DETERMINATION OF EXPRESSION SIGNATURE AND PROPORTION OF mtDNA IN PLASMA FRACTIONS IN PATIENTS WITH RENAL CELL CARCINOMA.

Elena Arance-Criado¹, Fernando Vázquez-Alonso, M^a Yarmila García-Iglesias³, Rocío López-Cintas³, Sara Martín-Esteban⁴, Ginesa López-Torres⁵, Ana Isabel Cortés-Valverde⁶, María Jesús Álvarez Cubero^{1,7} and Luis Javier Martínez-Gonzalez^{2*}

¹GENYO. Centre for Genomics and Oncological Research. Pfizer / University of Granada / Andalusian Regional Government. ²Urology Department, Hospital Virgen de las Nieves. Av. De las Fuerzas Armadas 2, 18014, Granada, Spain. ³Gran Capitán Health Center. Calle Gran Capitán, 10, 18002 Granada. ⁴Peligros Health Center. Calle Valencia, s/n, 18210 Peligros, Granada. ⁵Salvador Caballero Health Center. Calle Dr. Azpitarte, 6, 18012 Granada. ⁶Casería Montijo Health Center. Carr. de Jaén, s/n, 18013 Granada. ⁷Department of Biochemistry and Molecular Biology III and Inmunology, Faculty of Medicine, University of Granada. Av. de la Investigación, 11, 18016, Granada, Spain.



Renal cell carcinoma (RCC) is the third most common urologic malignancy¹, and its incidence is increasing globally. Currently, the main issue is the high stage in diagnosis of this disease. This aspect reinforces the need of identifying novel predictive biomarkers for diagnosis, progression



and prognosis of RCC^2 .

mtDNA exists as a circular, double-stranded nucleic acid with a high copy number. It is known the susceptibility of mtDNA to oxidative stress and mutation³ and there are some studies that found the relationship between mtDNA copy number (mtDNAcn) and risk of developing several cancer^{4,5}.

In this study the main objective is to determine the concentration of exosomal mtDNA present in plasma fractions of controls and patients as a potencial non-invasive biomarker of RCC in liquid biopsy.

Figure 1: Diagram of the formation of exosomes and emitted vesicles



MATERIALS AND METHODS



0.001	2	4	6	8	10	12	14	16	18	20 Cyc	22 Ie	24	26	28	30	32	34	36	38		
HV1 Mitochondrial hypervariable region HBB Beta nuclear																					
	CYB Apocytochrome B of complex III								hemoglobin												

RESULTS AND DISCUSION

After making a statistical analysis of the samples to each of the study phases, we observed that in phase F appears the only significant data regarding the difference in DNA concentration between cases and controls (Figure 3).



In that fraction, with a higher content of exosomal mtDNA, p value shows statistically significant differences in mitochondrial genes HV long and CYB long. While the nuclear gene (both short and long) shows high p values, short fragments for the mitochondrial genes (HV short and HV long) show interesting p values even though they are not significant (Table A3).

It is observed a little increase in the mitochondrial long genes of this study (HV and CYB), due to the protective role of the lipid bilayer of the exosomes (Figure A1).

Gene	P value	Mean of controls	Mean of cases
HV-Short	0,074871	0,003839	0,01005
HV-Long	0,021676	0,002671	0,007255
CYB-Short	0,058973	0,002189	0,004655
CYB-Long	0,049960	0,002233	0,004892
HBB-Short	0,263638	0,003772	0,008355
HBB-Long	0,410166	0,003674	0,01176

Gene

Figure 3: DNA concentration of each gene fragment of controls and patients. The black bars represent the group of the controls, and the grey ones the group of the patients.

CONCLUSION

Table 1: Means of each gene fragment of controls and cases in each of the phase F. P value regarding the difference in DNA concentration between cases and controls applying a t-test (t-student).

This study allowed us to analyze the fragment size distribution pattern of different regions of interest in each plasma fraction, and we have confirmed the high mtDNA content in exosomes as a powerful biomarker. Therefore, application of liquid biopsy in the clinical scenario is a promising non-invasive technique for prediction, diagnosis and monitoring of cancer treatment. We affirm that it would be quite interesting to study how the amount of mitochondrial DNA varies in controls and patients with RCC in fractions rich in exosomal content, allowing the preservation of mtDNA thanks to its lipid bilayer structure.



REFERENCES

- 1. Capitanio, U. et al. Epidemiology of Renal Cell Carcinoma [Figure presented]. European Urology vol. 75 74–84 (2019).
- 2. Oto, J. et al. Urinary microRNAs: Looking for a New Tool in Diagnosis, Prognosis, and Monitoring of Renal Cancer. Current Urology Reports vol. 21 (2020).
- K. et al. Pre-diagnostic leukocyte mitochondrial DNA copy number and colorectal cancer risk. Carcinogenesis 40, 3. 1462–1468 (2019)
- 4. Yu, M. Circulating cell-free mitochondrial DNA as a novel cancer biomarker: Opportunities and challenges. Mitochondrial DNA vol. 23 329–332 (2012).
- 5. Gentiluomo, M. et al. Mitochondrial DNA copy-number variation and pancreatic cancer risk in the prospective EPIC cohort. Cancer Epidemiol. Biomarkers Prev. 29, 681–686 (2020).
- 6. Figure 1 and 2 created with Biorender.com