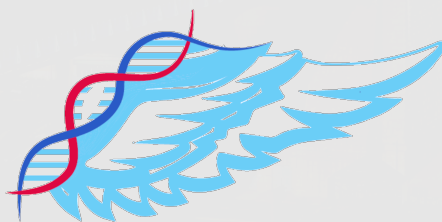


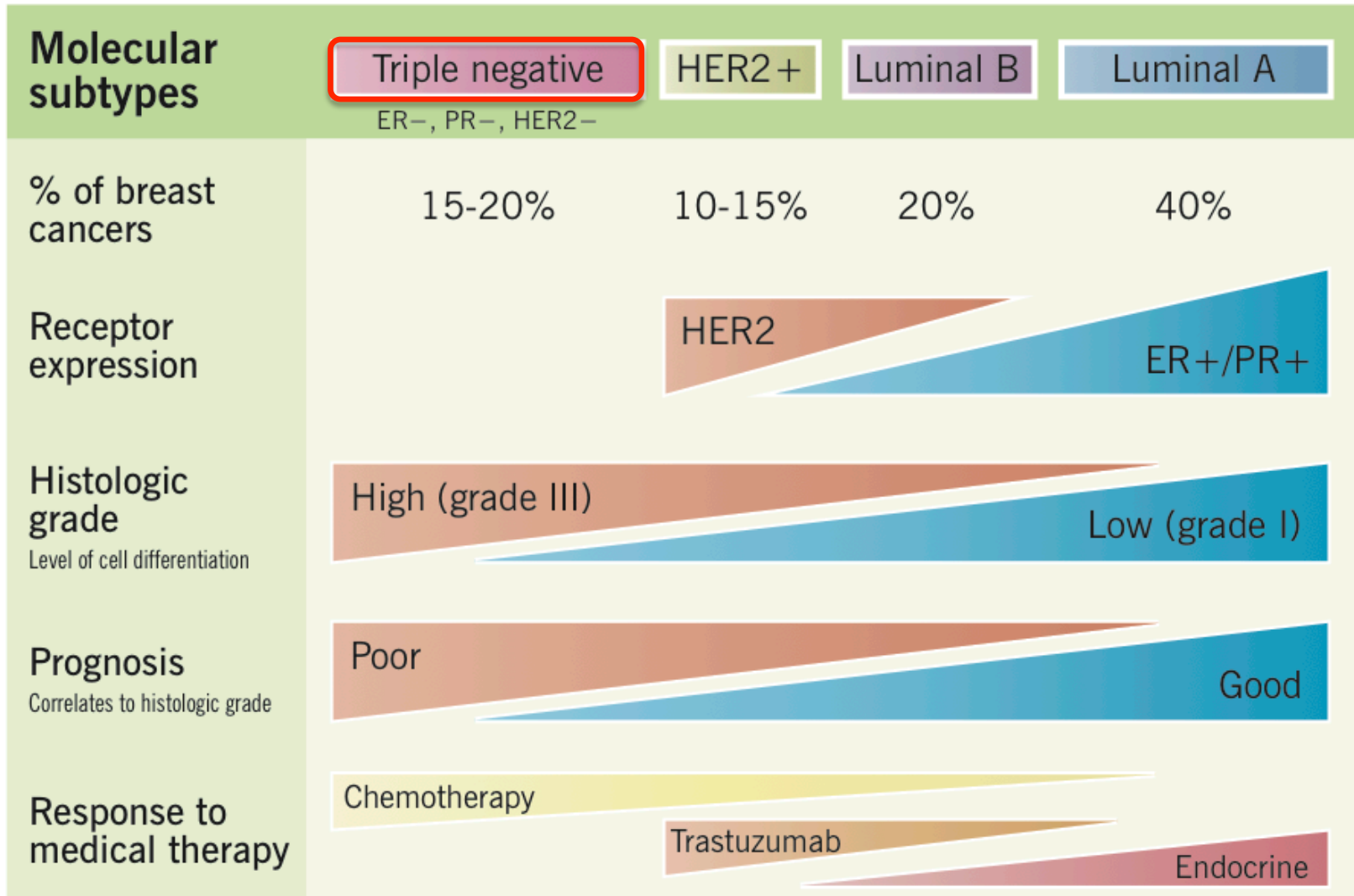
Triple Negative Breast Cancer Therapy by microRNA Blockade with PNA-peptides, without Passenger Strand Side Effects

XXII International Roundtable on Nucleosides, Nucleotides
and Nucleic Acids



Bound Therapeutics LLC

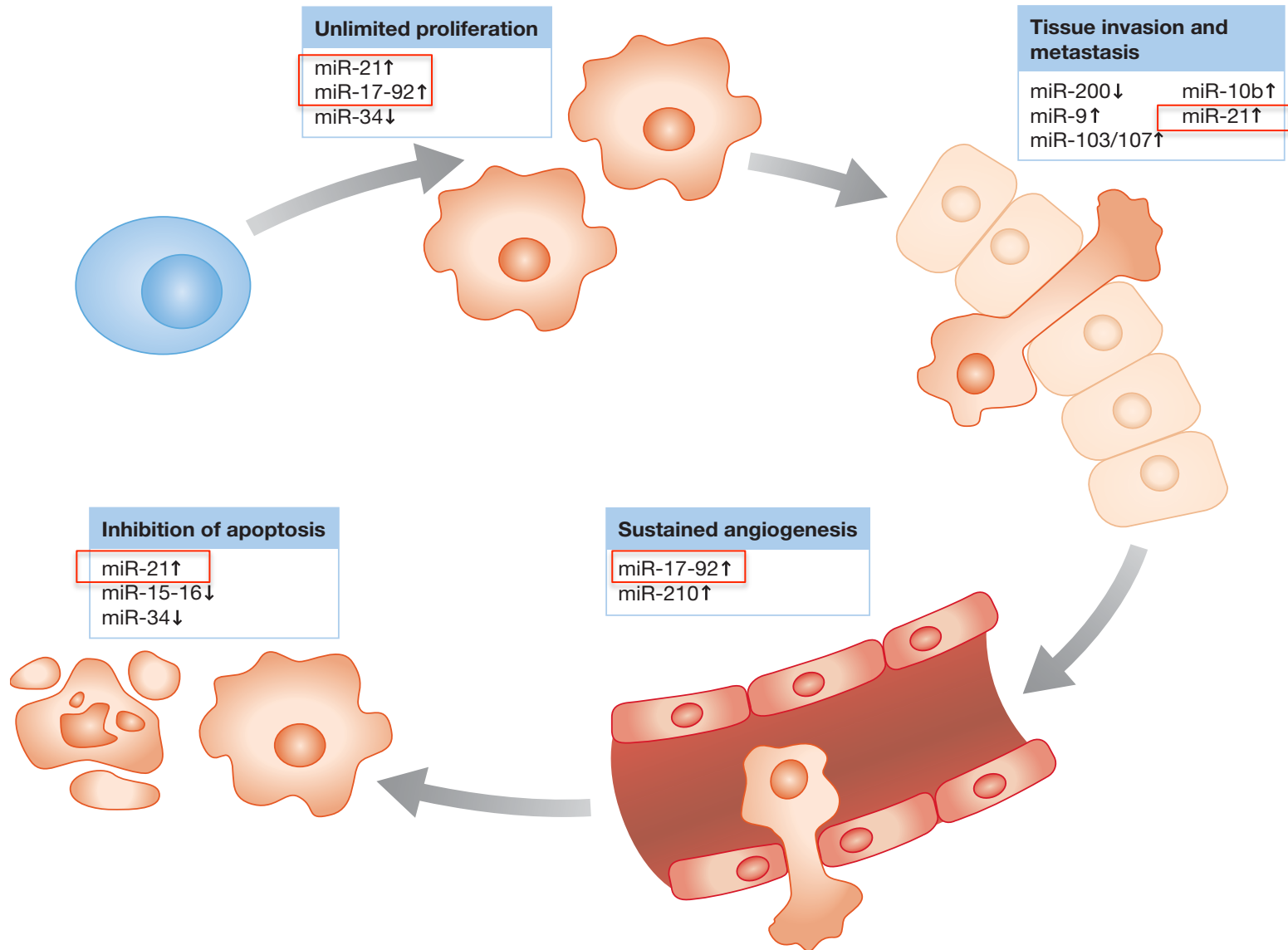
Breast cancer subtypes



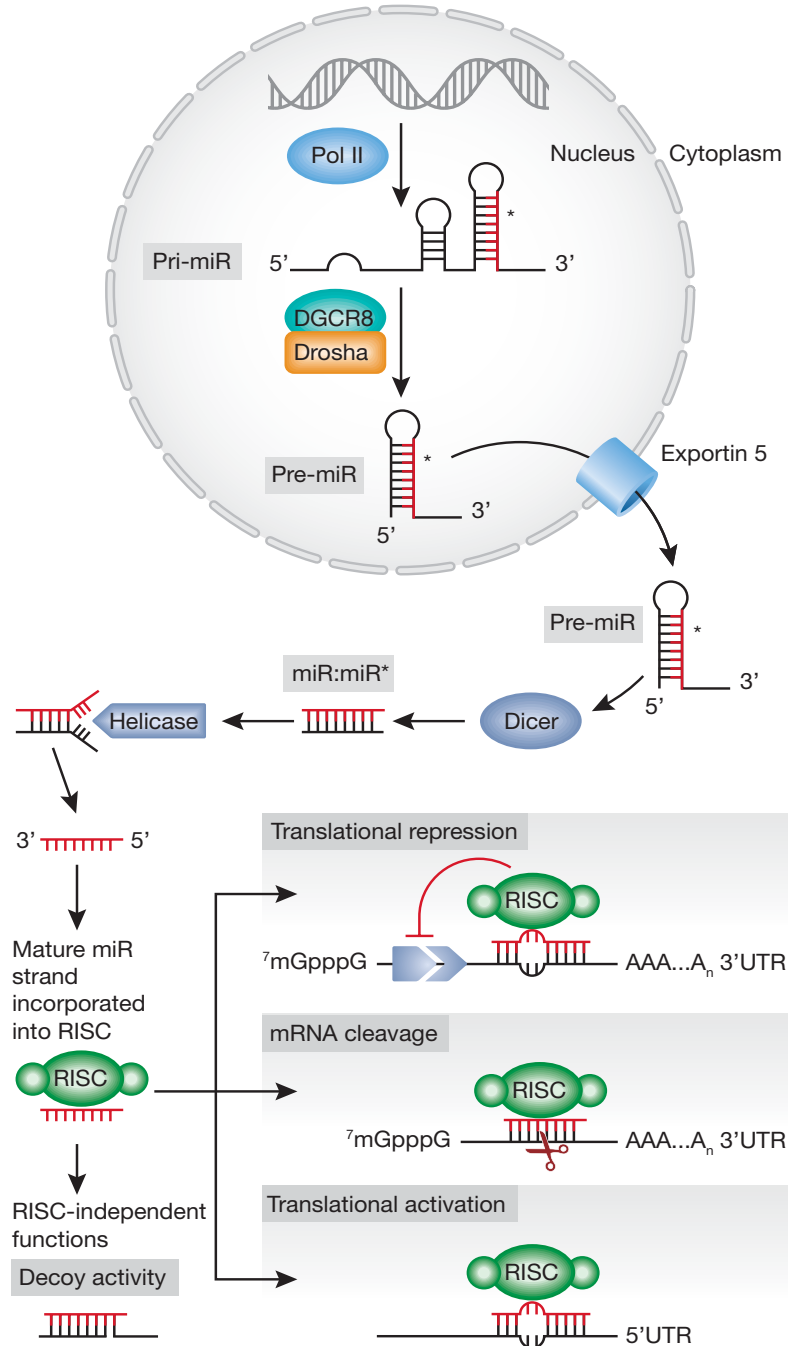
Triple negative tumours respond best to chemotherapy, similar to other aggressive cancers.

Luminal A tumours respond best to endocrine therapy, e.g. antiestrogen or aromatase inhibitor.

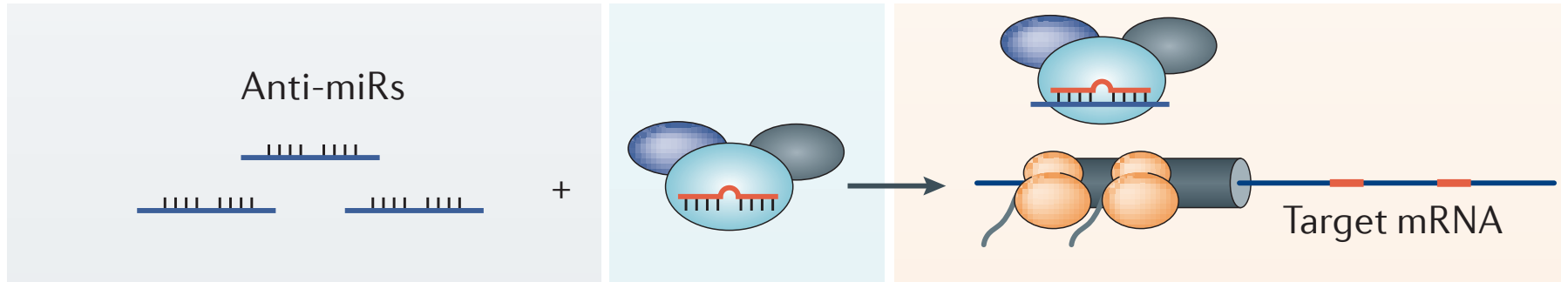
OncomiRs in cancer



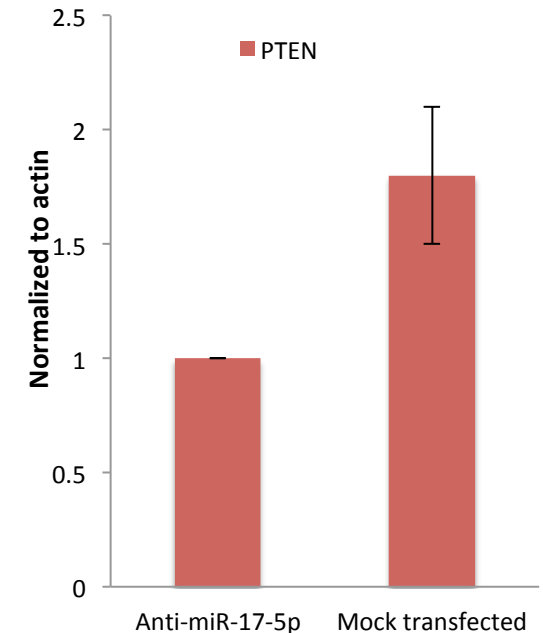
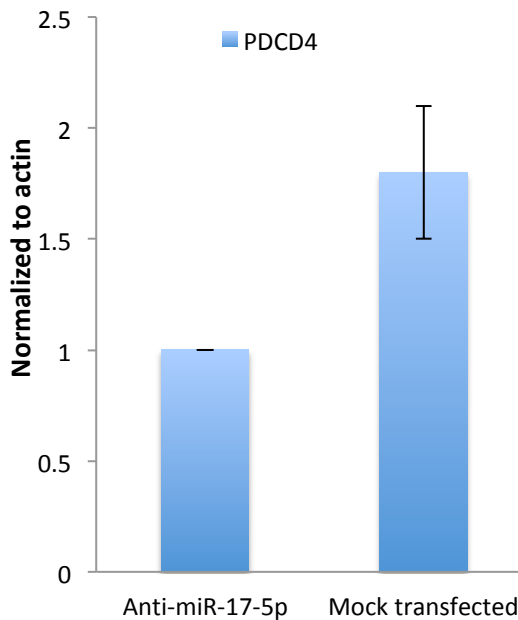
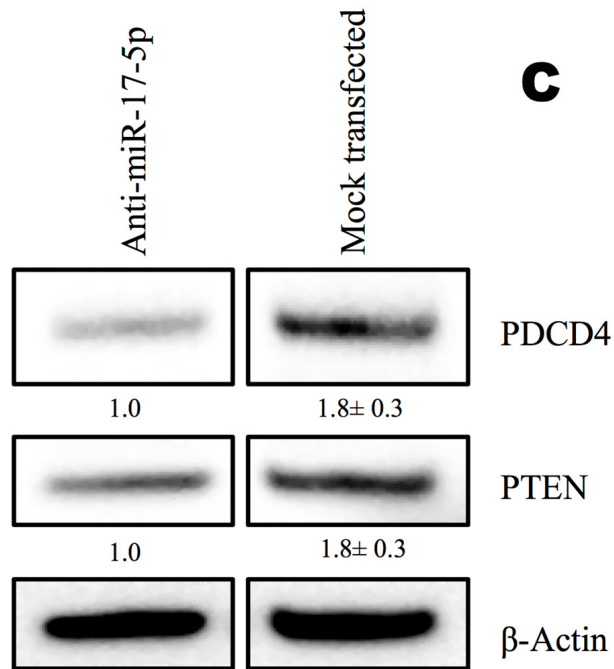
miRNA Biogenesis and Mechanisms of Action



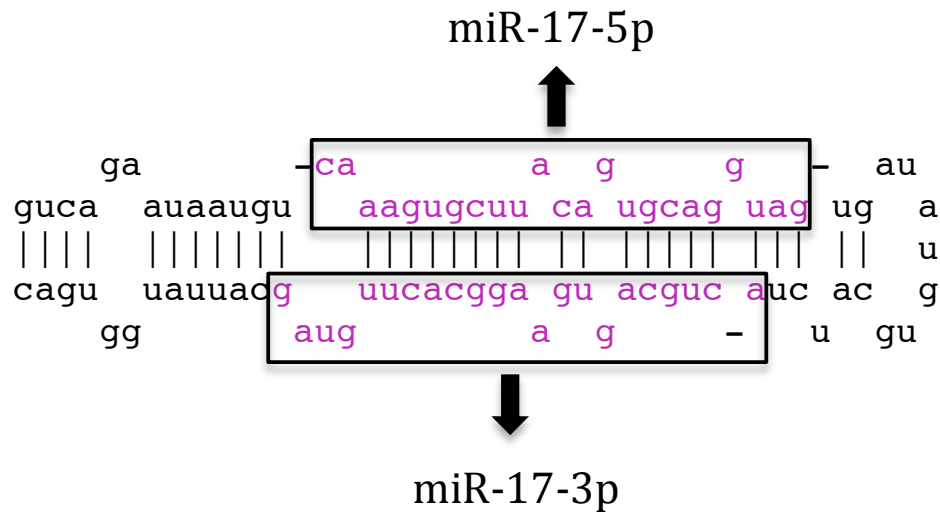
miRNA inhibition by modified oligonucleotides



miR-17-5p knockdown by DNA-LNA chimera unexpectedly decreased PDCD4 and PTEN protein in MDA-MB-231 cells.

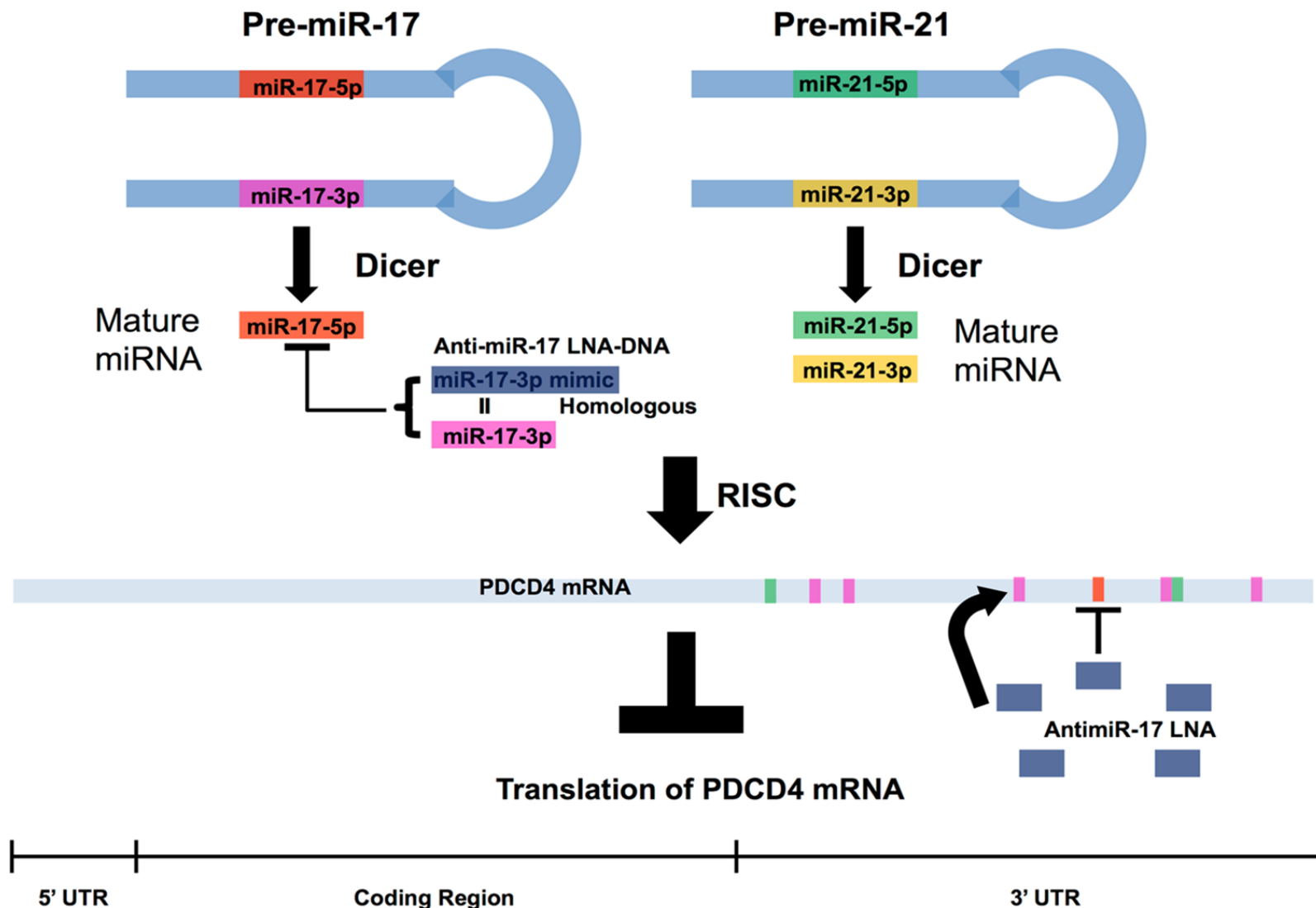


pre-miRNA structure of miR-17 revealed sequence similarity between DNA-LNA chimera and miR-17-3p passenger strand.

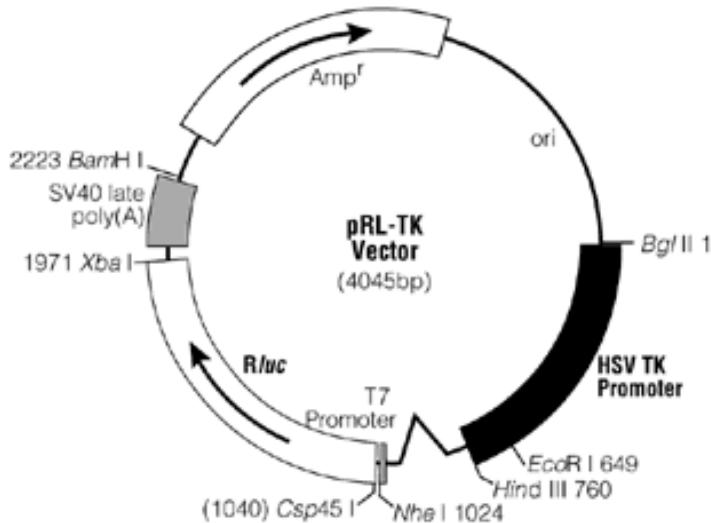
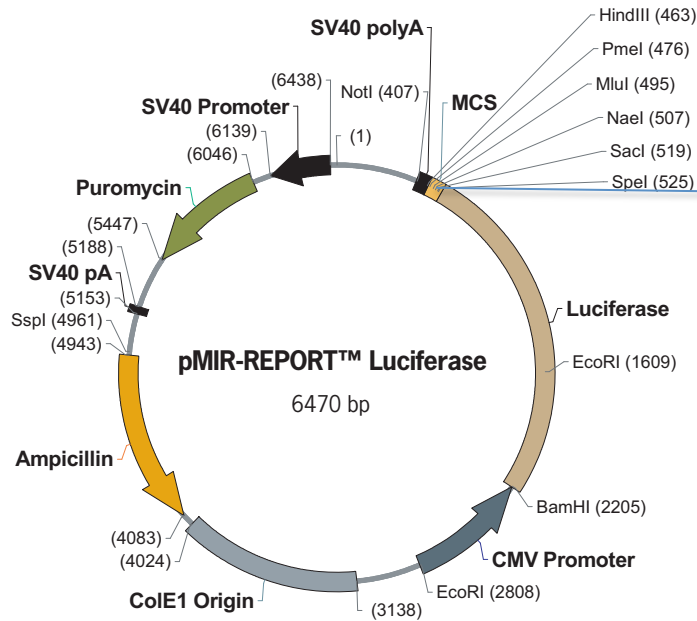


5' A-CUGCAGUG-AAGGCAC-UUGUAG 3' miR-17-3p
 5' ACCTGCACCTGTAAG-CACTTTG 3' Anti-miR-17-5p LNA

Competition between anti-miR-17-5p and miR-17-5p for inhibition of *PDCD4* mRNA

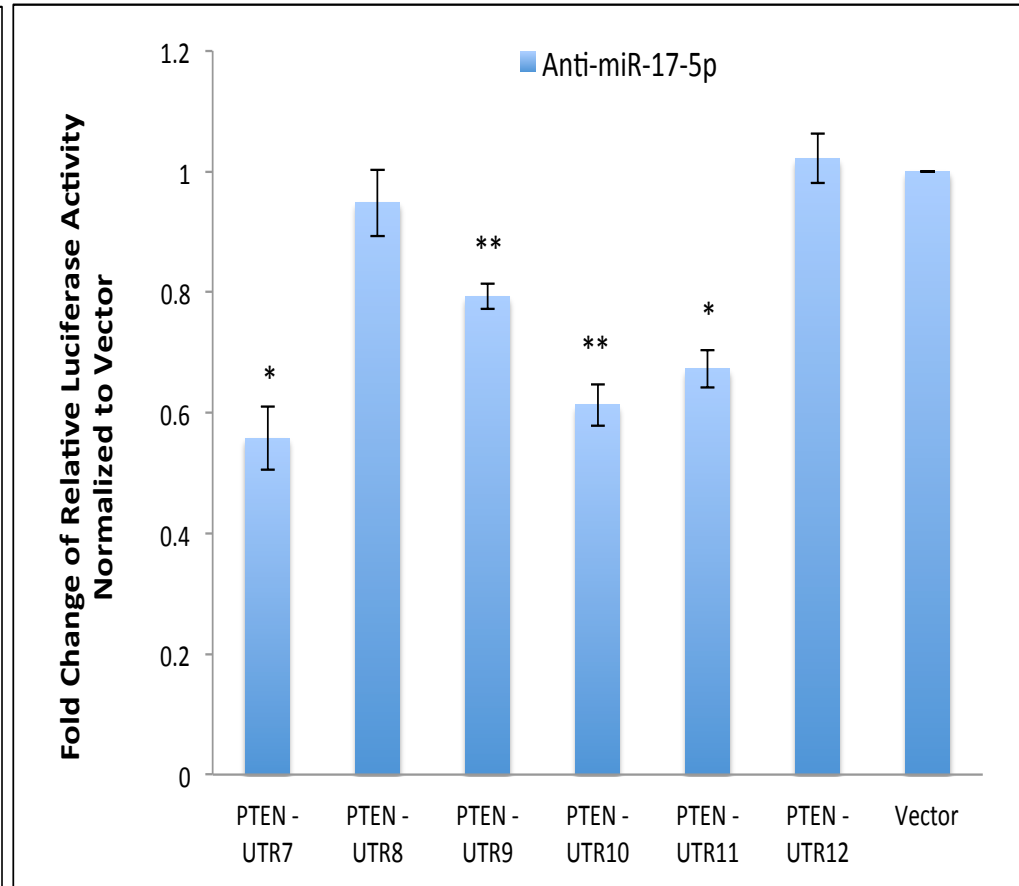
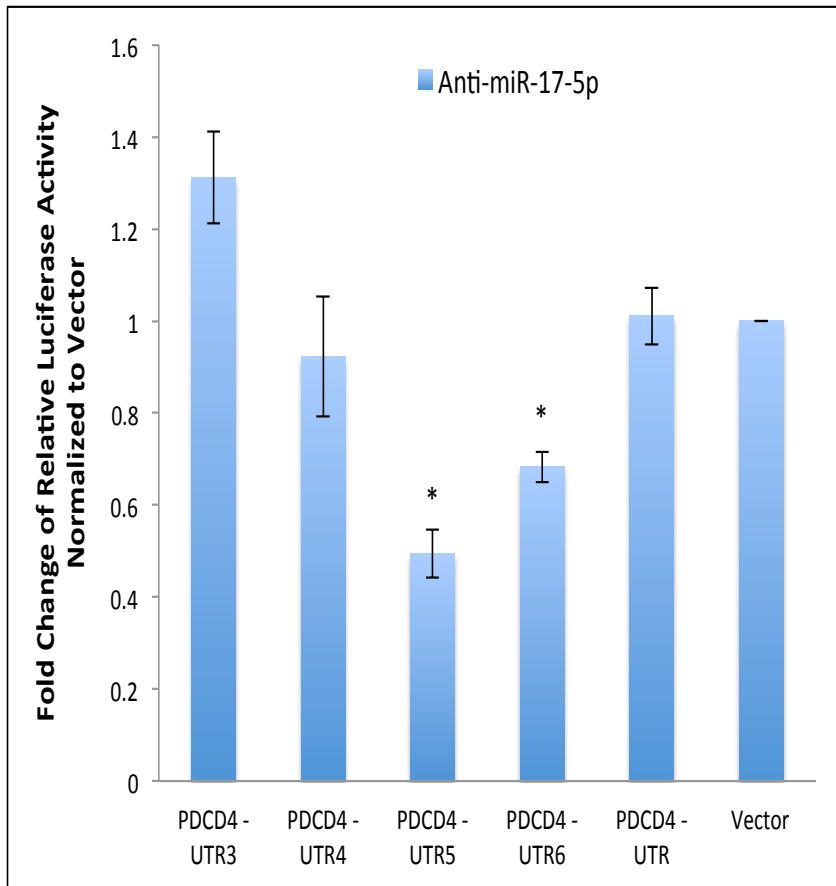


Luciferase assay system to test anti-miR-17 – mRNA interaction



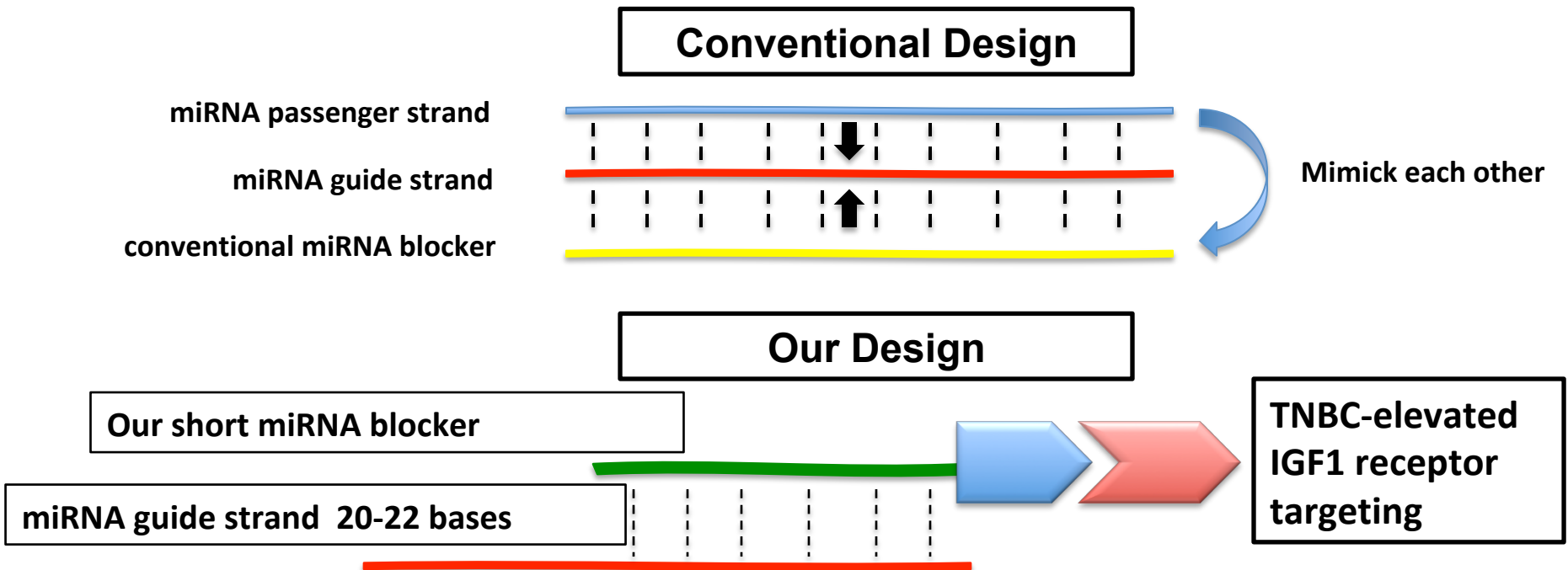
miRNA binding sites
PDCD4 3'UTR -3 – 19
PDCD4 3'UTR 325 - 346
PDCD4 3'UTR 1110 - 1136
PDCD4 3'UTR 1631 - 1652
PDCD4 3'UTR
PTEN 3'UTR 1199 - 1220
PTEN 3'UTR 5075 - 5096
PTEN 3'UTR 5828 - 5849
PTEN 3'UTR 5871 - 5892
PTEN 3'UTR 5908 - 5928
PTEN 3'UTR 6059 - 6080

Anti-miR-17-5p DNA-LNA lowered the expression of luciferase vectors containing several predicted *PDCD4* and *PTEN*'s 3'UTR target sites for miR-17-3p.



miRNA blocker design strategy

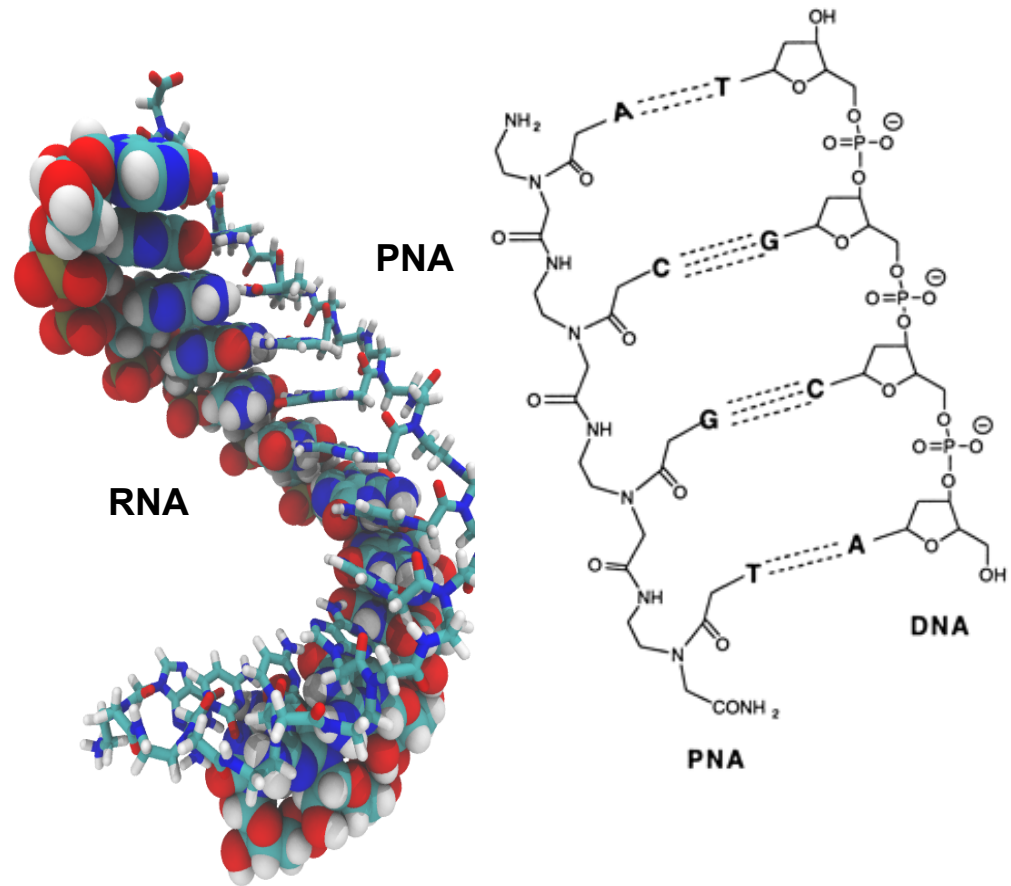
- Eliminate extra side-effects of conventional microRNA blockers
- TNBC cell-specific delivery method
- No complicated formulation, soluble in saline, intravenous route
- Next generation RNA backbones (FANA & NC-BNA vs. PNA) will elevate efficacy and potency



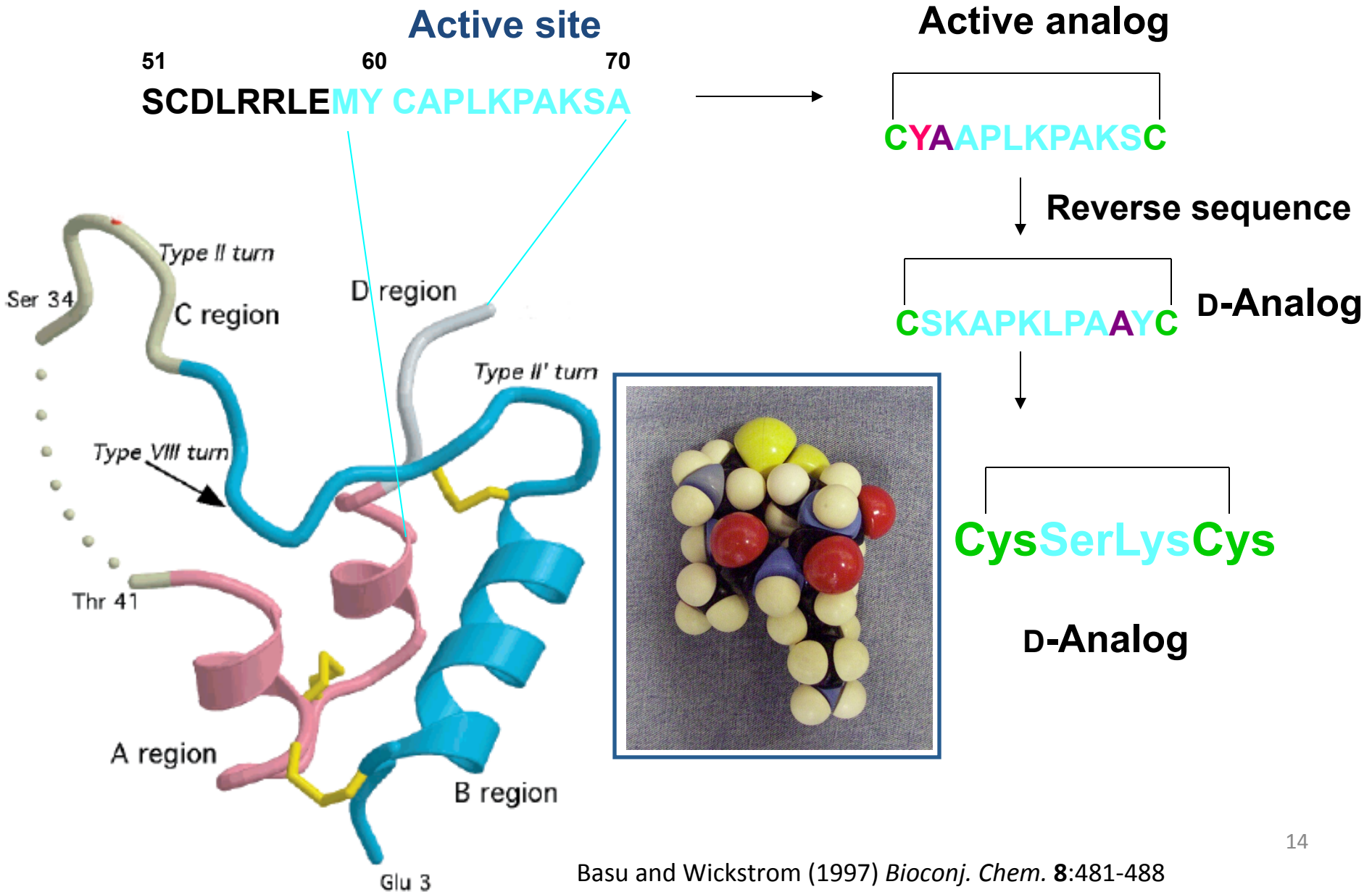
Nucleotide Analog - Peptide Nucleic Acids

Increasing stability, binding affinity and specificity

- High binding affinity to complementary DNA/RNA.
- Differentiation of single-base mismatch by high destabilizing effect.
- High chemical stability to temperature and pH.
- High biological stability to nuclease and protease.
- Good uptake via basic peptides or receptor-specific ligands
- Mice given up to 100 mg/kg dose of PNA-peptide conjugate daily did not show any irreversible toxicity (Chaubey et al., 2008).



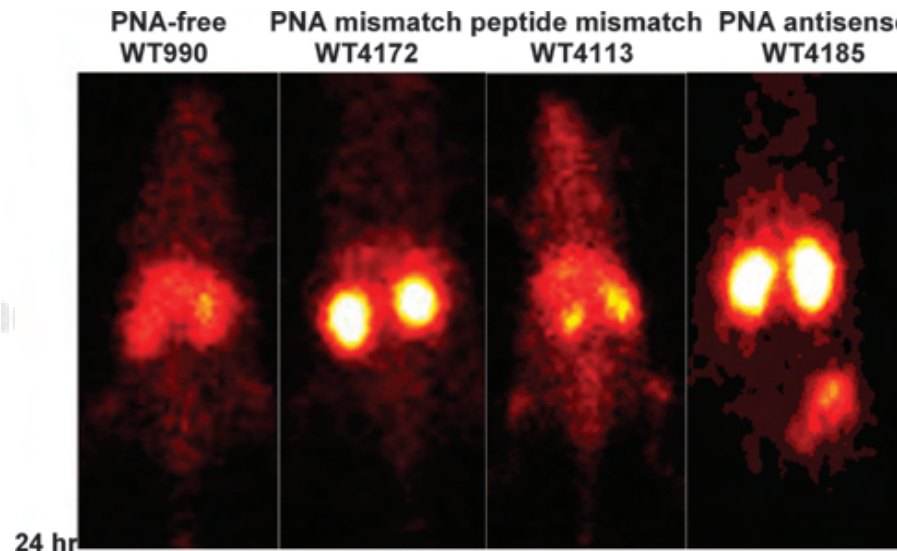
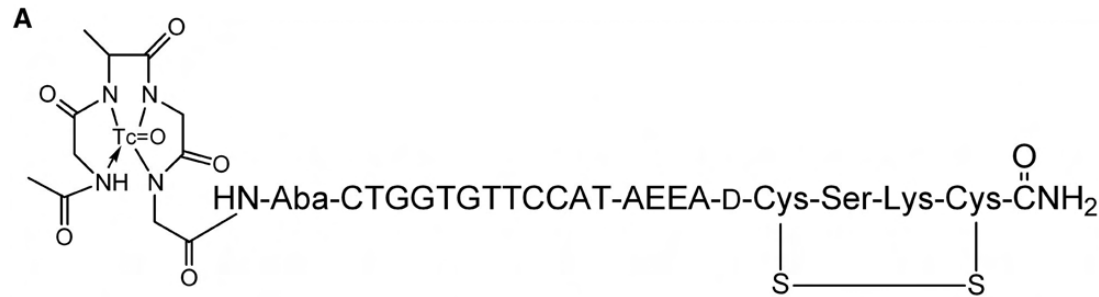
Delivery - IGF1 retro-inverso analog



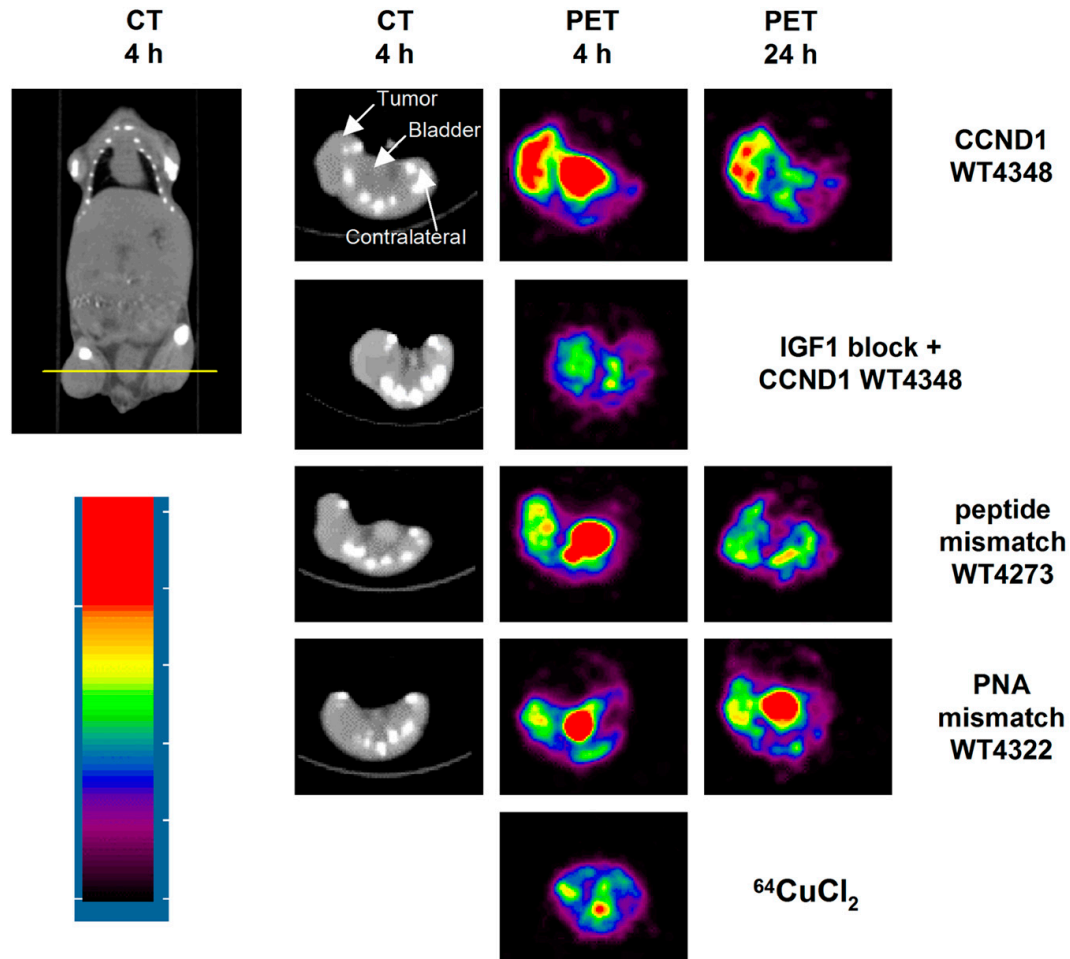
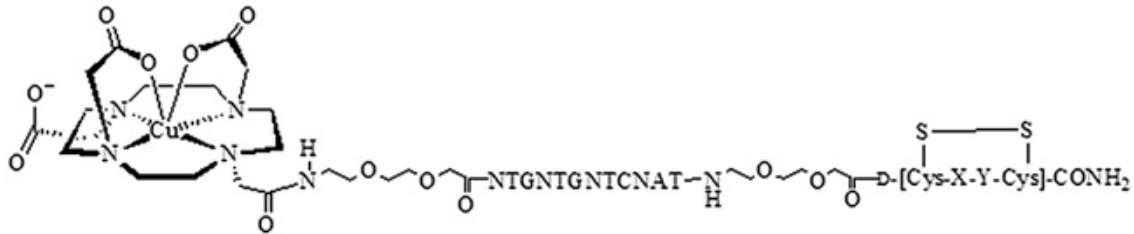
In vivo specificity of 12-mer PNA-IGF1 tetrapeptides

External Imaging of CCND1 Cancer Gene Activity in Experimental Human Breast Cancer Xenografts with ^{99m}Tc -Peptide-Peptide Nucleic Acid-Peptide Chimeras

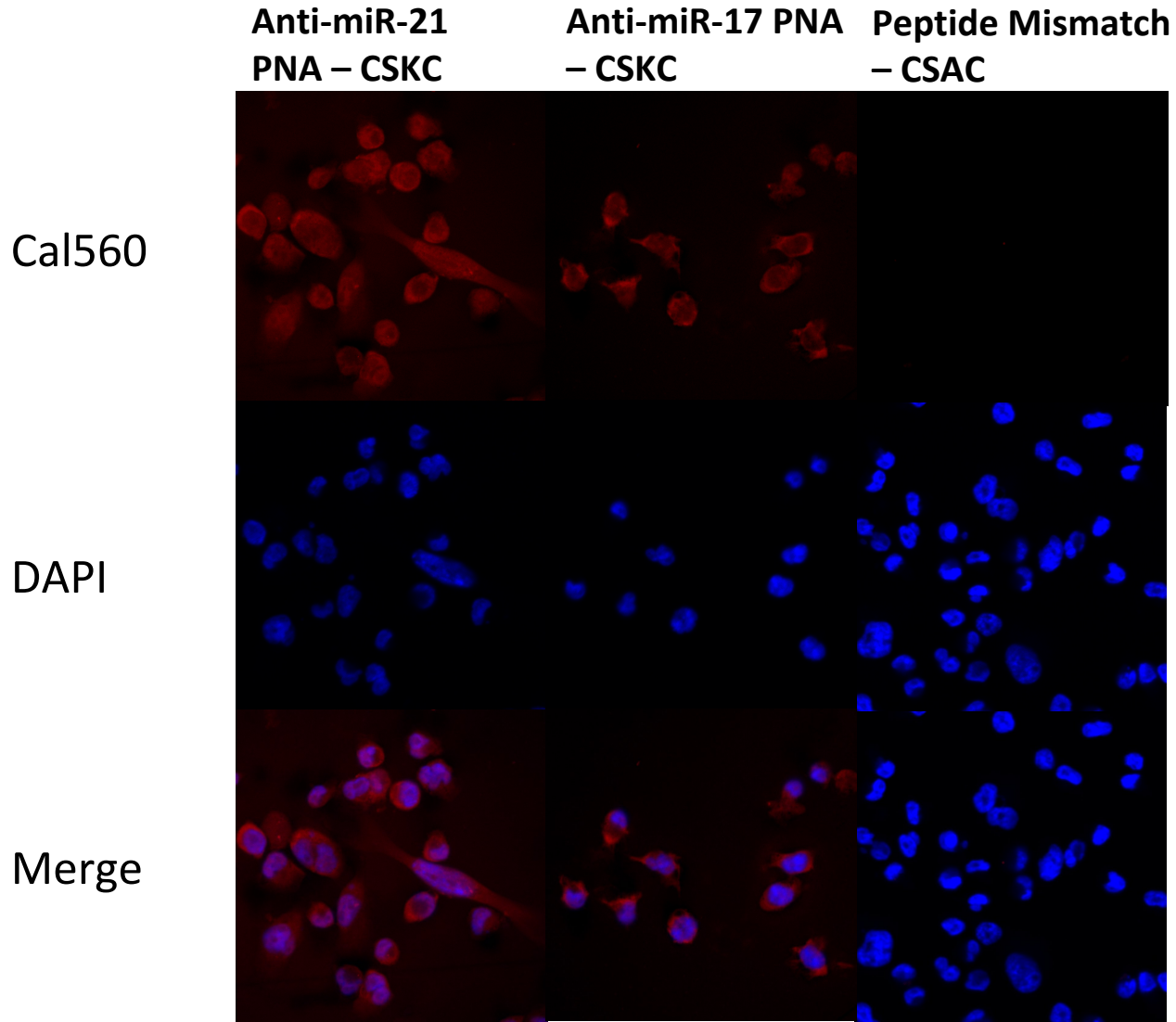
Xiaobing Tian, PhD¹; Mohan R. Aruva, PhD²; Wenyi Qin, MD³; Weizhu Zhu, MD³; Kevin T. Duffy, MBA¹; Edward R. Sauter, MD³; Mathew L. Thakur, PhD^{2,4}; and Eric Wickstrom, PhD^{1,4}



In vivo specificity of 12-mer PNA-IGF1 tetrapeptides



MDA-MB-231 cell uptake of Cal560-Anti-miR PNA-IGF1 tetrapeptide



Cells were incubated in 200 nM of Cal560-Anti-miR PNA-IGF1 tetrapeptide and negative controls for 4 hours at 37°C in complete medium. Ex: 543 Em: 560

1 μ M anti-miR PNA-IGF1 tetrapeptide elevated the expression of PDCD4 and PTEN

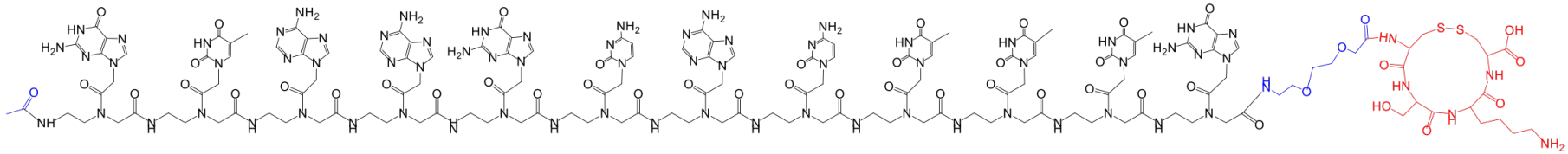
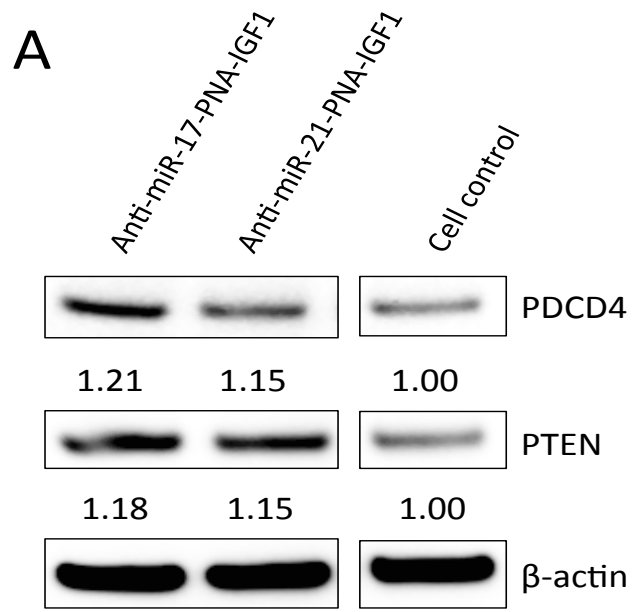
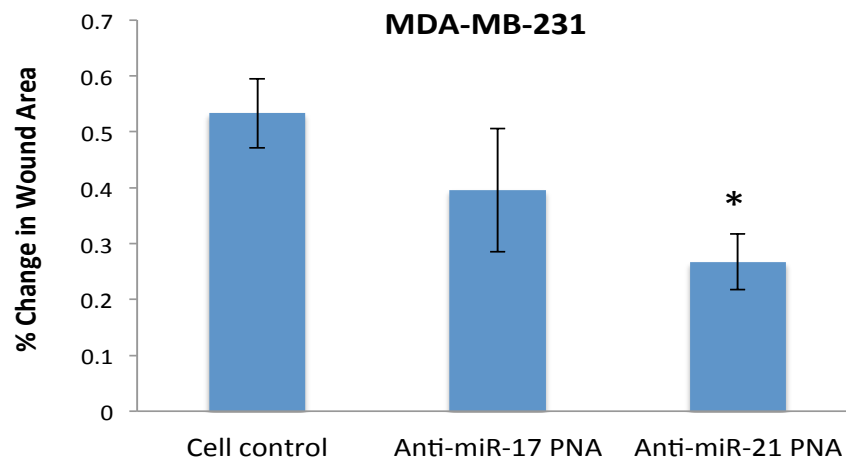
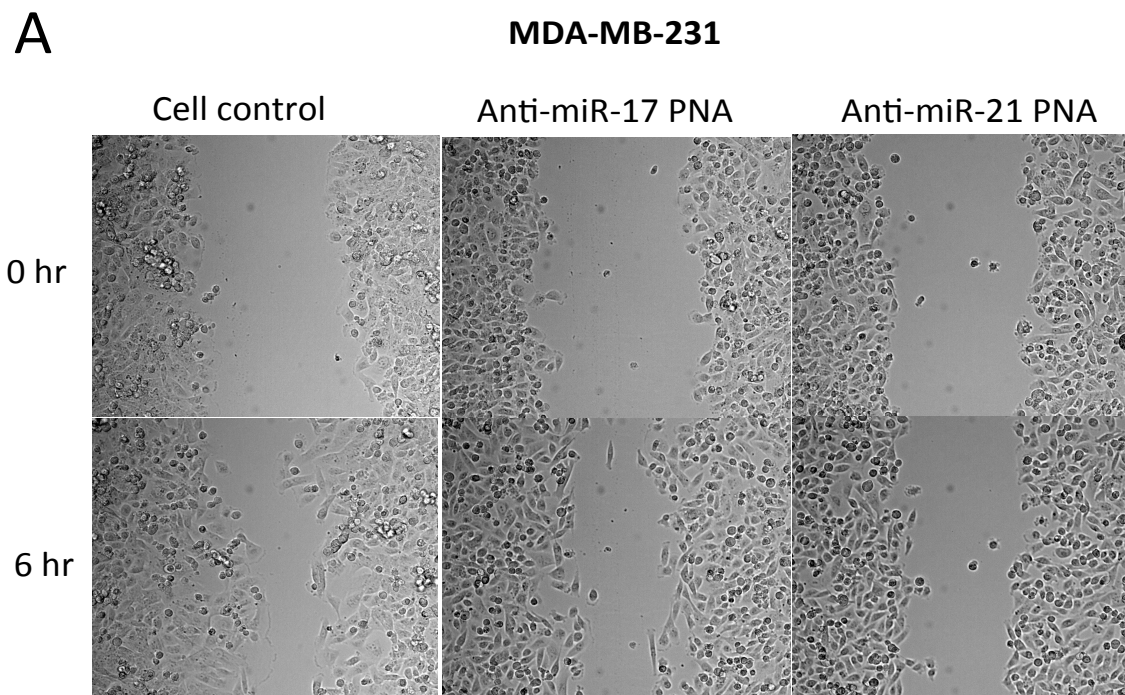


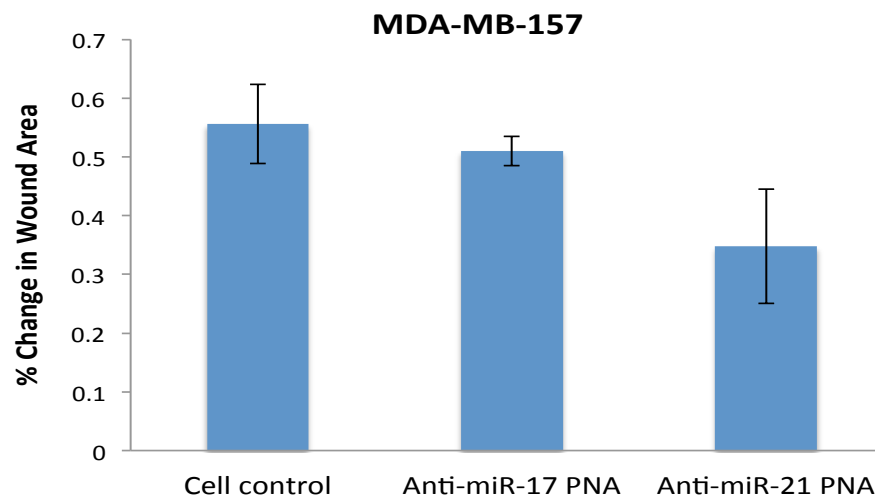
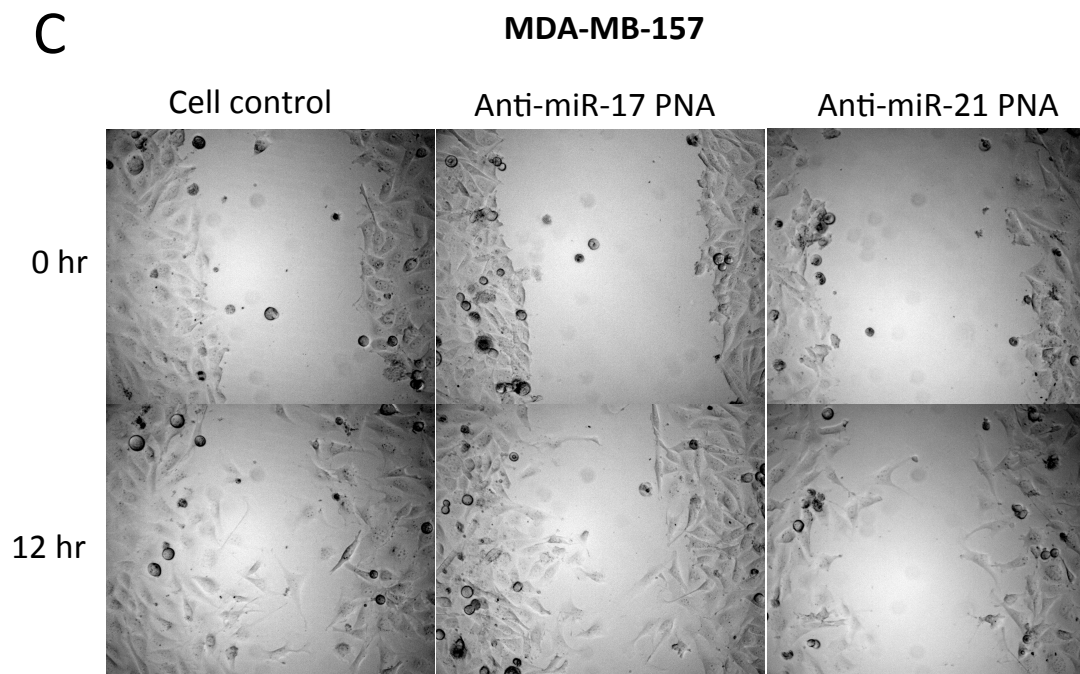
Fig. 1. PNA-AEEA-cyclo-D(Cys-Ser-Lys-Cys) blocker of miR-17-5p.



Blocking miR-21 with anti-miR-21 PNA-IGF1 tetrapeptide slowed down MDA-MB-231 cell migration.

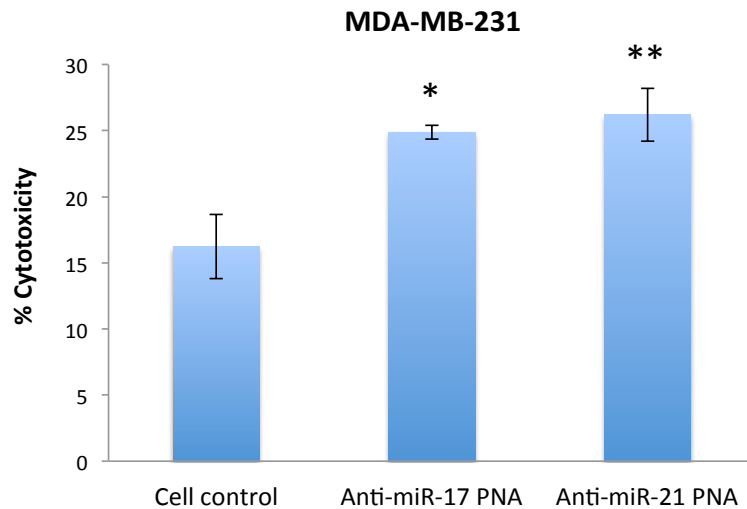


Blocking miR-21 with PNA-IGF1 tetrapeptide slowed down MDA-MB-157 cell migration.

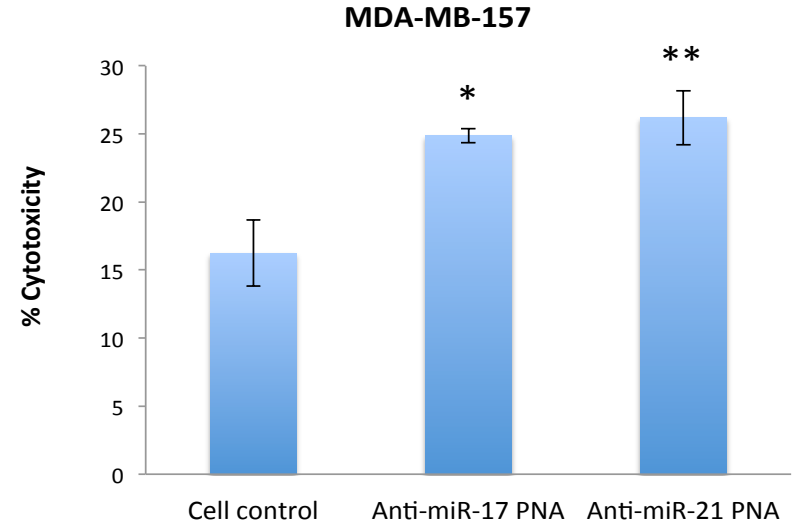


Blocking miR-21/17 with PNA-IGF1 tetrapeptide induced apoptosis in MSL type MDA-MB-231 and MDA-MB-157 cells.

A



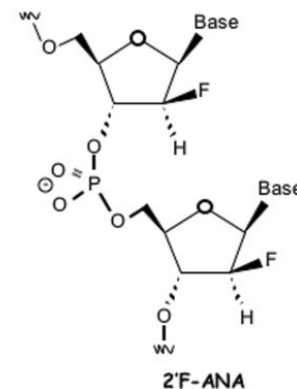
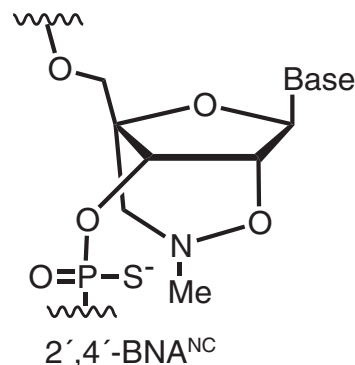
B



Summary

- The functional changes as a result of 1 μM PNA-IGF1 peptide treatment are modest, indicating low efficacy.
- TNBC cell lines that rely on PI3K/AKT/mTOR pathway are likely to respond to miR-21/17 blockage.
- Future antagomiRs can be optimized by:

- Alternative oligonucleotide analog that triggers RNase H (NC-BNA, FANA)



- Increasing the length of antagomiRs without mimicking passenger strand
- Better delivery target



Acknowledgements

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Bound Therapeutics LLC, NJ, USA

Dr. Eric Wickstrom

Henan Normal University, Xinxiang, Henan, China

Dr. Chang-Po Chen

