

Gene Expression Biomarkers for Identifying Vigilance Impairment from Total Sleep Deprivation

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Fatigue is a concern in transportation safety, and health in general. Yet how do we objectively measure fatigue?



The image is a poster from the National Transportation Safety Board (NTSB) titled "NTSB 2016 MOST WANTED TRANSPORTATION SAFETY IMPROVEMENTS". The poster features the NTSB logo on the left, which includes an eagle with a shield and the text "NATIONAL TRANSPORTATION SAFETY BOARD". The main text reads "NTSB 2016 MOST WANTED TRANSPORTATION SAFETY IMPROVEMENTS" in blue, followed by "REDUCE FATIGUE-RELATED ACCIDENTS" in black on a yellow background. Below the text are three photographs: a man in a suit shouting with his hand on his forehead while driving, a close-up of a man resting his head on his hand, and a pilot in a cockpit wearing a headset and looking out the window.

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Experimentally impose acute sleep deprivation

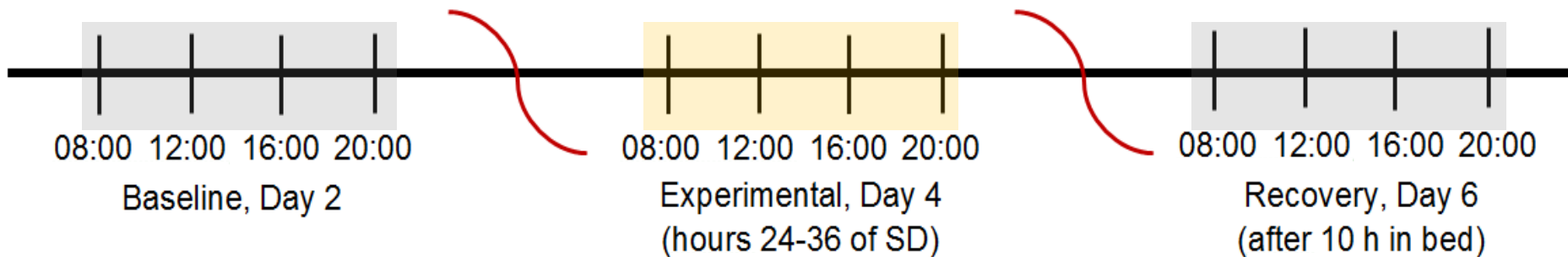
- **Wash. State Univ. sleep lab for 7 days**
 - 6 Control persons: 10 h in bed each night
 - 11 Sleep Deprived persons (SD): 62 h wakefulness
- **Data collection**
 - Blood for gene expression microarrays
 - Psychomotor Vigilance Test (PVT) for cognition
- **Test for differential gene expression**
 - Treatment effect: Control vs. SD individuals
 - Cognitive performance: PVT lapses

For cognition measurements and overview of study design, see Whitney et al. 2015 (<http://dx.doi.org/10.5665/sleep.4668>)



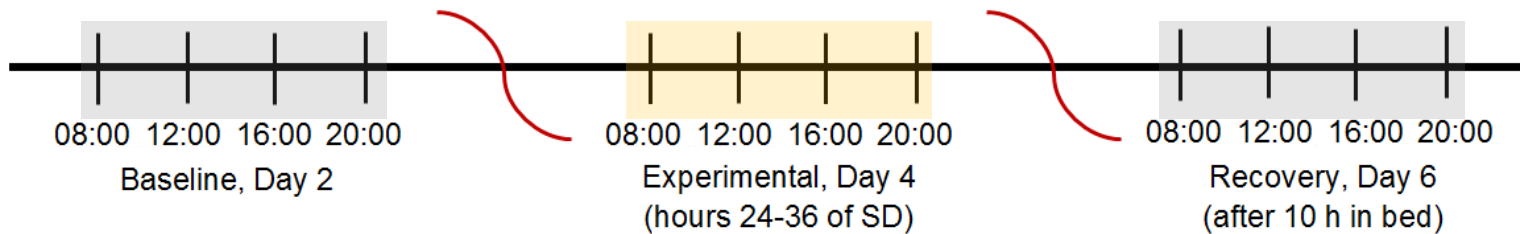
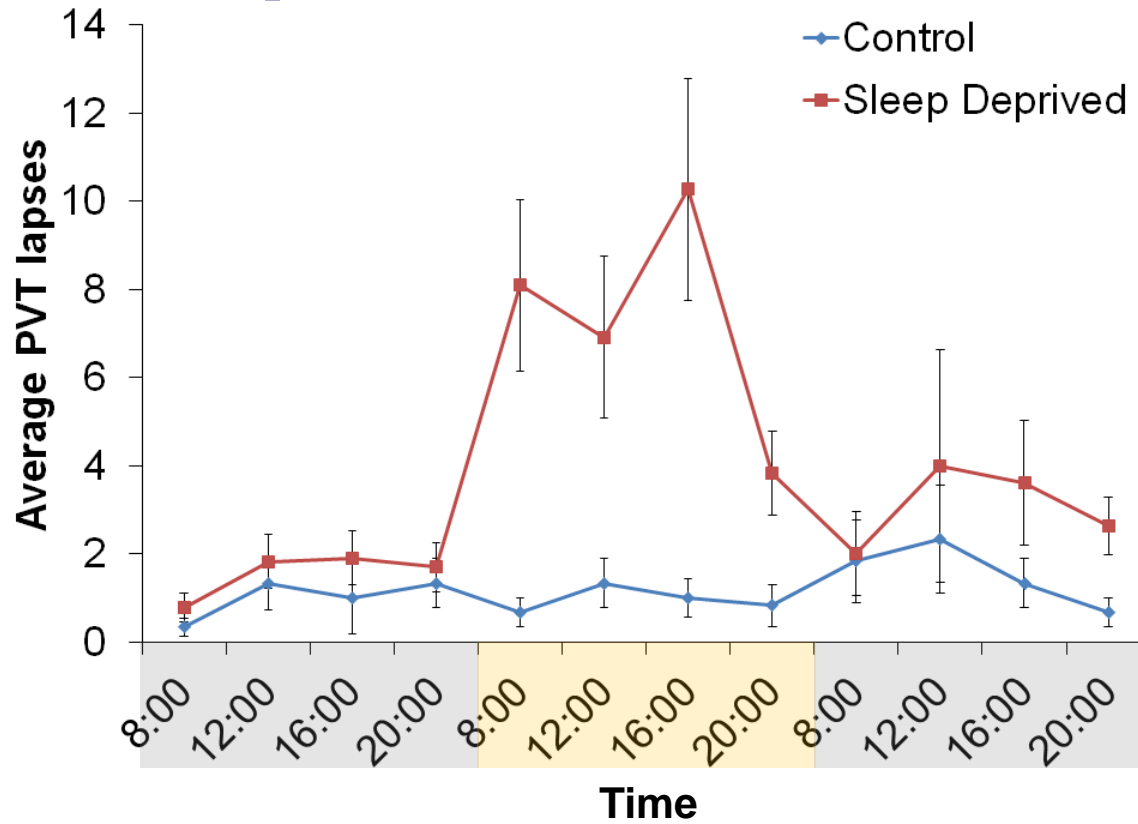
Blood draws for gene expression every other day

- **Control:** 10 h in bed every night
- **Sleep Deprived:** no sleep from 08:00 Day 3, until 22:00 Day 5



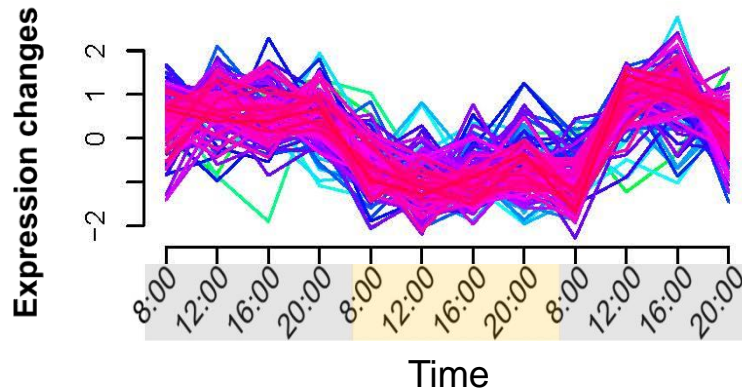
Informed consent and IRB approval were obtained for this research.

PVT lapses increase with SD

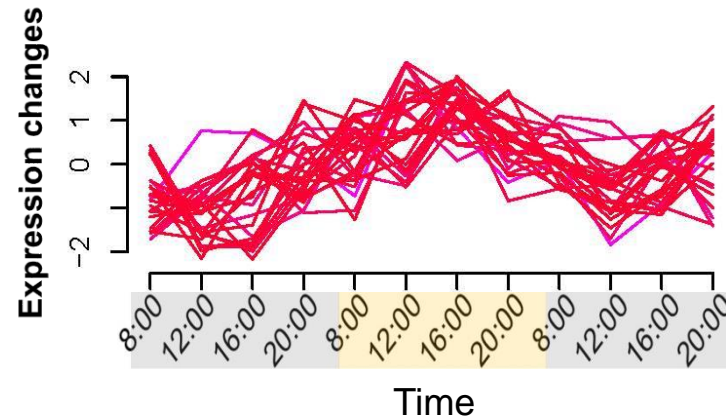


Clustering ~212 treatment effect genes by temporal expression in SD subjects

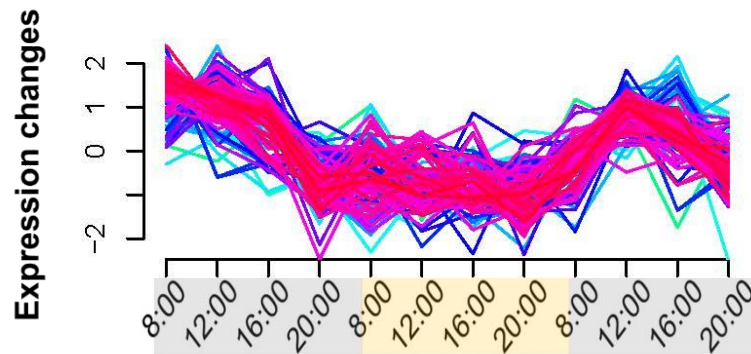
Cluster 1



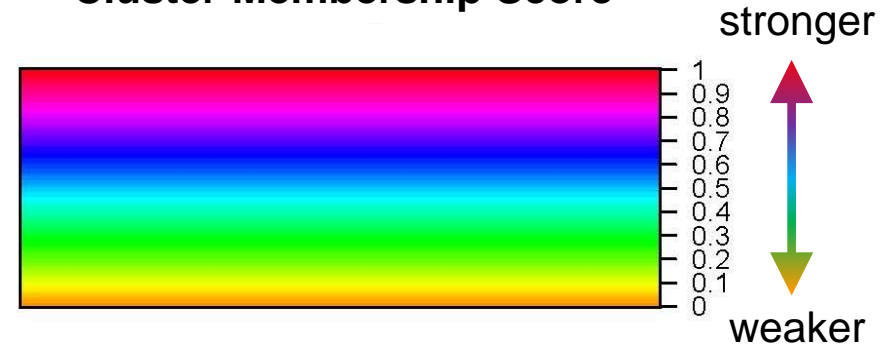
Cluster 2



Cluster 3



Cluster Membership Score



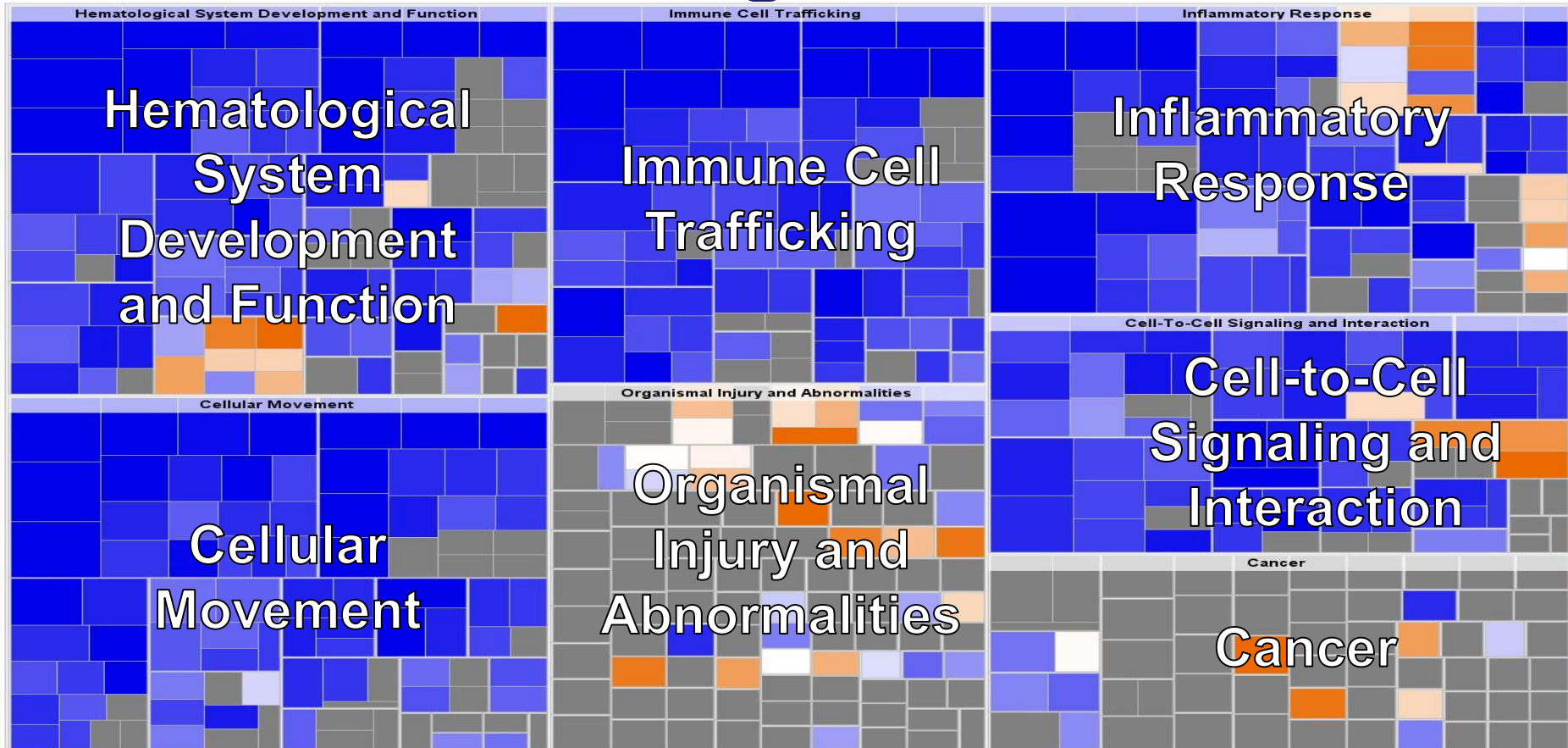
Functional enrichment of the 212 genes related to SD

Functional Group	Enrichment Score	Number of Genes	Description
1	4.43	99	membrane
2	2.82	14	immunoglobulin
3	2.74	16	cell adhesion
4	2.34	38	cell motility; inflammatory response
5	2.24	13	cell junction
6	2.03	12	coagulation
7	1.96	21	cell fraction
8	1.76	3	metal ion-binding site:calcium
9	1.44	15	vesicle
10	1.43	3	sushi; complement control module
11	1.33	13	lipoprotein

Functional analysis performed with the DAVID bioinformatics tool at <https://david.ncifcrf.gov>



Diseases and functions for the 212 SD genes



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Blue = inhibition. Orange = activation.

Z-score: <-3.32

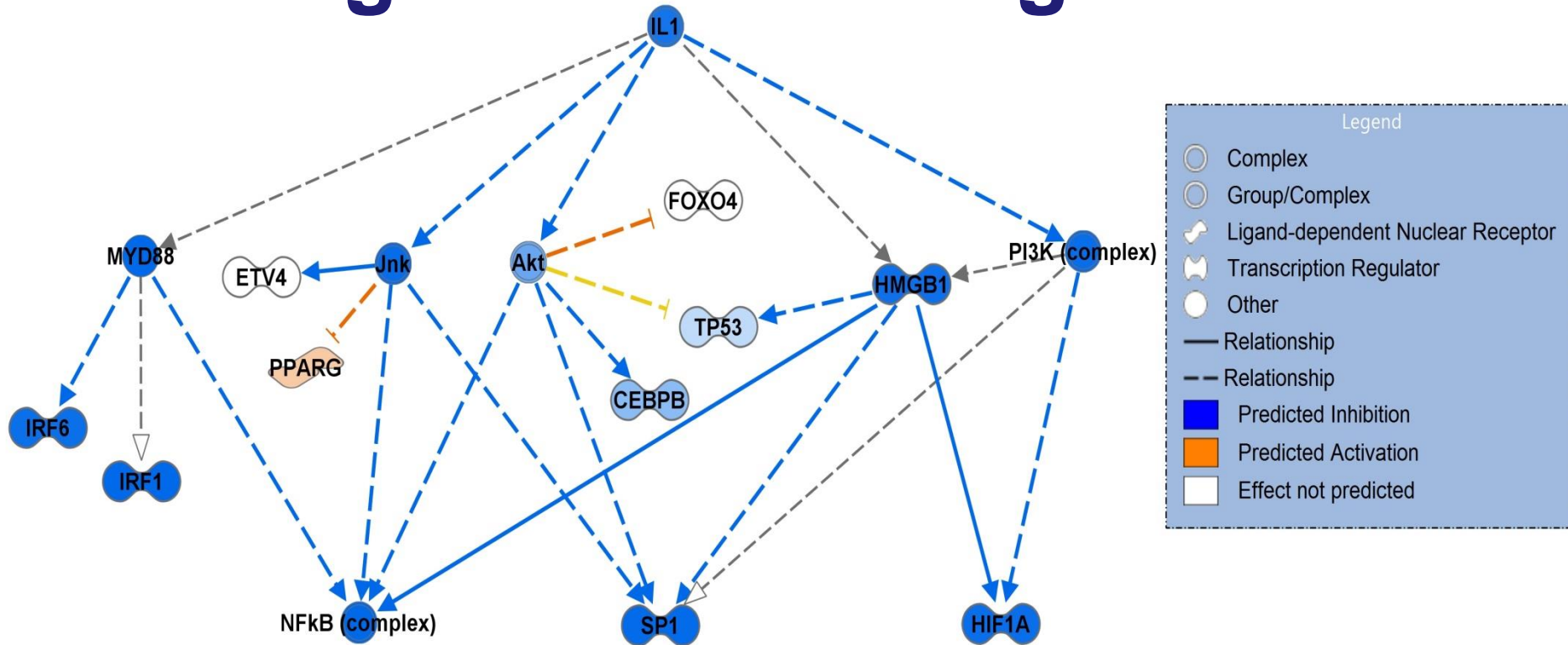


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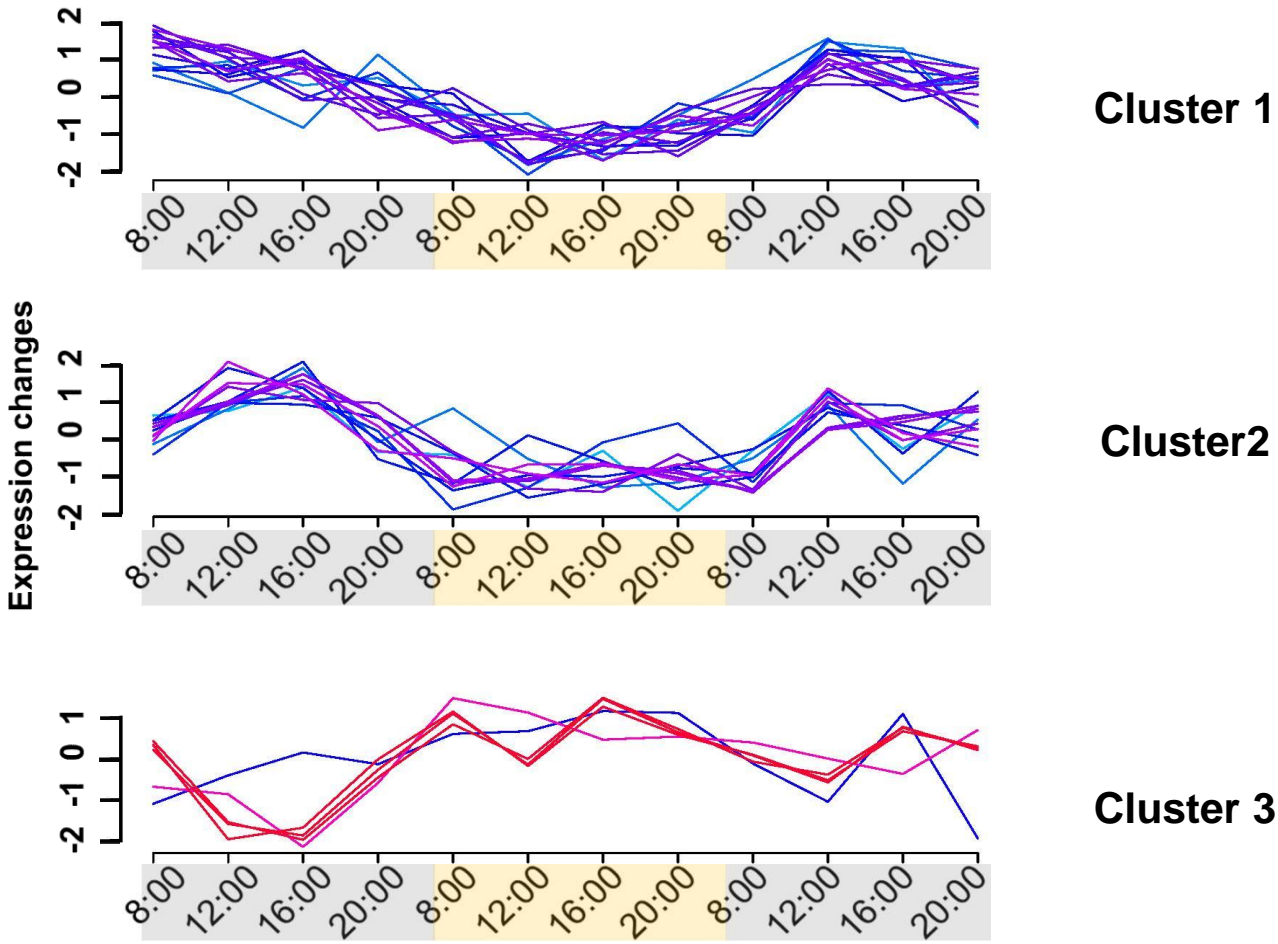
Mechanistic network of upstream regulators for SD genes



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59 downstream SD genes used to predict this network
(all but 4 of which are down-regulated by SD)

Clustering 28 PVT cognition genes by their temporal expression in SD subjects



Functional enrichment of the 28 genes related to PVT scores

Functional Group	Enrichment Score	Number of Genes	Description
1	1.65	9	cell motility; inflammatory response
2	1.37	9	vesicle-mediated transport; intracellular signaling cascade
3	1.37	19	membrane
4	1.35	5	calcium-binding EF-hand

Functional analysis performed with the DAVID bioinformatics tool at <https://david.ncifcrf.gov>



Summary of findings

- **Measurable effects on cognition and gene expression (e.g., cytokines) with one night of sleep deprivation**
- **Overall, we see a reduction in gene expression with sleep deprivation**
- **Future needs:**
 - Fine-tune cognitive measures
 - Increase sample size
 - Assess circadian component

Acknowledgments

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