

Modelling wnt signaling and colon cancer

Antony W Burgess

Melbourne Branch, Ludwig Institute for Cancer Research, Australia, 3050

MOST colon cancers are associated with the truncation of the tumor suppressor gene *apc*. Although the cell physiology and biochemistry associated with *apc* are still far from clear, initial measurements indicated that *apc* regulates the intracellular concentration of β -catenin and consequently, its sub-cellular distribution. Indeed the canonical wnt signaling involves receptor mediated inhibition of GSK3 β , a reduction in phosphorylation of β -catenin, and so a reduction in *apc*/axin mediated proteosomal degradation of phospho- β -catenin, and a consequential increase in cytosolic and nuclear β -catenin. Many of the initial experimental reports have studied transfected proteins and cell lines not closely related to colonic cells. We have studied the concentrations and distribution of E-cadherin, β -catenin (and its phospho-isoforms), GSK3 β , axin and *apc* in several epithelial and colonic cell lines. We are measuring the responses of these proteins to wnt signaling. Our initial data indicate wnt induced changes to the E-cadherin/ β -catenin complex at the membrane and relocalization of a sub-population of β -catenin to cytosolic structures. In colonic epithelial cells we have not detected a significant wnt induced change in the cellular levels of β -catenin nor an increase in destruction of β -catenin in the absence of wnt signaling. Significant wnt induced increases in the nuclear localization of β -catenin only occur in the absence of cell surface cadherins and involve processes other than the truncation of *apc*. We are developing a quantitative model to simulate the time dependent effects of wnt signaling and *apc* truncation on the concentration and distribution of β -catenin isoforms in normal and neoplastic colonic cells.