Dissociating the signaling mechanisms underlying addiction vulnerability from the consequences of drug use Stephanie M. Groman, Becky Carlyle, Rashaun Wilson, Angus Nairn, and Jane R. Taylor Department of Psychiatry, Yale University

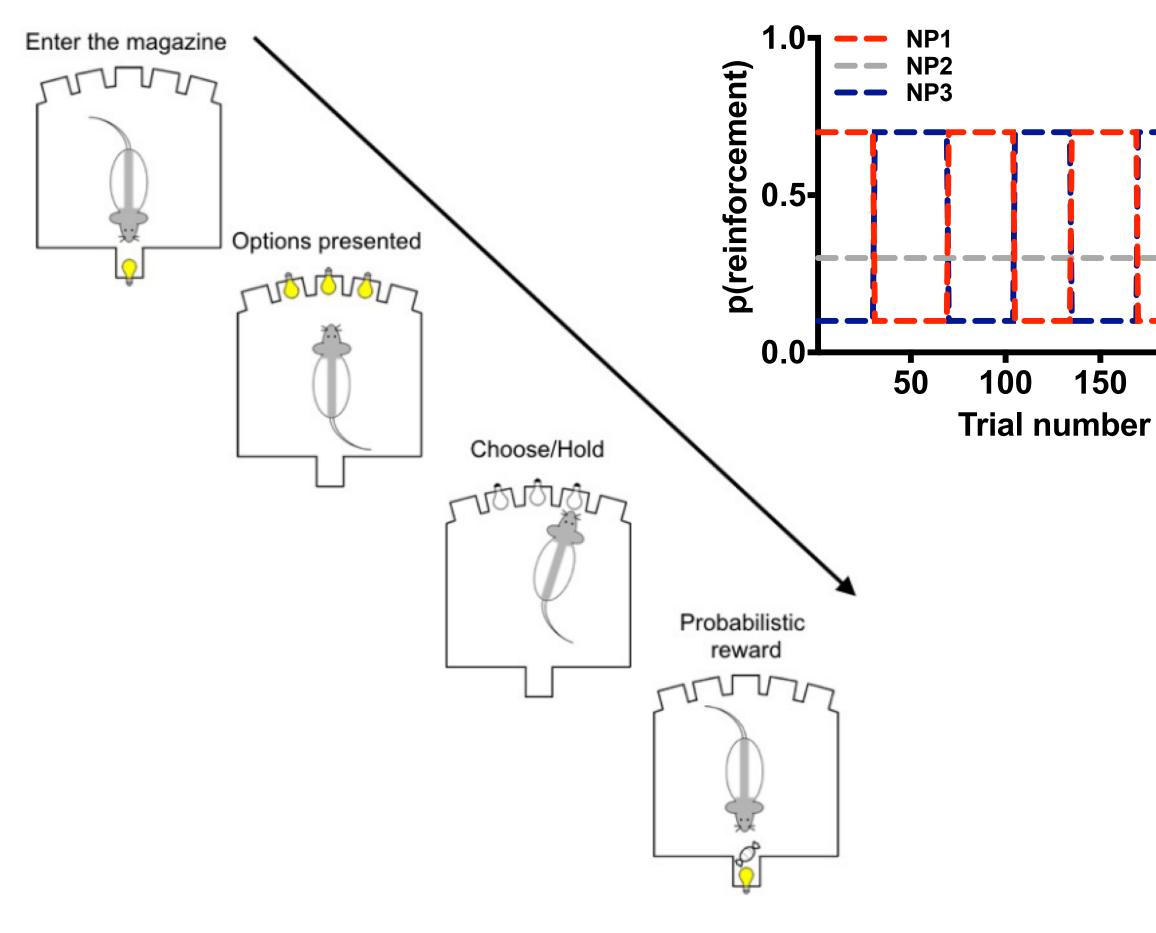
INTRODUCTION

Methamphetamine self-administration Adaptive, flexible decision-making is disrupted in addicted After PRL testing, rats (N=40) were implanted with intra-jugular catheters and individuals and believed, in part, to be a consequence of trained to self-administer methamphetamine (0.05 mg/kg/infusion) or saline in chronic drug use. Recent studies, however, have suggested 6 h long-access sessions for 14 days. that pre-existing alterations in decision-making might reversals d in PRL influence future drug-taking behaviors. Decision-making may, therefore, be a critical biomarker for understanding the neural mechanisms of addiction. Here, we investigated in rats the role of decision-making in methamphetamine selflun. administration to isolate the proteins involved in addiction Saline Math susceptibility from those involved in addiction consequence.

METHODS

Probabilistic reversal learning task

Adult, male Long-Evans rats (N=80) were trained on a three-choice, probabilistic reversal-learning (PRL) task. Reinforcement probabilities for each noseport were assigned at the beginning of each session. These probabilities remained stable until rats met a performance criterion (24 correct in last 30 trials completed) at which point the probabilities between two choices reversed and remained stable until the performance criterion was met again. Rats could complete up to 8 reversals each session



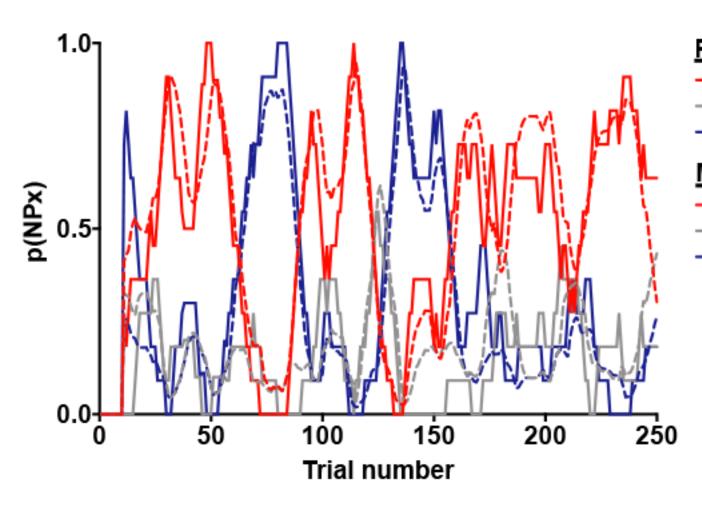
Computational analysis

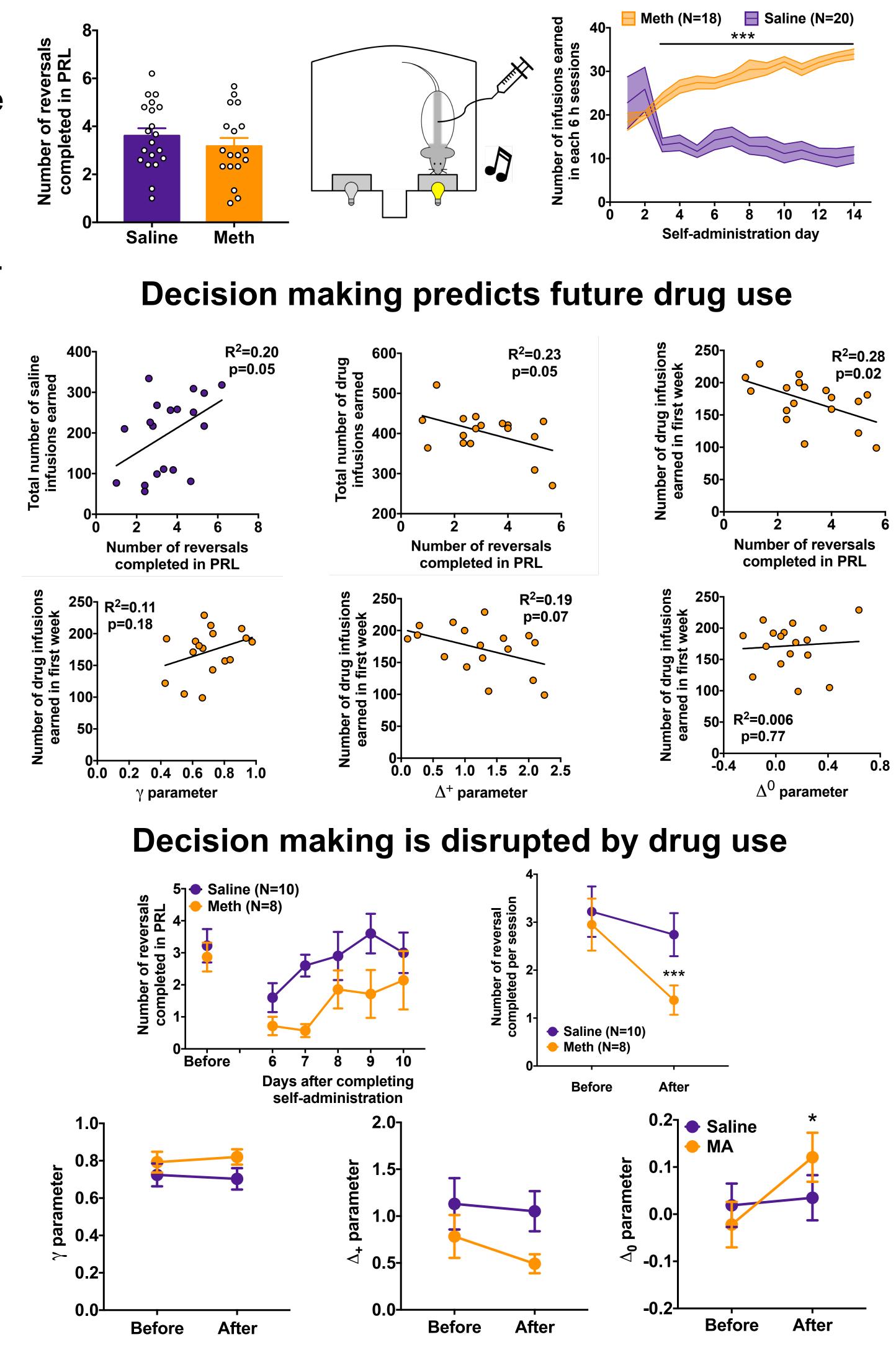
Choice data was analyzed with a reinforcement-learning algorithm. Action values for each option were updated according to the following equation: $Q(t+1) = \gamma Q(t) + \Delta_i$

where the decay rate γ determines how quickly the action values decay and Δ_i indicates the change in the action value that depends on the outcome from the chosen noseport. If the outcome was reward, then the value function of the chosen noseport was updated by Δ_{+} , the appetitive strength of reward. If the outcome was no reward, then the value function of the chosen noseport was updated by Δ_0 , the aversive strength of no reward. Decay of action values for unchosen options was determined by the γ parameter.

Three free parameters:

- γ decay rate
- Δ_+ appetitive strength of rewards
- Δ_0 aversive strength of no rewards



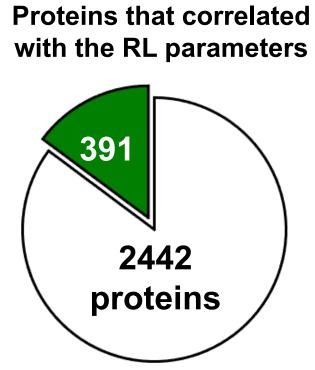


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Rat choices — Port 1 — Port 2 — Port 3 Model choices -- Port 1 -- Port 2 -- Port 3

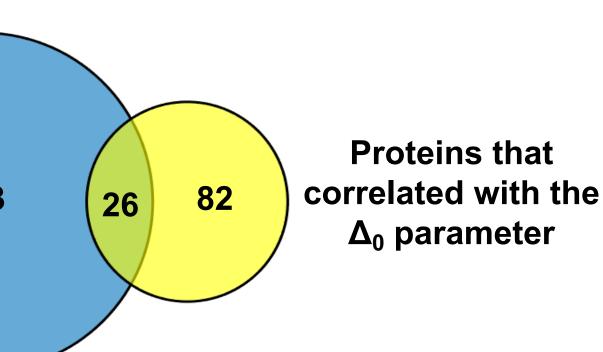
Proteomics

Tissue from the ventral striatum was collected from rats tested on the PRL task who were either drug-naïve (N=18) or had self-administered meth for 14 days (N=16). Proteins were extracted and purified, and peptides fractionated for liquid chromatography mass spectrometry (LC-MS/MS). Expression of each protein was correlated to the individual reinforcement-learning parameters for each rat to identify proteins that co-varied with separable aspects of reinforcement learning.

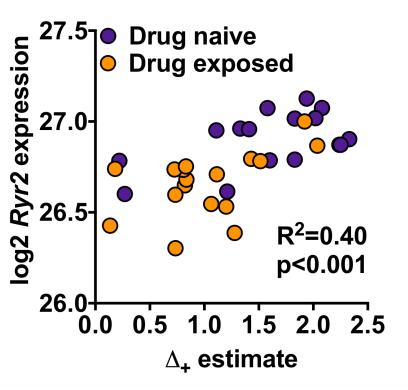


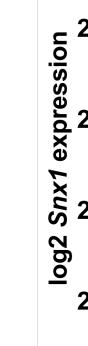
Proteins that correlated with the Δ_+ parameter

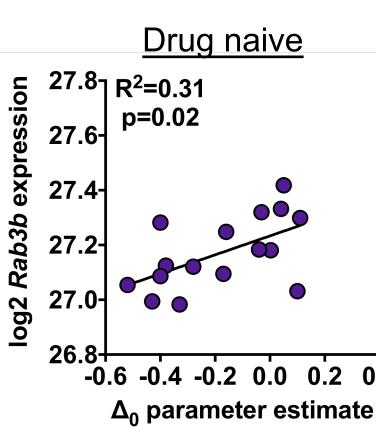
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Identifying protein targets mediating addiction susceptibility			
		2240	Proteins that were not significantly different between drug-naïve and drug-exposed rats
	Proteins that correl with the ∆ ₊ paramet drug-naïve rats	er in 300 217	Proteins that correlated with the ∆ ₊ parameter in drug-exposed rats
Gene	Protein	Function	Previously linked to addiction?
Ndufb10	NADH: ubiquinone oxidoreductase subunit B10	Subunit of mitochondrial membrane respiratory	Altered in alcohol preferring rats (McClintick et al., 2017)
Dpp10	Inactive dipeptidyl peptidase 10	Promotes surface expression of KCND2	
Setd7	Histone-lysine N- methyltransferase SETD7	Monomethylates Lys-4 of histone 3 (methylates nkkb and histones – wb hlk4); histone extraction; histone here repssive at lysine9	Genetic association with smoking behaviors (Thorgeirsson et al., 2010)
Sort1	Sortilin	Sorting receptor in the Golgi compartment	Low expression in high novelty seeking rats (Kabbaj et al., 2004)
Ryr2	Ryanodine receptor 2	Channel that mediates Ca2+ release from sarcoplasmic reticulum	Genetic association with impulsivity and gambling (Khadka et al., 2014; Lind et al., 2012)
Snx1	Sorting nexin-1	Intracellular trafficking	Reduced following meth CPP (Yang et al., 2008)
Gamt	Guanidinoacetate N- methyltransferase	Converts guanidoacetate to creatine	Reduced in alcohol dependent individuals (Sokolov et al., 2003)
Naa15	N(alpha)- acetyltransferase 15	Subunit of NatA complex; important for neuron growth	
Atxn2l	Ataxin 2-like	Involved in stress granule and P-body formation	Genetic association with lifetime THC use (Pasman et al., 2018)
oissa 27.0- 26.5- 26.0-	Drug naive Drug exposed	27.5 Drug naive Drug exposed 27.0 26.5 26.0 26.0 0.0 0.5 1.0 1.5 2.0 2.5 Δ ₊ estimate	27 Drug naive Drug exposed 26 Drug exposed 25 $R^2=0.47$ P<0.001 24 0.0 0.5 1.0 1.5 2.0 2.5 Δ_{+} estimate
Ident	•	•	g addiction consequence
Proteins that correlated with the Δ_0 parameter in drug-exposed rats Proteins that correlated with the Δ_0 parameter in Proteins that correlated with the Δ_0 parameter in			
<mark>ج 27</mark> .	drug-naïve rats <u>Drug naive</u> ⁸ 7 R ² =0.31	Drug expos	<u> </u>
log2 <i>Rab3b</i> expression 27. 27. 27. 27. 27. 26.	6- 4- 2- 0-		0.2 0.4







CONCLUSIONS

These data indicate that the protein-behavior correlates mediating addiction susceptibility differ from those that are disrupted by drug use. Future studies will manipulate expression of these proteins to demonstrate causal evidence for these correlations. Our innovative platform highlights the potential of decision-making biomarkers to isolate protein targets that could be manipulated to promote addiction resilience or treat addiction.

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