

THE DEVELOPMENT OF mGluR₈ PAM AGONISTS

A. S. Hogendorf,^{1,2} P. Brański,³ G. Burnat,³ R. Bugno,¹ A. Hogendorf,¹ B. Chruścicka,³
A. Stankiewicz,¹ M. Trela¹, A. J. Bojarski¹

- 1) Department of Medicinal Chemistry, Institute of Pharmacology, Polish Academy of Sciences, 12 Smętna Street, 31-343 Cracow, Poland
- 2) Department of Organic Chemistry, Jagiellonian University, 3 Ingardena Street, Cracow, Poland
- 3) Department of Neurobiology, Institute of Pharmacology, Polish Academy of Sciences, 12 Smętna Street, 31-343 Cracow, Poland



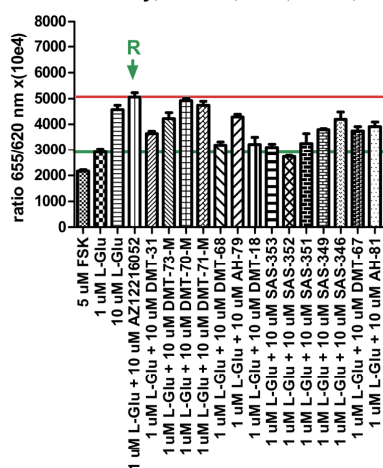
INTRODUCTION

Glutamate is the main excitatory neurotransmitter in the central nervous system (CNS). It is an essential molecule, e.g. for cognitive functions such as memory formation and learning.¹

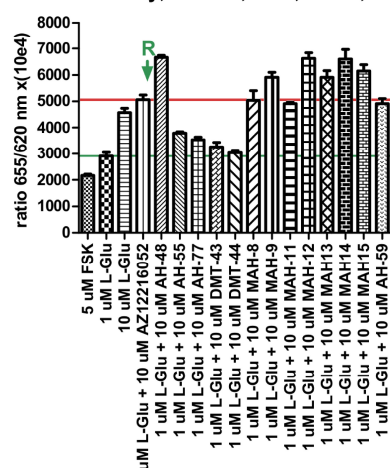
Group III metabotropic glutamate receptors (mGluR₄, mGluR₆, mGluR₇ and mGluR₈) are considered promising drug targets for treatment of neurological disorders e.g. Parkinson's disease, schizophrenia, major depressive disorder and pain.² Apart from the traditional concept of targeting orthosteric site, mGluR allosteric modulation is considered a very promising approach.³ Due to little differences in the aminoacid sequences of orthosteric binding sites of mGluR₄, mGluR₇ and mGluR₈, finding selective ligands is notoriously difficult.

mGluR₈ receptor, which is positively coupled to G_{α(i/o)}, functions as a presynaptic autoreceptor. GRM8 polymorphism may be involved in pathogenesis of schizophrenia.⁴ Activation of mGluR₈ can elicit both hyperalgesic and analgesic effects. Behavioural experiments suggest that mGluR₈ plays role in regulation of anxiety. mGluR₈ knockout (KO) mice exhibit an anxiety phenotype further implying involvement of this receptor in mood regulation.⁵

mGluR₈, 10 uM Group II Cmd. + EC20 L-Glu, potential PAM activity, column, ratio, cAMP, 05-08-2015

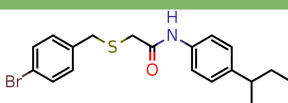


mGluR₈, 10 uM Group I Cmd. + EC20 L-Glu, potential PAM activity, column, ratio, cAMP, 05-08-2015



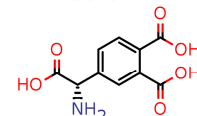
CHEMISTRY

Compound libraries (Group I and II) were synthesized, purity examined by LC-MS, structure confirmed by ¹H NMR and ¹³C NMR. Elemental analysis is consistent with calculated values.



AZ 12216052

mGluR8 PAM



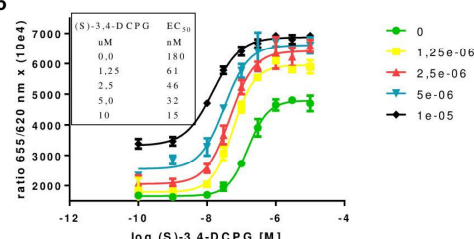
(S)-3,4-DPCG

mGluR8 agonist

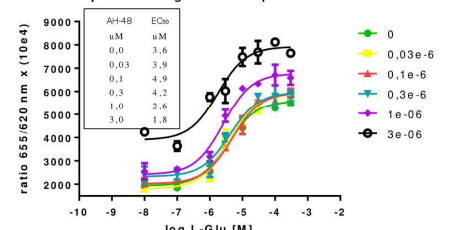
BIOLOGICAL ASSAYS

Cell line: Host: T-RexTM-293 cell line
Final cell lines: HEK293_T-REX-293_hmGluR8
Direct quantitative determination level of cAMP by Homogenous Time-Resolved Fluorescence
AZ 12216052 - mGluR8 Positive Allosteric Modulator has been used in the experiments as a reference.

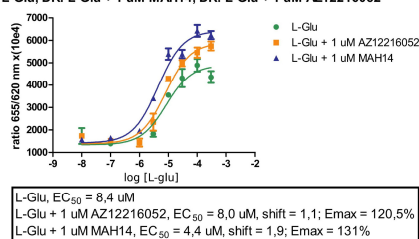
Response to the agonist (S)-3,4-DPCG in presence of AH-48



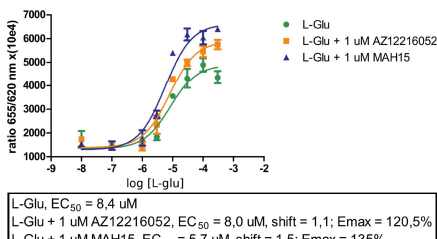
Response to the agonist L-Glu in presence of AH-48



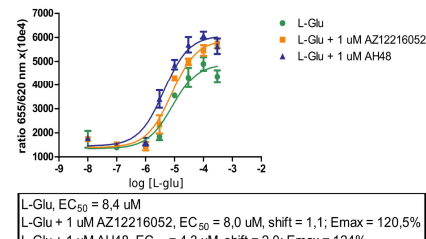
mGluR₈, DR: L-Glu, DR: L-Glu + 1 uM MAH14, DR: L-Glu + 1 uM AZ12216052



mGluR₈, DR: L-Glu, DR: L-Glu + 1 uM MAH15, DR: L-Glu + 1 uM AZ12216052



mGluR₈, DR: L-Glu, DR: L-Glu + 1 uM AH48, DR: L-Glu + 1 uM AZ12216052



Tab.1 Allosteric binding sites sequences of mGluRs⁶

PIN	mGlu ₄	mGlu ₅	mGlu ₆	mGlu ₇	mGlu ₈	mGlu ₄	mGlu ₅	mGlu ₆	mGlu ₇	mGlu ₈	mGlu ₄	mGlu ₅	mGlu ₆	mGlu ₇	mGlu ₈	Conservation	quality**	Consensus
2.45	624	G	G	G	G	G	G	G	G	G	G	G	G	G	G	11	17	100
2.46	625	I	I	V	V	I	I	I	I	I	I	I	I	I	I	9	16.7	77
2.48	628	G	G	C	C	S	C	I	C	C	C	G	S	S	S	5	7.4	55
3.36	651	I	V	L	L	L	L	L	L	L	L	I	I	I	I	9	16.2	66
3.37	652	G	G	G	G	G	G	G	G	G	G	G	G	G	G	11	17	100
3.39	654	S	S	A	S	F	M	G	G	G	G	G	G	G	G	8	13.9	55
3.40	655	P	S	F	F	F	M	T	M	M	M	F	A	A	A	4	8.4	33
3.43	658	S	C	C	C	S	S	S	S	S	S	I	S	S	S	5	9.4	55
3.44	659	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	Y	11	17	100
5.43	743	P	P	S	S	S	L	C	S	S	S	S	S	S	S	4	10	55
5.44	744	L	L	L	L	L	L	L	L	L	L	L	L	L	L	7	13.2	88
5.47	747	N	N	N	D	S	S	S	S	S	S	N	Q	Q	Q	7	13.9	44
5.51	751	I	I	I	V	M	M	M	M	M	M	I	I	I	I	8	15.2	44
6.46	781	T	T	T	T	T	T	T	T	T	T	T	T	T	T	11	17	100
6.49	784	I	I	I	V	I	V	I	V	I	V	I	I	I	I	9	16.4	77
6.50	785	W	W	W	W	W	W	W	W	W	W	W	W	W	W	11	17	100
6.53	788	F	F	F	F	F	F	F	F	F	F	F	F	F	F	11	17	100
7.32	802	M	T	M	M	L	L	L	L	L	L	L	L	L	L	6	14.6	55
7.35	805	S	A	S	S	S	S	S	S	S	S	S	S	S	S	8	16.4	88
7.36	806	V	V	V	V	V	V	V	V	V	V	V	V	V	V	8	15.5	55
7.39	809	S	S	S	S	S	S	S	S	S	S	S	S	S	S	11	17	100
7.40	810	A	V	G	G	A	A	A	A	A	A	A	A	A	A	8	13.4	66
7.43	813	A	A	V	V	S	S	A	S	A	S	A	S	A	S	6	13.1	44

CONCLUSIONS

We have developed two distinctive chemotypes with 22 entities exhibiting PAM properties. Six compounds from group II are more potent than the reference compound **AZ 12216052**, which is the only commercially available mGluR₈ ago-PAM till date. Lead compound AH-48 has the following characteristics:

- it activates mGluR₈ as an agonist (EC₅₀ = 2.6 uM),
- it acts as a Positive Allosteric Modulator (EC₅₀ = 4.3 uM in the presence of 1 uM L-Glu),
- AH-48 acts as mGluR₈ full PAM-agonist in contrast to benchmark compound **AZ 12216052** which activates the receptor only partially.

The transmembrane binding sites sequences between different group III mGluRs differ very slightly (tab.1). Thus obtaining selective ligands is very challenging. Herein presented group I and II ligands are not mGluR₄ selective nor is **AZ 12216052**. A further investigation of binding modes in order to exploit binding site differences may potentially yield selective ligands.

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