

March 2023

Loss of Connective Tissue Growth Factor Expression Promotes Remodeling of the Extracellular Matrix and Epithelial-to-Mesenchymal Transition in Ovarian Cancer

Mc Harry Ramos
Wayne State University, he3507@wayne.edu

Sandra Galoforo
C.S. Mott Center for Human Growth and Development

Colton Morris
C.S. Mott Center for Human Growth and Development, morris2@kenyon.edu

Gil Mor
C.S. Mott Center for Human Growth and Development, gmor@med.wayne.edu

Ayesha Alvero
C.S. Mott Center for Human Growth and Development, ayasha.alvero@wayne.edu

See next page for additional authors

Follow this and additional works at: https://digitalcommons.wayne.edu/som_srs



Part of the [Medicine and Health Sciences Commons](#)

Recommended Citation

Ramos, Mc Harry; Galoforo, Sandra; Morris, Colton; Mor, Gil; Alvero, Ayesha; and Gogoi, Radhika, "Loss of Connective Tissue Growth Factor Expression Promotes Remodeling of the Extracellular Matrix and Epithelial-to-Mesenchymal Transition in Ovarian Cancer" (2023). *Medical Student Research Symposium*. 277.

https://digitalcommons.wayne.edu/som_srs/277

This Research Abstract is brought to you for free and open access by the School of Medicine at DigitalCommons@WayneState. It has been accepted for inclusion in Medical Student Research Symposium by an authorized administrator of DigitalCommons@WayneState.

Authors

Mc Harry Ramos, Sandra Galoforo, Colton Morris, Gil Mor, Ayesha Alvero, and Radhika Gogoi

Loss of Connective Tissue Growth Factor Expression Promotes Remodeling of the Extracellular Matrix and Epithelial-to-Mesenchymal Transition in Ovarian Cancer

Harry Ramos¹, Sandra Galoforo², Colton Morris², Gil Mor², Ayesha Alvero², Radhika Gogoi^{1,2}

¹Wayne State University School of Medicine, Detroit, MI

²C.S. Mott Center for Human Growth & Development, Detroit, MI

Background: Ovarian Cancer (OC) is the leading cause of death from gynecologic malignancies in the United States largely due to the advanced stage at the time of diagnosis. Epithelial-to-mesenchymal transition (EMT) is a key biological process implicated in the pathophysiology of the metastatic spread of OC. Discovering the “trigger/s,” its downstream targets, and therapeutic targeting are essential to substantively improve the survival of women with OC. The objective of our study is to evaluate the role of Connective Tissue Growth Factor (CTGF) in EMT in OC.

Methods: R182 and R2615 are well-described epithelial OC cell and MR182 and MR2615 are the mesenchymal counterparts. R182/R2615 CTGF knock outs (KO) were derived utilizing a Cas9/CRISPR-Cas9 lentivirus plasmid vector and verified by indel sequencing. Invasion, anoikis resistance, and chemosensitivity assays were performed in wild-type (WT) and KO cells. RNA sequence analysis was performed and analyzed using iPathway guide. Top five upregulated and downregulated genes involved in ECM organization pathway were validated by quantitative PCR (qPCR). Immunofluorescence was performed for F-actin.

Results: CTGF was expressed in the epithelial and not in the mesenchymal OC cell lines. Loss of CTGF was associated with anoikis resistance, where KO and WT cells displayed 75% and 10% viability, respectively. KO cells were significantly more invasive than WT cells. Administration of exogenous CTGF in KO cells decreased invasion in a dose dependent manner. No change was seen in chemosensitivity to Cisplatin in KO cells. RNA seq analysis identified ECM organization as the biologic process most affected by loss of CTGF. Upregulated (FREM2, LAMC2, ITGB4) and downregulated (SPP1, SV2A, RELN, COL6A3, COL4A6) extracellular matrix genes were validated by qPCR. Immunofluorescence staining of F-actin demonstrated increased cytoskeleton expression of F-actin in CTGF KO cells.

Conclusion: Our data suggests that CTGF expression maintains the epithelial phenotype in OC. Loss of CTGF may be one of the early triggers of EMT in OC through extracellular matrix remodeling affecting anoikis and adhesion characteristics, thus acquiring a more migratory and invasive phenotype.