

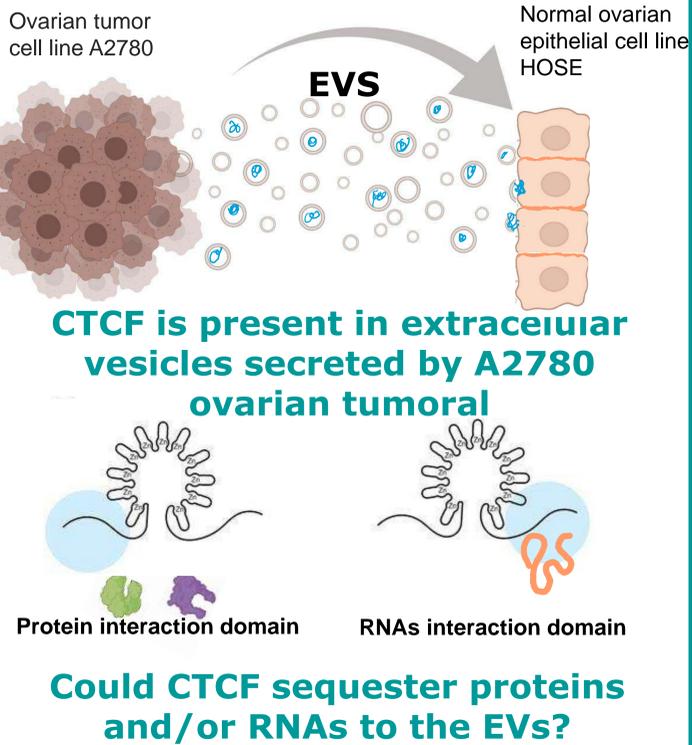
# Studying the implication of the presence of master waiver of the genome in extracellular vesicles secreted by the ovarian tumor cell line A2780 when received by normal epithelial ovarian cells



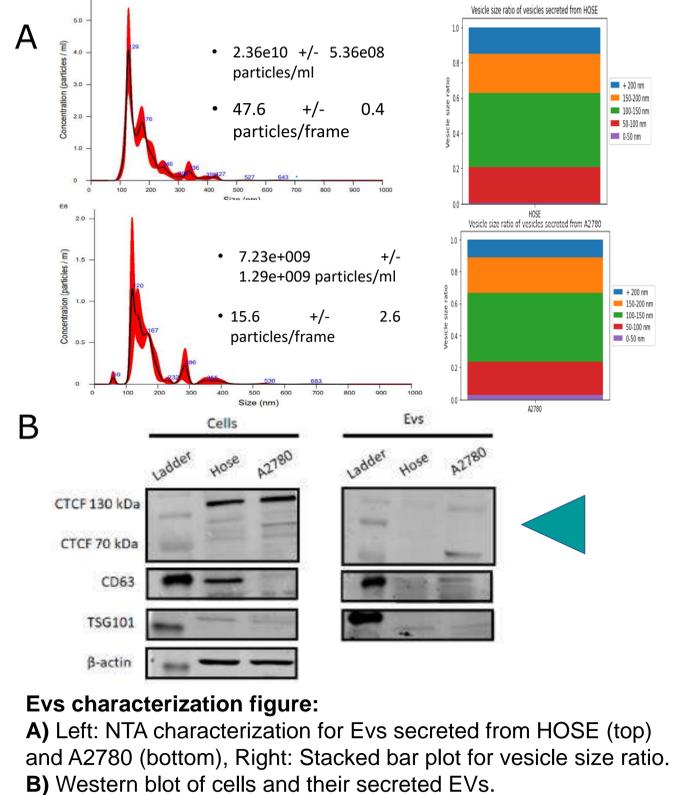
Matías del Campo-Smith, Eduardo Duran, Sebastián Urrejola, Carmen Romero Osses, Lorena Lobos-González

The master waiver of the genome CTCF modulates the differentiation and heterogeneity of eukaryotic cells through its interaction with the genome. Its functions include the structuring of the genome in loops to regulate the physical interaction between enhancer and promoter sequences, direct promotion and inhibition of gene expression, and even modulates nucleosome positioning. For this reason, the study of its role in cancer becomes crucial for understanding the progression of this disease. One of the fundamental mechanisms for cancer development is the control of the tumor microenvironment, and in this category extracellular vesicles (EVs) are crucial. To date, its presence and role in EVs have not been described. In this study we characterize the presence of this protein and its transcript in EVs secreted by A2780 ovarian cancer cells. EVs were characterized through western blot, NTA and electron microscopy CTCF protein was detected by western blot. It has recently been discovered that CTCF has a region capable of binding RNAs, which makes us suspect that this protein may be capable of sequestering RNAs and incorporating them into micro vesicles. Our bioinformatics studies showed that the RNAs with the highest probability of being sequestered in patients with ovarian cancer are MALAT1, NEAT1, PLEC, CPNE1, PNN, HGS and ADRM1. The next steps are to study the capacity of these CTCF+ EVs in modulating crucial tumor-promoting or cancer-related properties/characteristics such as invasion (Wound and transwell assays, clone, and spheroids) in the normal ovarian epithelium cells.

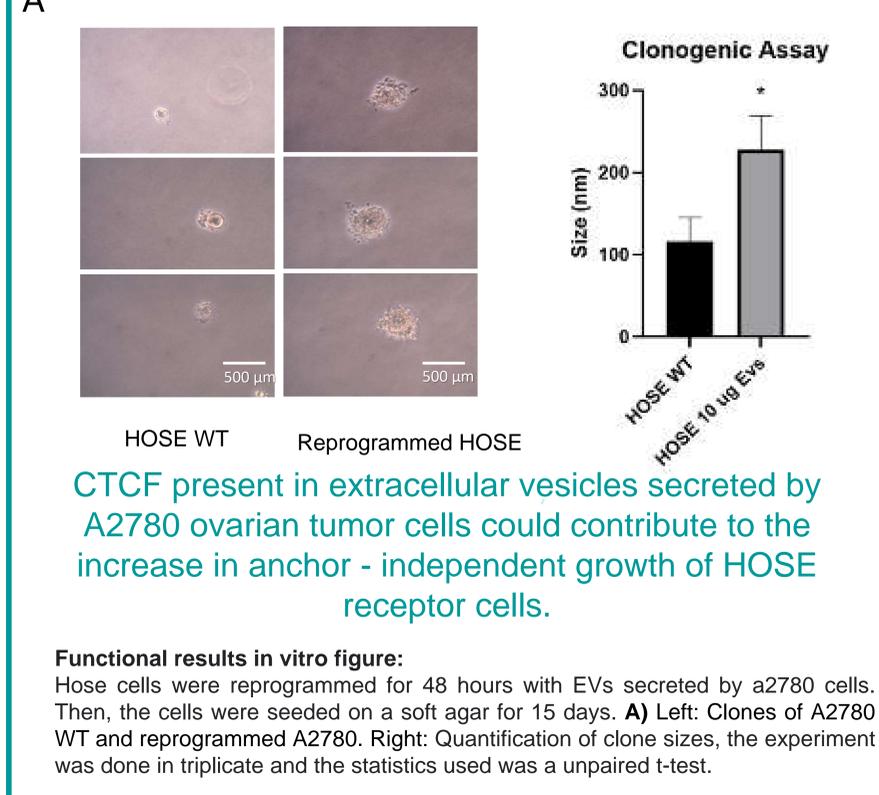
## Experimental Set-up



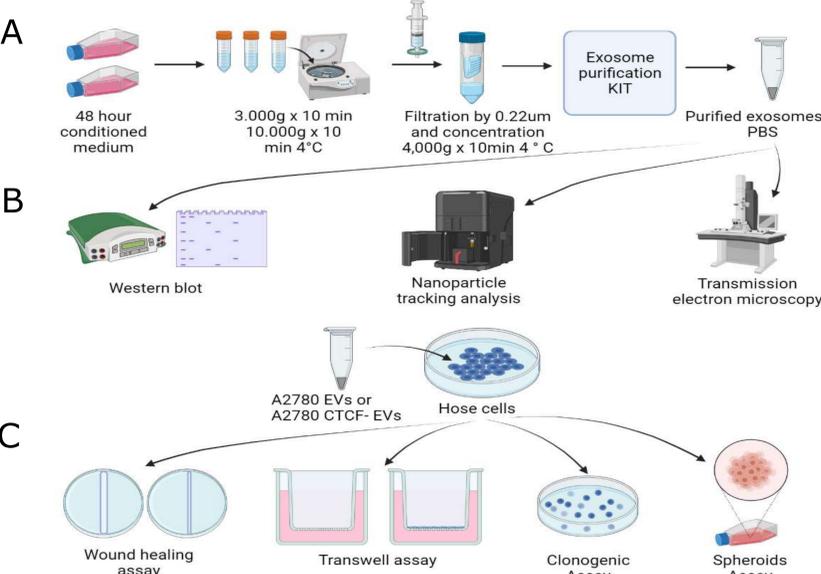
## Evs Characterization



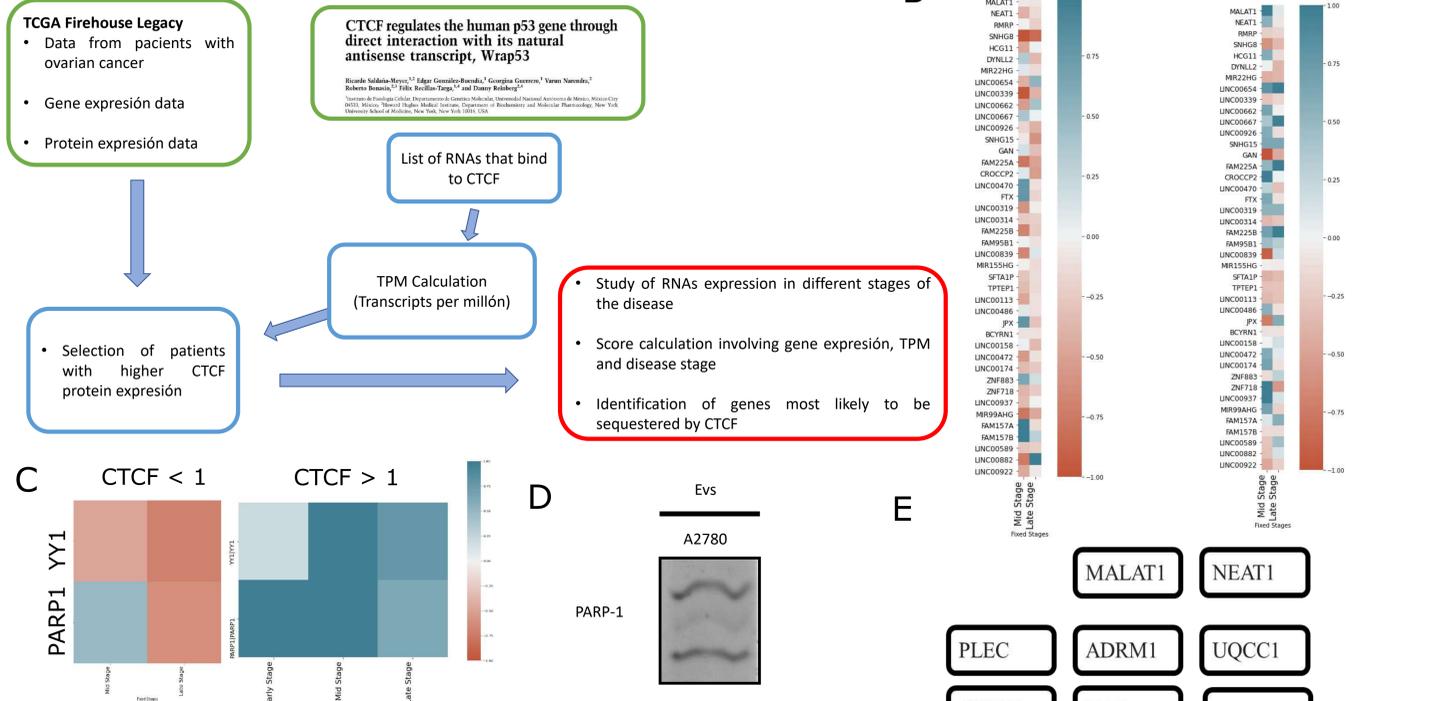
## Functional Results In Vitro



## Materials and methods



## In Silico results



## Conclusions

- CTCF is present as a cargo in EVs secreted by the a2780 cell line.
- CTCF is a promising agent in promoting metastasis when loaded to evs due to the large number of functions in the regulation of genetic expression

## Discussions

Since functional experiments have not yet been carried out, it remains to be demonstrated what is the contribution of CTCF as cargo in EVs in increasing the metastatic properties of receptor cells.

## Acknowledgment

1211223 Fondecyt  
1160139 Fondecyt  
15130011 Fondap

- Phillips JE, et al (2009). CTCF: master weaver of the genome. Cell 2009; 137: 1194-1211
- Lintao Zhao, et al (2017). CTCF promotes epithelial ovarian cancer metastasis by broadly controlling the expression of metastasis-associated genes. Oncotarget, 2017, Vol 8, (No. 37), pp: 62217-62230