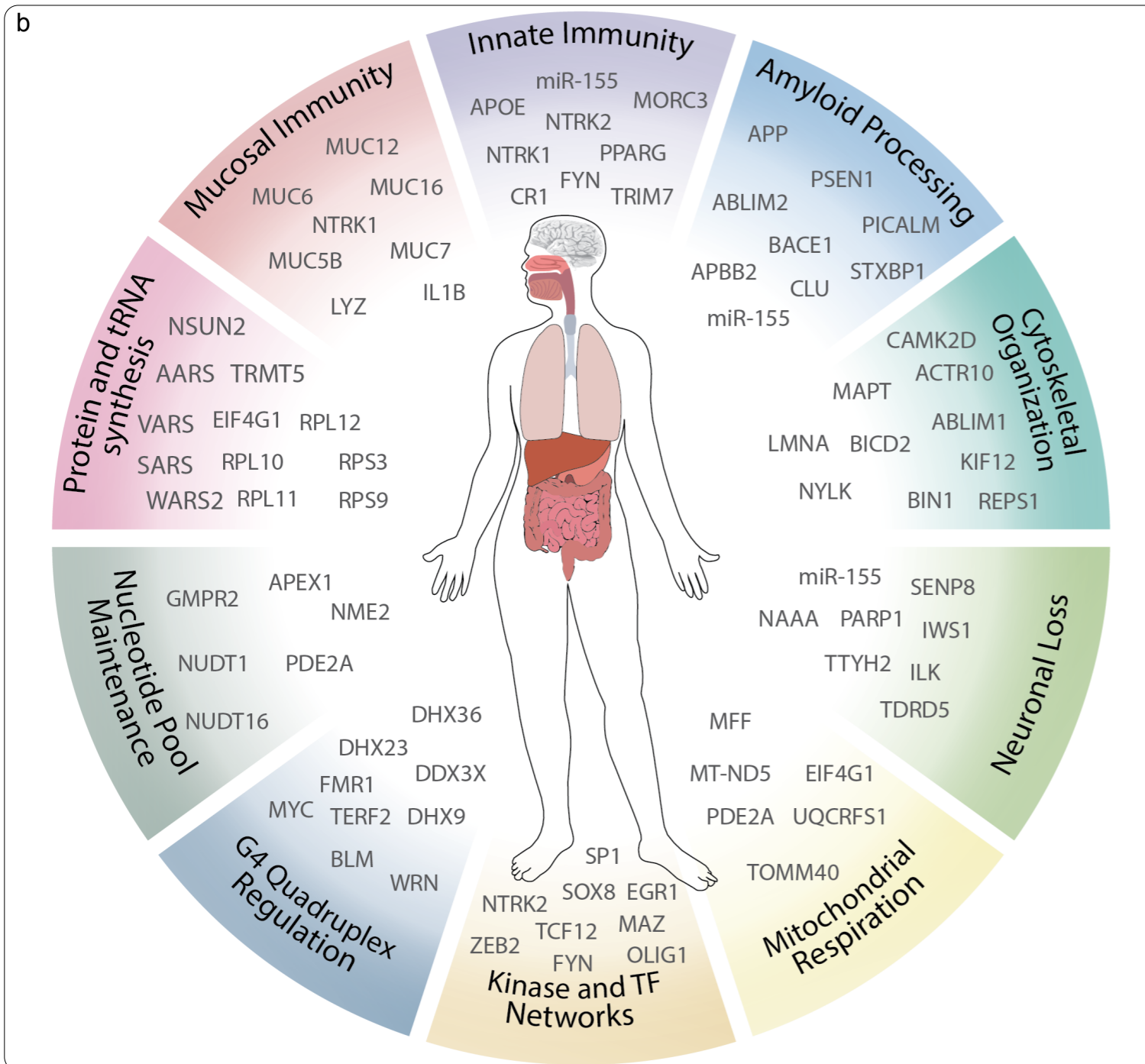
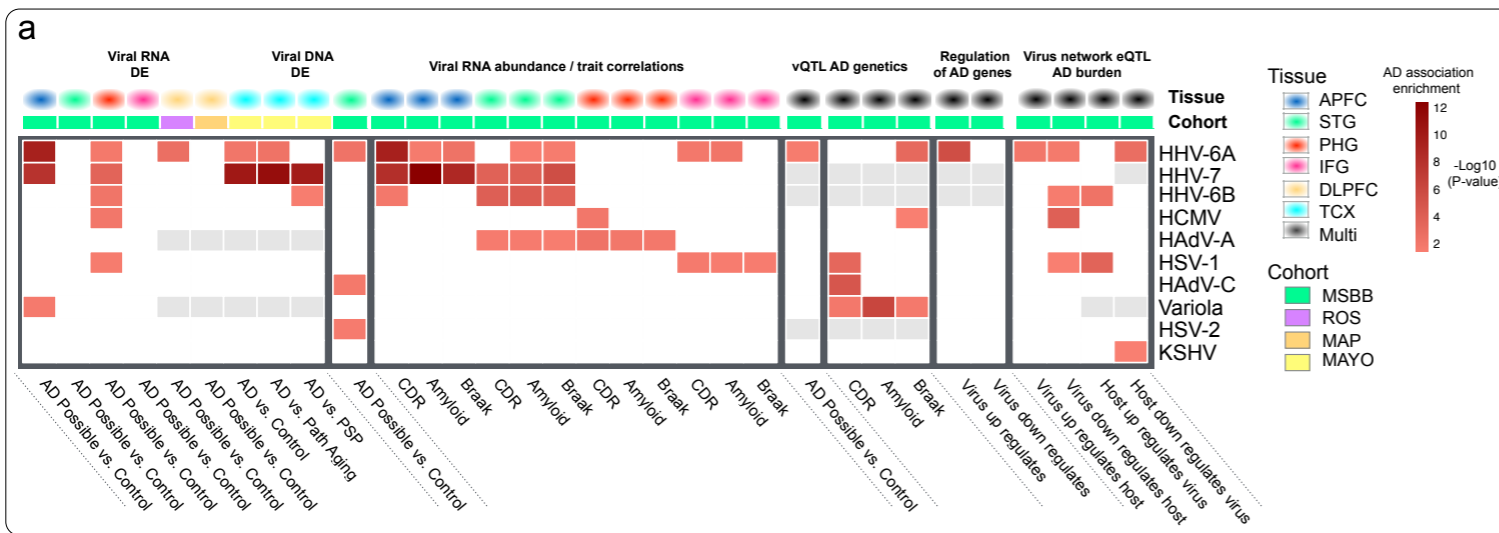


# Drugs and Bugs in Alzheimer's Disease

**Joel Dudley, PhD**

Director of the Institute for Next Generation Healthcare  
Associate Professor of Genetics and Genomic Sciences  
Mount Sinai Professor of Biomedical Data Science

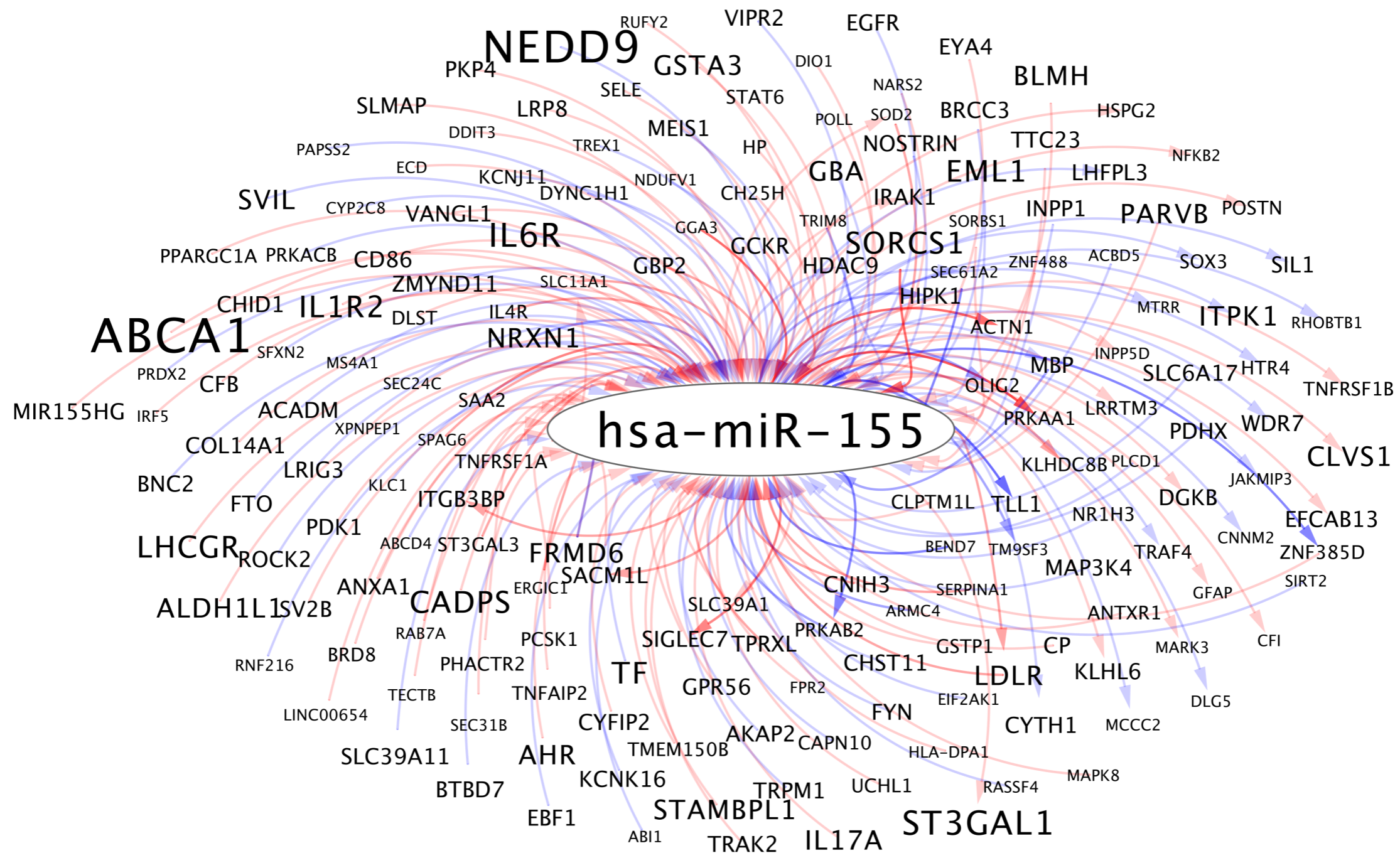




**Host-Virus interaction networks in AD identify numerous potential therapeutic targets related to immune activation**

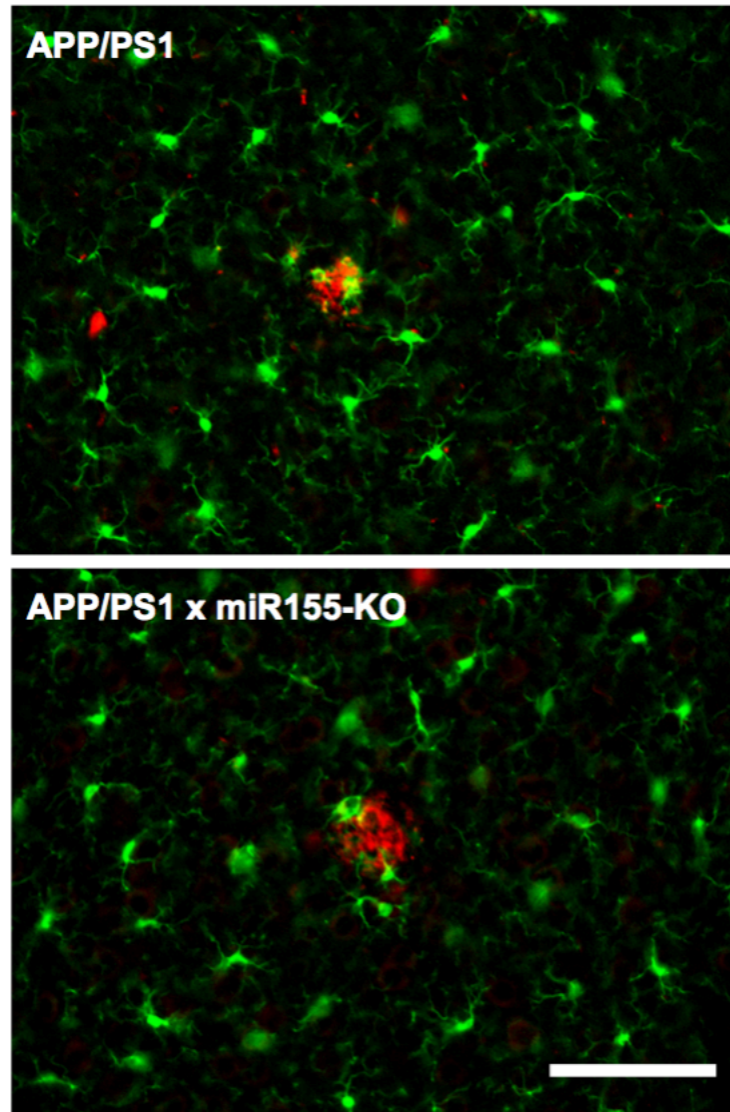
Readhead et al. *Neuron* 2018

# miR-155 is suppressed by HHV-6A, a regulator of preclinical and clinical AD networks

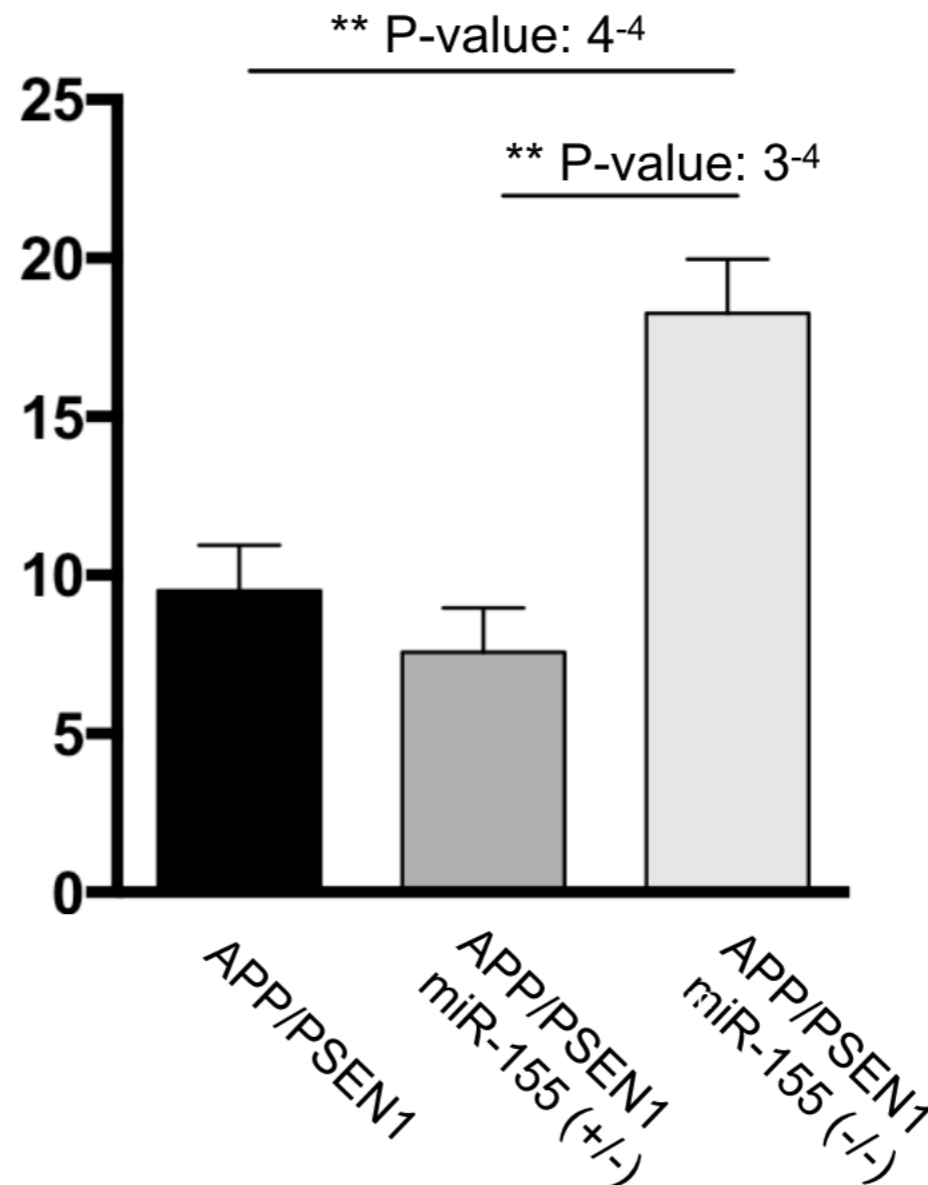


# miR-155 is suppressed by HHV-6A, a regulator of preclinical and clinical AD networks and alters $\beta$ -amyloid plaque and oligomer formation

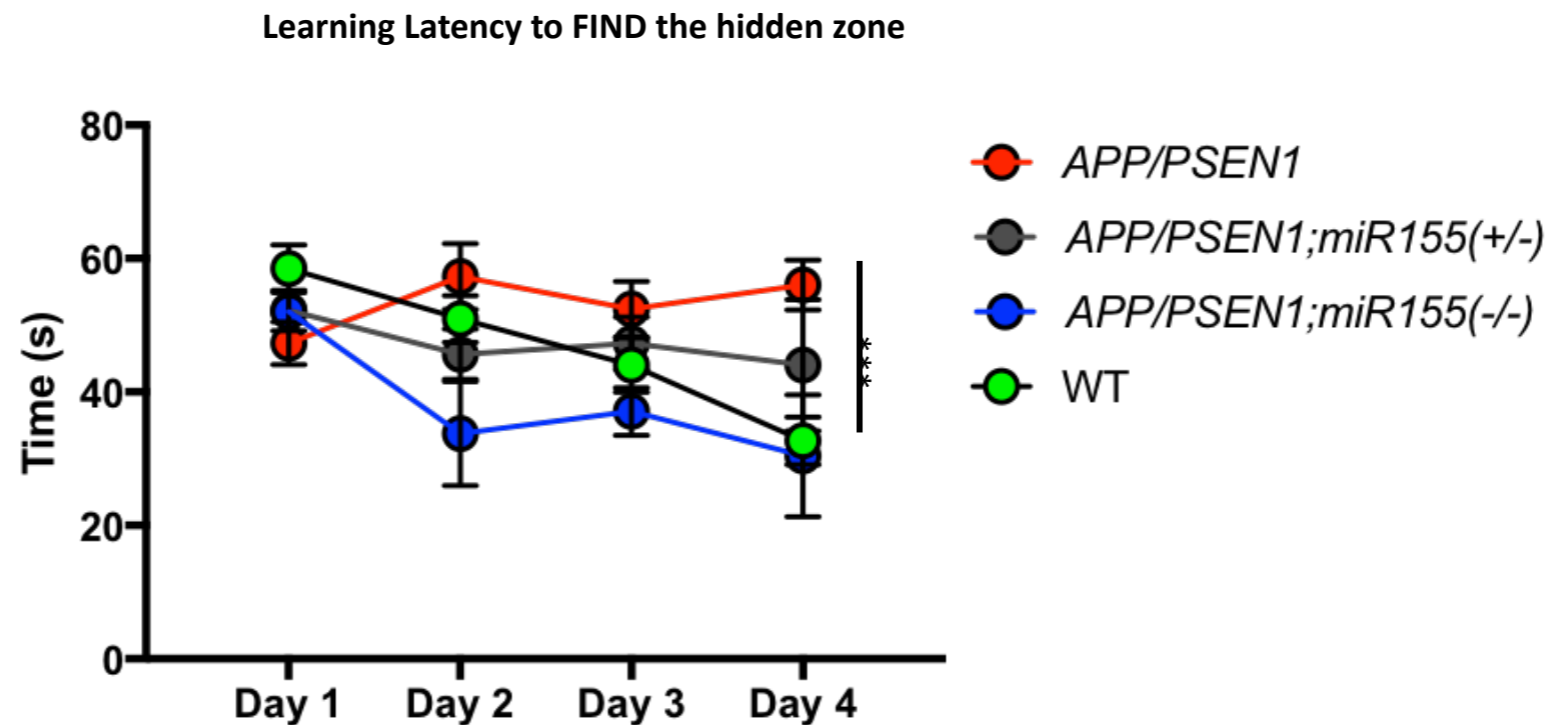
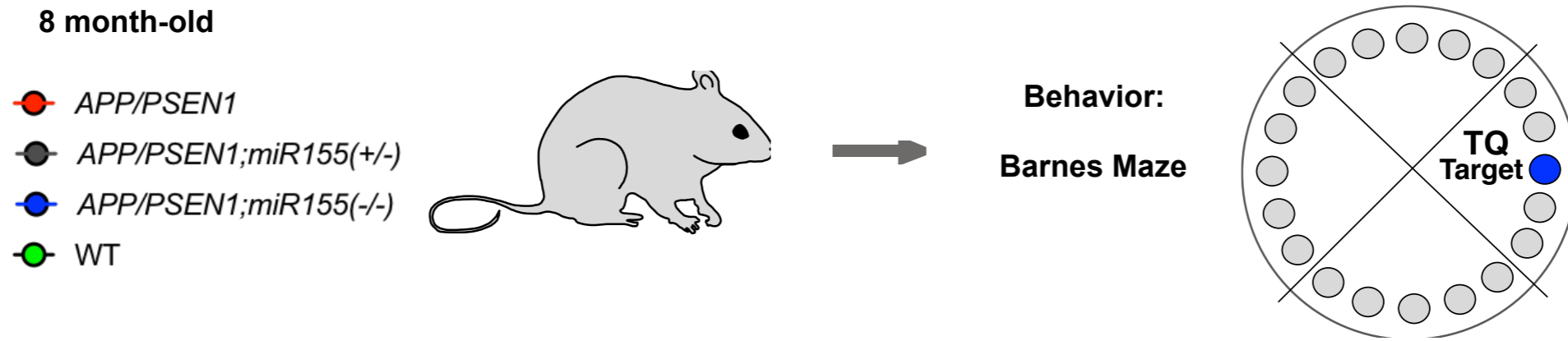
## APP/PS1 x MIR155-KO vs. APP/PS1 Frontal Cortex



### # $\beta$ -amyloid plaques

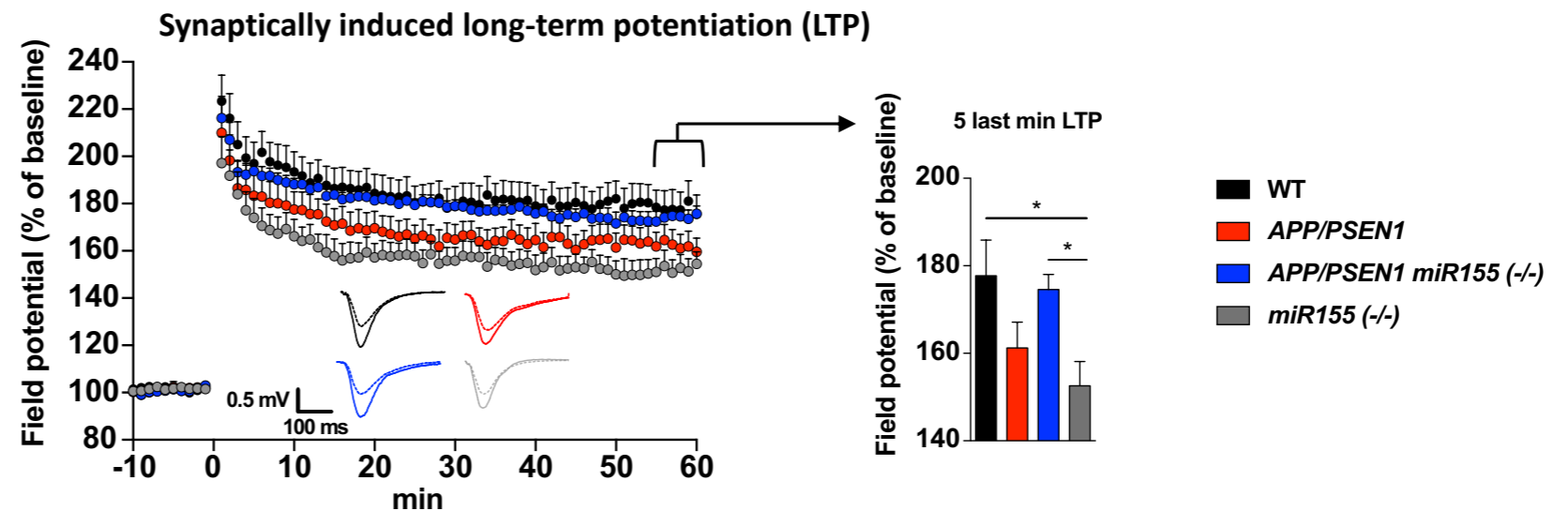
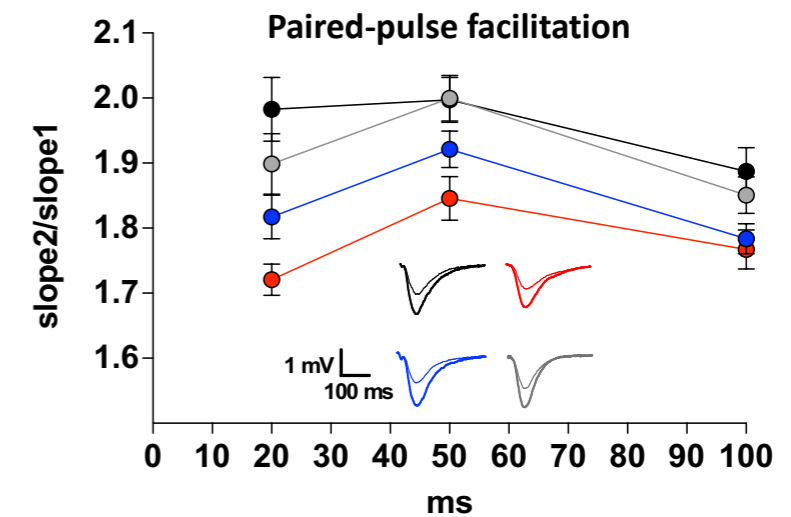
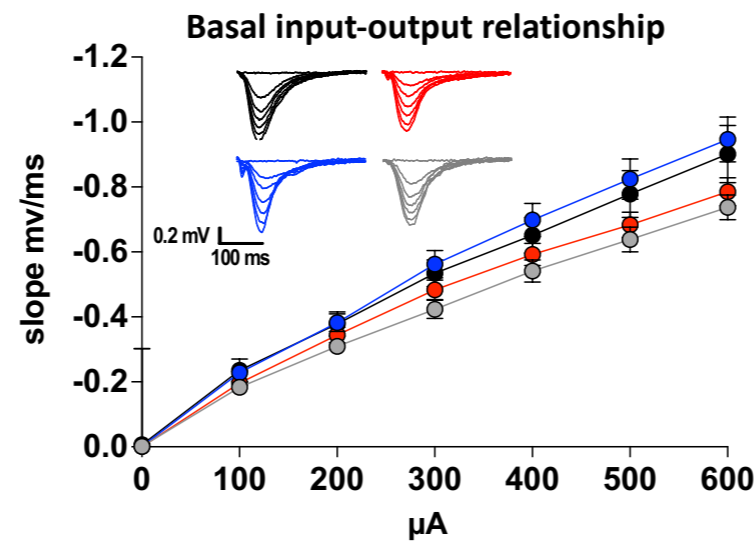
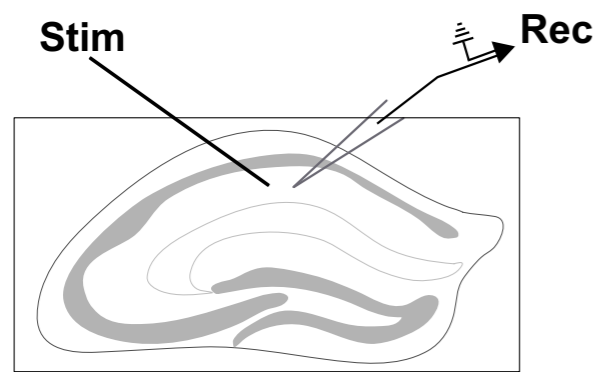


# Constitutive absence of *miR-155* in *APP/PSEN1* mice improves learning and memory performances in Barnes maze test



Unpublished work in progress

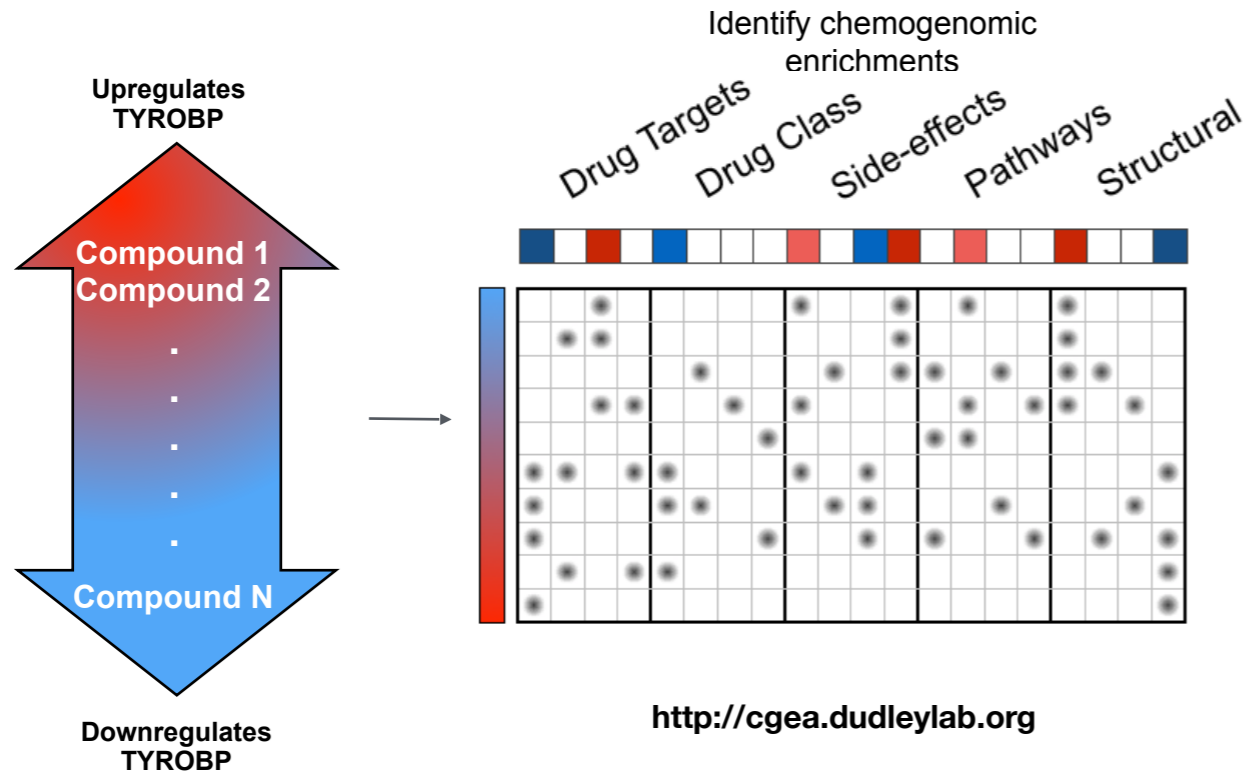
# Absence of mir-155 mice ameliorates synaptic plasticity defects in APP/PSEN1 mice But induces defect of synaptic plasticity in WT mice



Unpublished work in progress

# TYROBP repurposing use case

a



bi

Top CMAP compounds predicted to upregulate TYROBP

compound	ATC Level 3
thioguanosine	ANTIMETABOLITES
clorgiline	
methapyrilene	ANTIHISTAMINES FOR SYSTEMIC USE
estradiol	ESTROGENS
procainamide	ANTIARRHYTHMICS, CLASS I AND III
apigenin	
atropine	BELLADONNA AND DERIVATIVES, PLAIN
minaprine	ANTIDEPRESSANTS
clemizole	
salsolinol	
luteolin	
moxisylyte	PERIPHERAL VASODILATORS
alfuzosin	DRUGS USED IN BENIGN PROSTATIC HYPERTROPHY
tranylcypromine	ANTIDEPRESSANTS
vinpocetine	PSYCHOSTIMULANTS, AGENTS USED FOR ADHD AND NOOTROPICS

bii

Top CMAP compounds predicted to downregulate TYROBP

compound	ATC Level 3
myosmine	
imatinib	OTHER ANTINEOPLASTIC AGENTS
benzathine benzylpenicillin	BETA-LACTAM ANTIBACTERIALS, PENICILLINS
diphenylpyraline	ANTIHISTAMINES FOR SYSTEMIC USE
SC-58125	
lasalocid	
methyldopate	ANTIADRENERGIC AGENTS, CENTRALLY ACTING
harpagoside	
ketotifen	ANTIHISTAMINES FOR SYSTEMIC USE
cloxacillin	BETA-LACTAM ANTIBACTERIALS, PENICILLINS

c

Drug targets enriched in compounds that regulate TYROBP

Predicted Targets	Name
CHRND	cholinergic receptor, nicotinic, delta (muscle)
RIPK1	receptor (TNFRSF)-interacting serine-threonine kinase 1
KCNN2	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2
KYNU	kynureninase
CCR8	chemokine (C-C motif) receptor 8
CHKA	choline kinase alpha
ENPEP	glutamyl aminopeptidase (aminopeptidase A)
SELP	selectin P (granule membrane protein 140kDa, antigen CD62)
ADRA1B	adrenoceptor alpha 1B
CTSD	cathepsin D
AOC3	amine oxidase, copper containing 3

Unpublished work in progress

# Compounds that modulate TYROBP are enriched for drug targets that link to AD

Symbol	Name	Notes
CHRND	cholinergic receptor, nicotinic, delta (muscle)	
RIPK1	receptor (TNFRSF)-interacting serine-threonine kinase 1	Mediates AB induced TNF production by microglia in vitro (Zhou, 2014 for review)
KCNN2	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2	
KYNU	kynureninase	
CCR8	chemokine (C-C motif) receptor 8	Monocyte chemotaxis and localization of activated T-cells
CHKA	choline kinase alpha	
ENPEP	glutamyl aminopeptidase (aminopeptidase A)	
SELP	selectin P (granule membrane protein 140kDa, antigen CD62)	
ADRA1B	adrenoceptor alpha 1B	
CTSD	cathepsin D	Risk gene for AD (Schuur, 2011)
AOC3	amine oxidase, copper containing 3	

\* Targets enriched (FDR < 0.1) in CMAP compounds ranked according to TYROBP expression

Unpublished work in progress

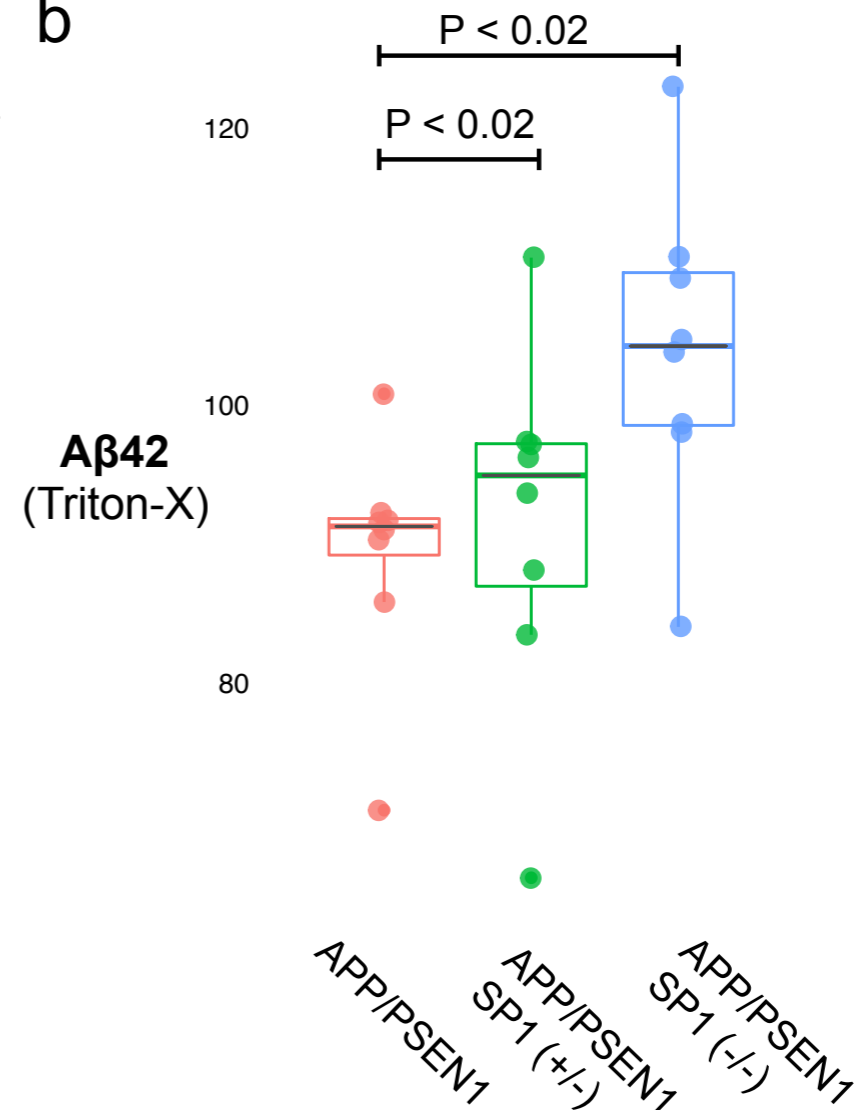


# APP/PSEN1 x SP1-KO vs. APP/PSEN1 RNA-sequencing Frontal Cortex

## a Differential gene expression

Symbol	Name	logFC	FDR
Amhr2	anti-Mullerian hormone type 2 receptor	3.60	3.52e-08
Sp1	trans-acting transcription factor 1	-0.90	4.25e-08
Pfn1	profilin 1	0.42	2.84e-03
Ppp1r9b	protein phosphatase 1, regulatory subunit 9B	0.26	5.02e-03
Mrps34	mitochondrial ribosomal protein S34	0.44	8.35e-03
Ptms	parathymosin	0.40	1.05e-02
Hist2h4	histone cluster 2, H4	0.56	1.05e-02
Hepacam	hepatocyte cell adhesion molecule	0.53	1.05e-02
Hnrnpa0	heterogeneous nuclear ribonucleoprotein A0	0.31	1.05e-02
Hnrnp3	heterogeneous nuclear ribonucleoprotein H3	0.74	1.05e-02

## b



## c Causal Inference Testing (CIT)

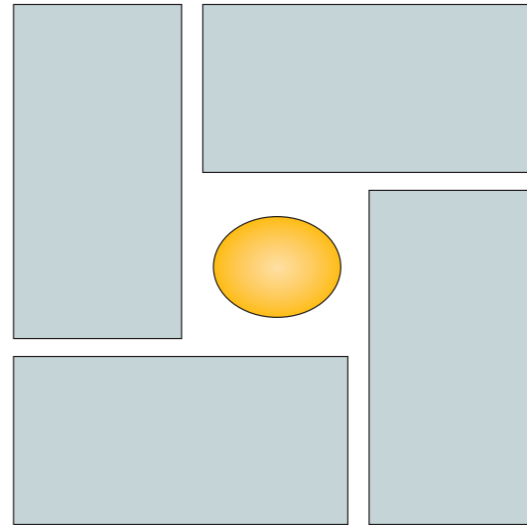
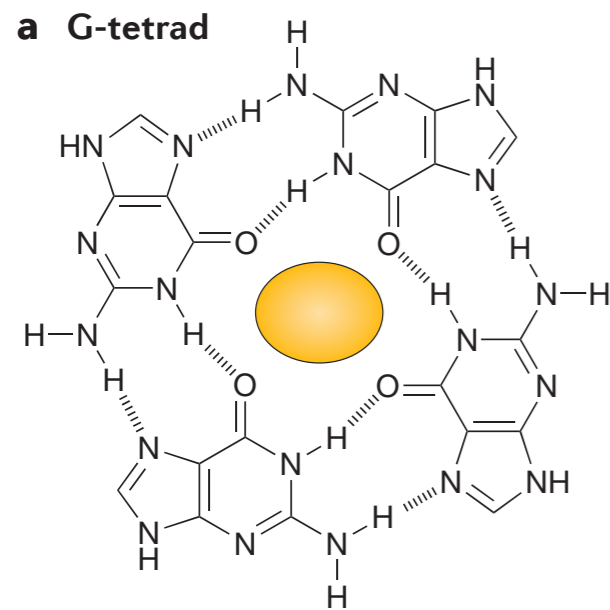


- SP1 and Aβ42 are associated
- SP1 and Aβ42-genes are associated
- SP1 associated with Aβ42-genes | Aβ42
- SP1 independent of Aβ42 | Aβ42-genes

## d

Symbol	Name	CIT (FDR)	Cor with SP1 dosage	Cor with Aβ42
Ppp1r9b	protein phosphatase 1, regulatory subunit 9B	9.99e-04	-0.71	0.71
Lrrk2	leucine-rich repeat kinase 2	9.99e-04	0.69	-0.67
Mettl7a1	methyltransferase like 7A1	9.99e-04	0.63	-0.76
Slc25a25	solute carrier family 25 (mitochondrial carrier, phosphate carrier), member 25	9.99e-04	0.73	-0.64
Edrf1	erythroid differentiation regulatory factor 1	9.99e-04	0.63	-0.80
Chd3	chromodomain helicase DNA binding protein 3	9.99e-04	-0.63	0.59
A230050P20Rik	RIKEN cDNA A230050P20 gene	9.99e-04	-0.71	0.58
Lgals9	lectin, galactose binding, soluble 9	9.99e-04	-0.71	0.58
Nbas	neuroblastoma amplified sequence	2.00e-03	0.72	-0.68
Tmem191c	transmembrane protein 191C	2.00e-03	-0.78	0.64

# G-quadruplex secondary structures associate with normal gene regulatory activity and viral replication

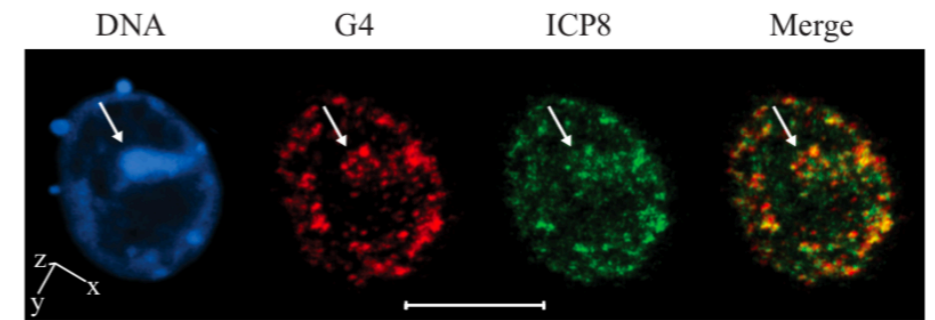
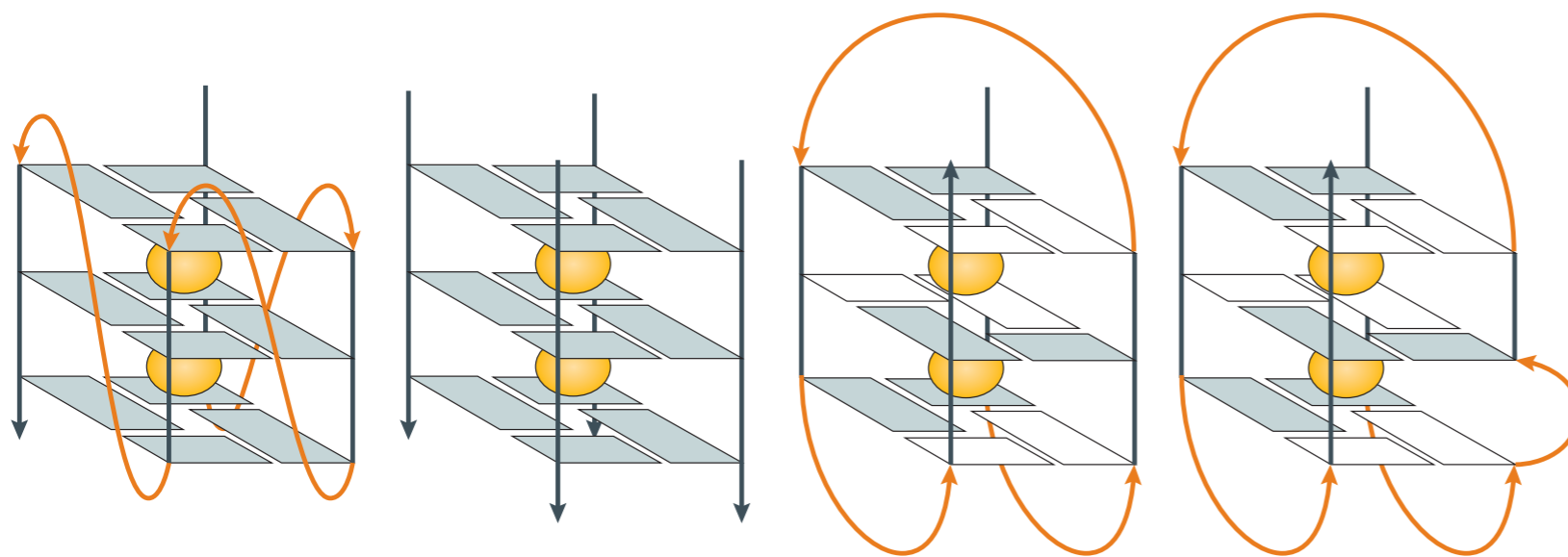


**b Unimolecular parallel**

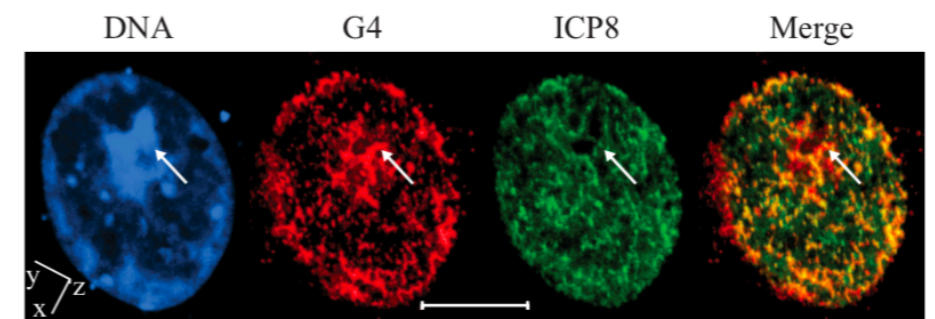
**c Tetramolecular parallel**

**d Unimolecular antiparallel**

**e Unimolecular bulged**



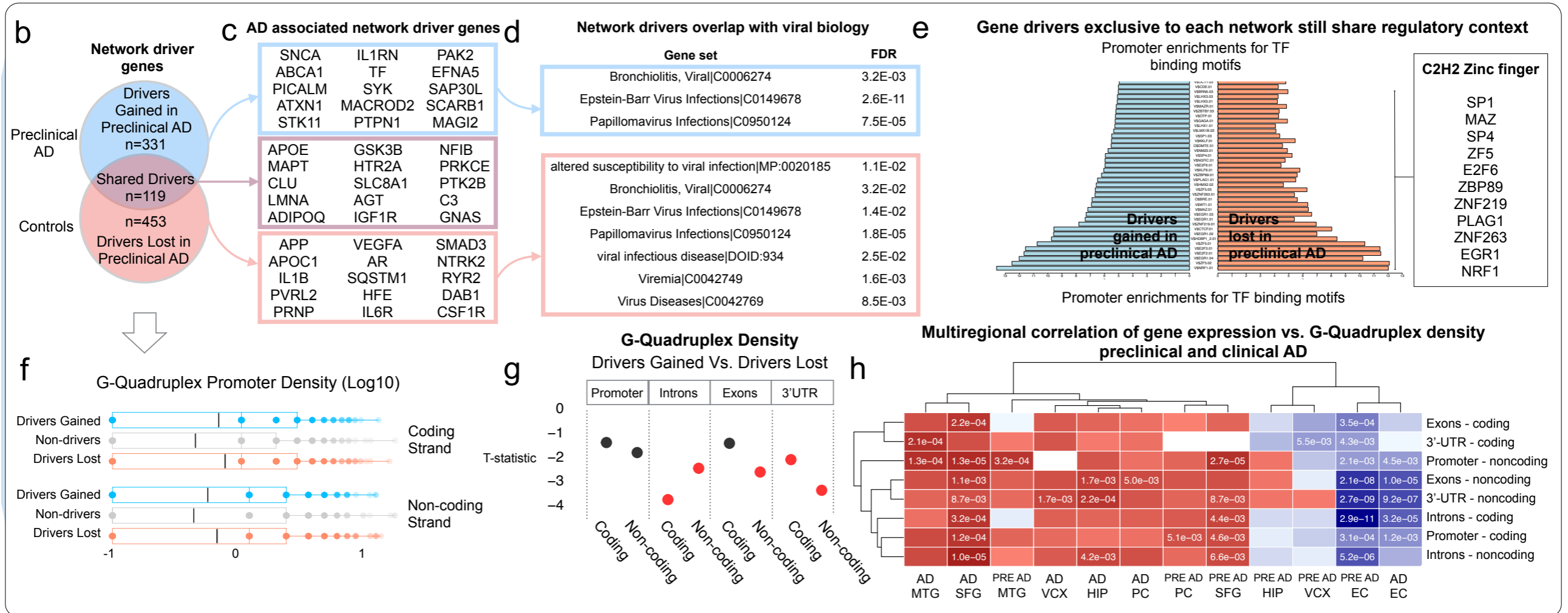
MOI 2.5



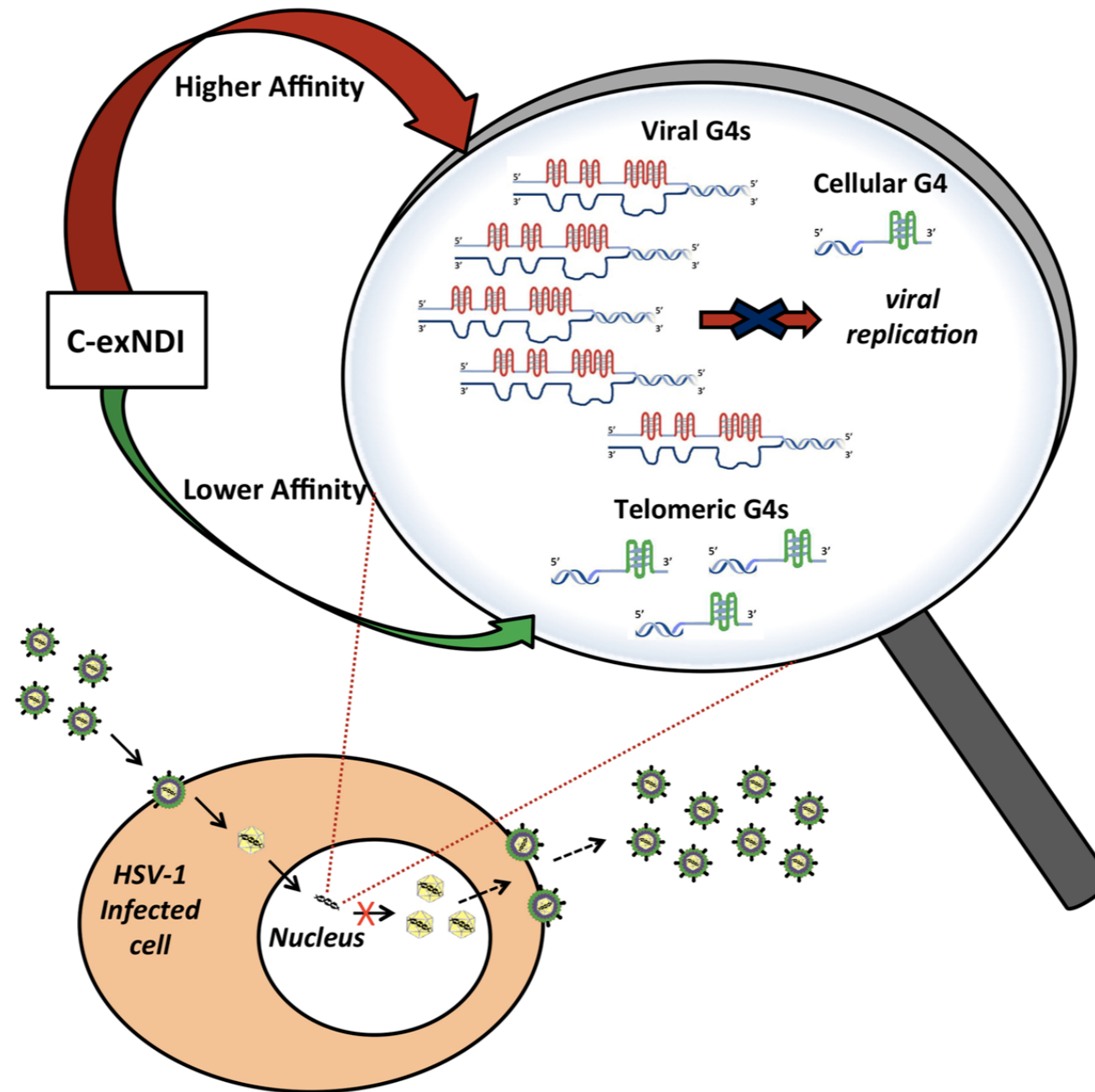
MOI 5

Artusi et al. *Nucleic Acids Research*, 2016, Vol. 44, No. 21

# Association between changes in G-quadruplex promoter density and network driver gain/loss

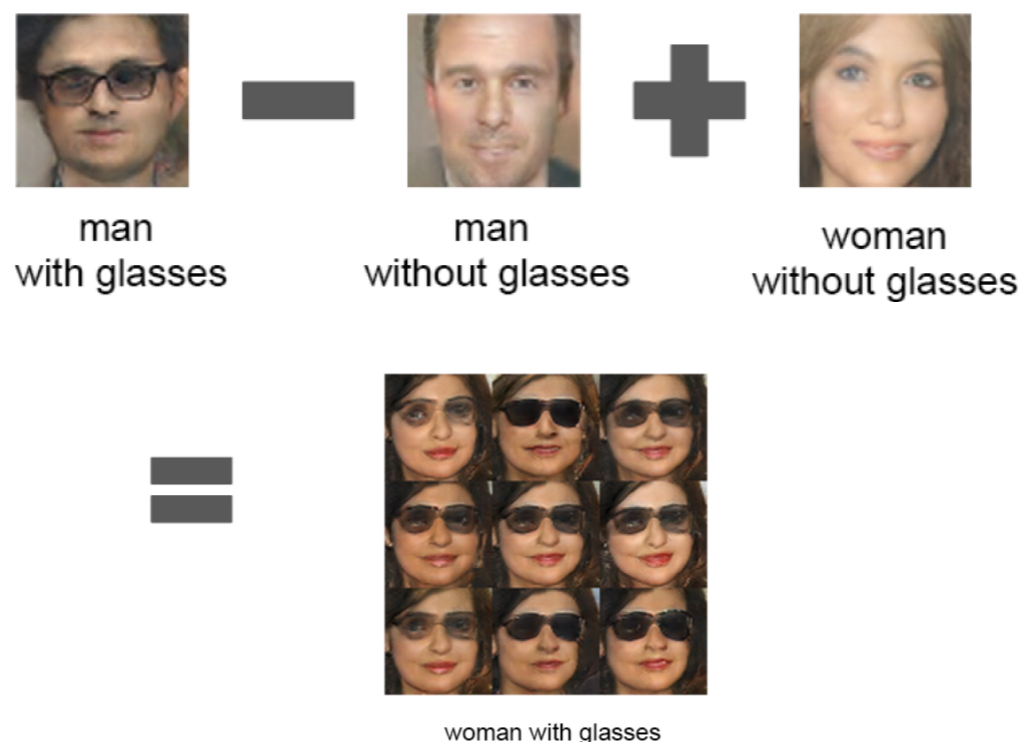


# A core extended naphthalene diimide G-quadruplex ligand potently inhibits herpes simplex virus 1 replication



**Figure 5.** Scheme of the proposed c-exNDI mechanism of anti-HSV-1 activity.

# Defining the “opposite phenotype” using Generative Adversarial Networks (GAN)

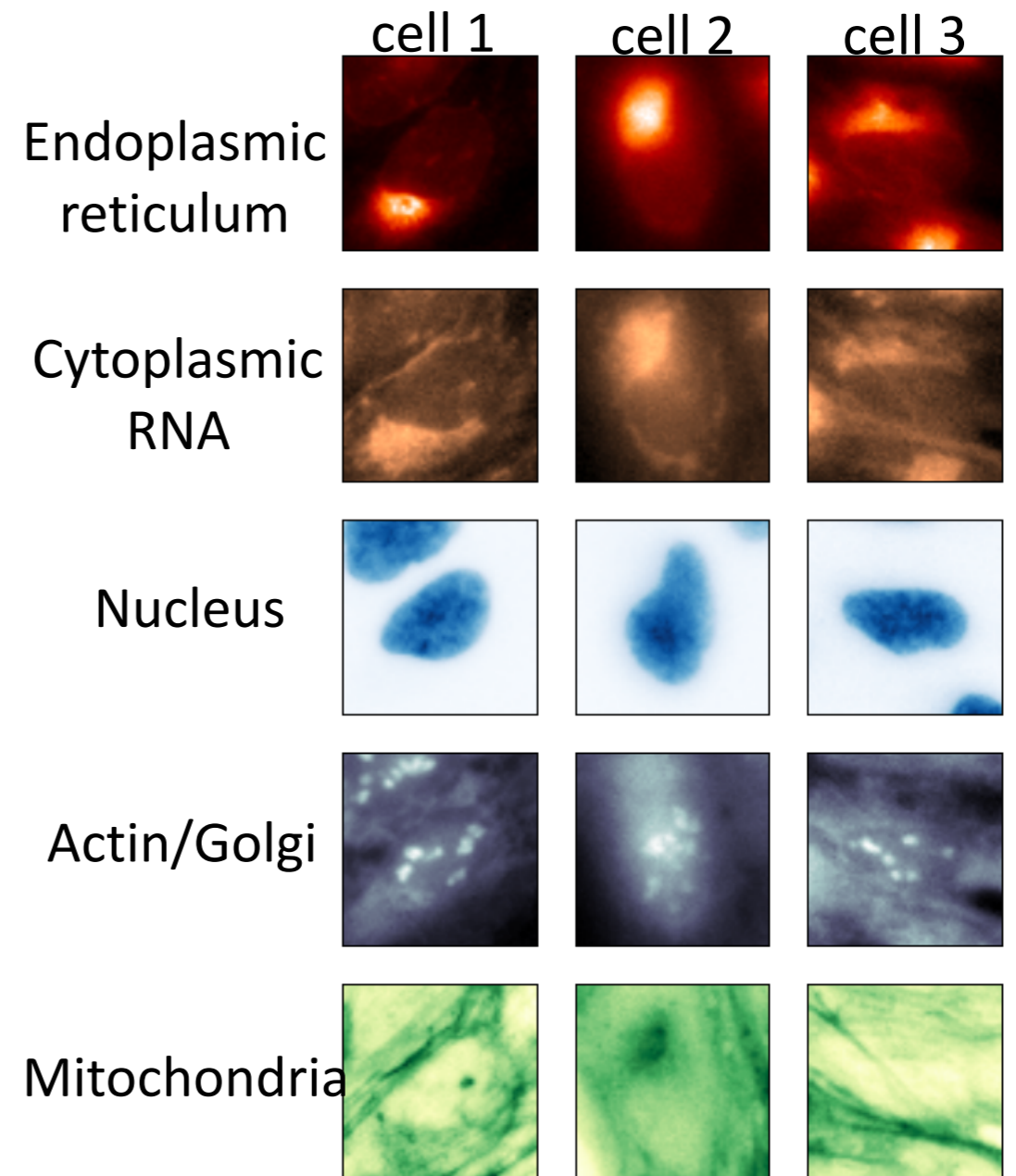


## Rationale for drug discovery:

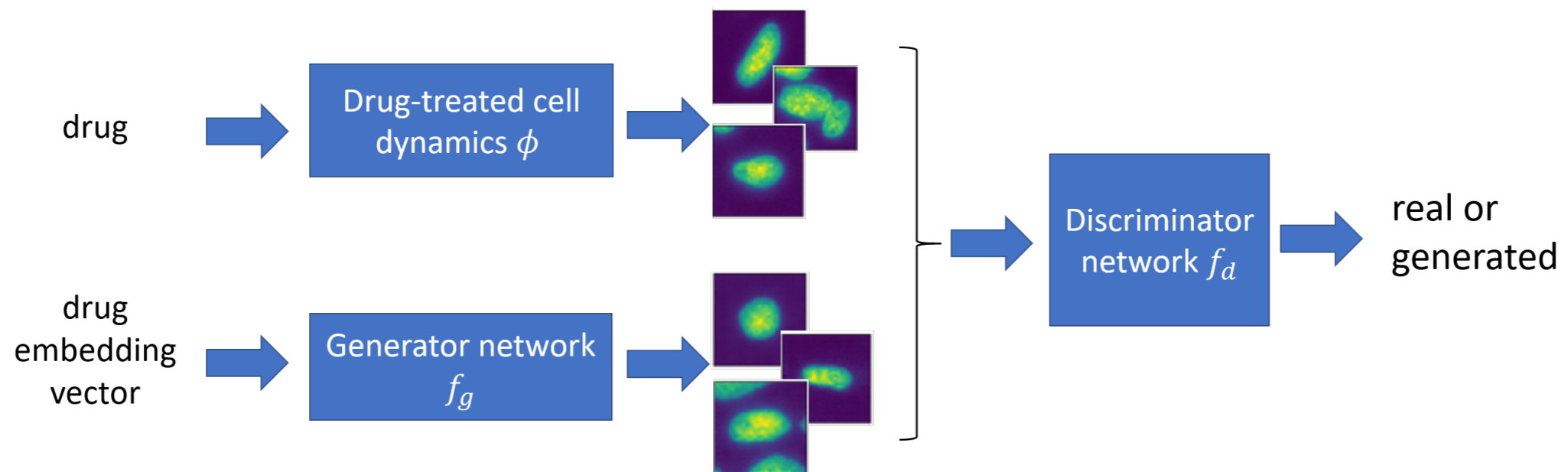
$$v_{without\ phenotype} - v_{with\ phenotype} \approx \text{action reversing the phenotype}$$

# Data source: drug-treated image “patches”

- On average 4 repeats per treatment
- 6 images form each repeat
- 10~50 cells from 1 images
- 2 TB of image patches (64x64)



# Using a conditional GAN to model high-throughput imaging data

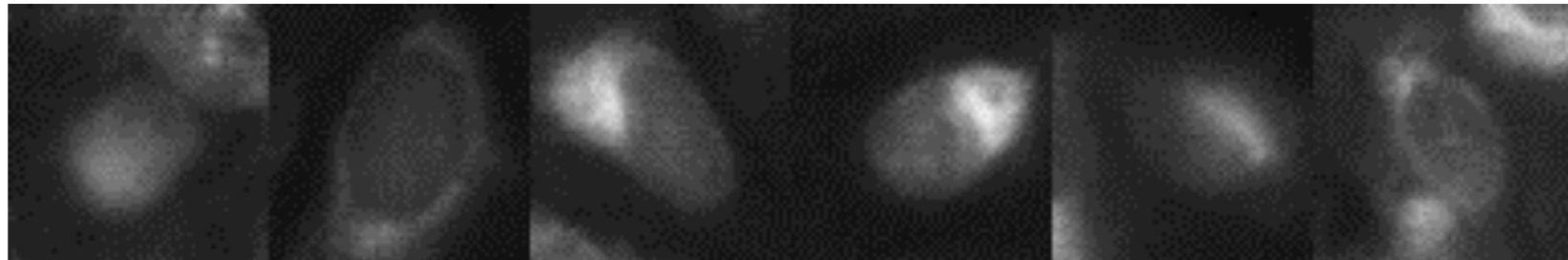


**Rationale for characterizing drug treated cell dynamics:**

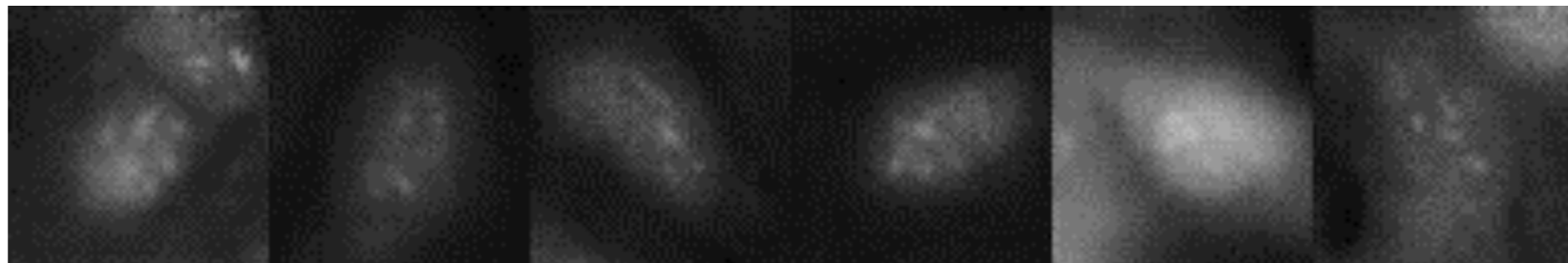
$$\phi \approx f_g \text{ if } f_d \text{ cannot tell the difference}$$

# Cellular phenotypes generated by GAN

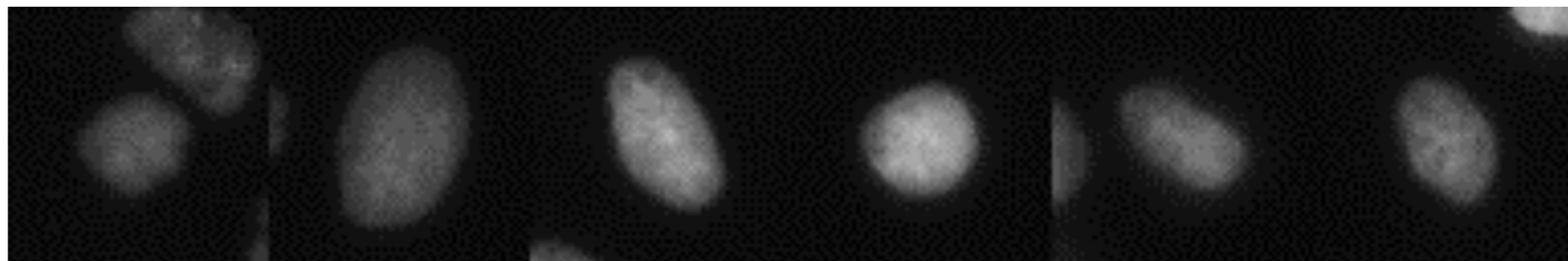
Endoplasmic  
reticulum



Cytoplasmic  
RNA



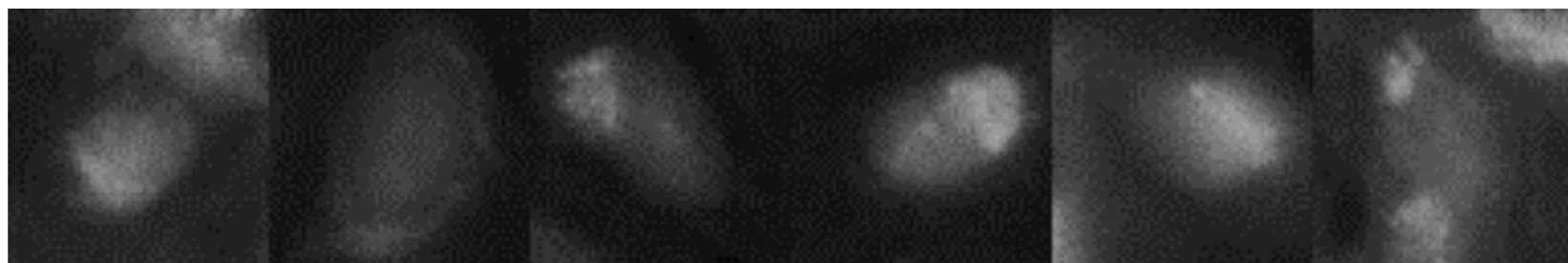
Nucleus



Actin/Golgi



Mitochondria



25  
epochs



# Thank you for your attention!

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