## Multivariate Perturbation of a Growth Factor-Cytokine Signalling Network Reveals Complex Systemic Responses in Glioblastoma Cells

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Abstract—Glioblastoma cells can evade **TRAIL-induced** apoptosis through various strategies involving the growth factor-activated **MEK-MAPK/ERK** PI3K-Akt/PKB and pro-survival signalling cascades. Although these signalling cascades have been studied extensively, our understanding of how they interact and participate in modulating apoptosis as part of a dynamic cell-wide network of signalling limited proteins is by traditional univariate experimental paradigms. Here. we study three human glioblastoma cell lines with differential response to TRAIL-induced apoptosis: LN229 (resistant), T98G, and A172 (both susceptible). We show that differential TRAIL susceptibility in these cell lines is unrelated to expression levels of agonist (DR4 and DR5) or antagonist (DcR1, DcR2, and OPG) receptors for TRAIL and thus TRAILinduced apoptosis in these cell lines is modulated at the intracellular signalling level. Serum, comprising multiple factors that regulate cellular activity, enhances TRAIL resistance in T98G but not LN229 and A172 cell lines. This protective effect against **TRAIL**induced apoptosis is recapitulated by the prototypical survival factor PDGF in T98G cells. Univariate inhibition of cell survival signalling cascades with

**U0126** MEK inhibitor and PI3K inhibitor LY294002 sensitized T98G cells to TRAIL but did not abrogate PDGF-mediated protection. However, further perturbation with inhibitors in combinatorial and multivariate a manner reveal synergistic effects and complex systemic responses which may be a basis for uncovering novel insights into the regulation of TRAIL-induced apoptosis.