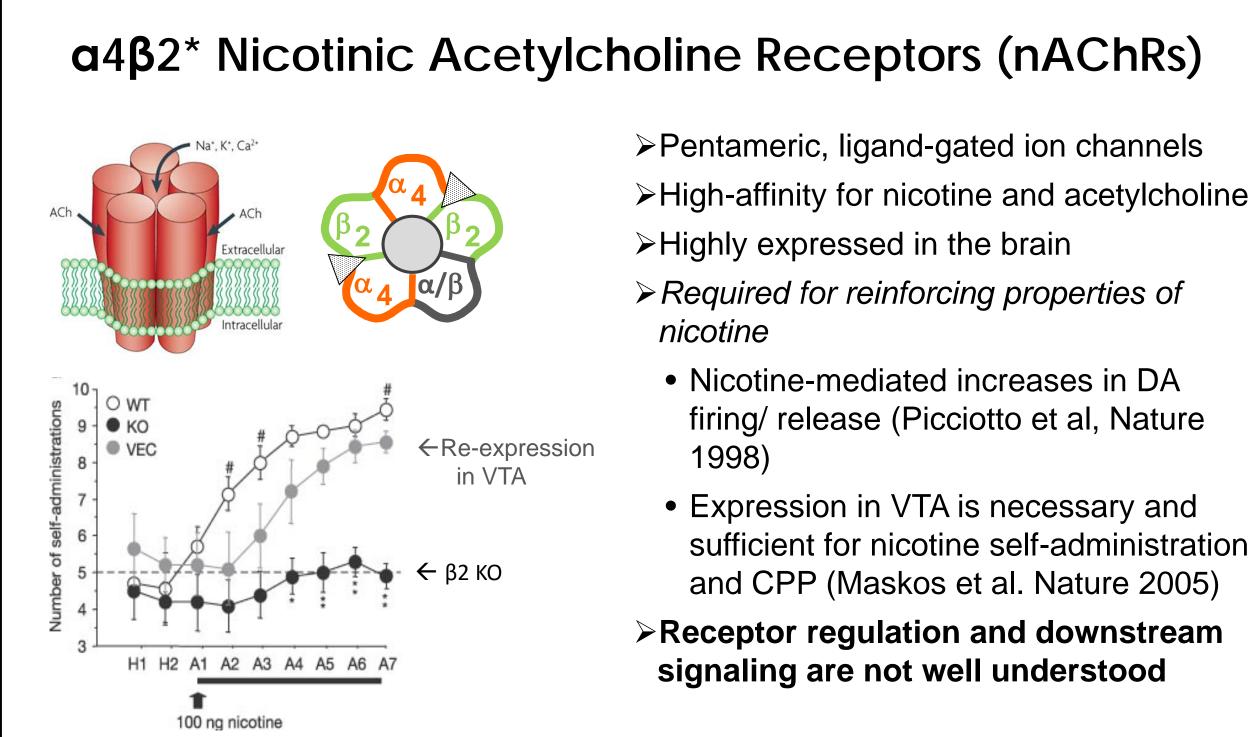


Background

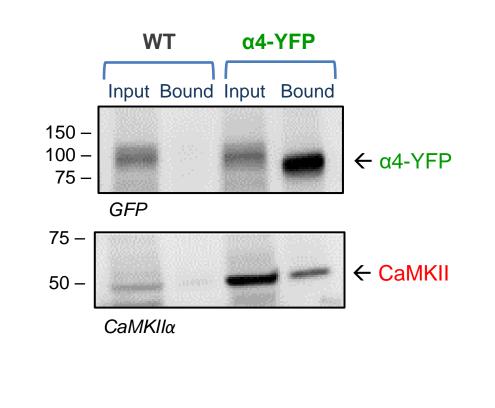
- > Tobacco use is the leading cause of preventable death worldwide
- > Nicotine binds to and activates nicotinic acetylcholine receptors (nAChRs)
- \geq Primary reinforcing properties of nicotine are mediated by $\alpha 4\beta 2^*$ nAChRs
- > Molecular mechanisms underlying regulation and downstream signaling of α4β2* nAChRs are not completely understood
- \succ CaMKII α , a highly expressed brain kinase with well-established roles in synaptic plasticity, was recently identified as a protein interactor of $\alpha 4\beta 2^*$ nAChRs in mouse and human brain
- \succ In this study, we aimed to elucidate the role of CaMKIIa in nicotinic receptor function, focusing on receptor phosphorylation and localization

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The a4β2* nAChR Protein Interactome

Correlation coefficient	N	F score	P value	Protein	UniProtKB accession No.	Cellular compartment	Previously identified?	Molecular function
0.748	12	20.268	0	Glial fibrillary acidic protein	P03995	Cytoplasm	No	Protein binding, structur
0.858	18	44.542	0	nAChR subunit a4	O70174	Cell junction	No	Neuro-transmitter recept
1	18	-	0	nAChR subunit β2	Q9ERK7	Cell junction	No	Neuro-transmitter recept
0.652	18 18	11.844	0.003	Neurofilament light poly peptide	P08551	Growth cone	No No	Protein binding, structur Nucleotide binding, prot
0.637	18	10.904	0.004	Calcium/calmodulin-dependent protein kinase type II subunit α	P11798	Cytoplasm	No	Transferase, nucleotide b
0.917	18	21.235	0.01	Calcium/calmodulin-dependent protein kinase type II subunit γ	Q923T9	Sarcoplasmic reticulum membrane	No	Transferase, nucleotide b
0.57	18	/.681	0.014	F-actin-capping protein subunit α2	P4//54	Cytoplasm	NO	Protein binding
0.562	18	7.386	0.015	Thyroid hormone receptor- associated protein 3	Q569Z6	Nucleus	No	Nucleotide binding, prot
0.665	12	7.933	0.018	Transcriptional activator protein Pur-α	P42669	Nucleus	No	Nucleic acid binding, tra regulator, protein bindir
0.539	18	6.563	0.021	Ectonucleotide pyrophosphatase/phosphodiesterase family member 6	Q8BGN3	Cell membrane	No	Catalytic activity, hydrol
0.519	18	5.884	0.027	Spectrin β chain, brain 1	Q62261	Cytoplasm	No	Protein binding, lipid bin structural molecule activ
0.856	6	11.009	0.029	Ras-related protein Rap-1A	P62835	Cell membrane	No	Hydrolase activity, prote
0.512	18	5.695	0.03	Myosin-10	Q61879	Cytoplasm	No	Protein binding, nucleoti
0.506	18	5.496	0.032	Myelin proteolipid protein	P60202	Cell membrane	No	Structural molecular, pro
0.502	18	5.378	0.034	Spectrin a chain, brain	P16546	Cytoplasm	Yes	Hydrolase, protein bindi
0.493	18	5.149	0.037	Tubulin β-3 chain	Q9ERD7	Cytoplasm	No	Hydrolase, nucleotide bi molecular, protein bindin



 \geq Immunoprecepitation from WT, β 2 KO or heterozygous mouse brain

- ➢Quantitative LC MS/MS for protein ID
- \geq <u>17 proteins</u> correlated significantly with β 2 nAChR gene dose, including CaMKII isoforms
- >CaMKII isoforms also found in nAChR interactome in human cortical tissue (McClure-Begley, eNeuro 2016)
- **CaMKII**α co-IP confirmed using lysates from α4-YFP transgenic mice

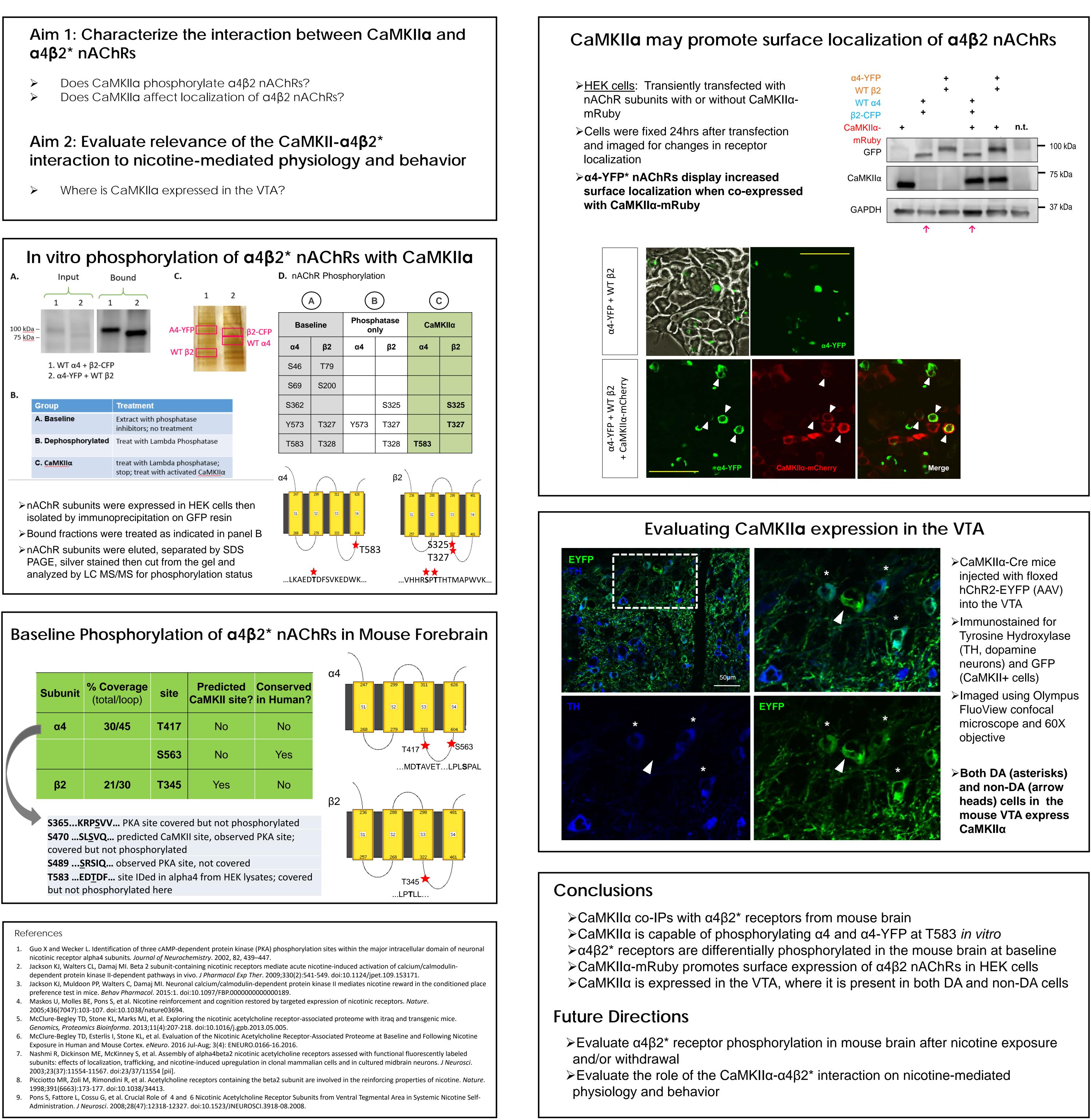
Examining the Role of CaMKIIa in a4β2* Nicotinic Receptor Function

Megan B. Miller, Wenliang Zhou, Jean Kanyo[†], TuKiet Lam[†], and Marina R. Picciotto Yale University School of Medicine, Department of Psychiatry; Yale/NIDA Neuroproteomics Resource[†]

- ral molecule iral molecule binding, protein binding binding, protein binding otein binding olase activity tein binding, nucleotide binding otide binding, hydrolase rotein binding ling, nucleotide binding oinding, structural ing, peptide McClure-Begley et al. GPB. 2013

a4β2* nAChRs

Does CaMKIIa phosphorylate a4β2 nAChRs?



	Subunit	% Coverage (total/loop)	site	Predicted CaMKII site?					
	α4	30/45	T417	No	No				
			S563	No	Yes				
	β2	21/30	T345	Yes	No				
S365KRP <u>S</u> VV PKA site covered but not phosphorylated S470SL <u>S</u> VQ predicted CaMKII site, observed PKA site; covered but not phosphorylated S489 <u>S</u> RSIQ observed PKA site, not covered T583EDTDF site IDed in alpha4 from HEK lysates; cover but not phosphorylated here									

500.01

