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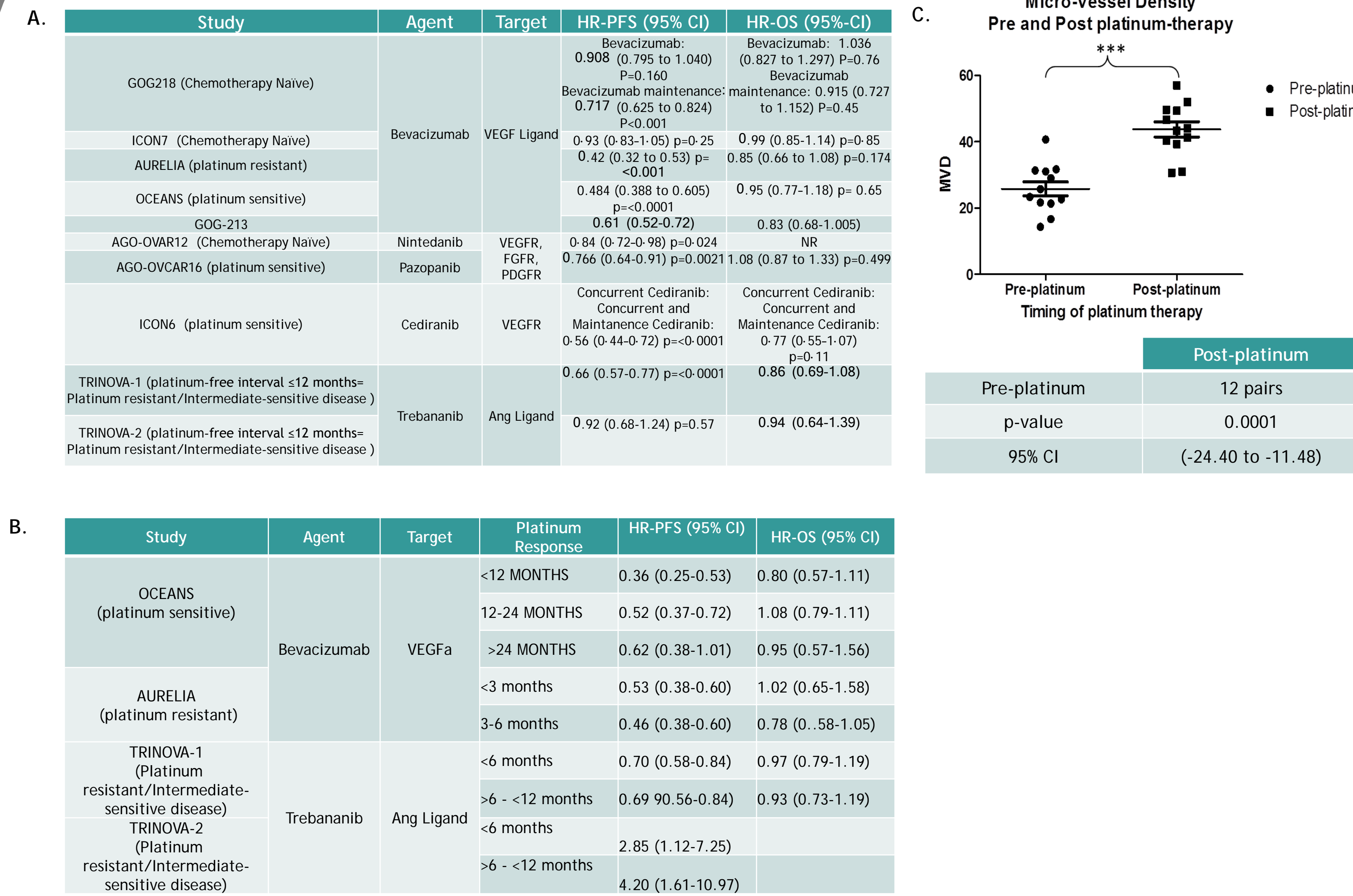
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Poster Number: 776

Background

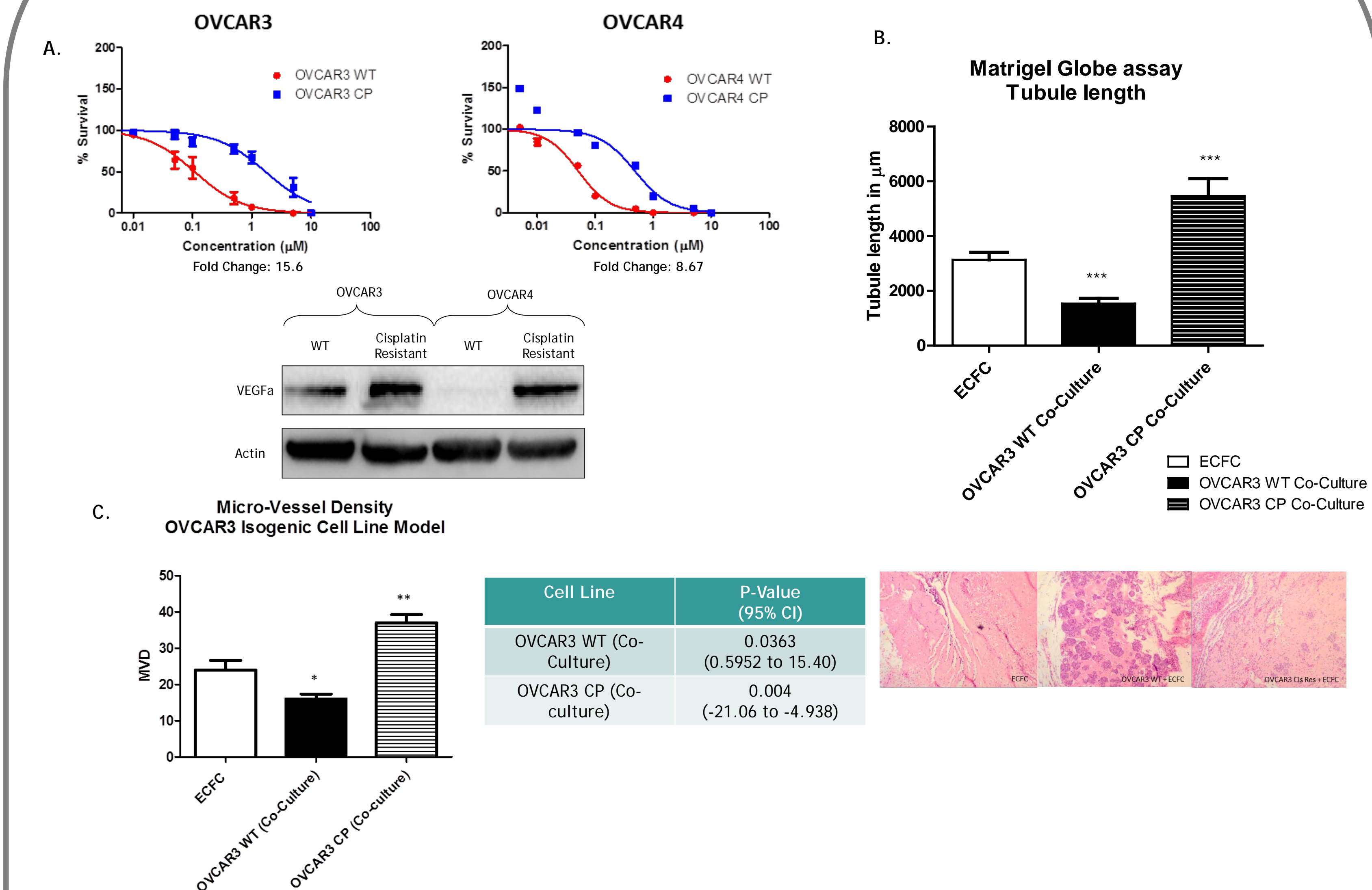
1. Angiogenesis is a key pathological feature of epithelial ovarian cancer (EOC) and anti-angiogenics have dominated the field of drug development in EOC
2. There are no clinically approved predictive biomarker to preselect the subgroup of high grade serous ovarian cancer (HGSOC) that will derive benefit from anti-angiogenic therapy
3. Relapse post platinum-based therapy is associated with a beneficial response to anti-angiogenics
4. Innate and acquired platinum resistance is characterised by upregulation of VEGFa and receptor tyrosine kinase expression that trigger an enriched angiogenic tumour micro-environment
5. In-vitro and in-vivo angiogenesis assays demonstrate that platinum resistant cell lines are sensitive to Cediranib and Nintedanib

I. Platinum therapy resistance is associated with response to anti-angiogenic agents in clinic trials.



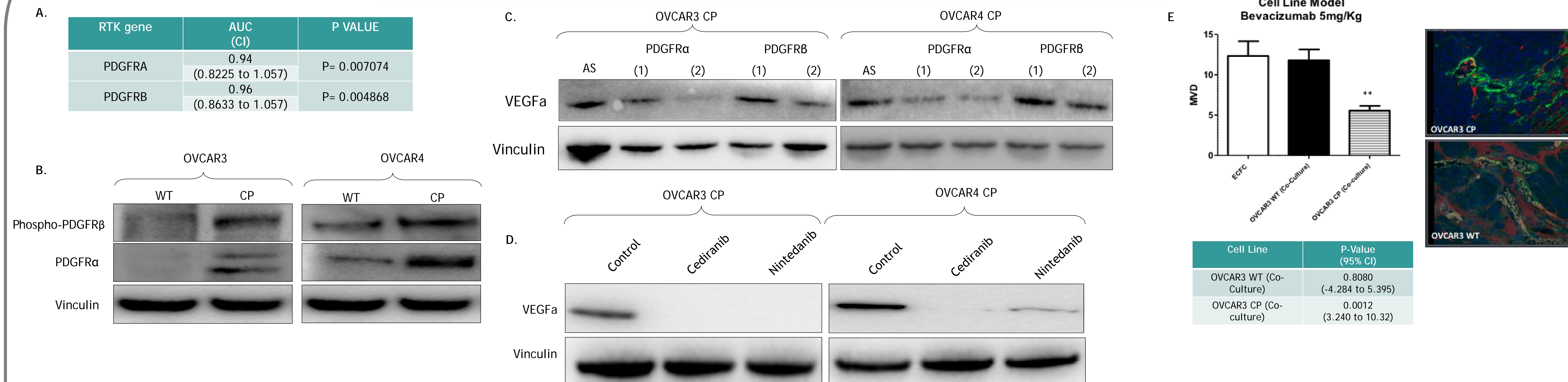
A. Review of 11 phase III clinical trials to determine the effect of antiangiogenics in the management of advanced ovarian cancer. B. Analysis of 4 phase III clinical trials to determine the effect of bevacizumab and Trebananib in platinum sensitive and resistant ovarian cancer. C. 12 paired patient samples pre and post platinum-based therapy. Quantification of MVD by CD31 IHC in the paired samples demonstrates that the post-platinum patient treatment samples have a higher MVD relative to their paired treatment-naïve pair (paired t-test; p-value: 0.0001).

II. Platinum therapy selects for an angiogenic phenotype in HGSOC and is associated with upregulation of VEGF expression



A. OVCAR3 and OVCAR4 HGSOC platinum-resistant cell lines have higher basal VEGFa expression relative to their parental platinum-sensitive cell lines. B. OVCAR3 platinum resistant cell demonstrate increased microtubule length (p=0.0009) when in co-culture with the Endothelial colony forming cells (ECFC). C. In-vivo matrigel-plug assay to determine the MVD in the OVCAR3 isogenic cell lines in co-culture with ECFC. The OVCAR3 platinum-resistant cell lines have a higher MVD than the OVCAR3 platinum-naïve pair relative to the ECFC cell lines (p-value: 0.004).

III. Platinum resistance is associated with response to anti-angiogenic agents in pre-clinical model systems



A. Gene expression profiling demonstrates a higher expression of PDGFRA (P=0.007074) and PDGFRB (P=0.004868) following platinum-based chemotherapy in the pre/post-treatment paired patient samples. B. Validation of the gene expression profiling data by western blot to demonstrate the upregulation of PDGFRA and PDGFRB in the isogenic platinum resistant cell line model (OVCAR3 and OVCAR4). C. siRNA of PDGFRA and PDGFRB in the isogenic platinum resistant cell line model (OVCAR3 and OVCAR4) leads to downregulation of VEGFa expression. D. Western blot following treatment of the OVCAR3 cisplatin-resistant and OVCAR4 cisplatin-resistant cells with Cediranib (VEGFR1-3 and PDGFRA/β inhibitor) and Nintedanib (VEGFR1-3, PDGFRA/β FGFR1-3 inhibitor) leads to downregulation of expression of VEGFa. E. In-vivo matrigel plug assay illustrating that bevacizumab has specificity for the OVCAR3 platinum-resistant cell line which is demonstrated by a reduction in MVD (p-value: 0.0012). Bevacizumab has no impact on MVD in the OVCAR3 platinum-naïve cell line (p-value: 0.8080).

Conclusions

1. The clinical and pre-clinical data discussed has potentially significant clinical implications in the management of treatment-relapsed HGSOC
2. Platinum-resistance (innate and acquired) in relapsed HGSOC is an indicator for response to anti-angiogenics
3. We have found that relapse post platinum-therapy in HGSOC is associated with an angiogenic biology
4. This clinical and pre-clinical data supports the use of anti-angiogenic agents in the first and second line setting in patients with innate and acquired resistance to platinum therapy, respectively

Acknowledgements:
All this work would not be possible without the dedication of patients and their families who have consented to take part in the primary cell generation study. This work was supported by Invest NI through the European Sustainable Competitiveness Programme 2007-2013, European Regional Development Fund (ERDF) and Almac Diagnostics Research and Development Department. The samples used in this research were received from the Edinburgh Cancer Research Centre. Ethical approval was obtained from Lothian Local Research Ethics Committee. (Ref: 07/S1102/33).

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