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A high throughput platform for the identification of inhibitors of integrin-mediated TGF β activation

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Integrins and Transforming Growth Factor beta (TGF β) regulate multiple cellular processes including adhesion, migration, proliferation, extracellular matrix (ECM) homeostasis and epithelial-mesenchymal transition. Integrin-mediated TGF β activation has been indicated in pathological conditions including fibrogenesis and immune suppression. Macrophages are a major cellular source of TGF β . Alteration of macrophage phenotypes and aberrant activation of TGF β lead to several fibrotic and immune diseases therefore inhibitors of TGF- β activation have potential values in multiple therapeutic areas. We have developed *de novo Homogenous Time Resolved Fluorescence* 1536-well binding assays using multiple integrins to identify inhibitors and to support mechanism of action studies. Further, we have developed a high throughput Detroit 562 cell co-culture assay to measure TGF β activation. In conjunction with these studies, we have also utilized a macrophage polarization model and developed cytokine and mRNA profiling assays to explore potential pharmacodynamic and/or disease biomarkers. These studies offer a novel platform to identify novel inhibitors of TGF β signaling associated with specific subsets of integrins.

Biography

Ruihua Chen has worked in the pharmaceutical and biotechnology industry for 18 years in the capacities of discovery and translational research. From the diverse roles he has played and various therapeutic areas he has worked on, he has come to a sense that drug targets can be shared in many diseases. While the specific interactions and pathways point to the direction of making certain biologics or chemotypes, translation is the key to the successful making of a drug to cure a disease. He is currently working in the areas of immunology and oncology, and one of his passions is to maximize the values of compounds by finding uses outside these areas.

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