

Introduction

The melanocortin system is comprised of five complex integral membrane protein isoforms (MC1R-MC5R), which mediate many key physiological functions and are a subclass of the GPCR superfamily. Activation of the MCRs is facilitated via binding of the endogenous melanotropin agonists (α -MSH, β -MSH, γ -MSH, and ACTH) in order to stimulate the cAMP signal transduction pathway.

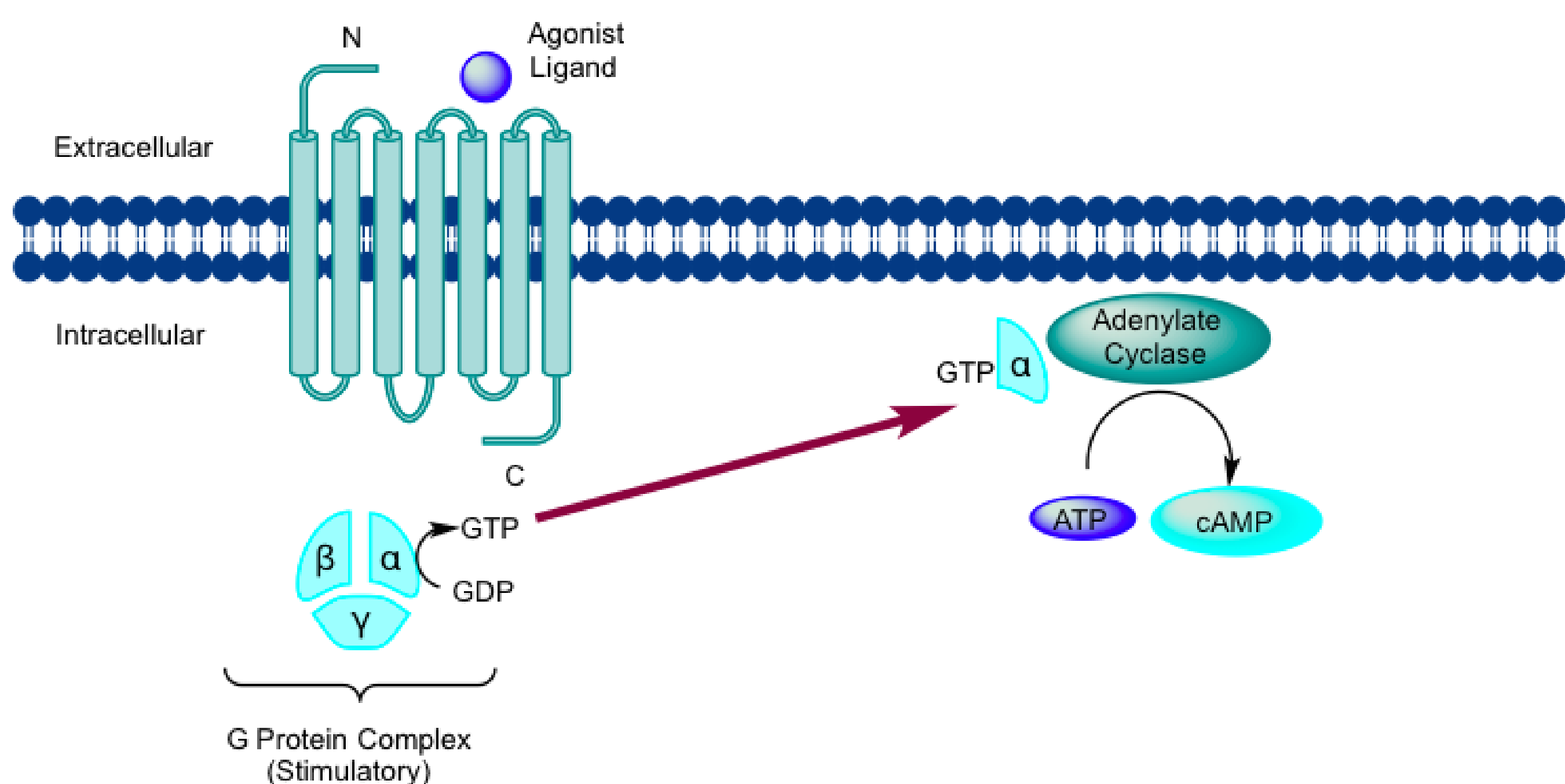


Figure 1.1—Activation of the heterotrimeric G_s protein and secondary messenger (cAMP) through agonist binding.

MC3R and MC4R have been shown to play an important role in energy metabolism and feeding behavior. They are also mediators of anti-inflammatory signaling in the brain. However, MC4R expression in the CNS is more abundant than MC3R; making this receptor an ideal target for suppressing brain inflammation. Guarini et al. showed that MC4R stimulation by NDP- α -MSH increases neuroprotection in a mouse model. Agonist analogs dubbed Anti-Inflammatory Melanotropins (AIMs) were synthesized in order to enhance ligand potency and selectivity at MC4R.

Design of AIM Drugs

Molecular recognition of agonists by the MCRs is based upon the conserved His-Phe-Arg-Trp (HFRW) pharmacophore sequence. AIMs contain multiple alterations to their structure to increase selectivity. Global stereochemical constraint is introduced via lactam bond formation, in addition β -amino acid substitutions allow for enhanced pharmacophore mobility. In order to increase stability in the binding pocket a constrained amino acid is introduced in the sequence. Halogenation of specific amino acids allows for ion induced dipole interactions in the binding pocket.

Table 1.1—Structural Sequence of Naturally-Occurring Melanotropin Agonists

Agonists	Amino Acid Sequence
α -MSH	Ac-SYSMEHFRWGKPV-NH ₂
β -MSH	H ₂ N-AEKKDEGPYRMEHFRWDRLF-OH
γ -MSH	H ₂ N-YVMGHFRWDRLF-OH
ACTH	H ₂ N-SYSMEHFRWGKPVGKKRRPVKYPNGAEDESAAFPLEF-OH

Table 1.2—Structural Sequence of Nonselective Exogenous Melanotropins

Drugs	Amino Acid Sequence
NDP- α -MSH	Ac-Ser-Tyr-Ser-Nle-Glu-His-D-Phe-Arg-Trp-Gly-Lys-Pro-Val-NH ₂
MT-II	Ac-Nle-cyclo[Asp-His-D-Phe-Arg-Trp-Lys]-NH ₂
SHU-9119	Ac-Nle-cyclo[Asp-His-D-Nal(2')-Arg-Trp-Lys]-NH ₂
SHU-9005	Ac-Ser-Tyr-Ser-Nle-Glu-His-(pI)-D-Phe-Arg-Trp-Gly-Lys-Pro-Val-NH ₂

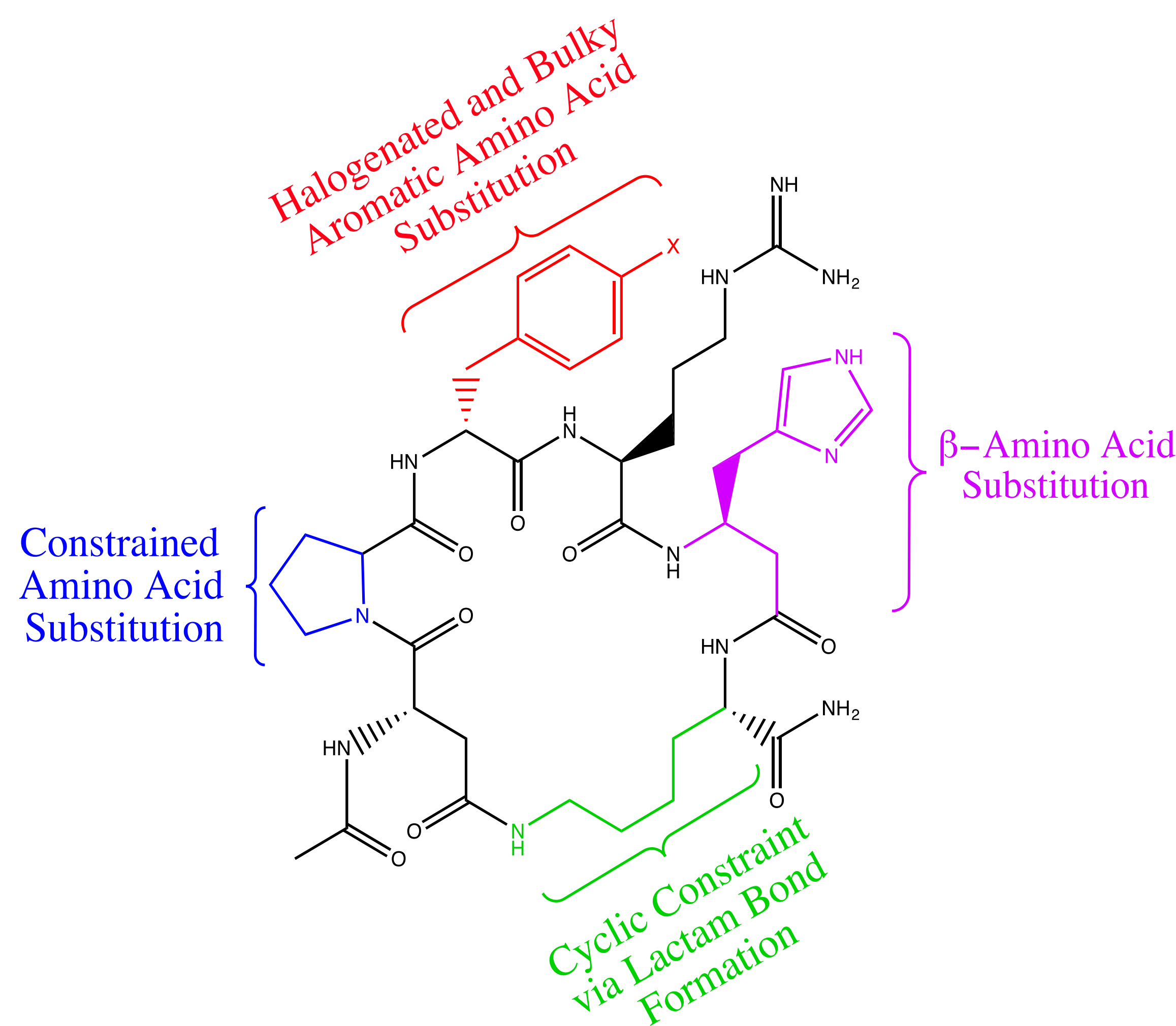


Figure 1.2—Principal features involved in the AIM design strategy.

Methods

The peptides were synthesized on low-loading Rink Amide resin using N^α-Fmoc chemistry and an orthogonal side chain protection strategy, which was executed through solid phase peptide synthesis. The linear chain was then acetylated at the N-terminus and cyclized via lactam bond formation. Each peptide was confirmed with Mass Spectroscopy and purified by HPLC.

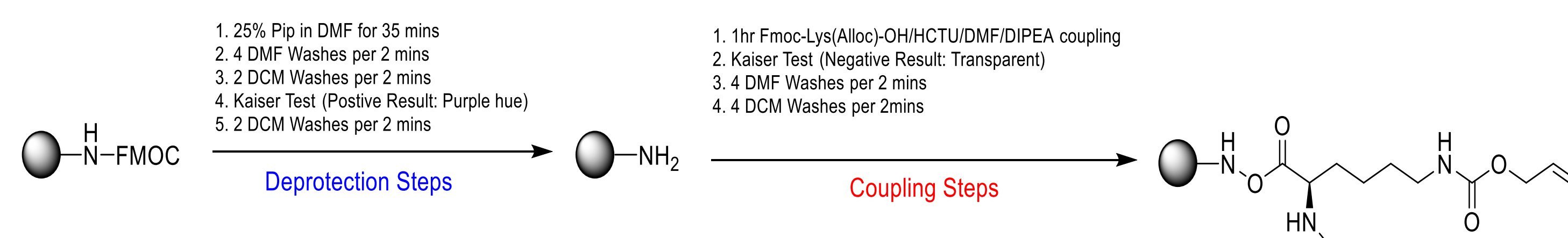


Figure 1.3—Solid phase peptide synthesis scheme.

Pharmacodynamics of AIMs were assessed by cAMP assays. This assay compares cAMP levels produced by receptor stimulation via binding of AIMs with a specific radiolabeled [³H]-cAMP concentration. The assay was ran on MCR transfected HEK293 cells and was evaluated on a MicroBeta² counter.

Biological Results

	hMC1R			hMC3R			hMC4R			hMC5R		
Drug	EC ₅₀ (nM)	Act%		EC ₅₀ (nM)	Act%		EC ₅₀ (nM)	Act%		EC ₅₀ (nM)	Act%	
AIM 1	120 ± 24	88		680 ± 25	88		>1000	63		580 ± 340	95	
AIM 2	220 ± 70	120		>1000	89		47±24	67		>1000	90	
AIM 3*	190 ± 90	82		120 ± 10	87		16±4	84		290 ± 50	82	
AIM 4*	64 ± 10	90		450 ± 830	76		630±175	46		530 ± 65	72	
AIM 5	340 ± 0	100		>1000	59		>1000	84		>1000	62	
AIM 6	>1000	61		>1000	49		>1000	120		>1000	51	
AIM 7	99 ± 0	67		>1000	24		240±10	57		>1000	21	
AIM 8*	130 ± 0	89		360 ± 25	46		110±90	66		>1000	38	
AIM 9**	240 ± 11	90		>1000	63		4±2	87		295 ± 165	76	
MT-II	32	100		3	100		10	100		2	100	

Table 1.3—cAMP Assay Data for AIMs.

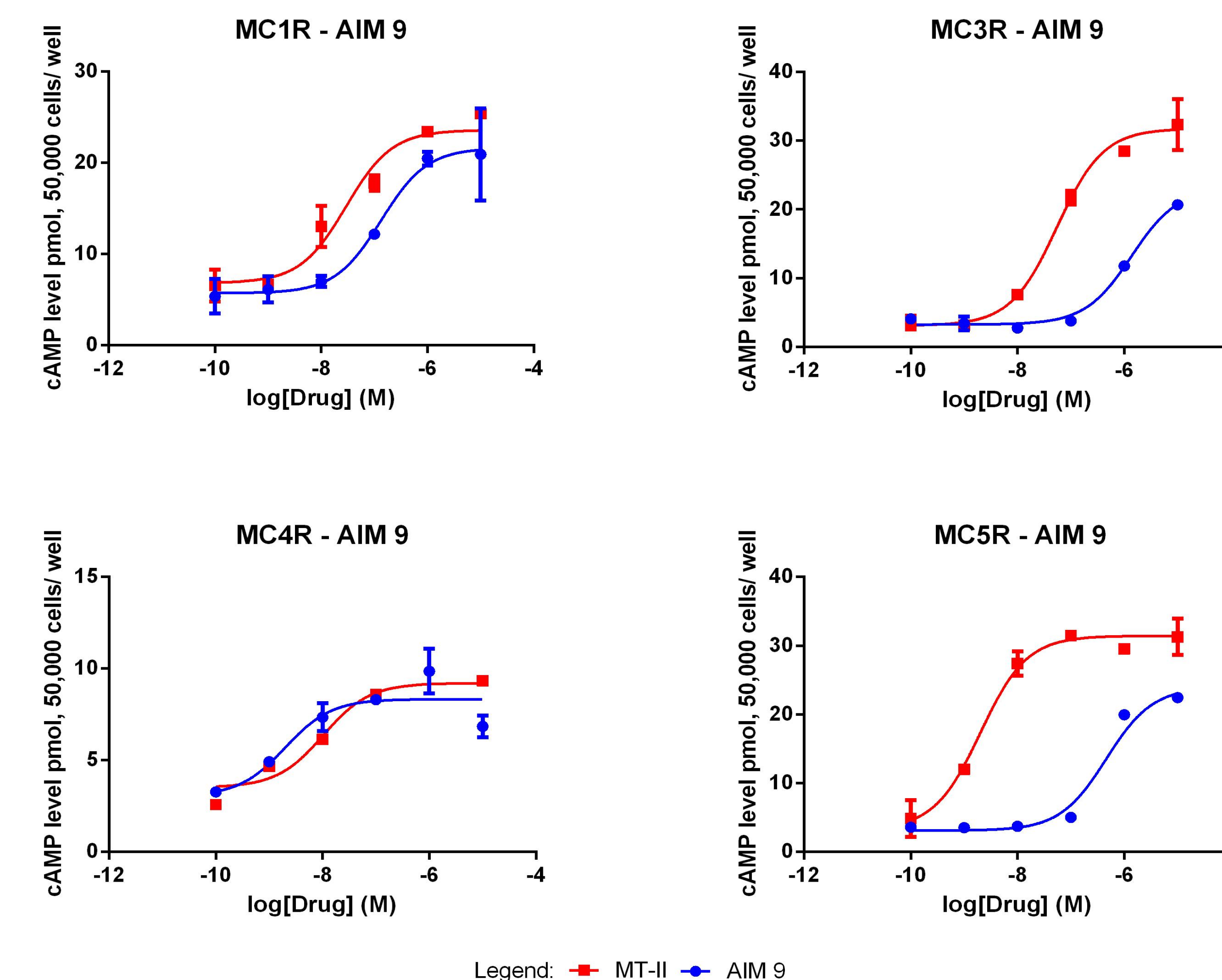
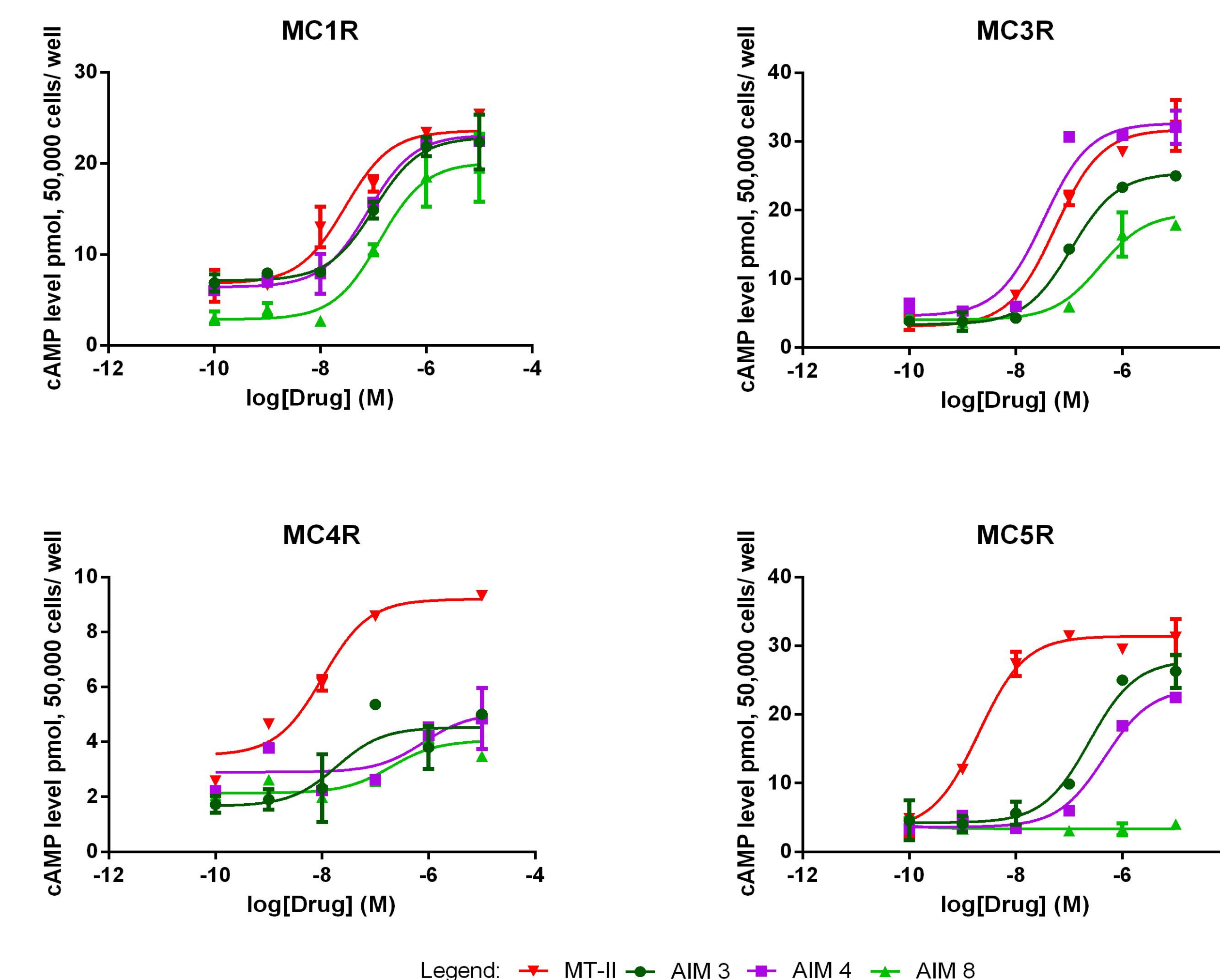


Figure 1.4—cAMP Assay Data for AIMs 3, 4, 5, 9, and the control MT-II on MCR's.

Conclusions

Current biological data suggests that all AIM molecules act as agonists on MCRs. Global constraint, β -amino acid substitutions, halogenation of the pharmacophore and a constrained amino acid addition all point towards selectivity for MC4R. This is best exhibited in AIM 9 – a peptide that contains all of the aforementioned modifications. Our future directions are to conduct detailed binding studies for AIMs on the MC4R and use the results to elucidate the structure of this receptor.

References

- [1] Cai M, Hruby VJ (2016) The Melanocortin Receptor System: A Target for Multiple Degenerative Diseases. *Cur. Pro. Pep. Sci.* 17: 488-496
- [2] Cai M, Hruby VJ (2016) Design of Cyclized Selective Melanotropins. *Pep. Sci.* 106: 876-883
- [3] Giuliani D, Guarini S (2013) Melanocortins protects against progression of Alzheimer's diseases in triple transgenic mice by targeting multiple pathophysiology pathways. *Neurobio. of Aging.* 35: 537-547

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