## **EDA/SDA/RCPE Delirium Teaching Day** 4 September 2019



# Delirium Association/Royal College of Physicians of Edinburgh Joint Conference on Delirium 5 and 6 September 2019



Poster 10







European Delirium Conference Edinburgh, 4-6 September 2019

EUROPEAN DELIRIUM ASSOCIATION

# Evaluation of circulating miRNA in serum samples of delirium patients in acute geriatric medical setting: insights on the inflammatory hypothesis of delirium

Authors: Patrícia Regueira 2, Ana Rita Silva 1, Ana Luísa Cardoso 1, Elisabete Albuquerque 2, Fabiana Ventura 2, Mário Carneiro 2, Inês Baldeiras 13, Isabel Santana 3, Joaquim Cerejeira 12 4Center for Neuroscience and Cell Biology, University of Coimbra, Coimbra, Portugal 3Department of Psychiatry, Coimbra University Hospitals, Coimbra, Portugal; 4Department of Neurology, Coimbra, Coimbra,

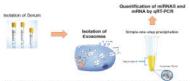
#### INTRODUCTION

MicroRNA (miRNA) are short non-coding RNA molecules usually composed of 18-25 nucleotides that play a role in the regulation of the neuroinflammation, however its association with delirium is not yet established. Acute medically-ill older patients are at high risk to develop delirium, which is frequently related with adverse outcomes. In this ongoing study we are testing the hypothesis of acute neuroinflammatory response as a disease marker of delirium, measured by serum biomarkers such as miRNA.

Aim: To determine whether elderly patients who develop delirium have a different pattern of inflammatory response; than patients without delirium.

#### **METHODS**

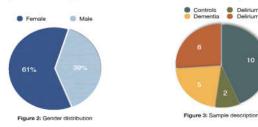
A case-control prospective study was developed on subjects aged 65 or over, unplanned admitted to Internal Medicine wards, with acute systemic infection. Patients are classified in four groups, according to their cognitive status: no delirium and no dementia (controls), delirium only, dementia only and delirium superimposed on dementia. We also collect blood samples to measure inflammatory mediators, namely levels of miRNAs in circulating monocytes and in exosomes. Serum levels of miRNA were determined by quantitative real-time PCR (qRT-PCR).



### **RESULTS**

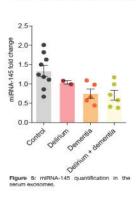
A total of 23 patients (61% female, mean age 85.3±6.1 years - min. 72 years, max. 96 years) (fig.1) were enrolled in this study: 10 controls, 2 delirium only, 5 dementia only and 6 delirium superimposed on dementia.

Cognitive impairment was present in 59% of the sample. Only controls differed significantly from the remaining groups in MoCA test (Mann-Whitney U = 0.0, Z = -2.167; -2.847; -3.277, p <. 05).



Total	11.9	6.6	10.5
Delirium+Dementia	6.1	2.6	5.0
Dementia	7.8	3.2	7.5
Delirium	5.5	0.7	5,.5
Control	18.3	3.2	18.0
Group	Mean	SD	Median

Mean Size Nº of particles 200 500000 of particles size 100 100000



Regarding the percentage of exosomes in serum, miRNA-145 diminished delirium+dementia group compared to controls (delirium+dementia miRNA-145 fold change=1.18±0.3; controls=2.1±0.56) (fig.5) and the mean size of exosomes in delirium alone is superior than the remaining groups (delirium 95.5±7.6, controls 71.5±10.6, dementia 55.4±5.6; dementia+delirium 68.5±10.8) (fig.4).

## CONCLUSION

These preliminary data suggest a down-regulation of miRNA-145 (an anti-inflammatory marker) in serum exosomes of patients diagnosed with delirium+dementia or dementia only. Delirium patients seem to have an increased size of extracellular vesicles when comparing with the other groups.





