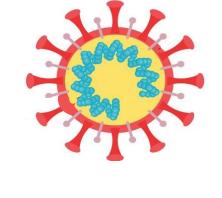


Addendum 4<u>a</u> Viruses





How To Defeat Viruses

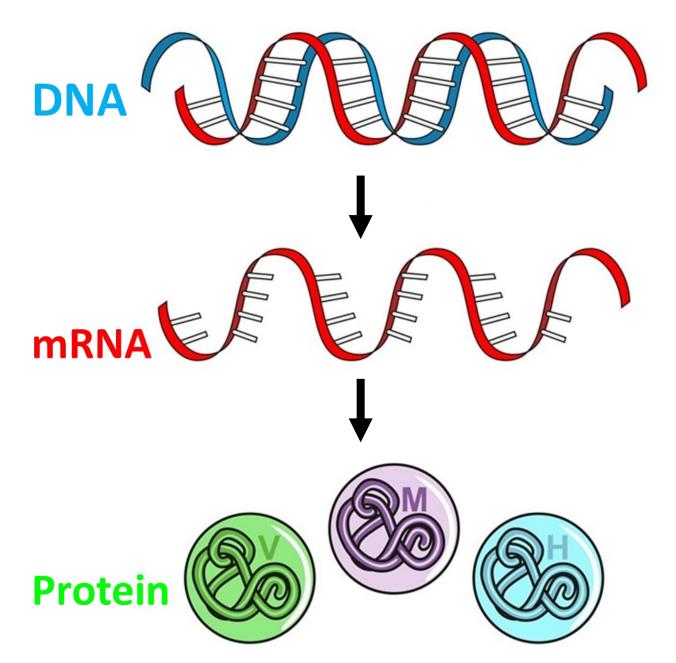
Presenter:

Dr. James Summerton Ph.D.

30 November 2022

This presentation was given to the Academy of Lifelong Learning at Oregon State University. Please see page 36 for the presentation on LK-1

Information flow in Cells

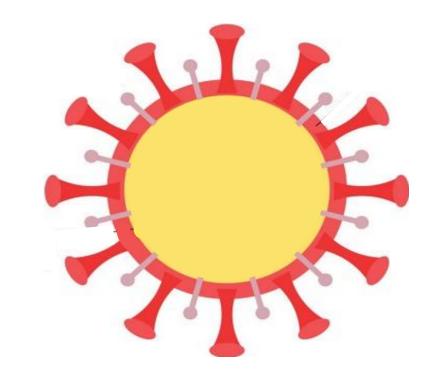


How to defeat COVID-19 and any other viruses

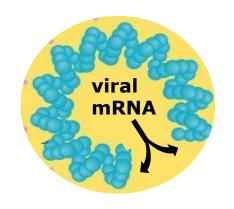
1) Select a suitable viral target

2) Develop a safe, effective and affordable therapeutic to prevent function of that target

The outer structures of COVID-19 serve to carry the viral payload (mRNA) from an infected cell to a new cell of the host, or to a new host.

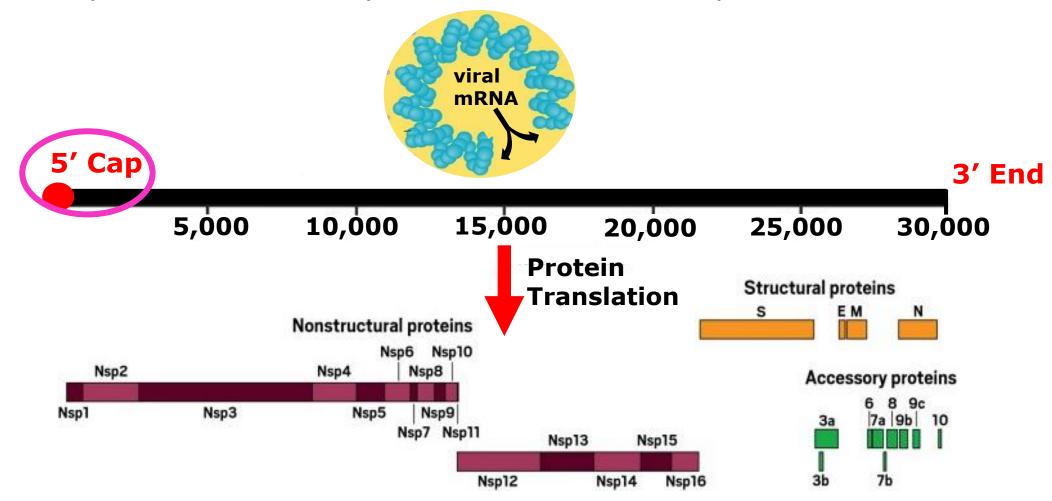


The payload (viral mRNA) comprises the genetic information for making <u>all</u> 29 viral proteins for viral function and reproduction



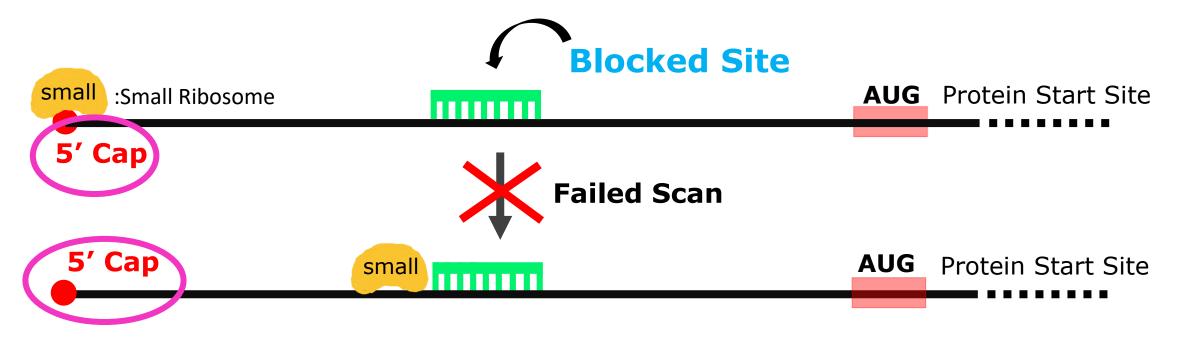
The 29 COVID-19 Proteins

The very-long viral mRNA (30,000 genetic letters) uses the cell's own protein synthesis machinery to make the virus' 29 proteins.



Targeting a Special Region: 5' cap to AUG start site for protein synthesis

Effective blocking of a site between the 5' cap site and AUG start site can prevent **ALL** function and reproduction of that COVID-19 mRNA.



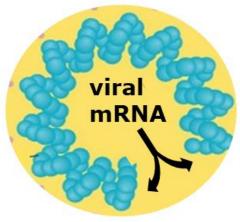
The virus remains **completely** nonfunctional as long as the blockage remains in place

Benefits of Targeting the Special Region

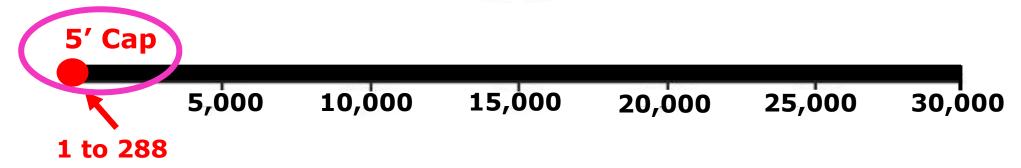
- Prevents synthesis of all 29 covid-19 proteins
- Prevents replication of entire covid-19 virus
- Sequence in special region is well conserved over decades

To Summarize

A) Select a suitable viral target. To date, we have settled on targeting the 30,000-genetic-letter viral messenger RNA (mRNA).



B) Target selection was further narrowed to the special region starting at the 5' cap and ending at genetic letter 288



Optimal Target Region:

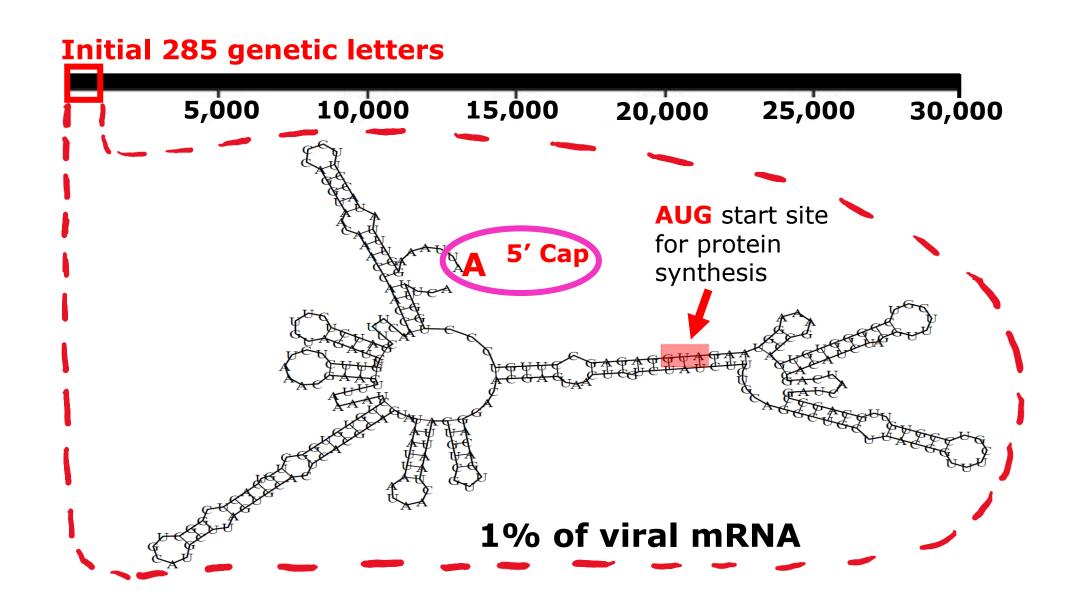
5' Cap to AUG start site

Simplified Optimal Target Region:



~1% of viral mRNA

Actual Optimal Target Region



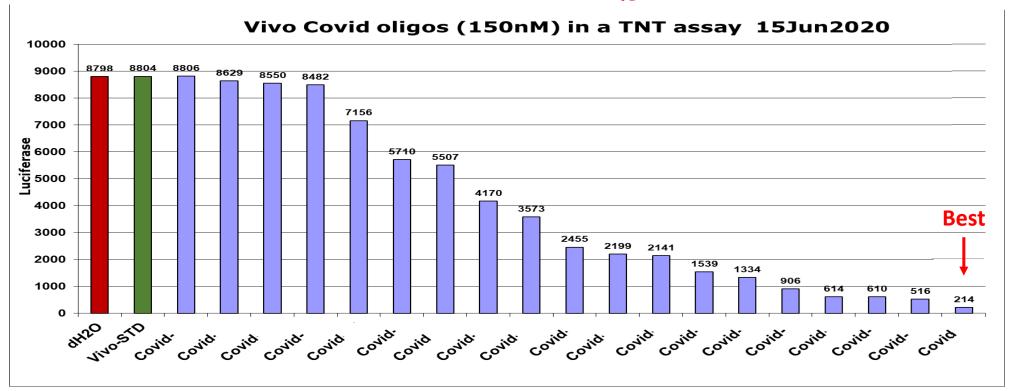
All Targetable Sites Are **NOT** Equal

5,000 10,000 15,000 20,000 25,000 30,000

AUG start site for protein synthesis

1% of viral mRNA

Preliminary target blockage results



How To Defeat COVID-19

- 1. Select a suitable covid-19 target
- Develop a safe, effective, affordable therapeutic to block the function of that selected covid-19 target

Structural design of covid-19 therapeutic

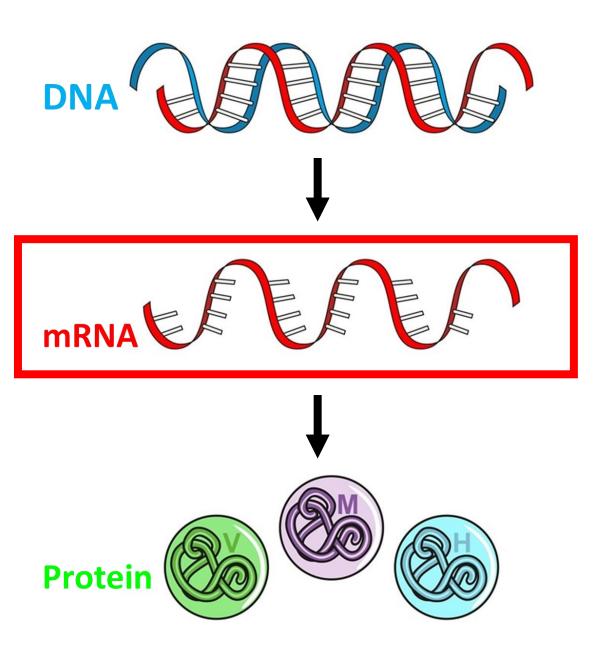
Morpholino

Precision blocking agent for selected RNA target

Delivery Component

To deliver Morpholino from the blood to the cytosol of cells

Information flow in Cells



Brief introduction to messenger RNA (mRNA)

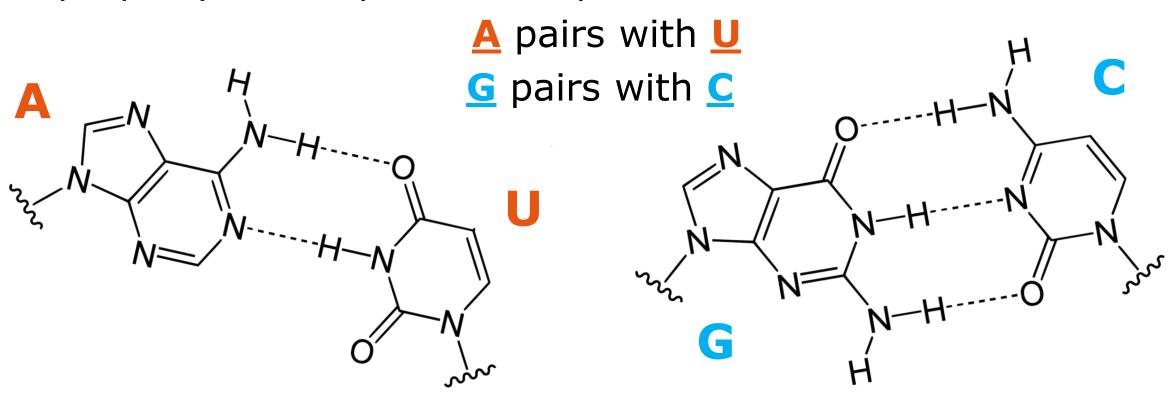
An mRNA, such as the one in covid-19, is made up of a long string of 4 genetic letters (A, C, G, U) in a specific sequence unique to the particular mRNA

The sequence of genetic letters in an mRNA can be read by a ribosome (the cell's protein synthesis machine) to assemble a corresponding protein that serves to carry out one or more functions in the body.

If the mRNA is from a **virus** the resultant protein(s) can instead carry out one or more nefarious functions for the invading virus.

Watson/Crick Pairing: A key interaction for all life as we know it

As noted earlier, a messenger RNA (mRNA) that codes for a specific protein is composed of a long sequence of 4 genetic letters. Those genetic letters have the special property of complementarity wherein:



<u>Antisense Therapeutics Strategy:</u> <u>Block a selected sequence in RNA</u>

In principle, specific destruction or long-term blocking of a selected mRNA can alter, halt or cure a wide variety of diseases and conditions.

An established technology for very specifically inactivating or altering a selected mRNA is commonly referred to as the "antisense therapeutics strategy."

Antisense Therapeutics

The antisense therapeutics strategy exploits this Watson/Crick pairing property to afford simple and efficient design and production of high-specificity **antisense** blocking agents for selectively blocking any of a wide variety of mRNA sequences (called **sense** strands because they carry the genetic information for making proteins).

Very-long RNA target sequence (sense strand):

Short blocking Agent (antisense strand)

Evolution of The Antisense Therapeutics Field

1. Initial **conceptions** of antisense strategy: **1967 - 1969**

2. First papers focused on antisense therapeutics strategy: 1978

ResearchersWork Done At:Summerton & BartlettBerkeleyZamecnik & StephensonHarvard

3. First **patent** issued in antisense field: **1978**

(Summerton & Bartlett, US Patent 4,123,610)

- First company founded to pursue development of antisense therapeutics: ANTIVIRALS, Inc., founded by Summerton in 1980 (In 2012 company was renamed Sarepta)
- By the mid-1980s the antisense therapeutics field was becoming wildly popular, with a great many researchers, granting agencies, and pharmaceutical companies becoming heavily involved.
- 6. In the late-1980s venture capitalists became involved by funding start ups of 4 more antisense companies in 1987 to 1989: Gilead, Genta, Hybridon, and lastly, Isis (recently renamed Ionis)

Many Technical Challenges

In the **1980s** it was becoming clear there were many technical challenges to overcome before the seemingly-simple antisense therapeutics strategy could provide treatments for the diseases and conditions for which that strategy appeared to hold promise

Major problems in the antisense field

While there have been many problems to solve in the antisense field, the most obvious first problems were :

- a) How to **avoid rapid degradation** of short DNA (or RNA) antisense oligos.
- b) How to achieve **highly-efficient inhibition** of the targeted mRNAs.
- c) How to achieve **highly-specific inhibition** of the targeted mRNAs.
- d) How to **deliver** the large, polar antisense oligos into the proper subcellular compartment of the patient's cells.

Problems, problems, and more problems

By the mid-1980s, the antisense therapeutics field somewhat resembled the arcade game: "Whack-a-mole"

In Whack-a-mole the problem is a toy mole that pops up. Your solution to that problem is to whack that toy mole with a mallet, causing it to disappear. But when that mole disappears one or more other moles pop up. When you whack those moles other moles pop up.

It seemed in the **1980s** antisense therapeutics followed a similar path – when one problem was fixed, that fix generated new problems.

Progress at Last?

During the **1980s** dozens of antisense structural types were made and tested – but it proved very difficult to devise a single structural type that solved most or all the most obvious problems deterring development of antisense agents suitable for use in patients.

However, in the mid-1980s a research group led by Cohen at the US National Institutes of Health (NIH), in collaboration with a research group led by Zon at the US Food and Drug Admin. (FDA), developed Phosphorothioate-linked DNA oligos (S-DNA) which showed substantial promise for antisense applications.

Solutions – but with problems

The new S-DNA structural type was quickly and widely adopted by much of the antisense research community. This was because **S-DNAs** offer:

- **A) Moderate resistance to degradation –** because their pendant sulfur on each intersubunit link slows attack by degradative enzymes;
- **B) Excellent efficiency** in inhibiting their targeted mRNA because the paired mRNA/S-DNA duplex is rapidly cleaved by the cell's RNase H;
- C) a rather limited level of specificity for their targeted mRNA limited because only a low amount of sequence information is recognized by the RNase H (only 6 or 7 base–pairs of mRNA/S-DNA duplex); and,
- **D)** A **ready-made delivery** capability **–** because delivery systems developed for transfecting DNA into cells can be used to deliver S-DNAs.

Fundamental flaws in S-DNA antisense oligos

1. Why S-DNAs are plagued with so many off—target effects (causing undesired biological effects)

The pendant sulfurs on the intersubunit linkages of S-DNAs strongly interact with a wide variety of proteins – which leads to a host of serious off–target effects that have been extensively documented starting in the 1990s

The Solution: Morpholinos

Not just a simple substitution of a sulfur for an oxygen

But instead

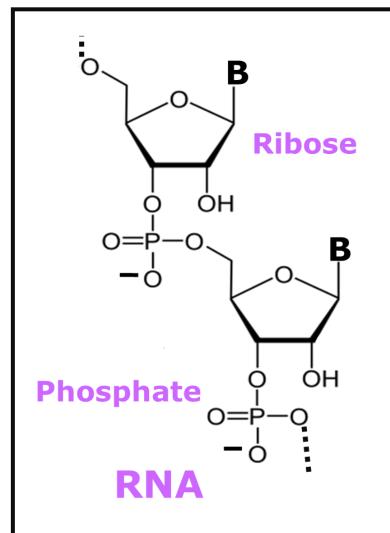
A radical re-design of genetic material

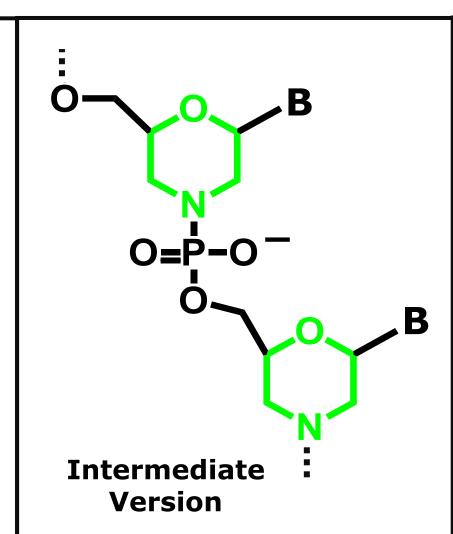
Radical Redesign of Genetic Material

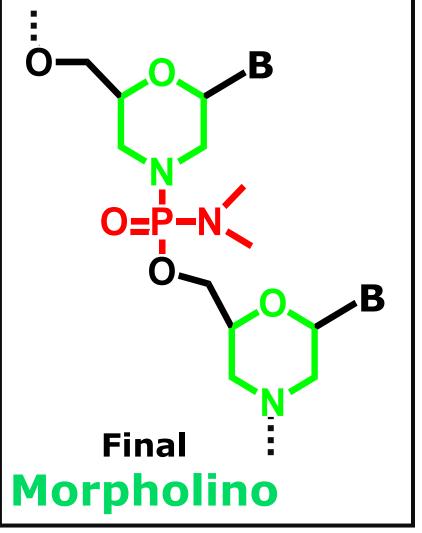
Start: 1985

B (Base): A, C, G, U

Finished: 1989







Functional Properties of "Magic Morpholinos"

- 1. Complete resistance to degradation in biological systems
- 2. Greatest sequence specificity of all antisense structural types
- 3. Highly effective without need for RNase H or RISC
- 4. Little or no non-antisense activity
- 5. Predictable targeting
- 6. Freely passes between cytosol and nucleus, and functions in both
- 7. Versatile: a) alter splicing in the nucleus
 - b) block translation in cytosol
 - c) and much more
- 8. Good aqueous solubility
- 9. Affordable: a) cheap starting materials
 - b) efficient assembly
 - c) easy workup

Morpholinos achieve these outstanding properties by virtue of their novel 6-membered morpholine backbone moieties joined by uncharged intersubunit linkages. These provide major advantages over the 5-membered ribose/deoxyribose moieties and negatively-charged intersubunit linkages characteristic of natural nucleic acids.

Our Therapeutic Designed For Preventing and Curing COVID-19

Morpholino

Precision blocking agent for selected RNA target

Delivery Component

To deliver Morpholino from the blood to the cytosol of cells

1985 - 1993

1993 - 2022

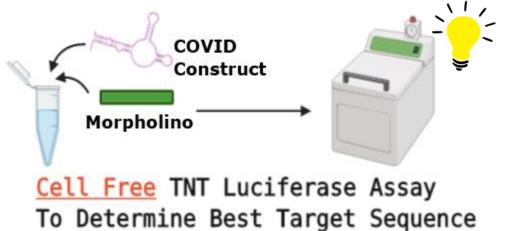
Therapeutic Applications

Morpholino

Delivery Component

By the year 2022

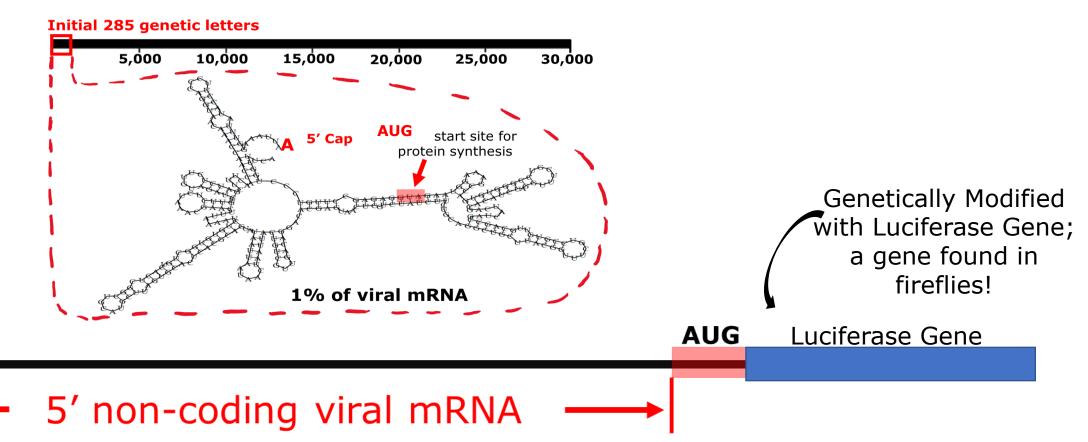
- Over 200,000 custom-sequence Morpholinos have been designed and produced for researchers around the world
- Over 11,000 scientific publications have described research with Morpholinos (Tabulated at : gene-tools.com)
- 3 FDA approved Morpholino therapeutics are being used to treat muscular dystrophy patients with more nearing FDA approval.



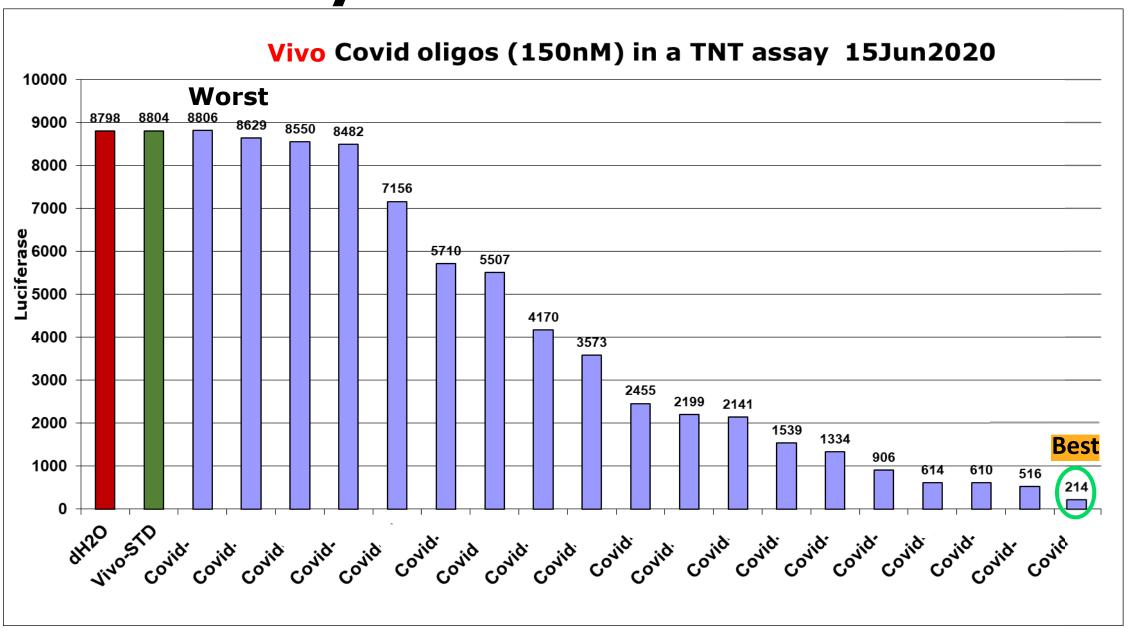
5' Cap

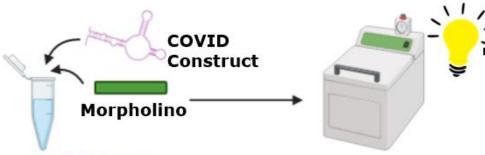
Cell Free Assay

Purpose: To determine the **BEST** targeted sequence for effective translating and blocking of Covid-19 virus.



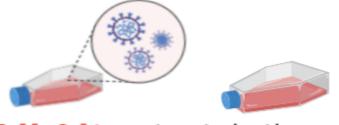
Preliminary Data From Cell Free Assay



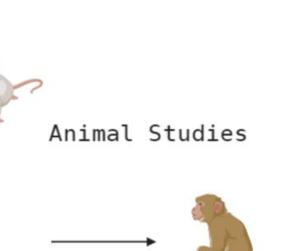


The Steps to Biological Testing

Cell Free TNT Luciferase Assay
To Determine Best Target Sequence



Cell Culture to study the efficiency of the COVID-19 target To study to efficiency of the delivery component



1. Cell Free Assay

2. Cell Studies

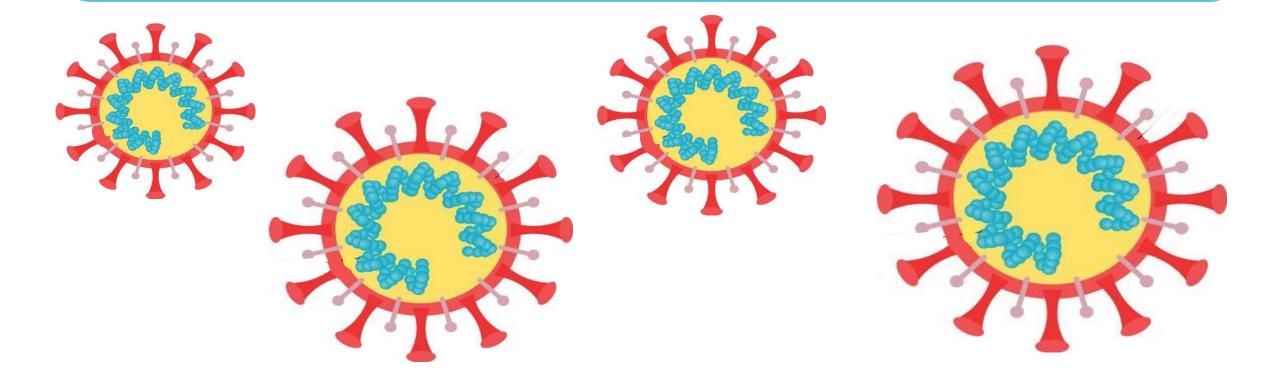
3. Animal Studies

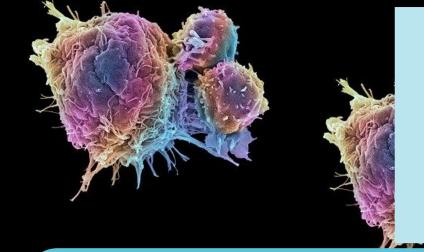
Compassionate Use

An alternative route to humans

Gilead's Remdesivir Precedent

Questions?





Addendum 4<u>b</u> Cancers



How To Defeat <u>Cancers</u>

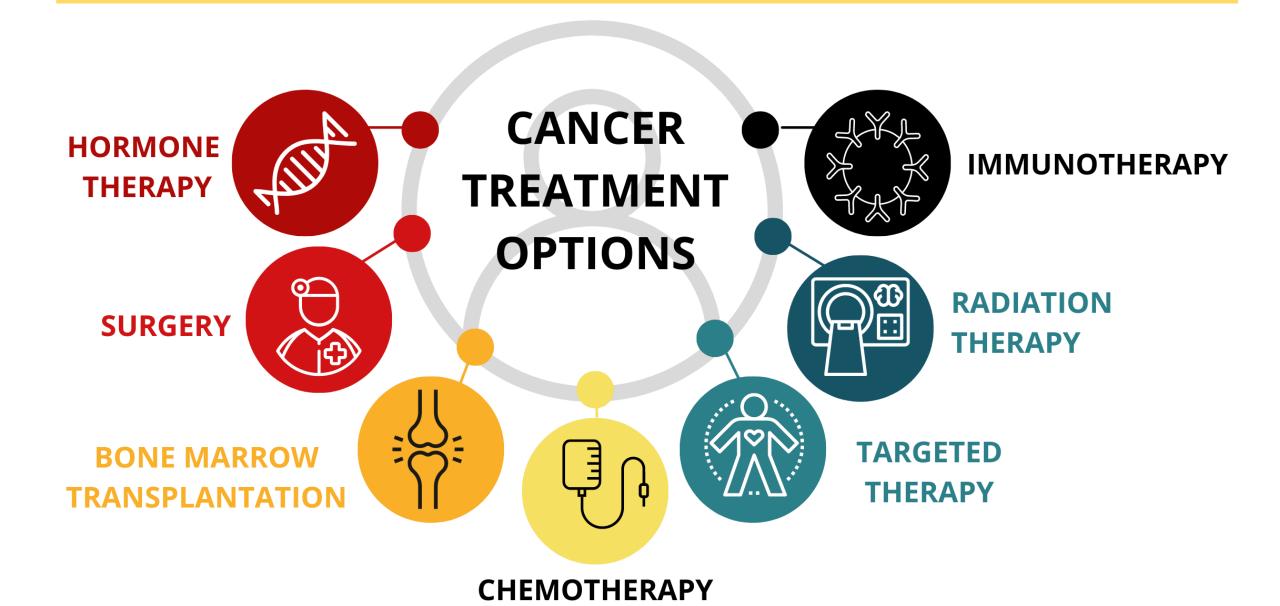
Presenter:

Lena Kinion (Ph.D. Candidate)

30 November 2022

This presentation was given to the Academy of Lifelong Learning at Oregon State University. Please see page 36 for the presentation on LK-1

Currently There Are No Reliable Cures For Cancer



How To Safely Defeat Cancer

1. Select a suitable target that will destroy the cancer

2. Avoid harm to the patient

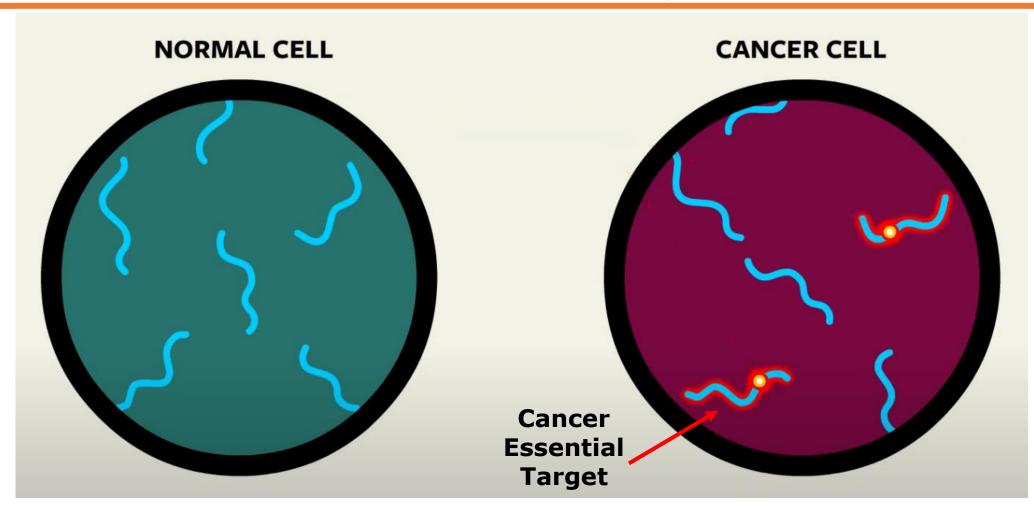
Invasion and metastasis Evading Sustaining growth proliferative suppressors signaling Reprogramming Evading energy immune metabolism destruction Enabling Resisting replicative immortality apoptosis Tumorpromoting Genomic inflammation instability Inducing angiogenesis

How To Find An Ideal Target That Will Destroy Cancer

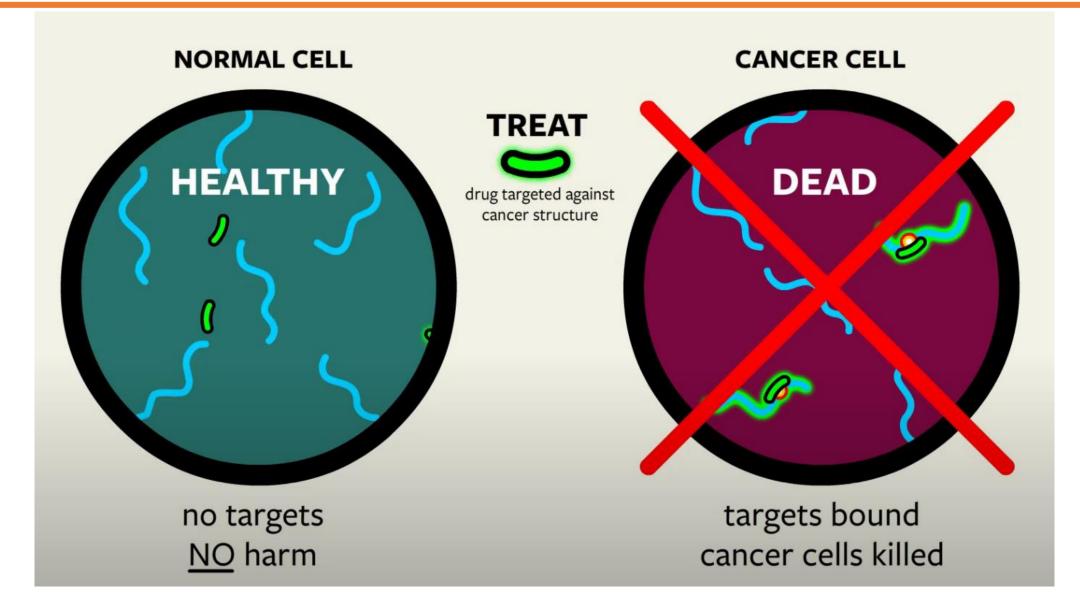
Weinberg identified multiple properties that are essential to cancers survival

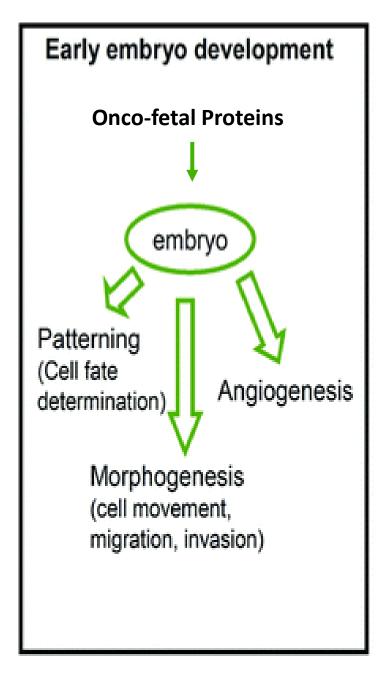


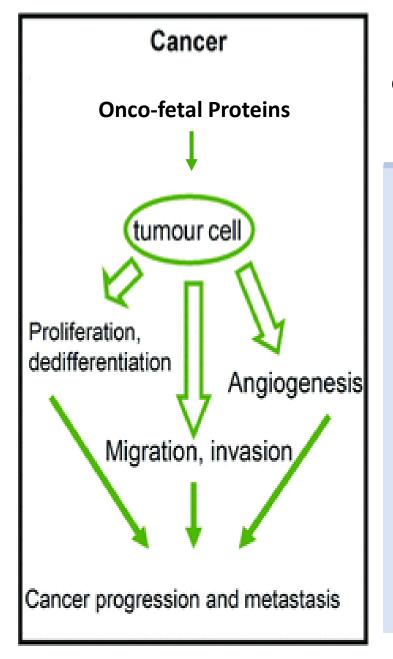
Strategy For Developing A Cure For Cancer Without Harming The Patient



Strategy For Developing A Cure For Cancer Without Harming The Patient





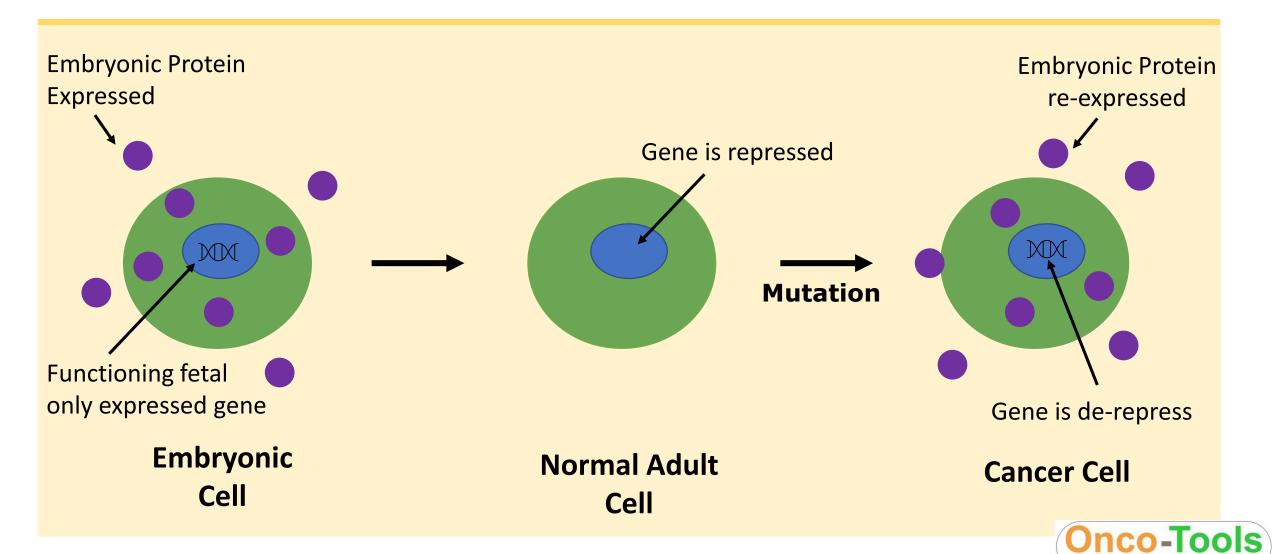


The relationship between early embryo development and tumorigenesis

- Similar signaling pathways in early embryo development and tumorigenesis.
- During tumorigenesis, these pathways are reactivated and contribute to cancer progression and metastasis.



Why Target Onco-Fetal Proteins?



Onco-Fetal Proteins

- These are proteins produced during fetal life
- They are present in high concentrations in fetuses & are disposed of after birth
- They reappear in individuals with cancer
- This demonstrates that certain genes are reactivated by mutation
- There are several oncofetal genes one being human chorionic gonadotropin-beta (hCG-b)

What is <u>Human Chorionic</u> <u>Gonadotropin (hCG)</u>

- hCG is used as a marker for at home pregnancy tests (known as pregnancy hormone)
- hCG is produced by the placenta during pregnancy
- Maintains other essential functions during pregnancy and protects the fetus from the mother's immune system



hCG Protein Structure

Identical to other Reproductive Proteins

a Subunit

β Subunit

Specific Region of the Protein

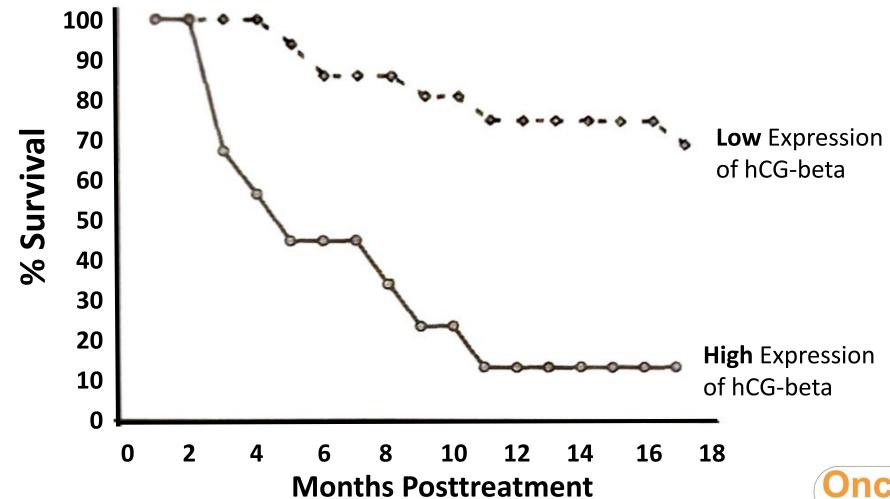


hCG-βeta has been linked to many cancers

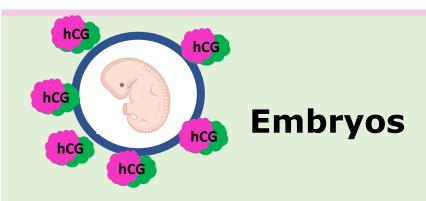
- When at home pregnancy tests were being developed, men were used as controls.
- Surprisingly, some of the controls came up positive!
- Researchers then, started looking at hCG and its influence in cancer



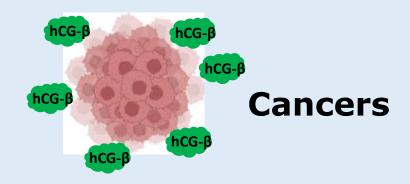
Low Prognosis For Patient's Who Express hCG-\(\beta \) In Cancers



hCG-β Activity in Humans



- Blocks programmed cell death
- Promotes <u>implantation</u> into uterus
- Hides the embryo from the <u>mother's</u> immune system
- Induces new blood vessels

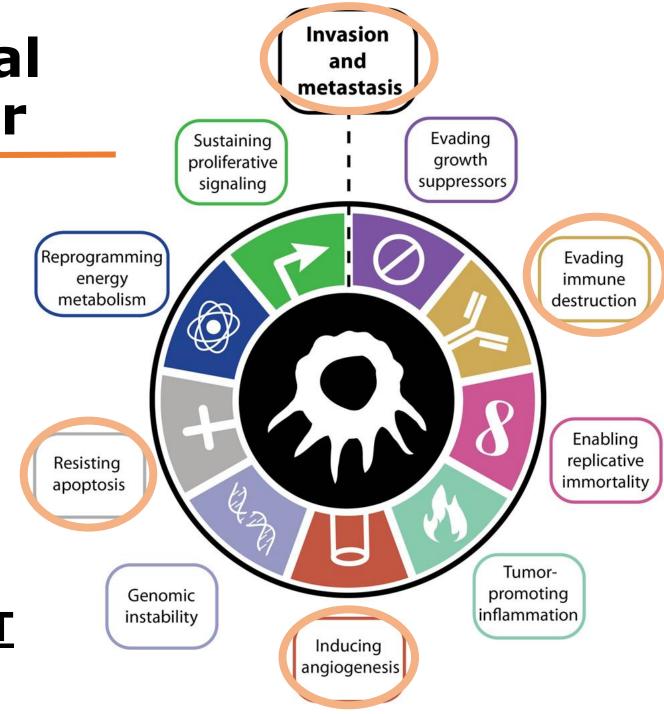


- Blocks programmed cell death
- Promotes <u>invasion</u> into normal tissues
- Hides the cancer from the <u>patient's</u> immune system
- Induces new blood vessels



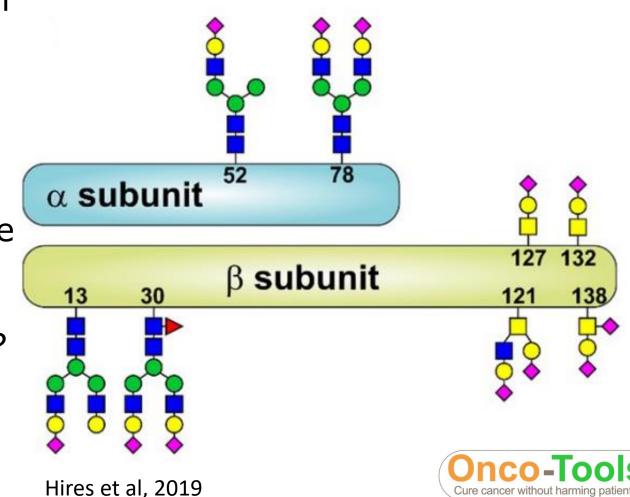
hCG-β Is The Ideal Target For Cancer

- hCG-β plays a major role in many different cancers
- Prevalent in many different cancers:
 - Prostate
 - Ovarian
 - Triple negative breast
 - Breast
 - Bladder
 - Skin, etc
- Essential to cancer but **NOT** essential in normal cell

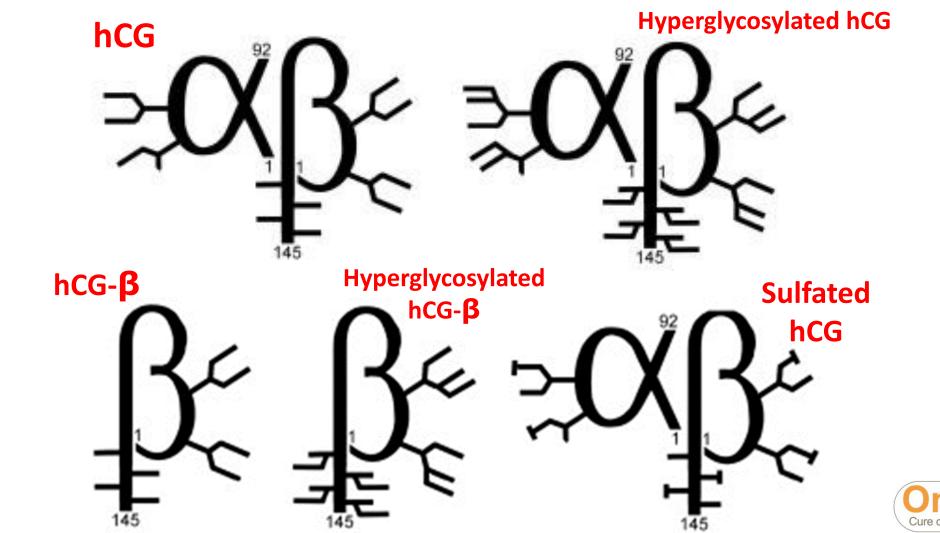


Why haven't Scientists been successful targeting hCG-\(\beta\)?

- hCG-β is a <u>very complicated</u> protein to target
- 3,000 out of 25,000 protein coding genes are druggable
- ~10% of the druggable proteins are FDA-approved drugs
- What does druggable mean to NIH?
 - Proteins with ability to bind to drug-like small molecules



Human Chorionic Gonadotropin Comes in Many Forms



Our Therapeutic Designed For Blocking Function of hCG-\(\beta \)

Delivery Component

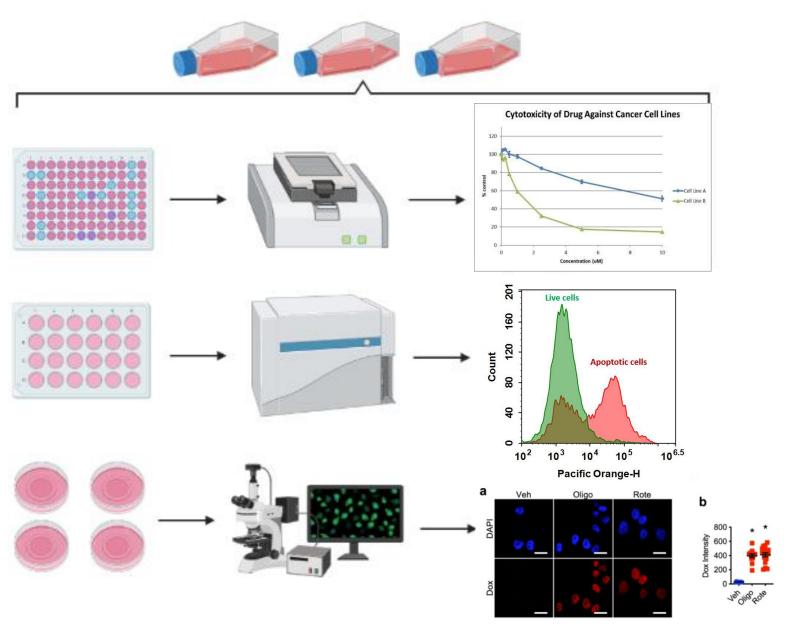
Precision blocking agent for selected target

To deliver Morpholino from the blood to the cytosol of cells

1985 - 1993

1993 - 2022

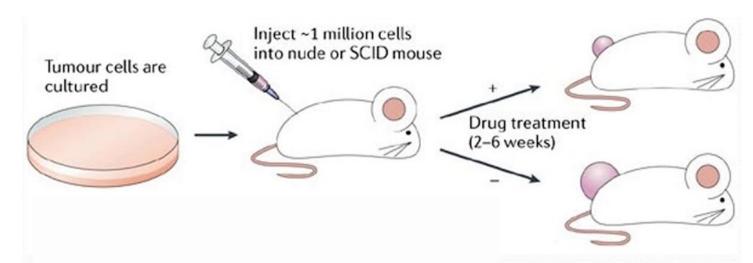
Various Cancer Types Tested



Steps to Develop a Cancer Therapeutic

- 1. Screen different treatment combinations
- 2. Identify dead cells
- 3. Validate best drug combination in several biological assays
- 4. Animal Studies





Xenograft studies can be designed to:

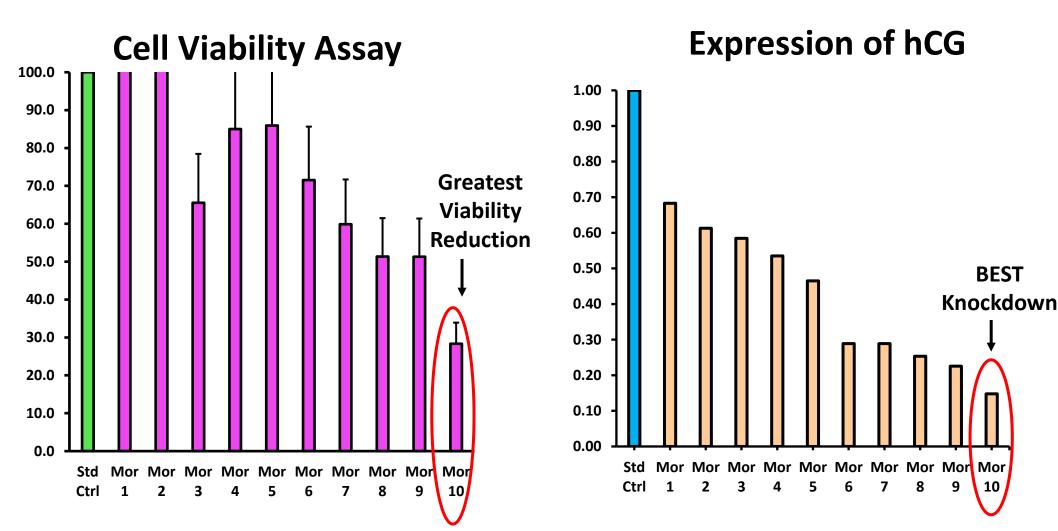
- Identify lead compounds
- Optimize dose schedules
- Identify combination strategies

Steps to Develop a Cancer Therapeutic

- 1. Screen different treatment combinations
- 2. Identify dead cells
- 3. Validate best drug combination in several biological assays
- 4. Animal Studies



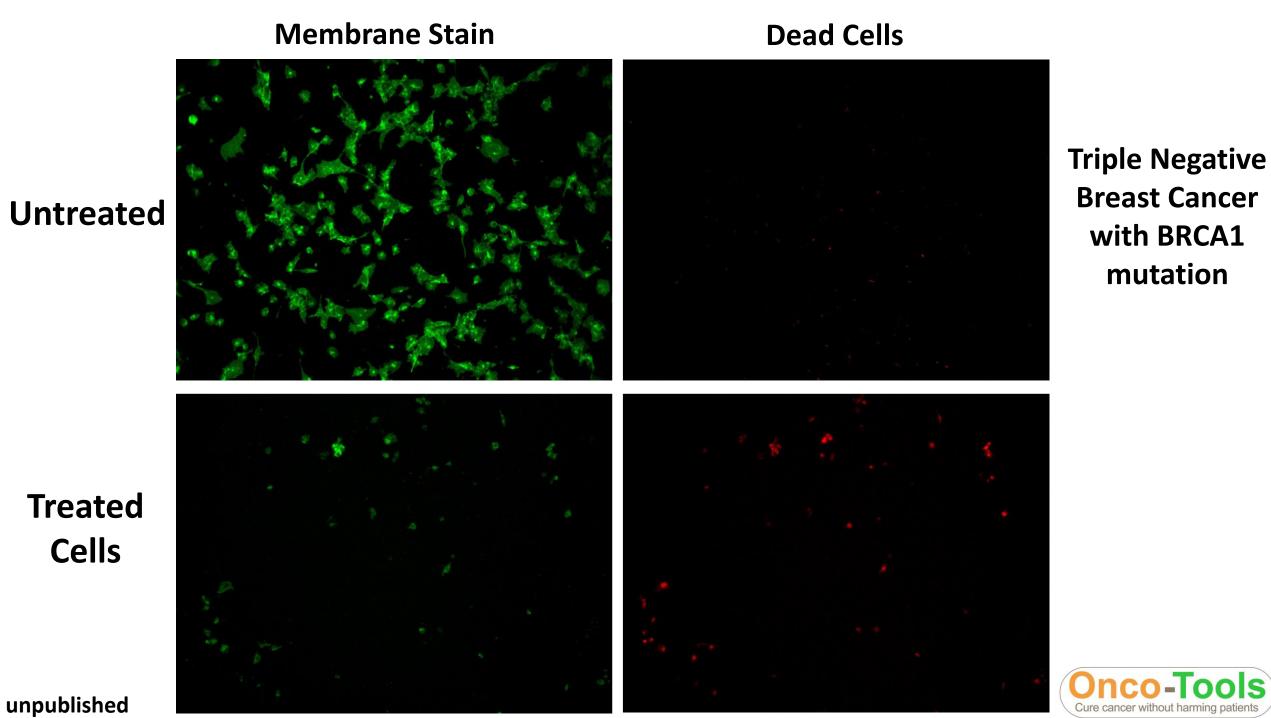
Various Morpholinos Designed to Block hCG- β Function

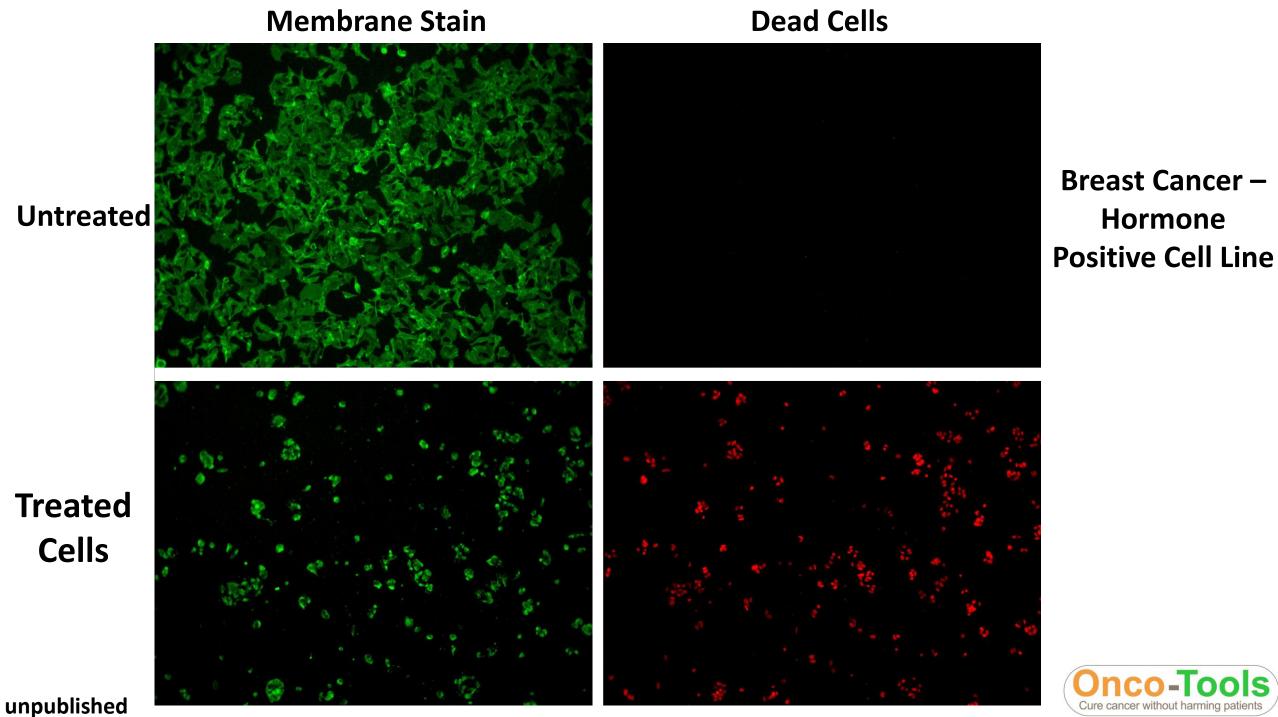


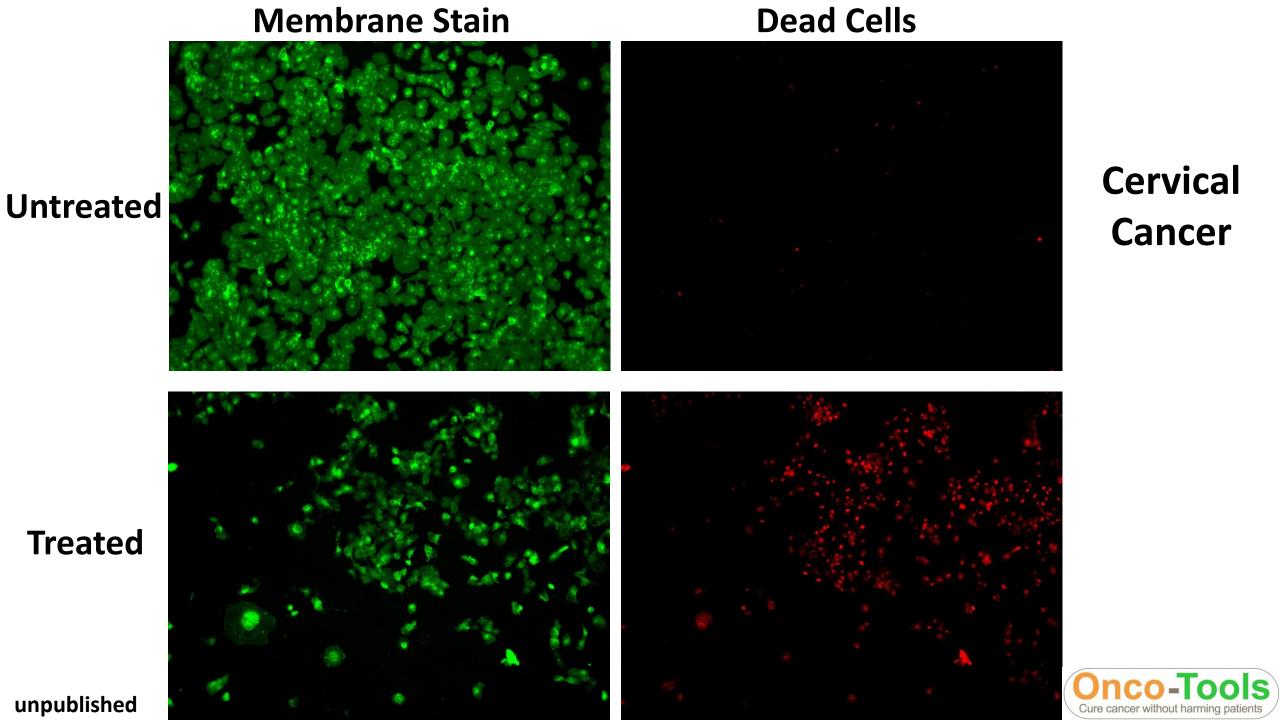
Membrane Stain Dead Cells Untreated Treated Cells

Triple Negative Breast Cancer Cell Line









Conclusion

hCG-β is a promising target for cancer therapeutics **because**:

- Only expressed in cancer, so will cause no harm to the patient
- Is ubiquitous in broad range of cancers, potentially a universal treatment for cancers
- Morpholino therapeutics allows hCG-β to go from an undruggable target to druggable target



