

Neurocognitive correlates of miRNA expression in the CNS of HIV positive subjects with a history of methamphetamine abuse

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Background

- A growing body of evidence indicates that microRNAs are important regulators of neuronal and brain function¹
- There may be links between changes in microRNA expression and neuronal function²
- Long-term infection with HIV leads to neurocognitive changes and associated cellular and gene expression changes in the central nervous system³
- Methamphetamine abuse among HIV-positive individuals represents a "double epidemic" affecting neurobehavioral outcomes⁴

Aims

- To assess differential expression of microRNAs in the Frontal Cortex of HIV-positive individuals with a history of Methamphetamine Abuse.
- Hypothesize that a set of microRNAs are differentially regulated and whose expression correlates to neurocognitive domains and clinical parameters.

Methods

Features of the Study Groups³

Table 1. Features of the Study Groups					
Group	N	Mean BDI (SD)	Mean PM hr (SD)	Mean Age yr (SD)	
Control	5		21.0 (11.9)	38.0 (14)	
HIV	6	11.6 (5)	15.2 (16.3)	42.1 (5.7)	
METH	5	15.6 (6.7)	19.6 (16.0)	42.1 (11.5)	

BDI - Beck's Depression Inventory, PM hr - postmortem interval, SD - standard deviation

Table 2. Neurocognitive Outcomes of the Study Groups											
Group	Mean GDS (SD)	Mean SIP_DS (SD)	VERBAL_DS (SD)	LEARN_DS (SD)	MEMORY_DS (SD)	EXEC_DS (SD)	WRKMEM_DS (SD)	MOTOR_DS (SD)			
Control											
HIV	1.23 0.79	1.78 (1.4)	0.83 (1.0)	0.50 (0.8)	0.58 (0.4)	1.67 (0.9)	1.08 (1.0)	2.00 (1.4)			
METH	0.68 0.58	0.73 (0.6)	0.20 (0.4)	0.60 (0.9)	0.70 (1.3)	0.50 (0.7)	0.60 (0.5)	1.60 (1.5)			

GDS - Global Deficit Score, SIP - Speed of information processing, DS - Deficit Score, EXEC - Executive functions, WRKMEM - Attention/Working memory, SD - standard deviation

Table 3. HIV Clinical Features of the Study Groups				
Group	CD4 (SD)	Mean NCD4 (SD)	Mean Log (PLA_RNA) (SD)	Mean Log (CSF_RNA) (SD)
Control				
HIV	53.5 (62.6)	30.67 (48)	3.76 (1.6)	1.94 (0.3)
METH	13.8 (11.5)	13.00 (12)	4.66 (1.3)	3.02 (1.2)

CD4 - CD4+ T-cells per mL, NCD4 - lifetime nadir CD4, PLA_RNA - Plasma viral RNA copies per mL, CSF - cerebrospinal fluid

Procedure

1. Isolate RNA from Frontal Cortex
2. TaqMan - based microRNA Array (380 microRNAs)
3. Quantify expression
4. Compare expression among the three groups (Mann Whitney U)
5. Test for correlation of expression with Neurocognitive and Clinical parameters

Results

Hypothesis Testing - Differential Expression by Group

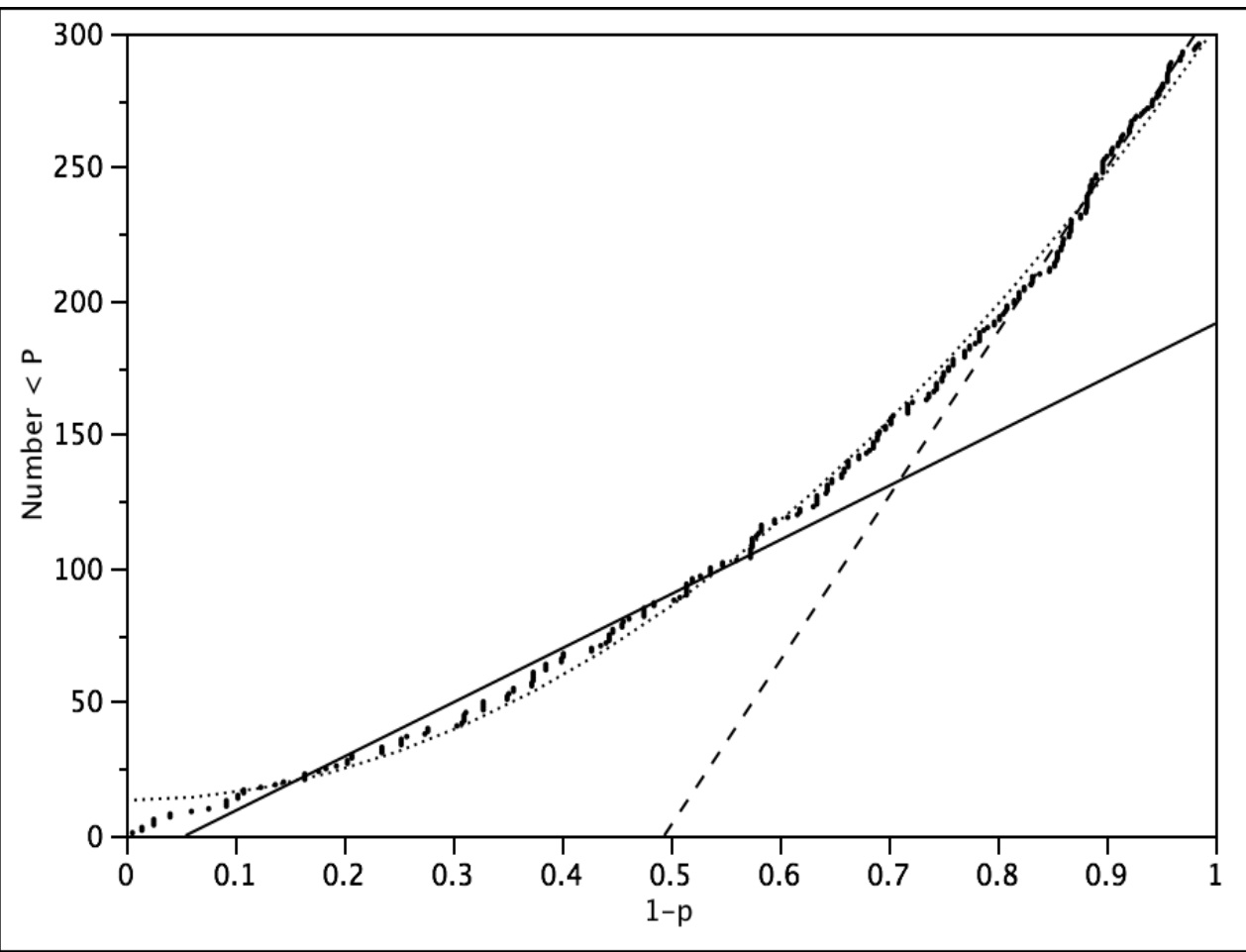


Figure 1. P-plot indicates non-random distribution of p-values from 296 hypothesis tests⁵

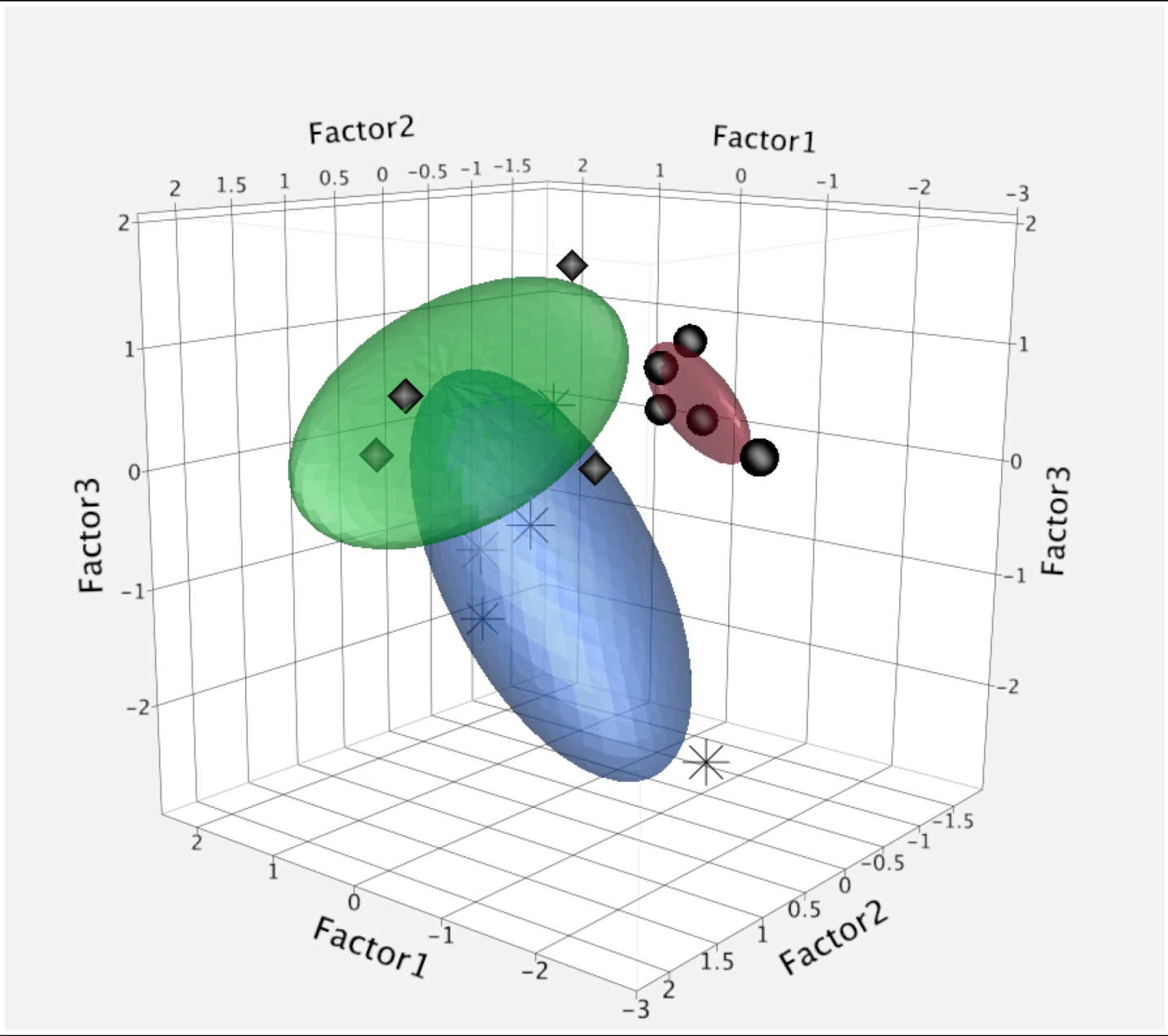


Figure 3. Principal components analysis indicates distinct miR profiles among the three groups.

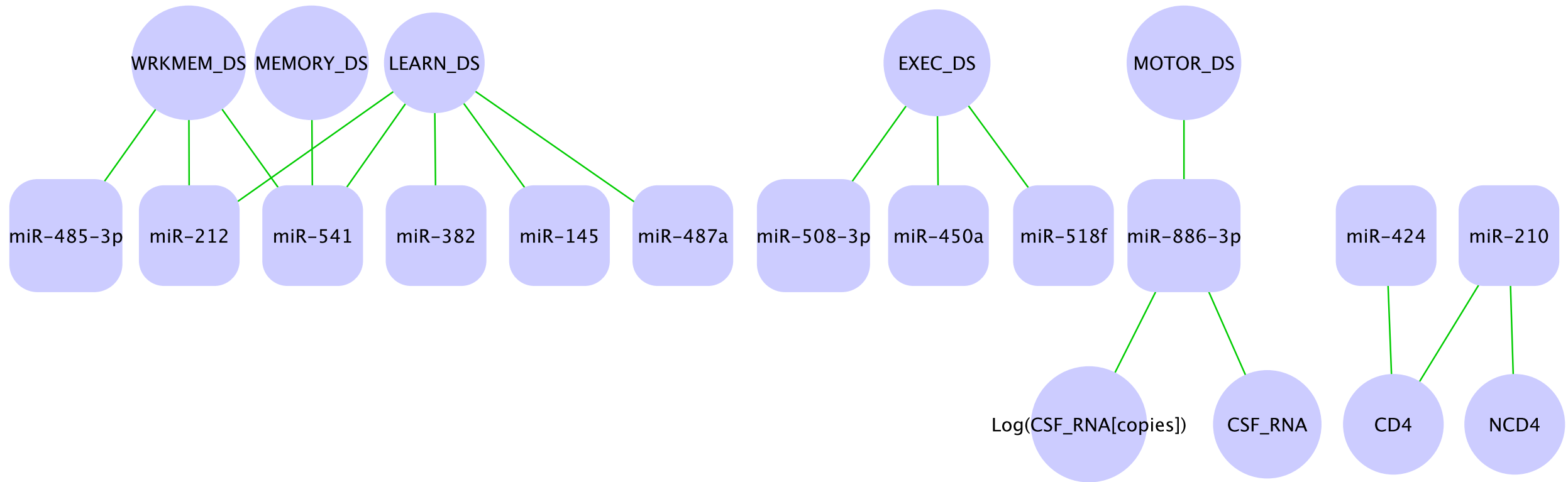


Figure 4. Visualization of microRNA and clinical correlates.

Conclusions

- Learning, Memory, Working Memory, and Executive Functioning Deficits correlate with microRNA expression in the CNS of HIV-infected individuals.
- MiR-212 and miR-541 may be potential molecular mediators of neuronal dysfunction in long-term HIV-infection. MiR-210 and miR-424 may be indicators of immune function.

References

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3. Carey, CL et al. "Predictive validity of global deficit scores in detecting neuropsychological impairment in HIV infection." J Clin Exp Neuropsychology. 26:3:307.
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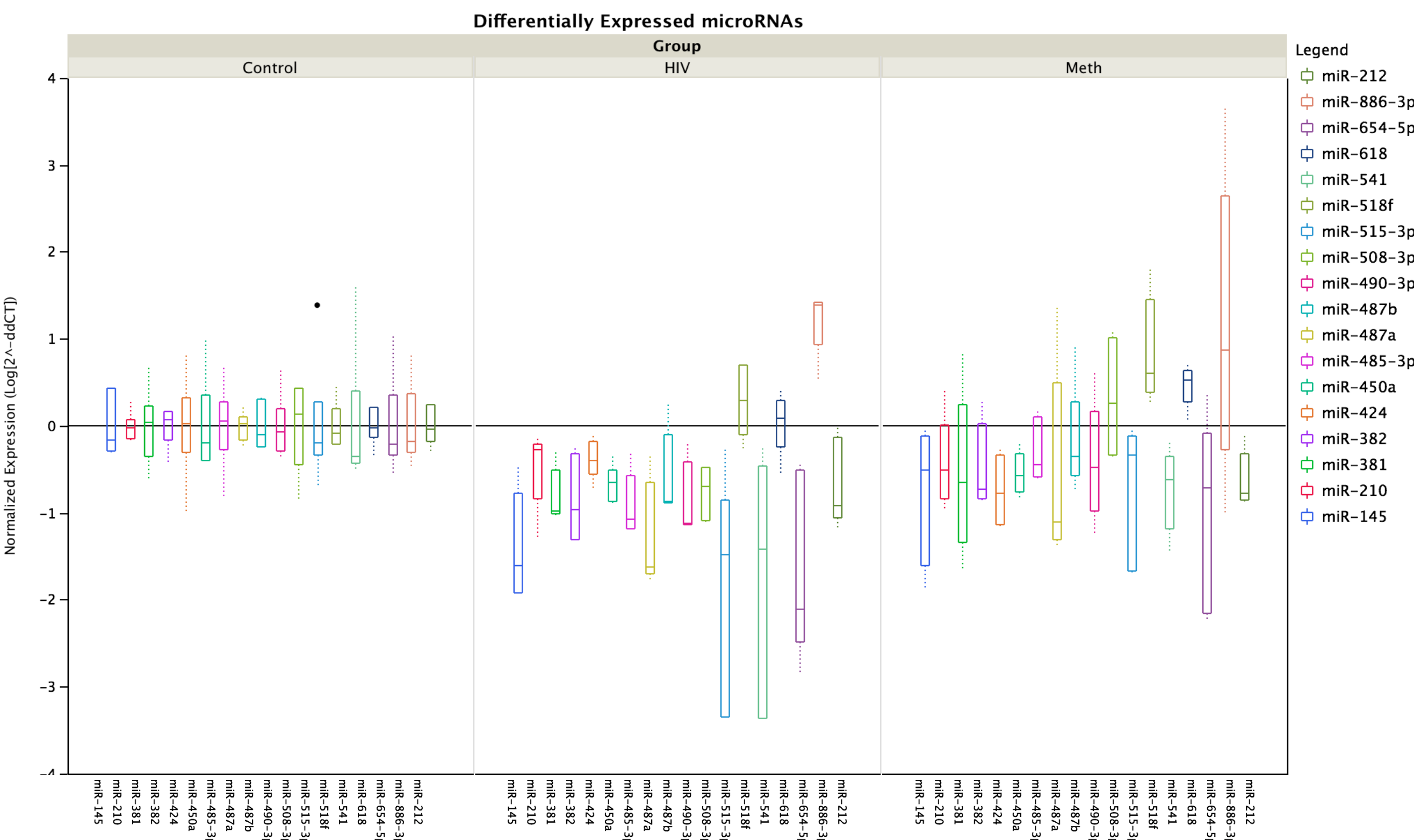


Figure 2. Generally, lower expression in HIV or Meth Groups if significantly different.

Correlation Analysis - MicroRNA Expression Compared to Clinical Variables

Table 4. Neurocognitive and Clinical Correlates to miR Expression in the Frontal Cortex			
Variable	by Variable	Spearman ρ	Prob> p
<i>Neurocognitive Variables</i>			
EXEC_DS	miR-450a	0.6523	0.0409
EXEC_DS	miR-508-3p	0.6461	0.0436
EXEC_DS	miR-518f	0.8697	0.0011
LEARN_DS	miR-145	-0.6806	0.0303
LEARN_DS	miR-212	-0.7838	0.0073
LEARN_DS	miR-382	-0.6394	0.0465
LEARN_DS	miR-487a	-0.8663	0.0012
LEARN_DS	miR-541	-0.8113	0.0044
MEMORY_DS	miR-541	-0.7694	0.0093
MOTOR_DS	miR-886-3p	0.7245	0.0178
WRKMEM_DS	miR-212	-0.702	0.0236
WRKMEM_DS	miR-485-3p	-0.6376	0.0473
WRKMEM_DS	miR-541	-0.7149	0.0201
<i>Clinical Variables</i>			
CD4	miR-210	-0.7927	0.0062
CD4	miR-424	-0.7683	0.0094
CSF_RNA	miR-886-3p	-0.8795	0.004
Log(CSF_RNA)	miR-886-3p	-0.7394	0.036
NCD4	miR-210	-0.7693	0.0093

MicroRNA expression was tested for correlation with neurocognitive and clinical variables using Spearman's rho test. Significant correlations are shown.