

# Understanding Oncogenic Transformation in Colon Cancer

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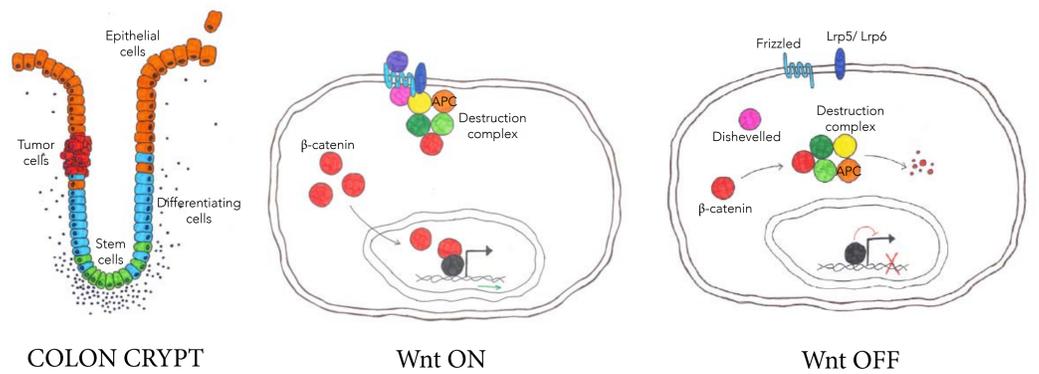
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## Dysregulation of the Wnt/ $\beta$ -catenin Signaling Pathway Can Lead to Tumor Formation

The Wnt/  $\beta$ -catenin signaling pathway is important for **stem cell maintenance** and **development**. Dysregulation of Wnt signaling can lead to **tumor formation** and cancer.

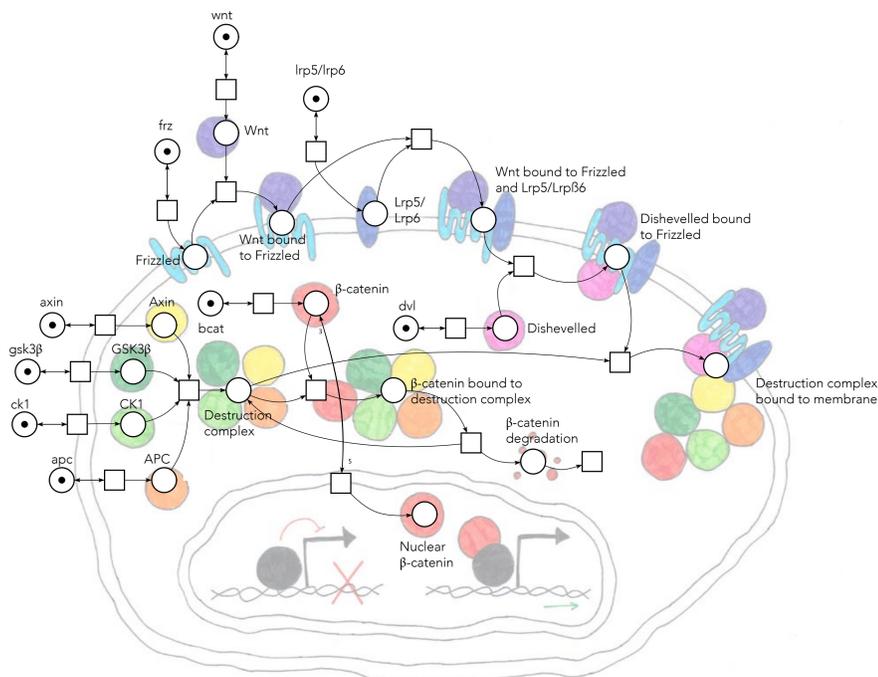
A **petri net model** of the Wnt/  $\beta$ -catenin signaling pathway was constructed in order to better **understand** the mechanism of **oncogenic signals**. The goal of this study is to:

**Recapitulate pathway responses from different perturbation of the Wnt/  $\beta$ -catenin signaling pathway**



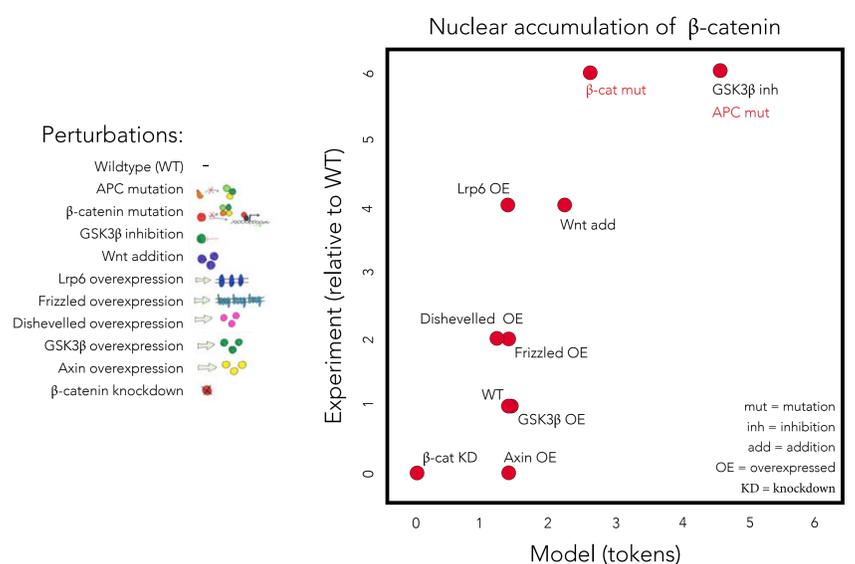
Stem cells are located in the base of the **colon crypt**, with active **Wnt signaling**. As the cells proliferate and differentiate they migrate towards the **top** where there is **no Wnt signal**. Progression into **tumor cells** can be caused by **oncogenic signaling**. The signaling pathway is active in stem cells (**Wnt ON**) and  $\beta$ -catenin translocates to the nucleus and **activates transcription** of  $\beta$ -catenin target genes. In epithelial cells (**Wnt OFF**)  $\beta$ -catenin is **degraded** in the cytoplasm by the destruction complex. Mutations in adenomatous polyposis coli (**APC**) and  **$\beta$ -catenin** are found in **~80** and **~15%** of colon cancer patients, respectively, and are among the most common oncogenes. Abberant Wnt signaling leads to both onset and progression of colon cancer.

## PETRI NET MODELING



A non-deterministic **petri net model** was constructed for the Wnt/  $\beta$ -catenin signaling pathway based on **current knowledge** of the system from literature and work performed experimentally. The places (round nodes) represent biological entities such as genes, proteins and complexes. The transitions (squared nodes) represent gene expression, protein interactions, protein degradation and protein translocation. The level of the **tokens range from 0 to 6** and represents the availability of the entity in the place (gene places are only 0 or 1). The arcs (edges) show the direction of flow and the **weights** indicate **requirements** for the transitions **from the input** place and the **effect** it has **on the output** place.

## REPRODUCING PATHWAY RESPONSES



Simulations of the model with different perturbations showed different levels of nuclear accumulation  $\beta$ -catenin, as seen by experiments. Next, recapitulation of more perturbation experiments will be done to adapt the model further. Further, **double experiments** will be done for **validating prediction** by the model. The model will then be used to explain **how  $\beta$ -catenin reaches steady state** in the different experiments.

It is also of interest to gain more insight into the regulation mechanism of the  **$\beta$ -catenin target genes** (e.g. **Axin2** and **Lgr5**) and the mechanism in which the **Human cytomegalovirus (HCMV)** disrupts the Wnt/  $\beta$ -catenin signaling pathway.

The extended model will hopefully lead to greater understanding of **how oncogenic signals propagate** through the network from disrupting the Wnt/  $\beta$ -catenin signaling pathway to transcription of target genes and regulation of gene transcription.