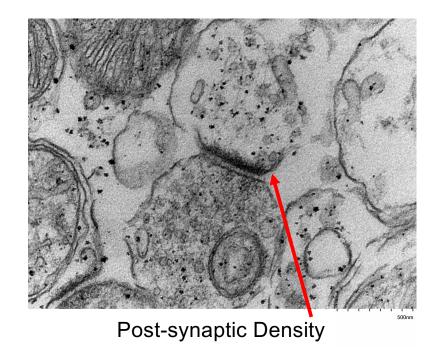
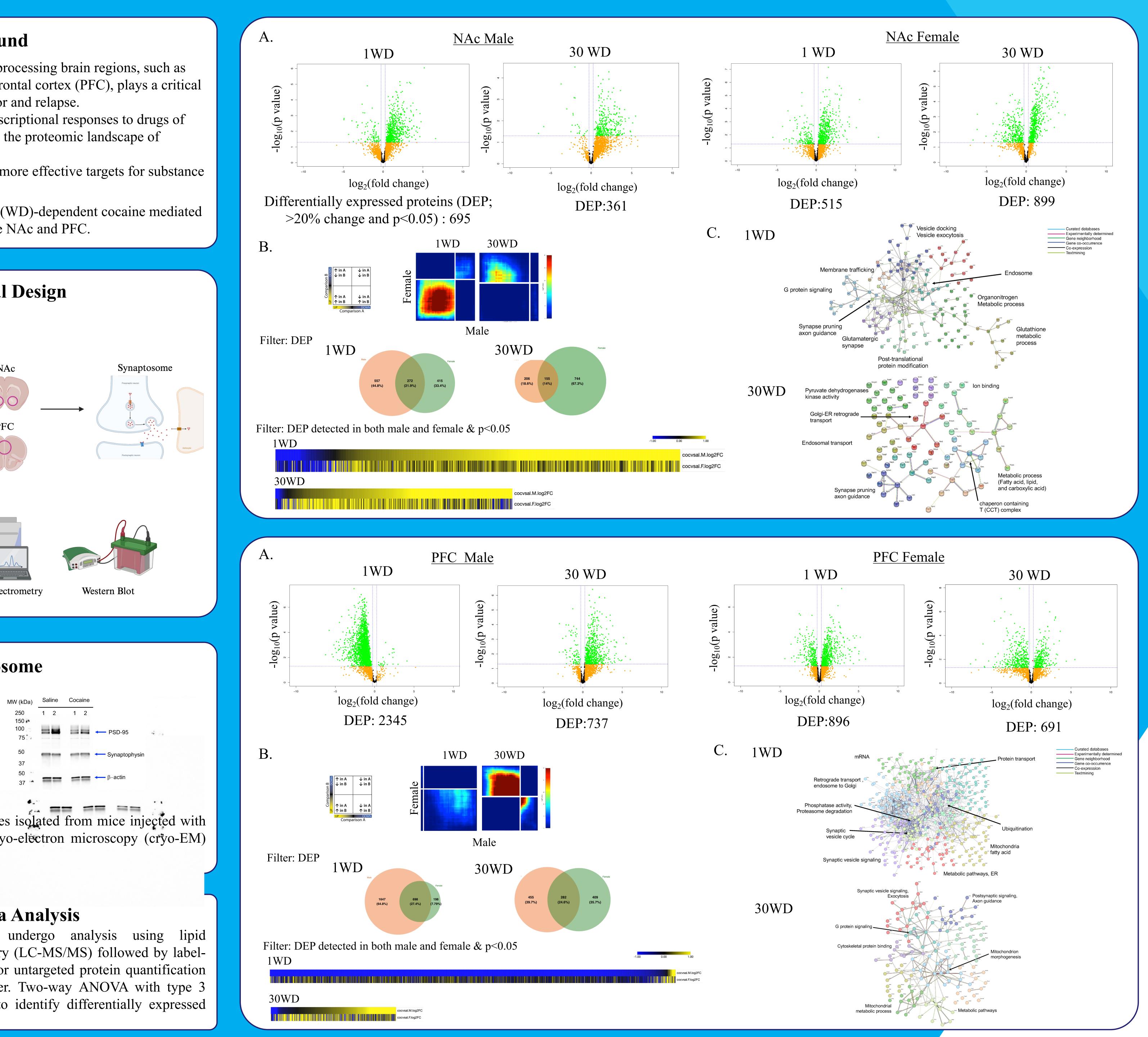
# **Sex- and Withdrawal-Dependent Proteomic Changes in Nucleus Accumbens and Prefrontal Cortex: Insights into Synaptic Adaptations in Substance Use Disorders**



Background		
<ul> <li>Dysregulated signaling within reward-processing brain regions, such as the nucleus accumbens (NAc) and prefrontal cortex (PFC), plays a critical role in promoting drug-seeking behavior and relapse.</li> <li>Compared to our understanding of transcriptional responses to drugs of abuse, our knowledge about changes in the proteomic landscape of synapses is limited.</li> <li>Identifying these changes could reveal more effective targets for substance use disorder (SUD) treatment.</li> </ul>		
• <u>Goal</u> : Identify the sex- and withdrawal (WD)-dependent cocaine mediate changes in the synaptic proteome of the NAc and PFC.		
<b>Experimental Design</b>		
A. Sample Preparation		
$C57BL/6J \qquad \qquad$	NAc OOOOO FFC	<image/>
B. Sample Analysis		
Electron Microscopy (EM)	Image: Addition of the sector of the secto	<image/>

## Synaptosome





The quality of NAc and PFC synaptosomes isolated from mice injected with saline and cocaine are assessed using cryo-electron microscopy (cryo-EM) and Western blot analysis.

### **Proteomic Data Analysis**

The NAc and PFC synaptosomes undergo analysis using lipid chromatography-tandem mass spectrometry (LC-MS/MS) followed by labelfree, data-dependent acquisition (DDA) for untargeted protein quantification at the Yale/NIDA neuroproteomics Center. Two-way ANOVA with type 3 sums of squares is conducted using R to identify differentially expressed proteins.

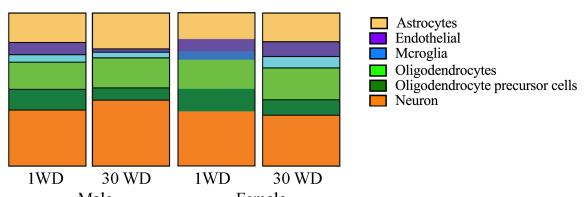
Yun Young Yim<sup>1</sup>, Arthur Godino<sup>1</sup>, Corrine Azizian<sup>1</sup>, Rita Futamura<sup>1</sup>, TuKiet Lam<sup>2</sup>, and Eric J.Nestler<sup>1</sup>

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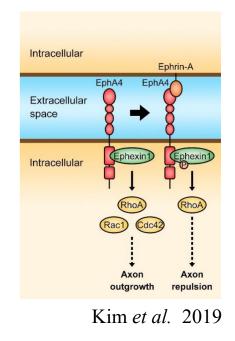


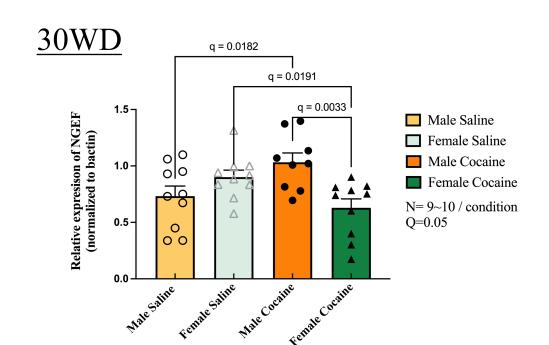
# **Conclusions & Future Directions**

- We conducted a whole-proteome analysis of synapses following the 7days injection model.
- We identified multiple synapse-enriched proteins that were either induced or repressed in a brain region-, sex-, and WD-timedependent manner.
- While most of these cocaine-regulated proteins are expressed by neurons, a significant subset is enriched in astrocytes or microglia, consistent with the involvement of these cell types in synaptic processes



- Interestingly, in the NAc after 30 days of WD, female mice exhibited  $\sim 2.5$  times more significant proteome changes than male mice. Conversely, in the PFC after 24 hours of WD, male mice displayed ~2.5 times more significant proteome changes, primarily repression, compared to female mice.
- We are currently validating these findings and characterizing particular synaptic proteins in NAc.





• Future goal is to identify novel synaptic protein targets regulated by drugs of abuse, such as cocaine and heroine, in a cell-type and circuit-specific manner, to better understand and develop treatment.

# Acknowledgements

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Data were analyzed through the use of R, GraphPad Prism (version 10.0.0 for Mac, GraphPad Software, Boston, Massachusetts USA, www.graphpad.com), and STRING database.

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