

Cellular Motility: A DIY Exercise

The information below has been extracted from Wikipedia for discussion only; not as a scientific resource.

“Adenomatous polyposis coli (**APC**) also known as deleted in polyposis 2.5 (DP2.5) is a protein that in humans is encoded by the **APC gene**.

The APC protein is a negative regulator that controls beta-catenin concentrations and interacts with E-cadherin, which are involved in cell adhesion.”

[https://en.wikipedia.org/wiki/Adenomatous_polyposis_coli#:~:text=Ade
nomatous%20polyposis%20coli%20\(APC\)%20also,are%20involved%20
in%20cell%20adhesion.](https://en.wikipedia.org/wiki/Adenomatous_polyposis_coli#:~:text=Ade%20nomatous%20polyposis%20coli%20(APC)%20also,are%20involved%20in%20cell%20adhesion.)

The following is provided for use as a DIY exercise based on verified science.

Cell Alignment: For Explanation and Discussion	
TNF-Alpha: TGF- Alpha (Calnexin) Density (CD-4)	
Calcium - threonine - magnesium (BRCA1)	p16
Calcium - serine - magnesium (BRCA2)	p18
Calcium - cysteine - magnesium (BRCA3)	p19
TNF-Beta: TGF- Beta (Calmodulin) Motility (CD-8)	
Calcium - phenylalanine - magnesium (HRas)	p21
Calcium - tyrosine - magnesium (KRas)	p27
Calcium - tryptophan - magnesium (NRas)	p57
TNF-Gamma: TGF- Gamma [VEGF] (Calcineurin) Modulatory Enzyme (CD-25)	
Iron - serine - Manganese	
Iron - cysteine - Manganese	
Iron - threonine - Manganese	

These are examples of the “enzymes” that have evolved with various designations; e.g. AKT, mTOR, PTEN, NF-kB, and MYC.

Any biologically astute individual can utilize bioinformatic search to correlate APC to cellular motility and calmodulin with supporting links from markers such as CD8, p21, KRas as well as TNF β and TGF β .

The Wikipedia comment refers to E-cadherin and cell adhesion. The same DIY exercise with E-cadherin as a search factor will result in results for density and correlation with the markers calnexin, CD4, p16 as well as TNF α and TGF α .

<https://phys.org/news/2019-11-achilles-heel-tumor-cells-inhibiting.html>

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Achilles heel of tumor cells: Inhibiting the mutated genes that keep cancer cells alive

by Robert Emmerich, Julius-Maximilians-Universität Würzburg

In almost all cases of colon cancer, a specific gene is mutated—this offers opportunities to develop broadly effective therapeutic approaches. Research teams in Würzburg have taken this a step further.

In 90 percent of all cases of colon cancer, the tumor cells have one thing in common: the APC gene is mutated. Research groups at Julius-Maximilians-Universität (JMU) Würzburg in Bavaria, Germany, were looking for targets in these cells that could be used to destroy the cells.

"We wanted to find genes that are only important for the survival of cells with APC mutations, but not for healthy cells," explains Dr. Armin Wiegering, head of a junior research group at the JMU Biocentre and physician in surgery at Würzburg University Hospital.

The search for a needle in a haystack was successful. The research teams now report this in the journal *Nature Cell Biology*. If they inhibited the gene called eIF2B5, the mutated colon cancer cells died of programmed cell death—a self-destruction program with which the organism normally disposes of damaged or aged cells. Healthy cells, on the other hand, were able to cope with the inhibition of the gene without any impairment.

Possible point of attack for treatment

"We have thus identified a very specific Achilles heel of APC-mutated tumors," says Professor Martin Eilers, cancer researcher at the Biocentre. "We now know of a site where newly developed antitumor drugs might be able to have a very targeted effect."

The efficacy of an eIF2B5 inhibition was shown in animal experiments. If the gene is not fully active in mice, they do not develop colon cancer so quickly and survive much

longer if they do. The researchers also experimented with organoids. These are miniature tumors that are cultivated in the laboratory from the cancer tissue of patients. If the amount of eIF2B5 was reduced, the organoids died.

Further genes will be investigated

Next, the researchers want to investigate further genes in colon cancer cells—because eIF2B5 is only one of five subunits of the larger eIF2B gene complex. "We also want to characterize the other subunits and see if we can also find a specificity here," Wiegering announces. We will then establish a method to degrade eIF2B5 in cancer cells. If this is successful, it might lead to a new option for therapies.

Colon cancer is one of the three most common tumor diseases. About six percent of all people in Germany fall ill with it in the course of their lives; about half of those affected die from the consequences of the tumor. Since more than 90 percent of all colon tumors show an APC mutation, research at JMU could lead to a very broad, new therapeutic approach.