

Genetic influences on human frontal cortical networks

Brian J. Mickey, MD, PhD



Disclosures

Speakers bureau:	none
Consultant:	none
Advisory boards:	none
Equity interest:	none
Salary support:	St Jude Medical, 2011-13 (5%)
Off-label uses:	none
Grant support:	National Institutes of Health University of Michigan Department of Psychiatry University of Michigan Depression Center Taubman Medical Research Institute

Acknowledgements



THE MOLECULAR & BEHAVIORAL
NEUROSCIENCE INSTITUTE
UNIVERSITY OF MICHIGAN



neuroscience graduate program

University of Michigan

Jon-Kar Zubieta

David Hsu

Marta Pecina

Tiffany Love

Mary Heitzeg

Tal Shafir

Heng Wang

Susan Kennedy

University of Pennsylvania

Falk Lohoff

NIAAA

David Goldman

Laura Bevilacqua

Zhifeng Zhou

Elizabeth Heinz

Pei-Hong Shen

Colin Hodgkinson

University of Illinois Chicago

Scott Langenecker

Sara Weisenbach

Overview

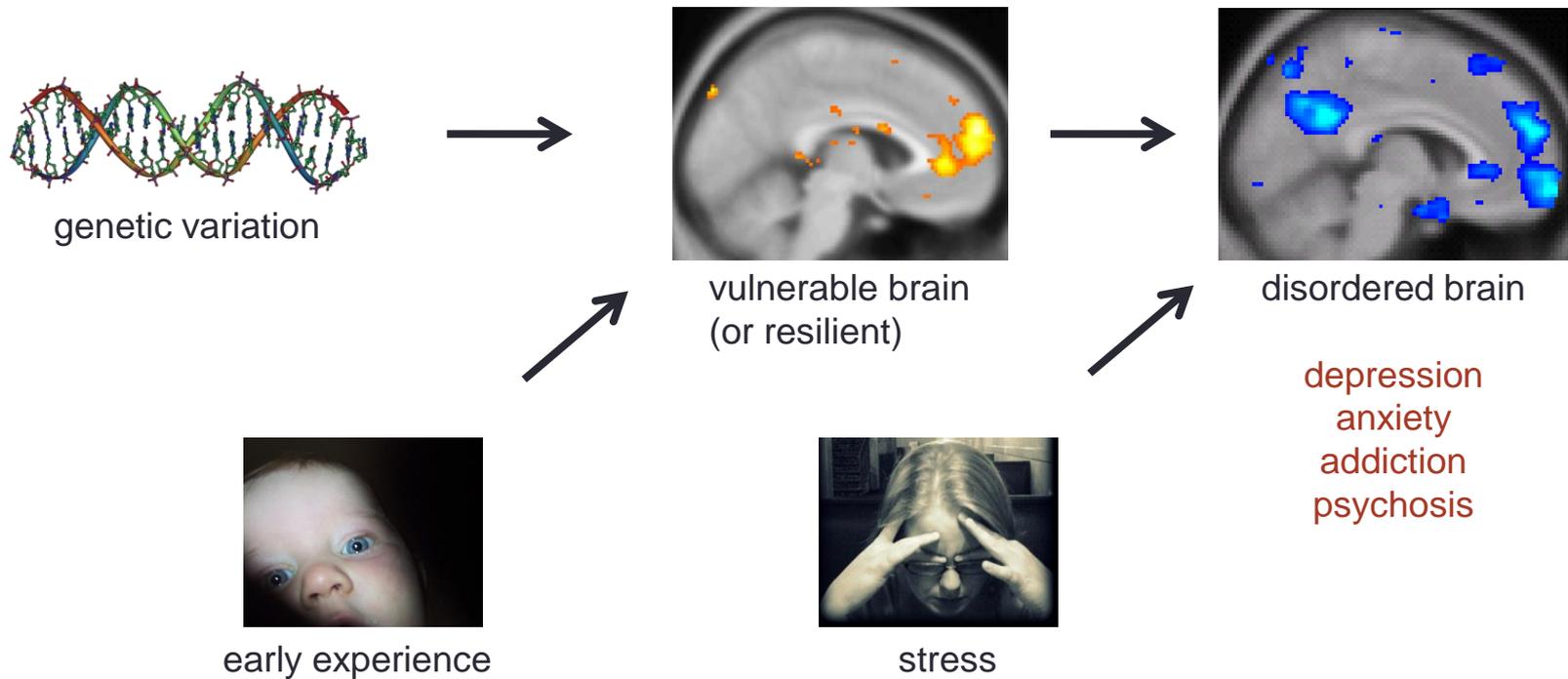
Genes, environment, risk, and the brain

Four genes that influence frontal-cingulate emotion circuitry

<i>NPY</i>	neuropeptide Y
<i>VMAT1</i>	vesicular monoamine transporter 1
<i>CRHR1</i>	corticotrophin releasing hormone receptor 1
<i>DRD2</i>	dopamine D2 receptor

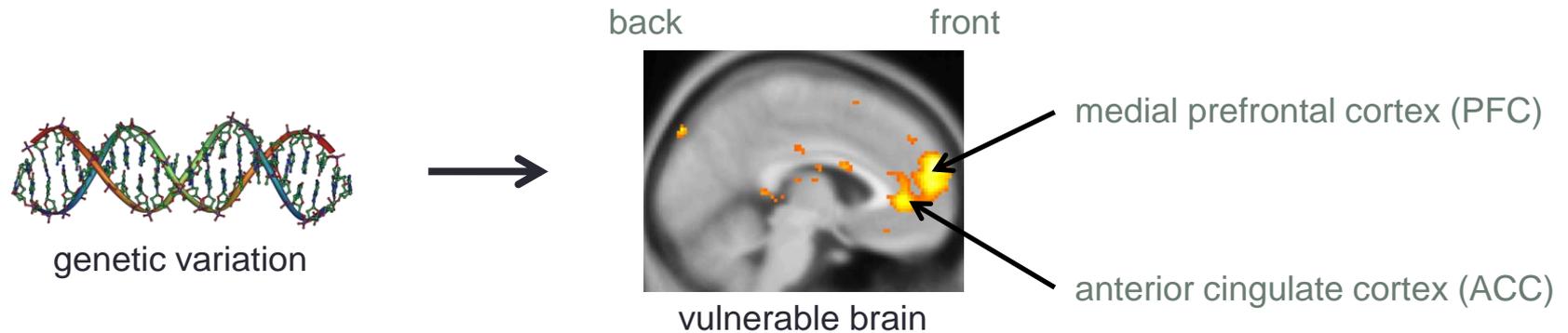
Summary and future directions

Genes, environment, and risk



Today's focus

Which variable genetic factors give rise to variation in emotional brain function?



Variant **X** is functional *in vitro* and has been implicated in psychiatric disorders

Is **X** associated with altered function of neural region **Y**?

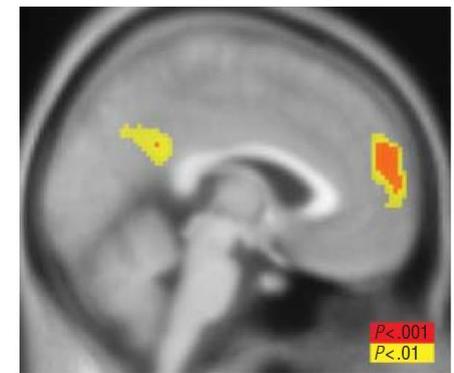
Altered **Y** = candidate intermediate phenotype

Imaging prefrontal emotion processing in humans

Functional magnetic resonance imaging (fMRI)
Blood oxygen level dependent (BOLD) response
Indirect measure of neural activity

Emotion word task
93 healthy adults and 18 with major depression

war	lost	burial	negative
...	iron	journal	neutral



Neuropeptide Y

A regulator of the brain's stress response

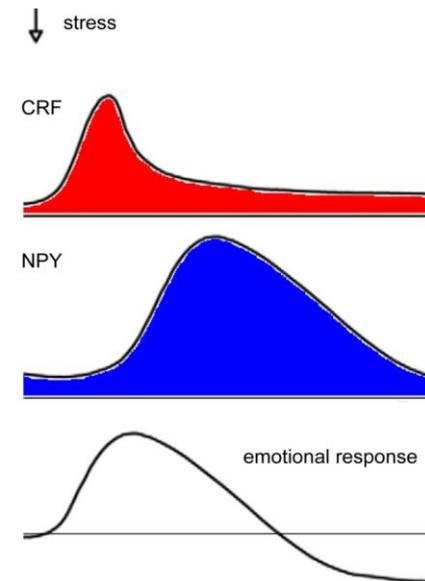
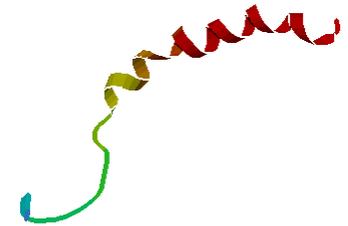
NPY is a 36-amino-acid peptide neurotransmitter
evolutionarily conserved
widely distributed in the brain
expressed at high concentrations

Neuropeptide Y is co-released with other neurotransmitters, including GABAergic interneurons in the cortex and striatum

(Kask et al., *Neurosci Biobehav Rev*, 2002)

Experiments in animal models indicate that stress increases expression and release of NPY in the amygdala and that NPY reduces anxiety-like behavior

(Heilig, *Neuropeptides*, 2004; Rhinehart et al., 2009)



Neuropeptide Y

Implicated in neuropsychiatric disease

Peripheral NPY has been positively associated with resilience to psychological stress

(Morgan et al., *Biol Psychiatry*, 2000, 2002, 2003; Yehuda et al., *Biol Psychiatry*, 2006)

Low NPY concentrations in plasma, cerebrospinal fluid, and postmortem brain have been variably associated with mood disorders

A single-nucleotide polymorphism in the *NPY* gene was linked with treatment-resistant major depressive disorder

(Heilig et al., *J Psychiatr Res*, 2004)

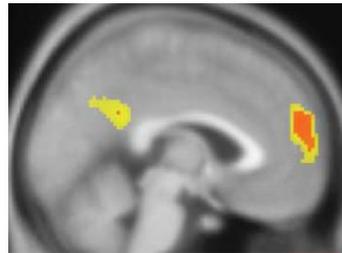
Variation in NPY expression is influenced by variation in the *NPY* gene

(Heilig et al., *J Psychiatr Res*, 2004; Zhou et al., *Nature*, 2008)

Does *NPY* variation affect prefrontal emotion processing in humans?



NPY Low



response to
negative emotion

...

mood and
anxiety
disorders

Functional genetic variation of NPY

Six-marker haplotype

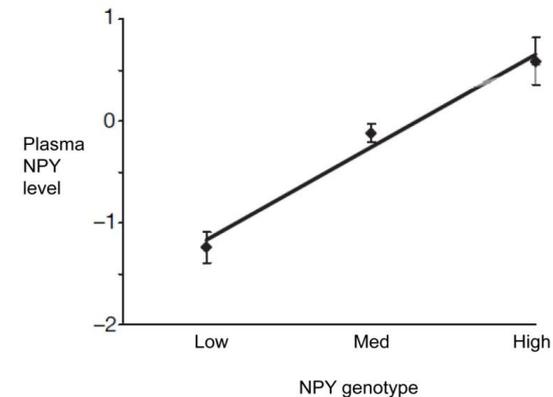
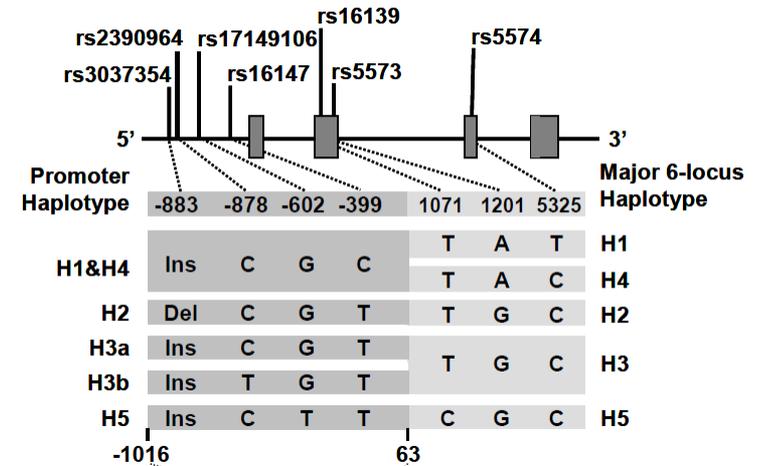
(Zhou et al., *Nature*, 2008)

Individuals with low-expression NPY genotypes exhibited:

lower endogenous opioid release during pain

greater amygdala responses to threat-related stimuli

greater trait anxiety



modified from Zhou et al., *Nature*, 2008

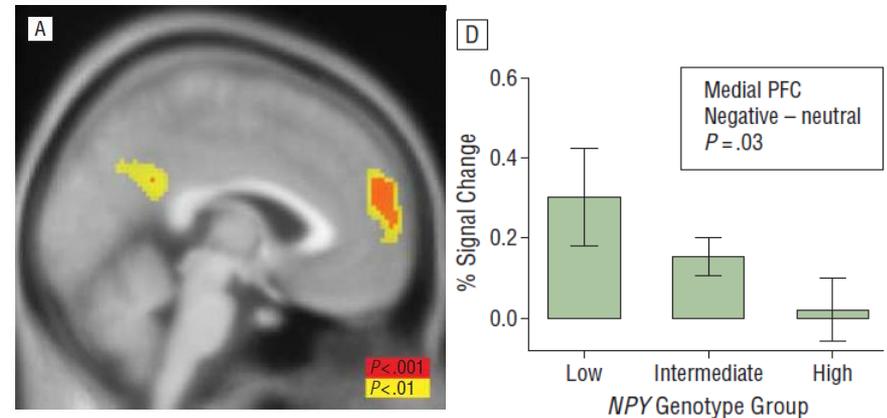
Effects of *NPY* variation on prefrontal function in humans

Increased response to negative words in the Low *NPY* group

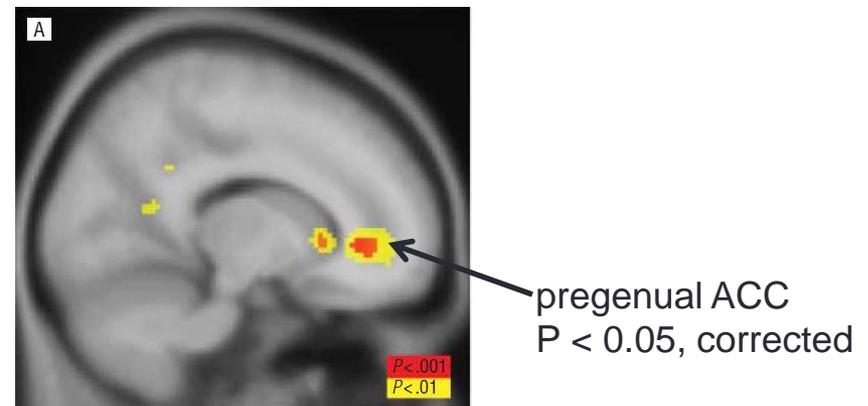
medial prefrontal cortex
anterior cingulate cortex

Gene effects persisted when adjusting for age, sex, and ancestry

Region-of-interest

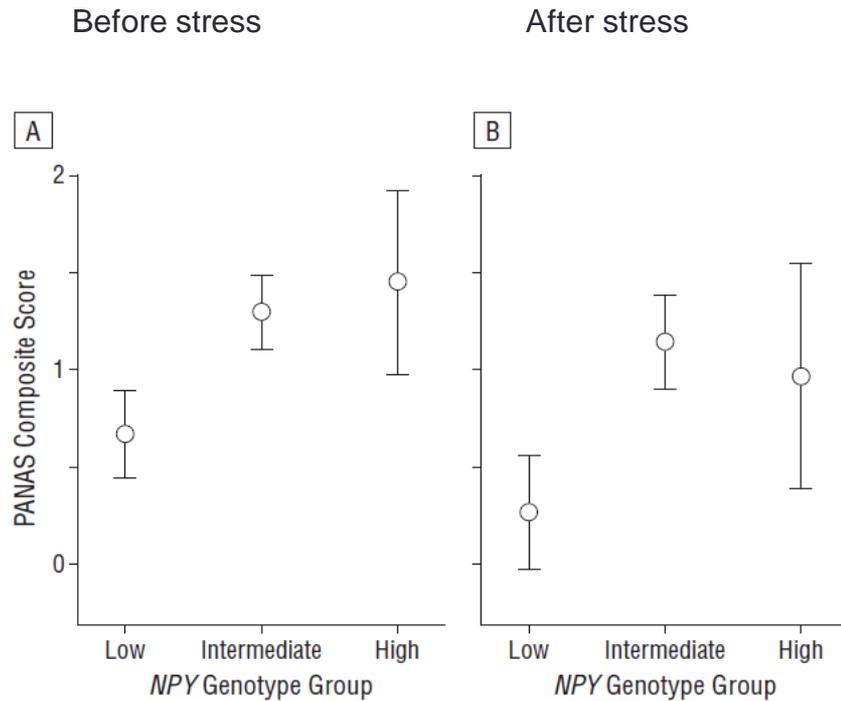


Whole-brain



NPY effects on emotion phenotypes

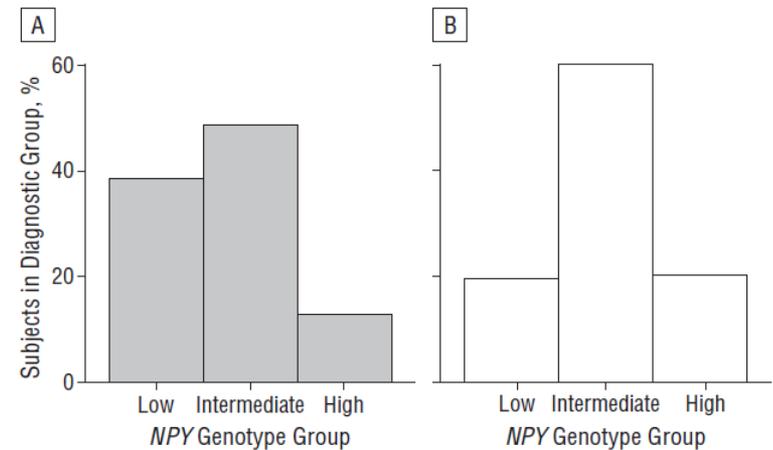
Affective state during pain-stress challenge (n = 78)



Association with depression

Major Depressive Disorder
n = 39

Controls
n = 113



Summary: *NPY* variation and neural processing of emotion

Low-expression variants of the *NPY* gene increase responsiveness of medial PFC and pregenual ACC to negative words

Preliminary data suggest effects of *NPY* on response of nucleus accumbens monetary loss

NPY-driven variation in prefrontal, cingulate, and accumbens response to negative stimuli is a potential endophenotype for mood and anxiety disorders

Low *NPY* expression by cortical and striatal interneurons may heighten the sensitivity to negative stimuli, thereby increasing risk of psychiatric illness

Vesicular monoamine transporter 1

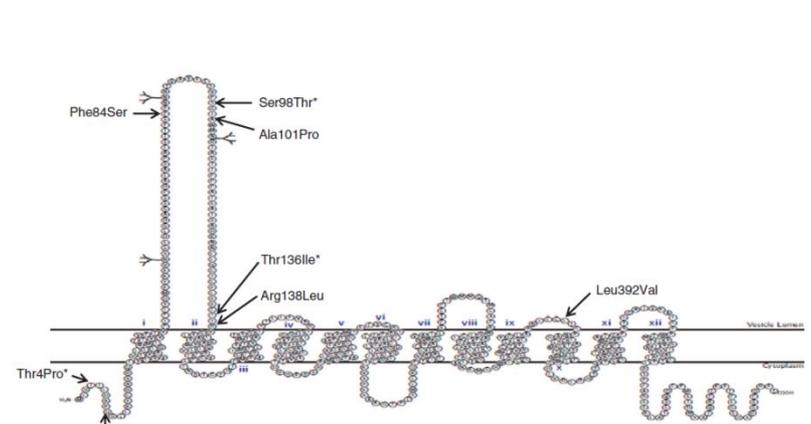
A key enzyme in neuronal monoamine storage

Packages monoamine transmitters into synaptic vesicles

serotonin (5HT)

dopamine (DA)

norepinephrine (NE)



Two vesicular monoamine transporters

VMAT1

VMAT2

VMAT1 relatively selective for serotonin

Vesicular monoamine transporter 1

Implicated in neuropsychiatric disease

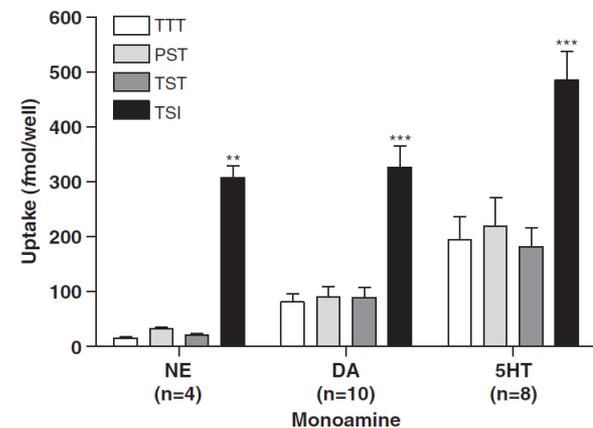
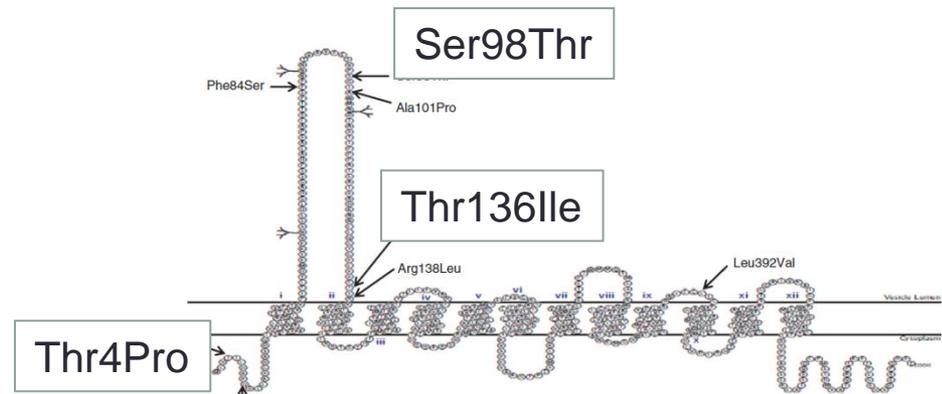
Pharmacologic blockade of VMAT causes depression in humans

Several coding variants in *VMAT1* gene (*SLC18A1*)

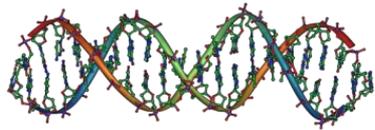
Thr136 associated with bipolar disorder (Lohoff et al., *Neuropsychopharm*, 2006)

Thr136 has lower activity *in vitro*

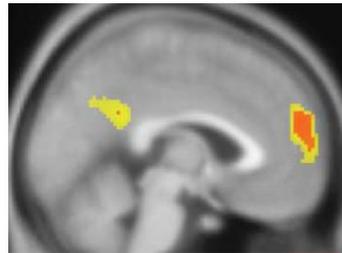
No effects of Thr4Pro or Ser98Thr



Does *VMAT1* variation influence emotion processing in humans?



VMAT Thr136



response to
negative emotion

...

mood
disorders

Effects of *VMAT1* variation on prefrontal function

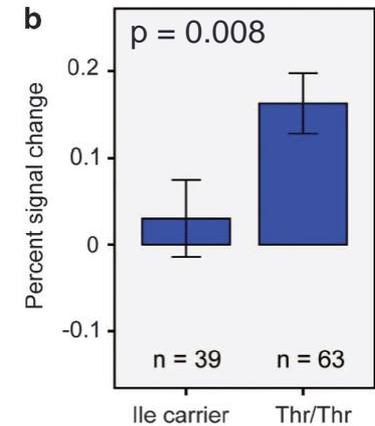
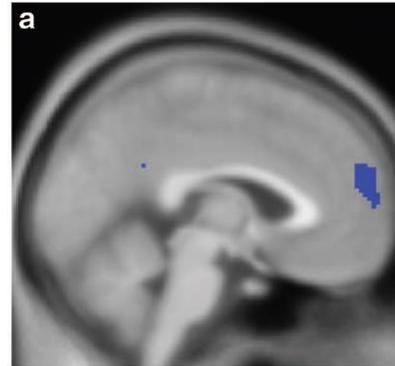
Greater activation among Thr136 homozygotes

medial prefrontal cortex
anterior cingulate cortex

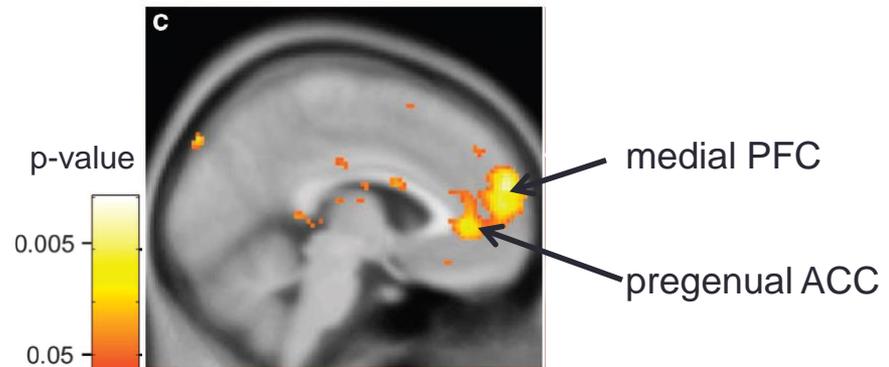
Gene effects persisted when adjusting for age, sex, ancestry, and diagnosis

No significant effects of putatively non-functional variants Thr4Pro and Ser98Thr

Region-of-interest



Whole-brain



Summary: *VMAT1* variation and emotion processing

VMAT1-driven variation in prefrontal activation with negative emotion is a potential endophenotype for psychiatric disorders

Thr136 may cause less efficient filling of monoamine vesicles, reducing frontal release of monoamines, increasing metabolic activity

Lower functioning allele Thr136 associated bipolar disorder, but rare hyper- and hypo-functional alleles may also be associated

(Lohoff et al., *Mol Psychiatry*, 2013)

VMAT1 genotype could contribute to variation in brain or clinical responses to monoaminergic drugs

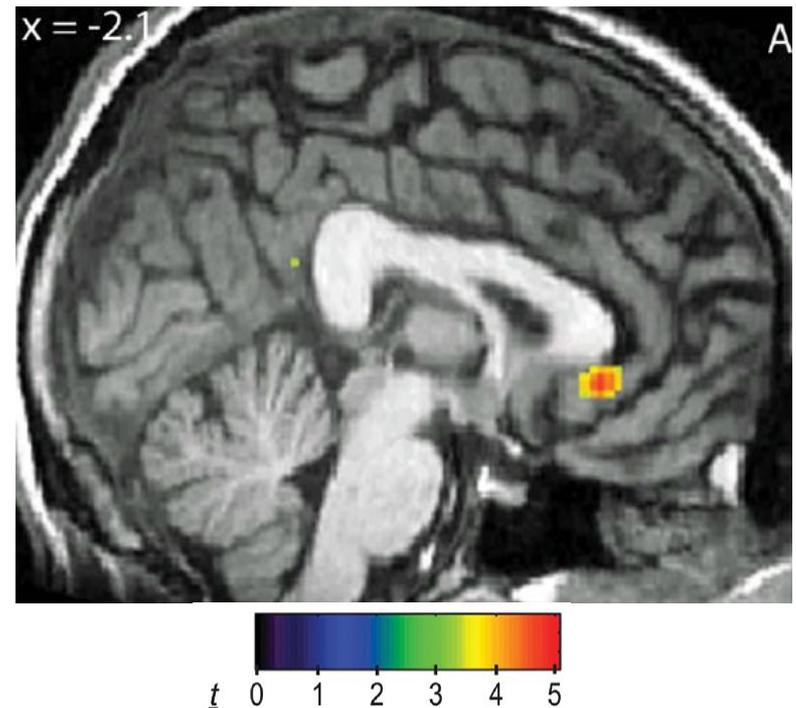
Corticotrophin releasing hormone receptor 1

Autonomic, hormonal, and behavioral response to stress

CRHR1 rs110402
single nucleotide variant

G allele associated with major depressive disorder (MDD) in context of childhood abuse
(Bradley et al., 2008; Polanczyk et al., 2009; Heim et al., 2009; Tyrka et al., 2009; Ressler et al., 2010)

Hyper-activation of subgenual anterior cingulate cortex in MDD patients among GG homozygotes but not A carriers

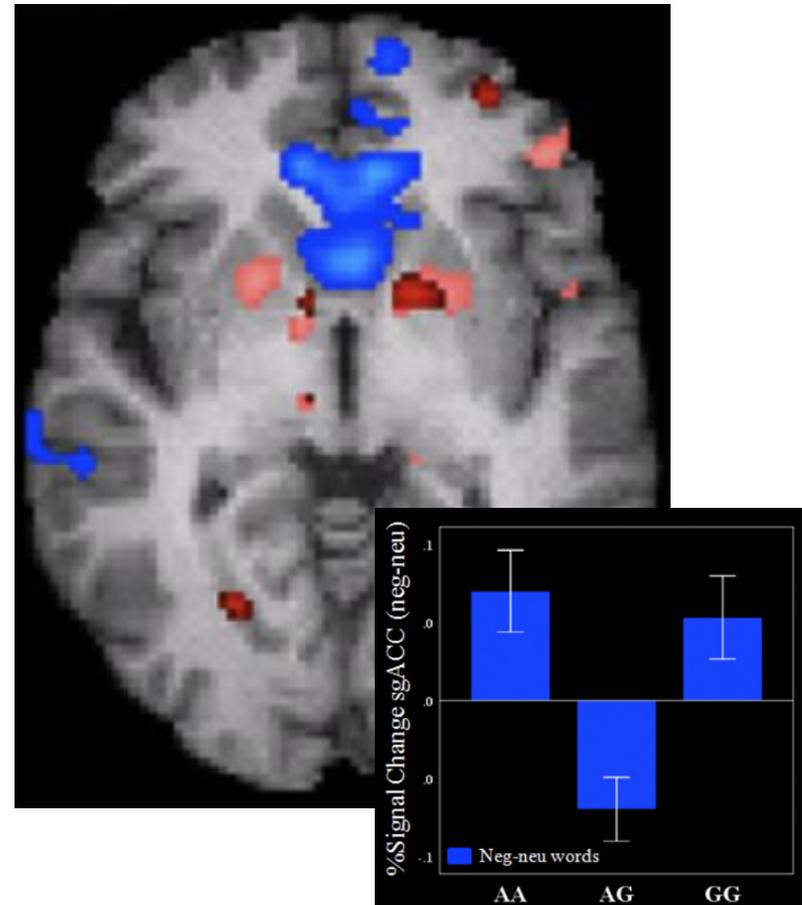


D2 dopamine receptor Processing of emotionally salient stimuli

DRD2 rs4274224
single nucleotide variant
- intronic, function unknown

Greater activation of pregenual
and subgenual anterior cingulate
cortex in GG and AA homozygotes

Similar genetic effects in
hemodynamic responses to
monetary reward and dopamine
release with pain



Pecina, Mickey, et al., *Cortex*, 2012

Overall summary

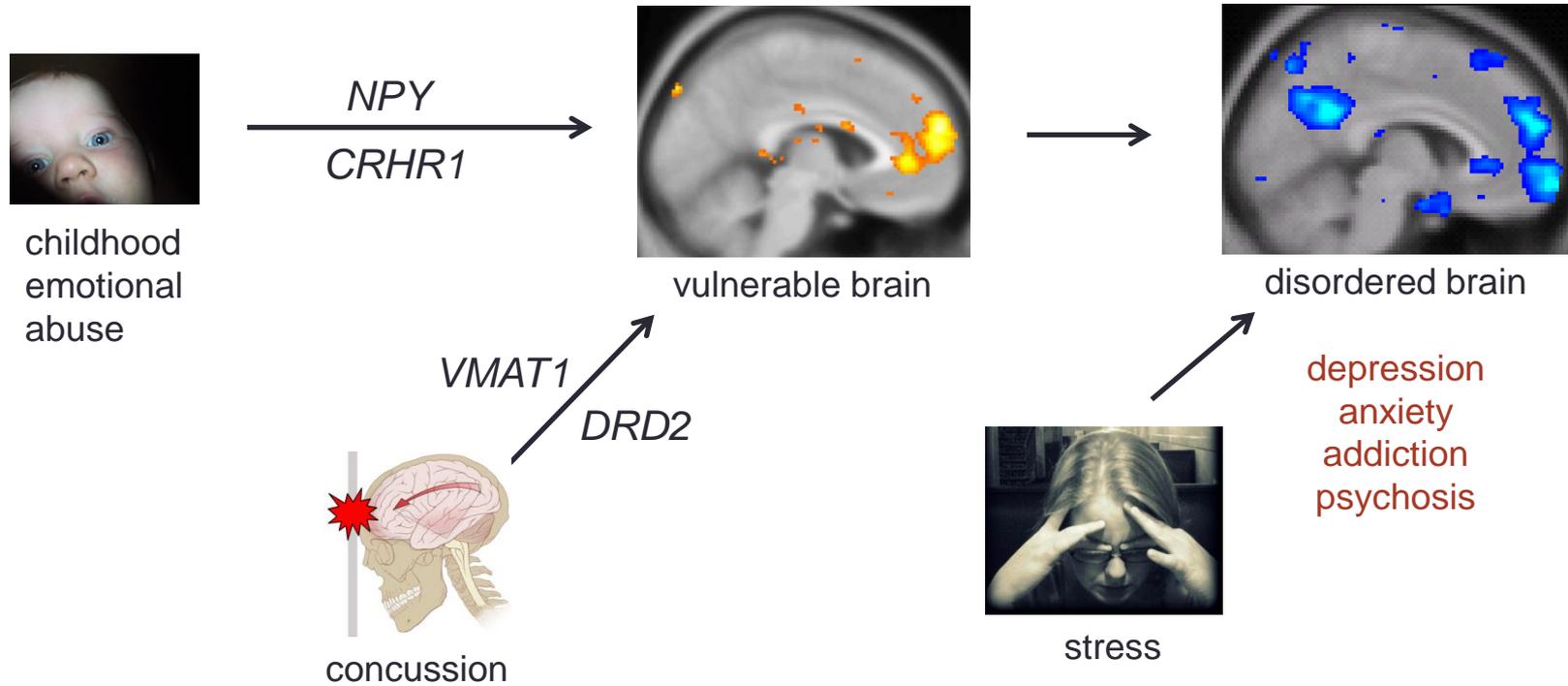
Common, putatively functional, genetic variants (*NPY*, *VMAT1*, *CRHR1*, *DRD2*) influence medial prefrontal and anterior cingulate responses to negative stimuli

Variation in stress- and emotion-sensitive circuitry across individuals may mediate risk for psychiatric disorders, and/or moderate response to specific treatments

These genetic causes of variation in brain function could be examined in experimental animals

Interactions of these genes with specific environmental factors (childhood adversity, recent stress) should be examined in humans and animal models

Speculative model framework



Acknowledgements



THE MOLECULAR & BEHAVIORAL
NEUROSCIENCE INSTITUTE
UNIVERSITY OF MICHIGAN



neuroscience graduate program

University of Michigan

Jon-Kar Zubieta

David Hsu

Marta Pecina

Tiffany Love

Mary Heitzeg

Tal Shafir

Heng Wang

Susan Kennedy

University of Pennsylvania

Falk Lohoff

NIAAA

David Goldman

Laura Bevilacqua

Zhifeng Zhou

Elizabeth Heinz

Pei-Hong Shen

Colin Hodgkinson

University of Illinois Chicago

Scott Langenecker

Sara Weisenbach