr aq. conc., $100^\circ C$, overnight, 94%; d) EDC*HCl, DMAP, DCM, r.t., overnight, 69%. aq. = aqueous; conc. = concentrated; DCM = dichloromethane; DME = Dimethoxyethane; DMF = N,N-dimethylformamide; EDC*HCl = N-(3-dimethylaminopropyl)-N'-ethylcarbodiimide hydrochloride; r.t. = room temperature.

BAY-6726 was synthesized in a convergent sequence of overall 4 steps

Adrenergic α_{2B} Antagonist iv Probe BAY-6096

SMOL X-ray BAY-6096

Fig. 1: Ortep-Plot (50 %) with labeling scheme (molecule A)

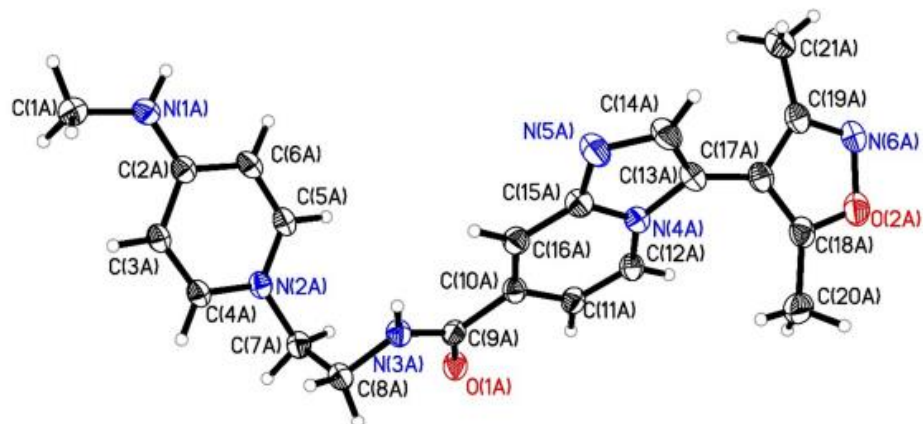
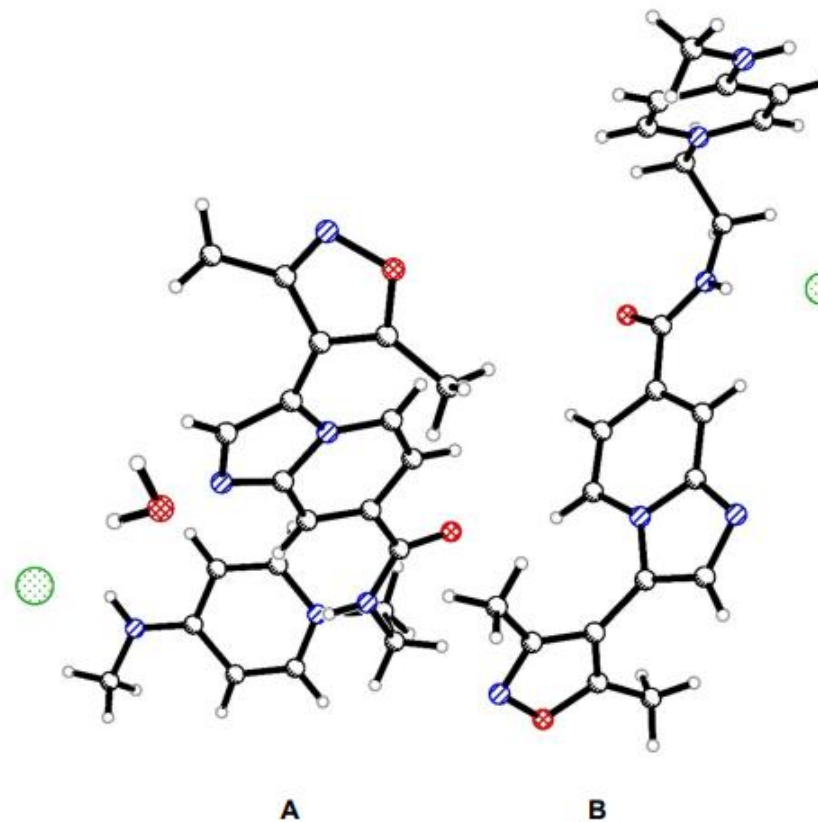


Fig. 2: Independent molecules in the asymmetric unit



BAY-6096 crystallized in the monoclinic P21/c space group