Nr4a1 Activation Suppresses Cocaine-induced Behavior via Epigenetic Regulation of Homeostatic Target Genes.

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A core feature of addiction is the propensity for relapse caused by the aggregation of neuroadaptations during abstinence (Shaham & Hope, 2005). To combat relapse, endogenous homeostatic mechanisms may restore and even reverse normal function to reward-related brain areas (Huang et al 2011; Keramati et al, 2017. To identify a master regulator of homeostatic gene expression, we profiled global transcriptomic changes in brain reward areas at early and late abstinence following cocaine self-administration in mice. Using this approach, we identified a key role for the transcription factor, nuclear receptor subfamily 4 group A member 1 (Nr4a1), in epigenetic regulation of homeostatic target gene expression and cocaine-evoked behavior.

Methods: Mice self-administered either saline or cocaine (0.7 mg/kg/infusion) on a fixed-ratio 1 schedule during daily 2-hour sessions for 21 consecutive days. Mice were sacrificed 1- and 28-days after the last cocaine session, brains were removed, and nucleus accumbens (NAc) was collected. Single-sample sequencing (S3EQ) (Xu & Heller, 2018) was used to analyze changes in gene expression and histone modifications. CRISPR/dCas9-mediated gene activation and repression was used to regulate *Nr4a1* in vivo. Cocaine conditioned place preference (i.p., 10 or 5 mg/kg) was used to assess cocaine reward behavior following CRISPR/dCas9-mediated Nr4a1 expression in NAc.

Results: Cocaine self-administration and abstinence regulated expression of the transcription factor, Nr4a1, and its target genes, *Cartpt* and *Vmat2*. Sustained activation of *Cartpt* and *Vmat2* at late abstinence was coupled with depletion of the repressive histone modification, H3K27me3, in the presence of the activating marks, H3K27ac and H3K4me3. CRISPR-mediated gene activation (dCas9-VP64) or repression (dCas9-KRAB) in NAc was sufficient to activate or repress expression of *Nr4a1* and target genes. Cocaine conditioned place preference was attenuated by Nr4a1 activation, and enhanced by its repression. Importantly, dCas9-VP64 activated Nr4a1 expression to the level evoked endogenously by cocaine self-administration.

Discussion: This work provides evidence of the relevance of *Nr4a1* and target genes, *Cartpt* and *Vmat*, in cocaine reward. We identified a molecular mechanism of action of Nr4a1. Specifically, Nr4a1 was sufficient to recapitulate cocaine-activation of Cartpt via loss of repressive histone modifications in the presence of extant activating modifications. Our findings provide evidence that epigenetic editing of endogenous homeostatic genes is a potential strategy to combat cocaine addiction.

References:

Shaham, Y. & Hope, B. T. Nat.Neurosci. 8, 1437–1439 (2005)
Keramati, M., Durand, A., Girardeau, P., Gutkin, B. & Ahmed, S. H. Psychol. Rev. 124, 130–153 (2017).
Huang, Y. H., Schlüter, O. M. & Dong, Y. Behav. Brain Res. 216, 9–18 (2011).
Xu, S. J. & Heller, E. A. J. Neurosci. Methods 308, 62–73 (2018).

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