

MiRacles for babies with abnormal lung development and Congenital Diaphragmatic Hernia

Richard Keijzer, MD, MSc, PhD, FACS

Thorlakson Chair in Surgical Research



Conflict of Interest Disclosure

I hold a patent application (PCT/CA2015/051028) containing technology described in the presentation

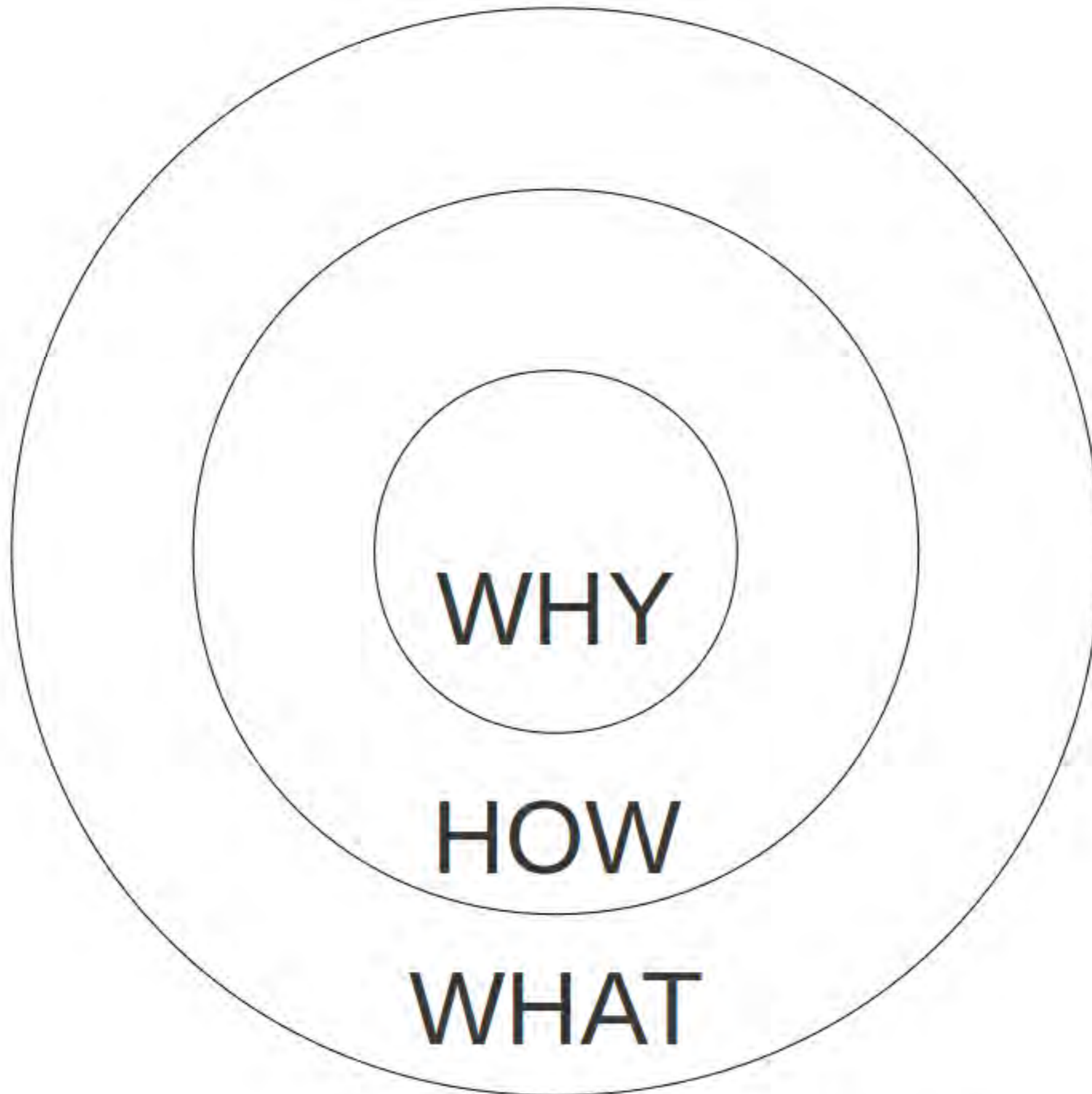
I have filed for a patent to use circular RNAs as biomarkers for abnormal lung development and
CDH

How it
began



We don't do chart reviews here, why don't you go in the lab for the next few months.





The 'golden circle' from Simon Sinek



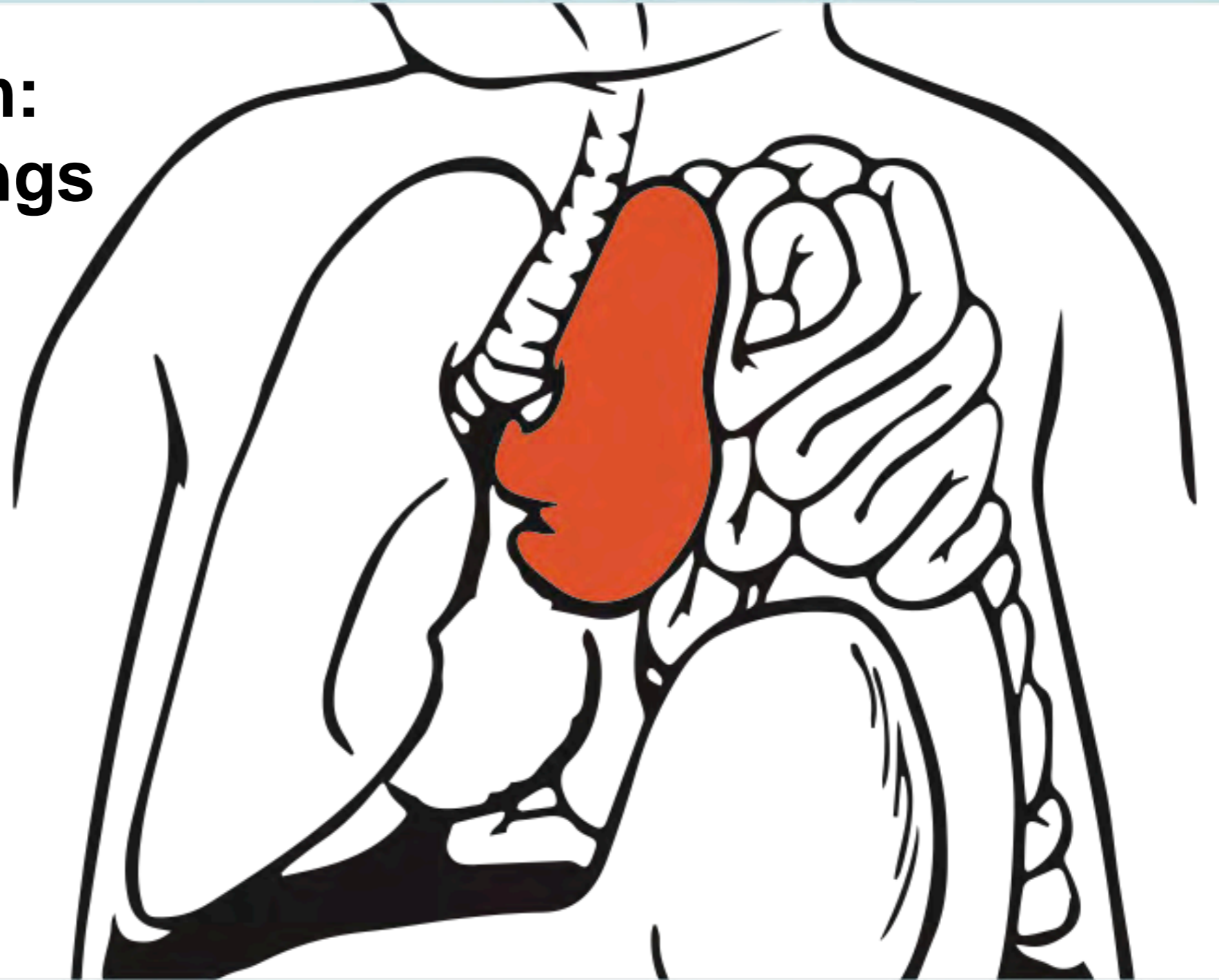
<http://store.winnipegfreepress.com/photostore/details/139975/>

Mortality: >400,000 since 2000



Image: Shutterstock

**Main problem:
Abnormal lungs**





MIRACLA

MicroRNAs • And • Congenital • Lung • Anomalies

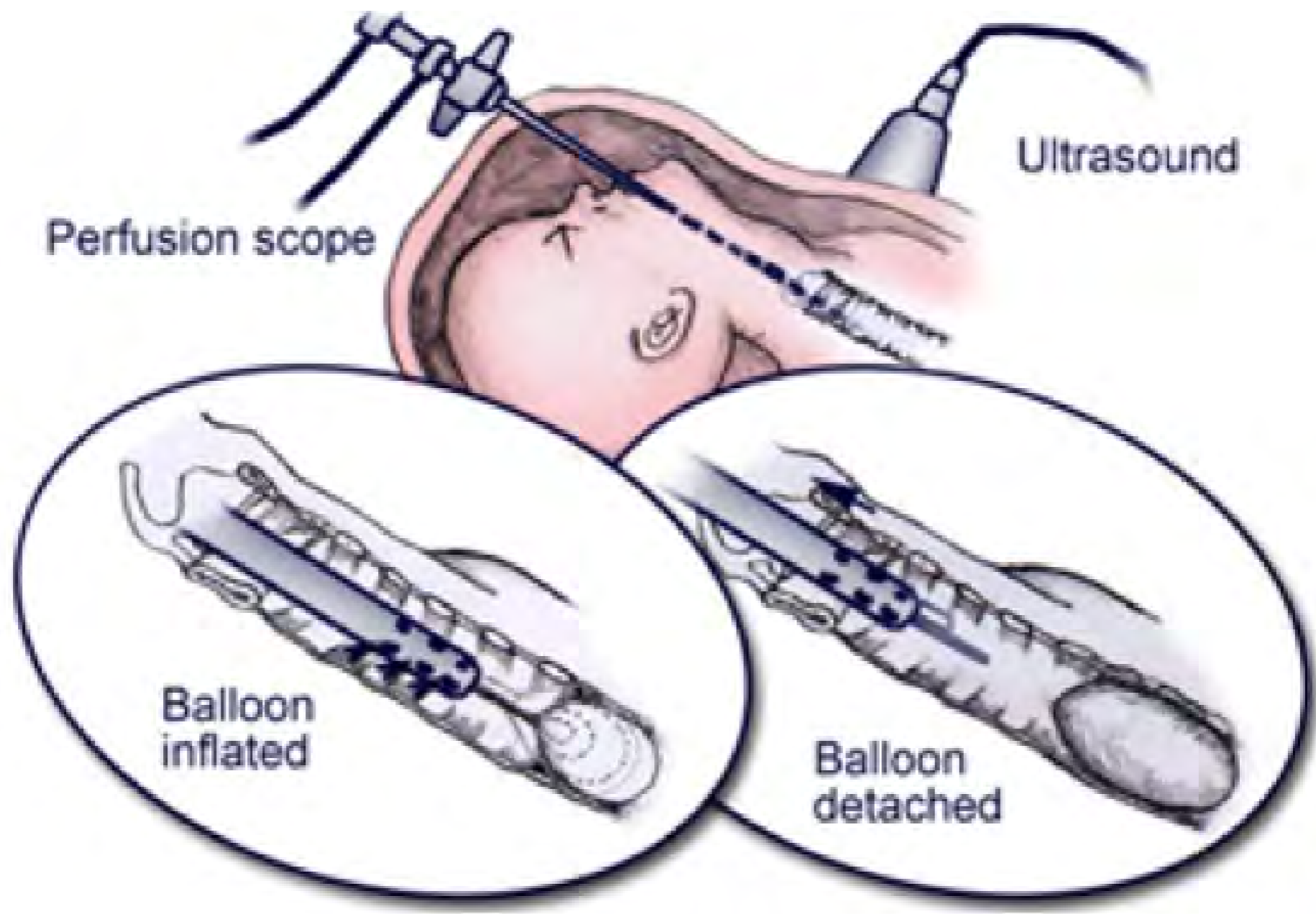
MicroRNAs • And • Congenital • Lung • Anomalies



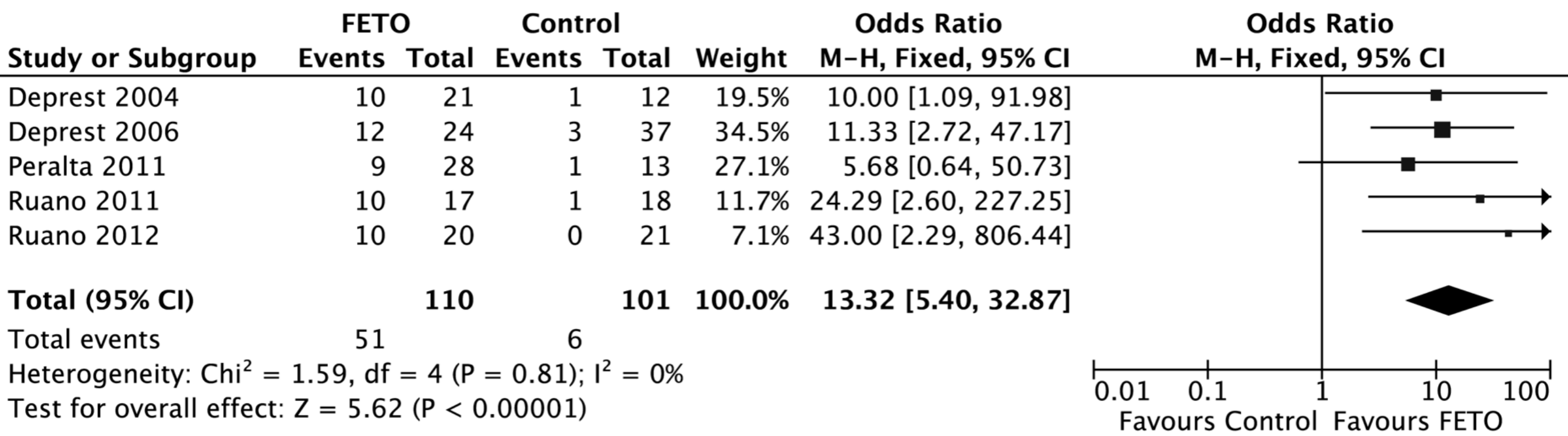
MIRACLA

What is wrong with CDH lungs?

Can we fix the lungs before birth?



Meta-analysis FETO improves survival in isolated CDH

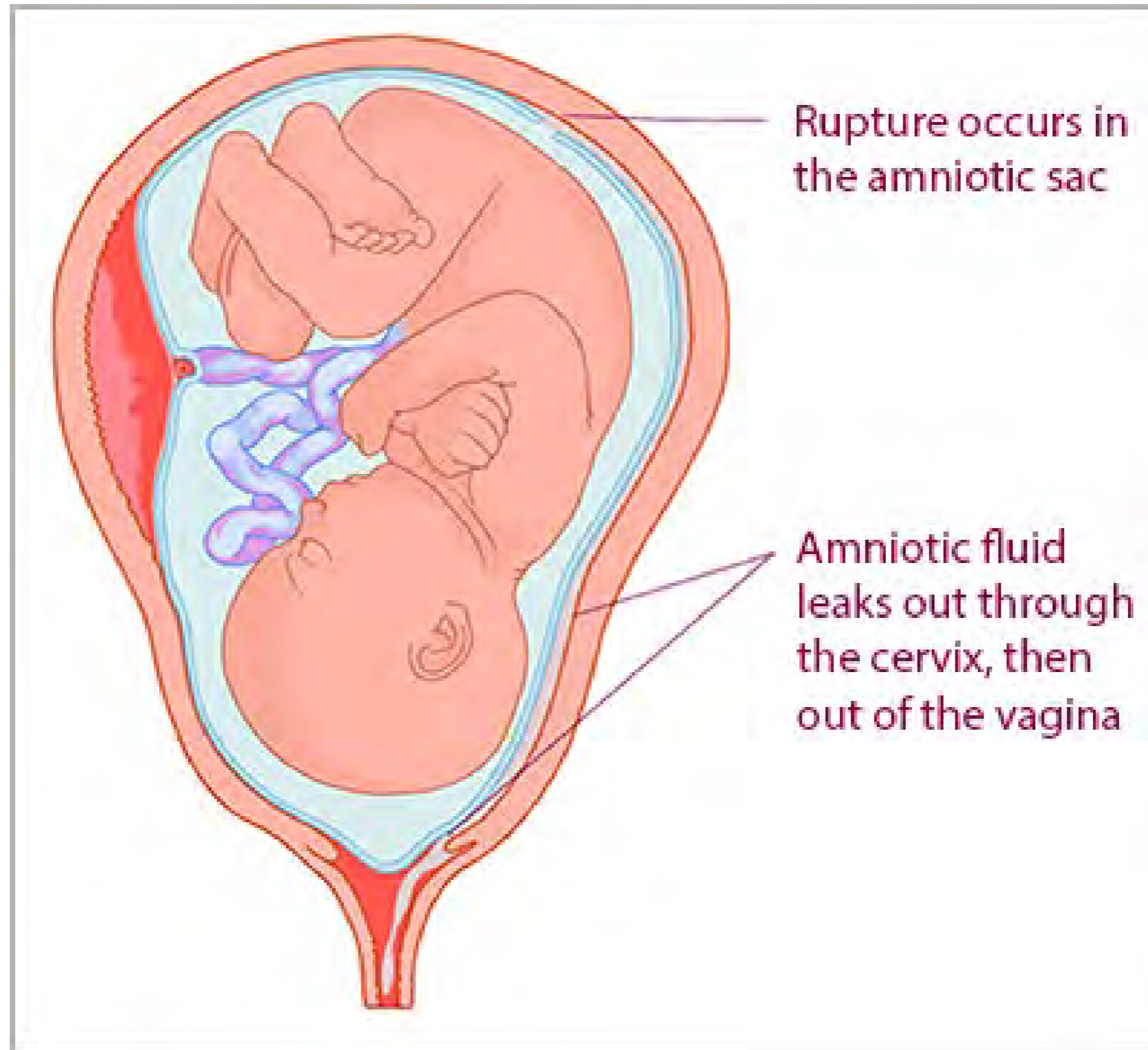


META-ANALYSIS

Fetal Tracheal Occlusion for Severe Pulmonary Hypoplasia in Isolated Congenital Diaphragmatic Hernia A Systematic Review and Meta-analysis of Survival

Jamila Al-Maary, MD,* Mary P. Eastwood, MBChB,† Francesca Maria Russo, MD,†
Jan A. Deprest, PhD,†§ and Richard Keijzer, PhD*‡

But, negative side effects!



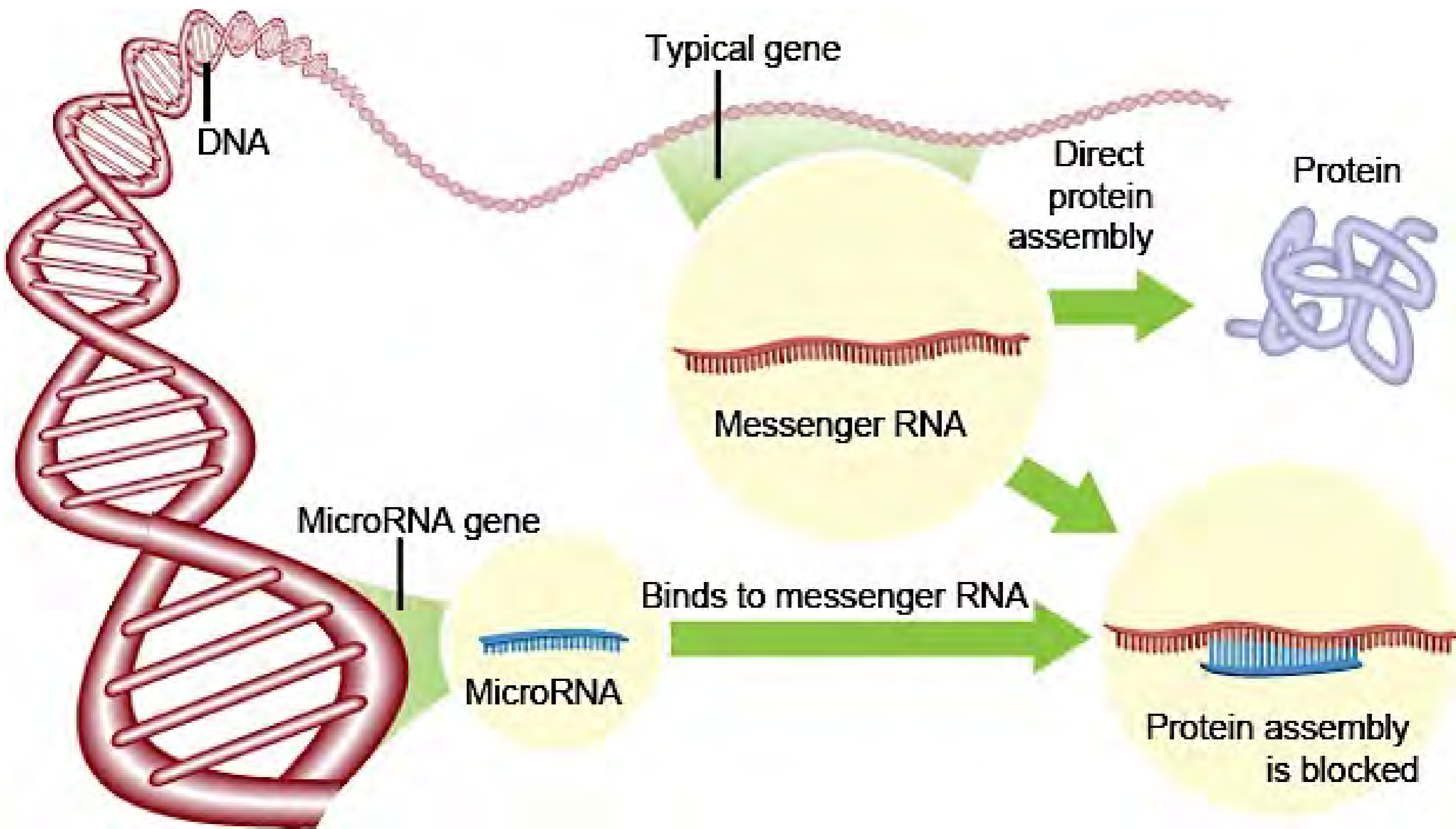
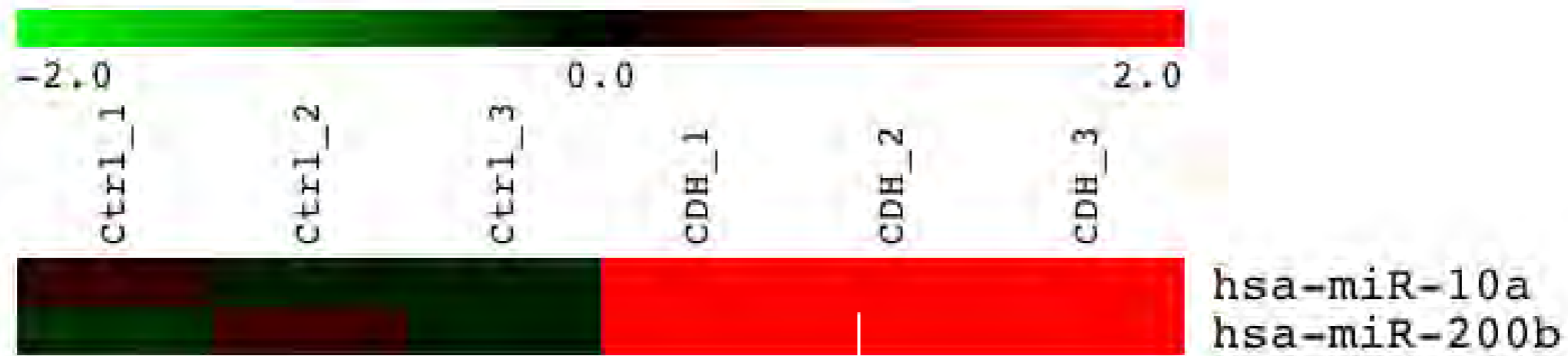


Image: Steve Karp, Discover Magazine

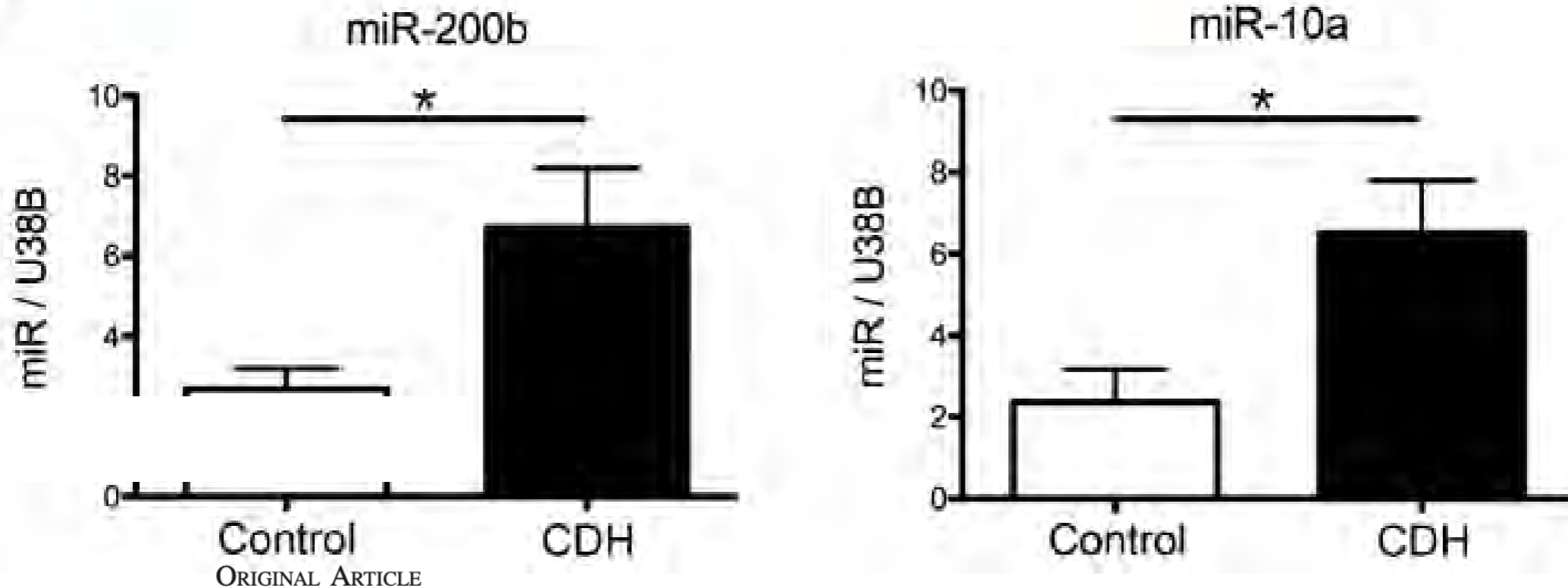


MicroRNAs and Congenital Diaphragmatic Hernia

A.

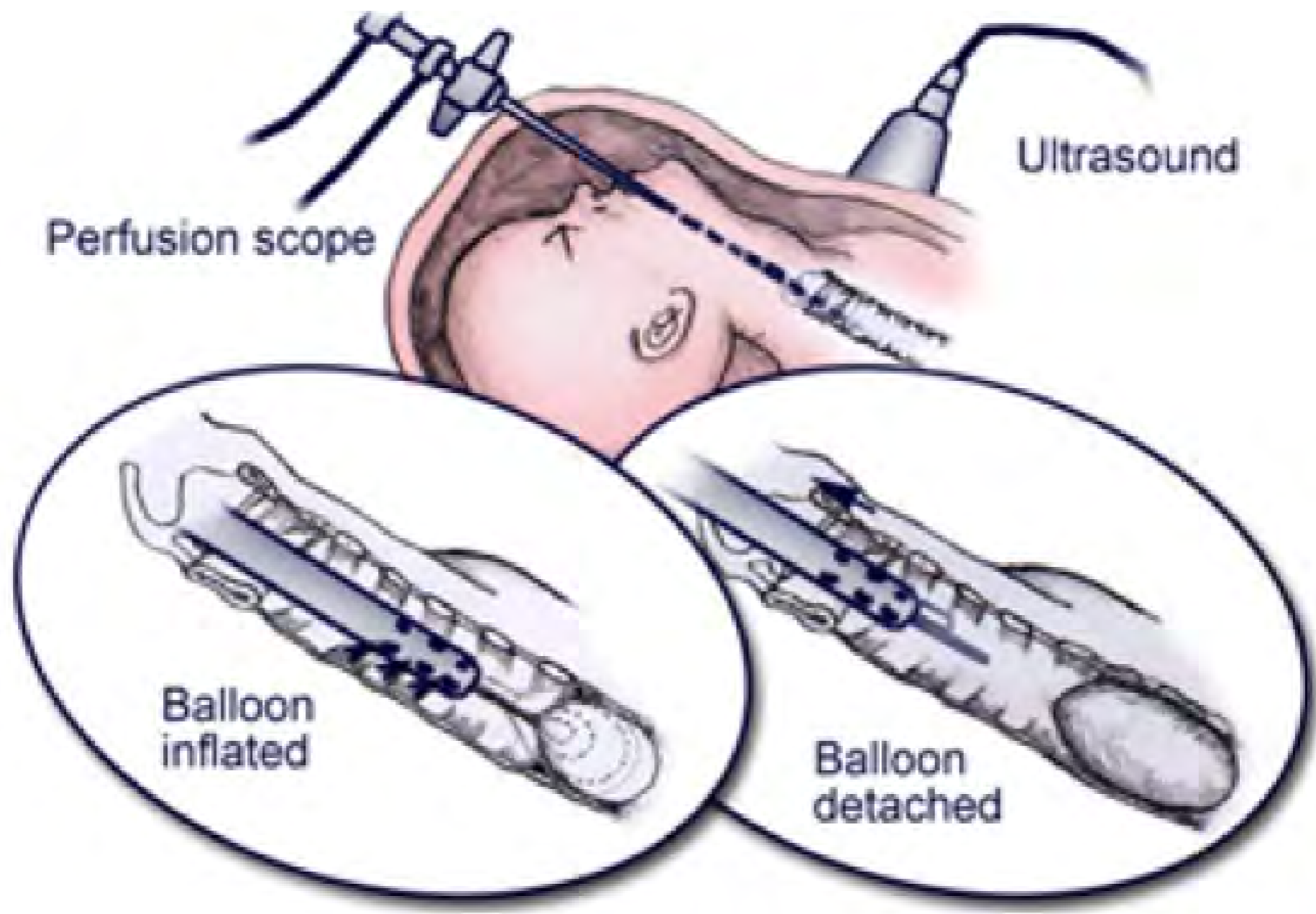


B.

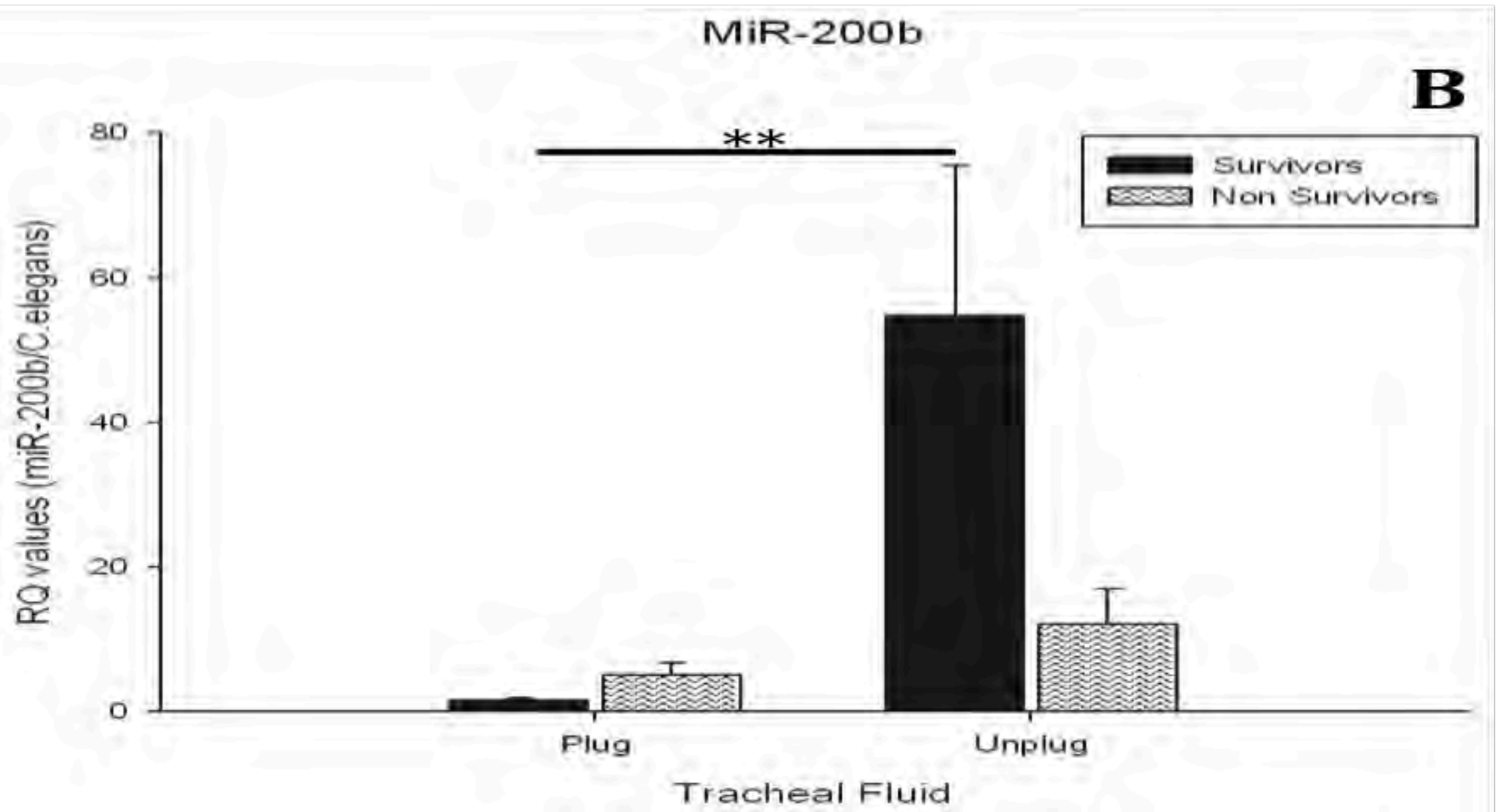


Unique Tracheal Fluid MicroRNA Signature Predicts Response to FETO in Patients With Congenital Diaphragmatic Hernia

Patrícia Pereira-Terra, MSc,*† Jan A. Depreest, MD, PhD,‡ Ramin Kholdebarin, MD, MSc,*
 Naghmeh Khoshgoo, MS,* Philip DeKoninck, MD, PhD,‡ Anne A. Boerema-De Munck,§ Jinxia Wang,¶
 Fuqin Zhu,* Robbert J. Rottier, PhD,§ Barbara M. Iwasiow, MSc,* Jorge Correia-Pinto, MD, PhD,†
 Dick Tibboel, MD, PhD,§ Martin Post, DVM, PhD,¶ and Richard Keijzer, PhD*



Higher miR-200b has better outcomes



Epithelial-to-Mesenchymal Transition (EMT)

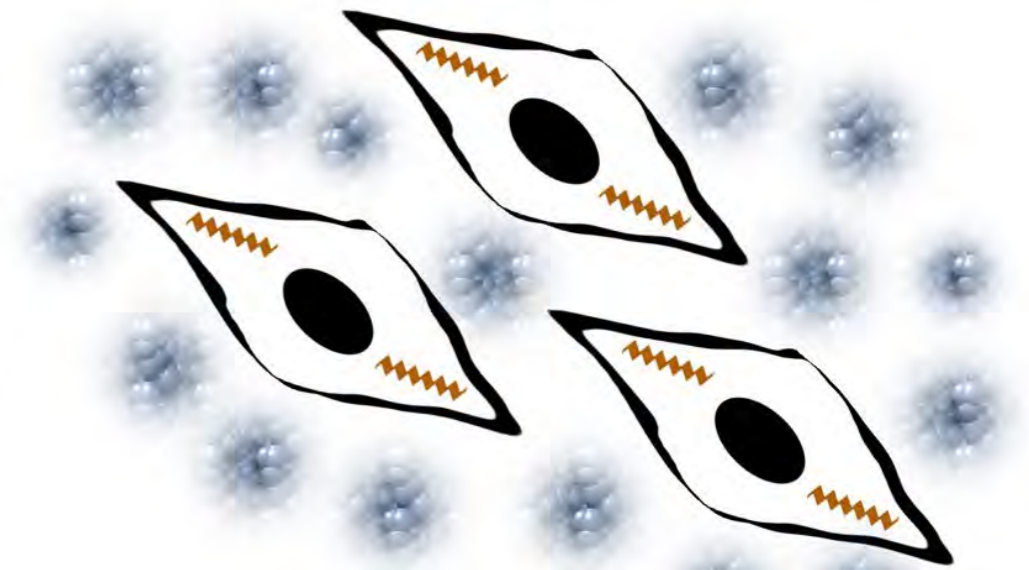


Epithelial cells

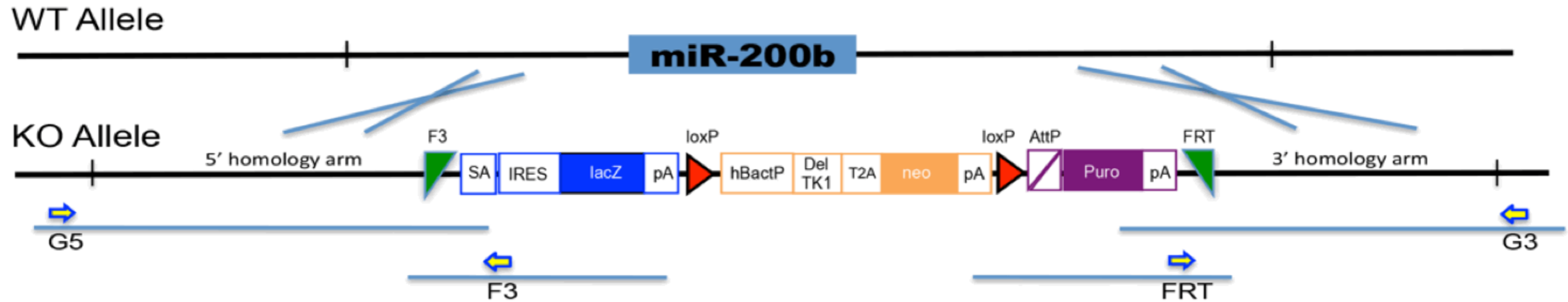
TGF- β /SMAD



miR-200b



Mesenchymal cells



SCIENTIFIC REPORTS

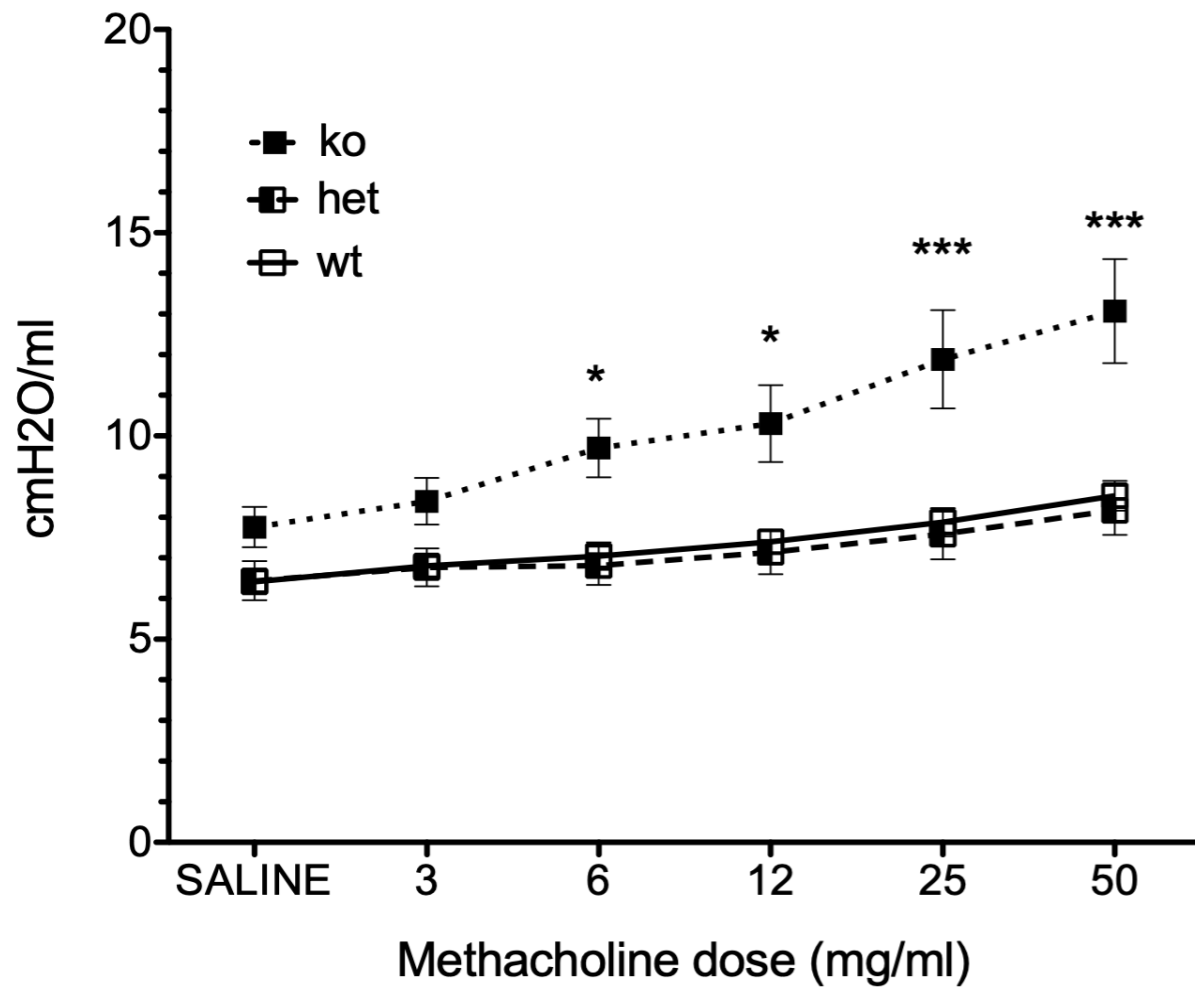
OPEN

MicroRNA-200b regulates distal airway development by maintaining epithelial integrity

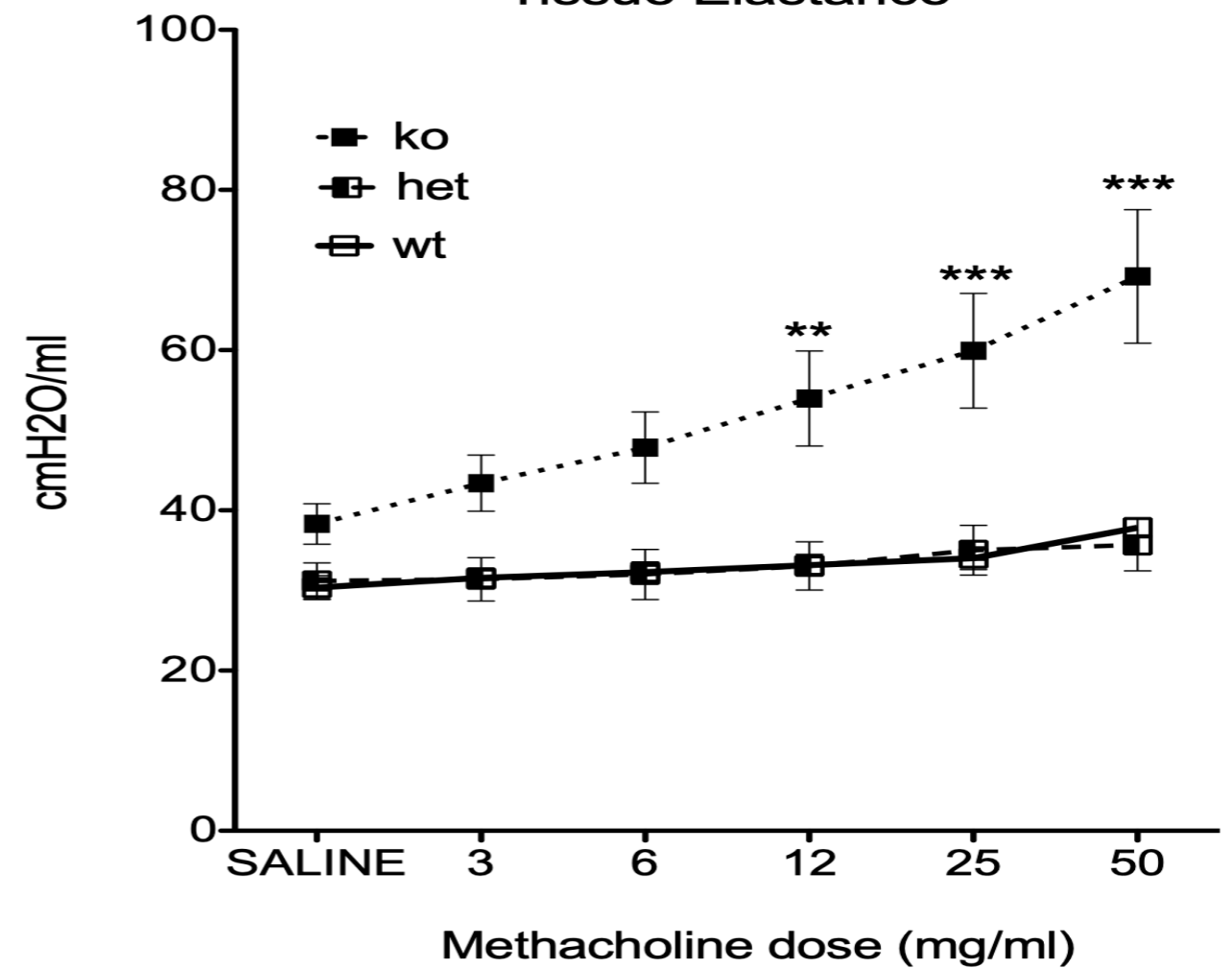
Naghmeh Khoshgoo^{1,2,3}, Robin Visser^{1,2}, Landon Falk^{1,2}, Chelsea A. Day^{1,2}, Dustin Ameis^{1,2}, Barbara M. Iwasiow^{1,2}, Fuqin Zhu^{1,2}, Arzu Öztürk^{4,5}, Sujata Basu^{1,3}, Molly Pind^{4,5}, Agnes Fresnosa^{4,5}, Mike Jackson⁶, Vinaya Kumar Siragam^{1,2}, Gerald Stelmack^{1,3}, Geoffrey G. Hicks^{4,5}, Andrew J. Halayko^{1,3} & Richard Keijzer^{1,2,3}

miR-200b -/- mice have higher lung tissue damping and elastance

Tissue Damping



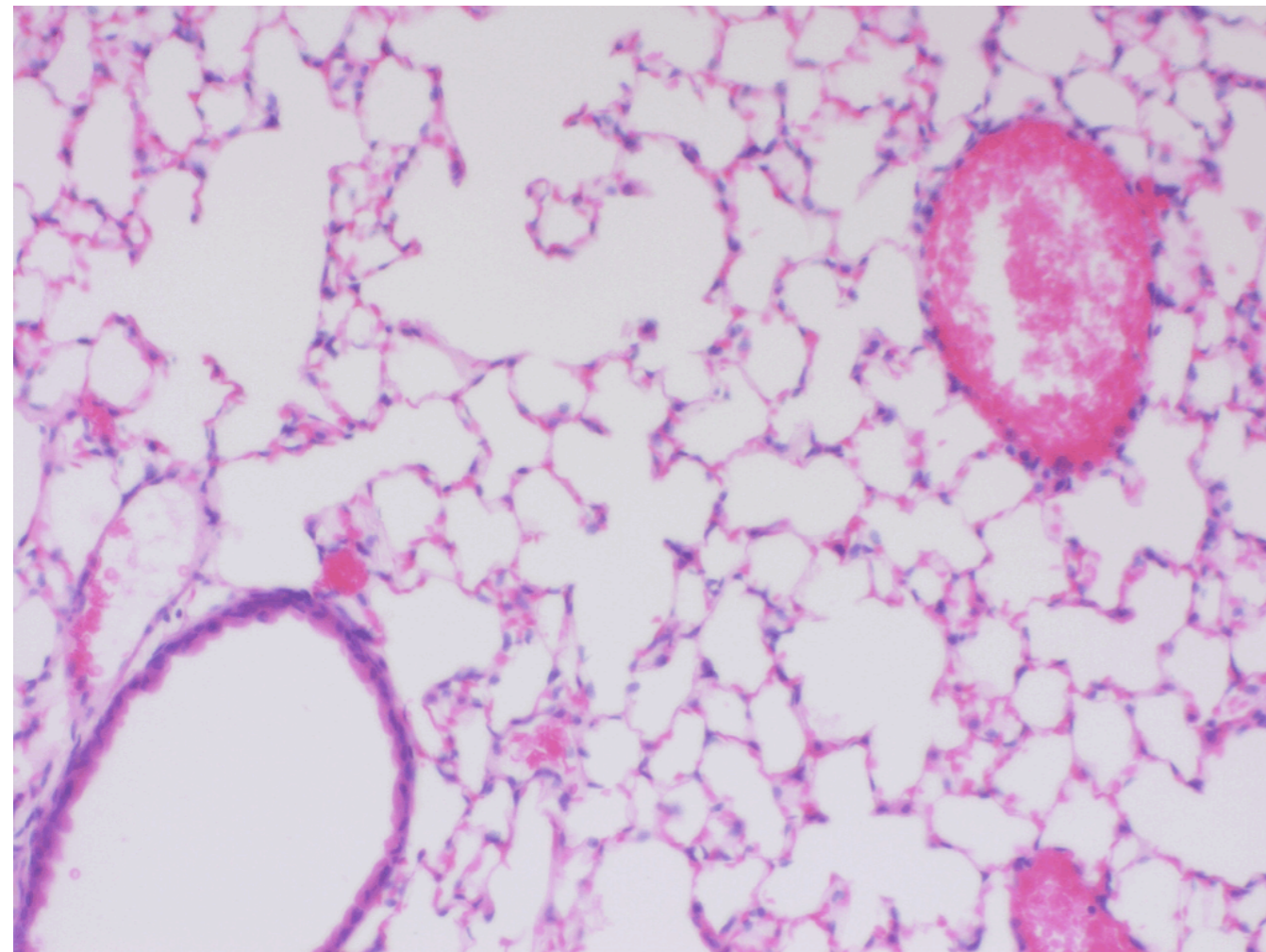
Tissue Elastance



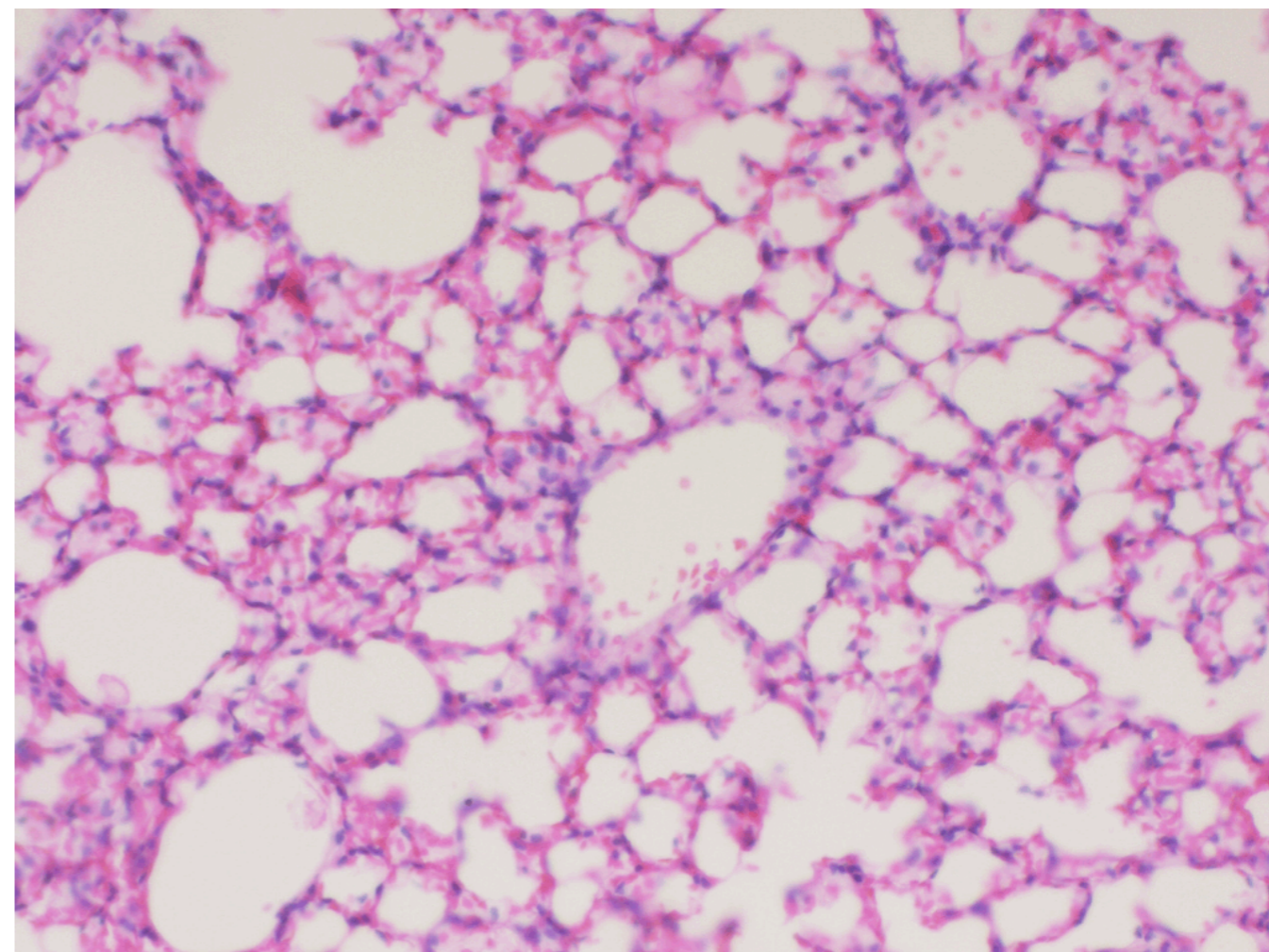
* P < 0.05, ** P < 0.01, *** P < 0.001

miR-200b $-/-$ lungs are hypoplastic

wt



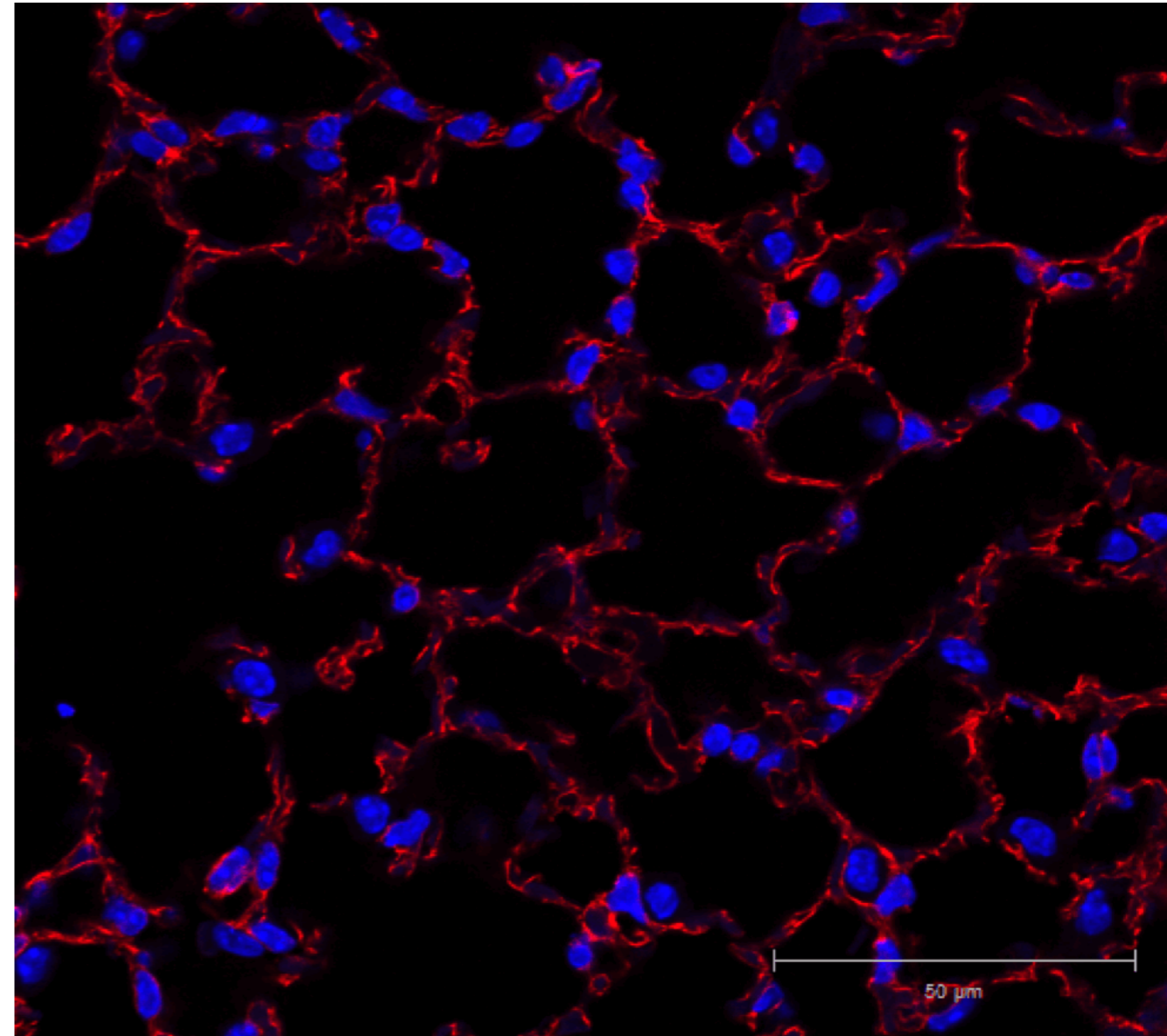
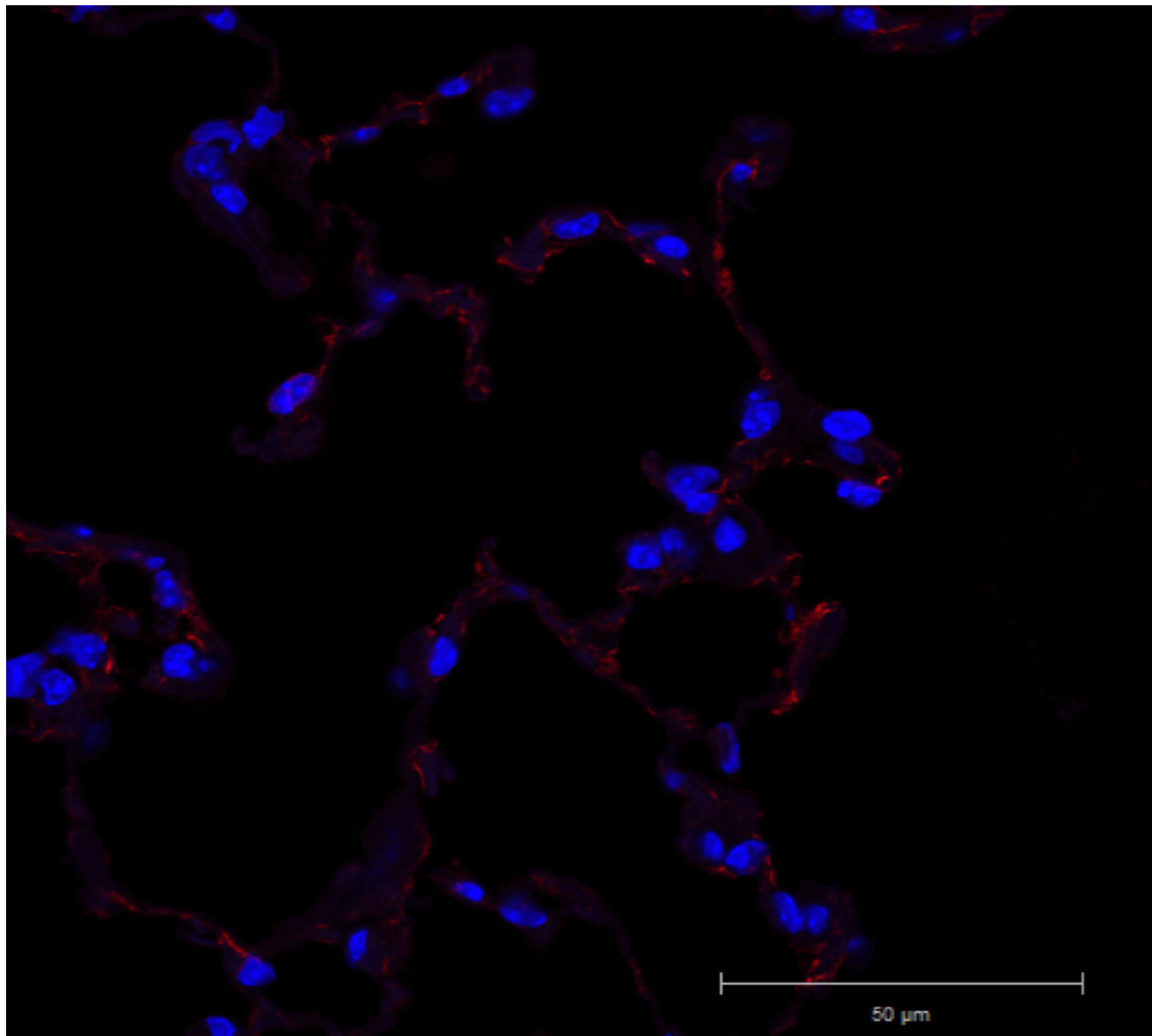
miR-200b $-/-$



miR-200b $-/-$ lungs have more vimentin

wt

miR-200b $-/-$

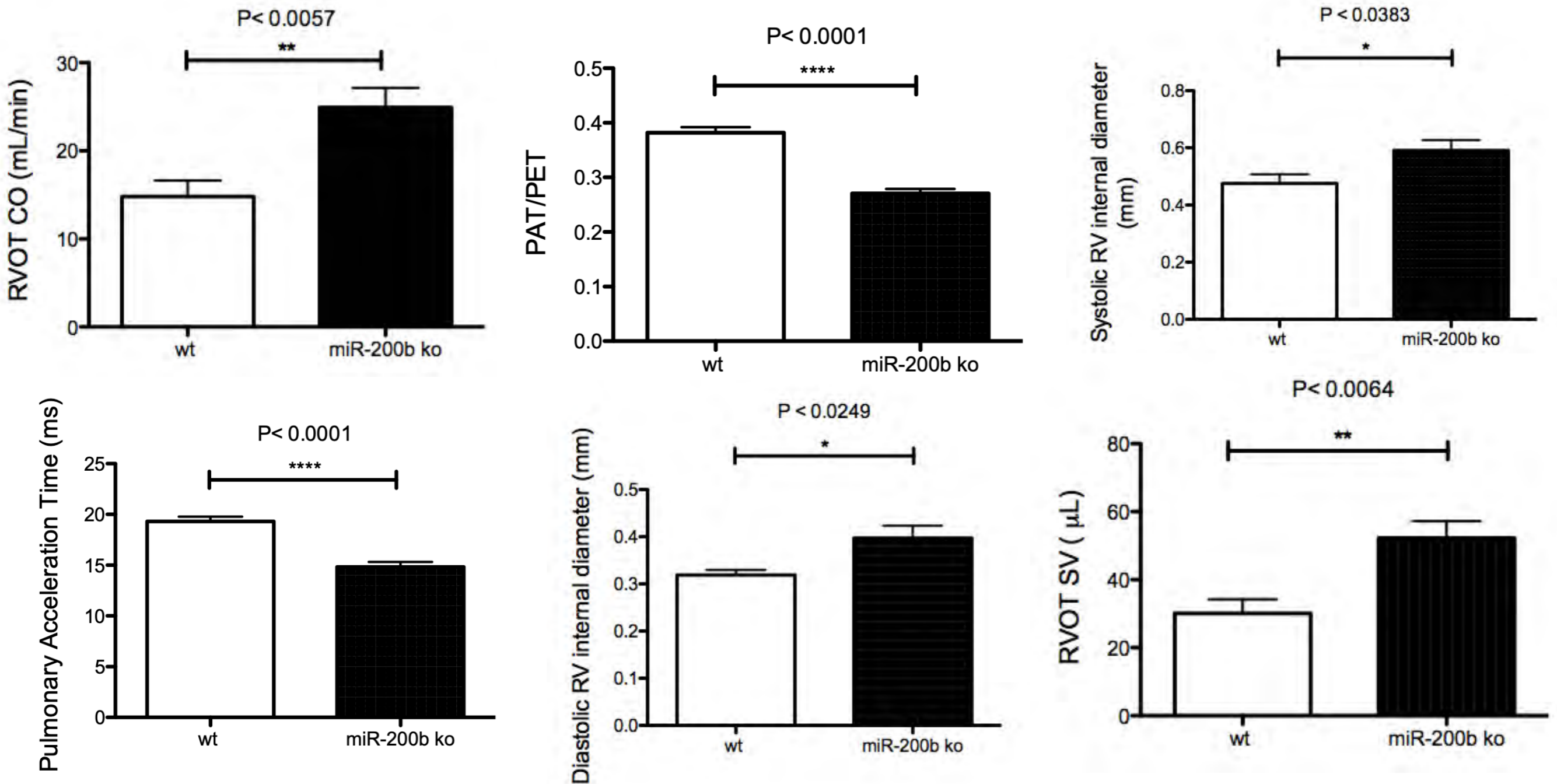


Wildtype

miR-200b -/-

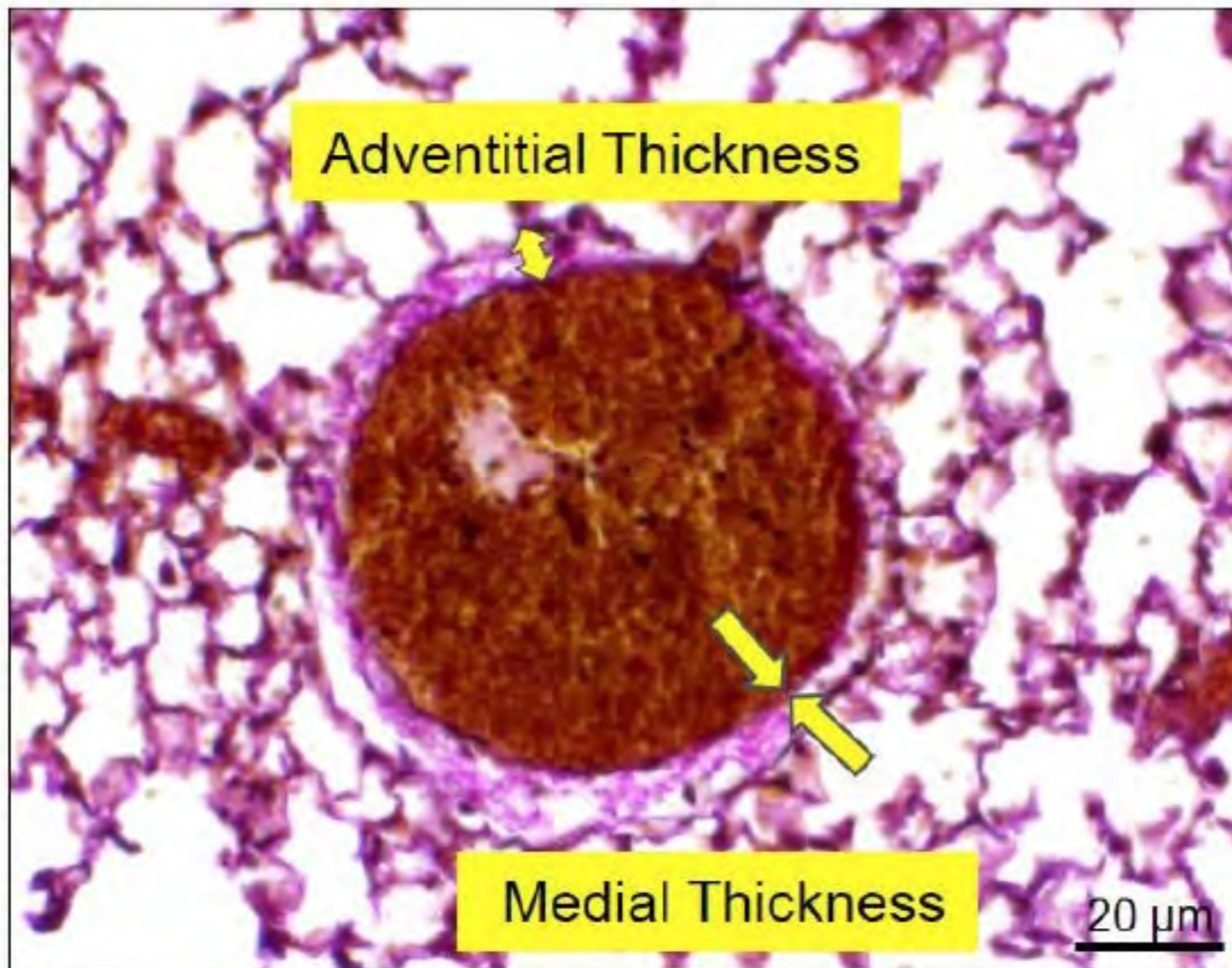


miR-200b knockout mice have pulmonary hypertension on cardiac echography

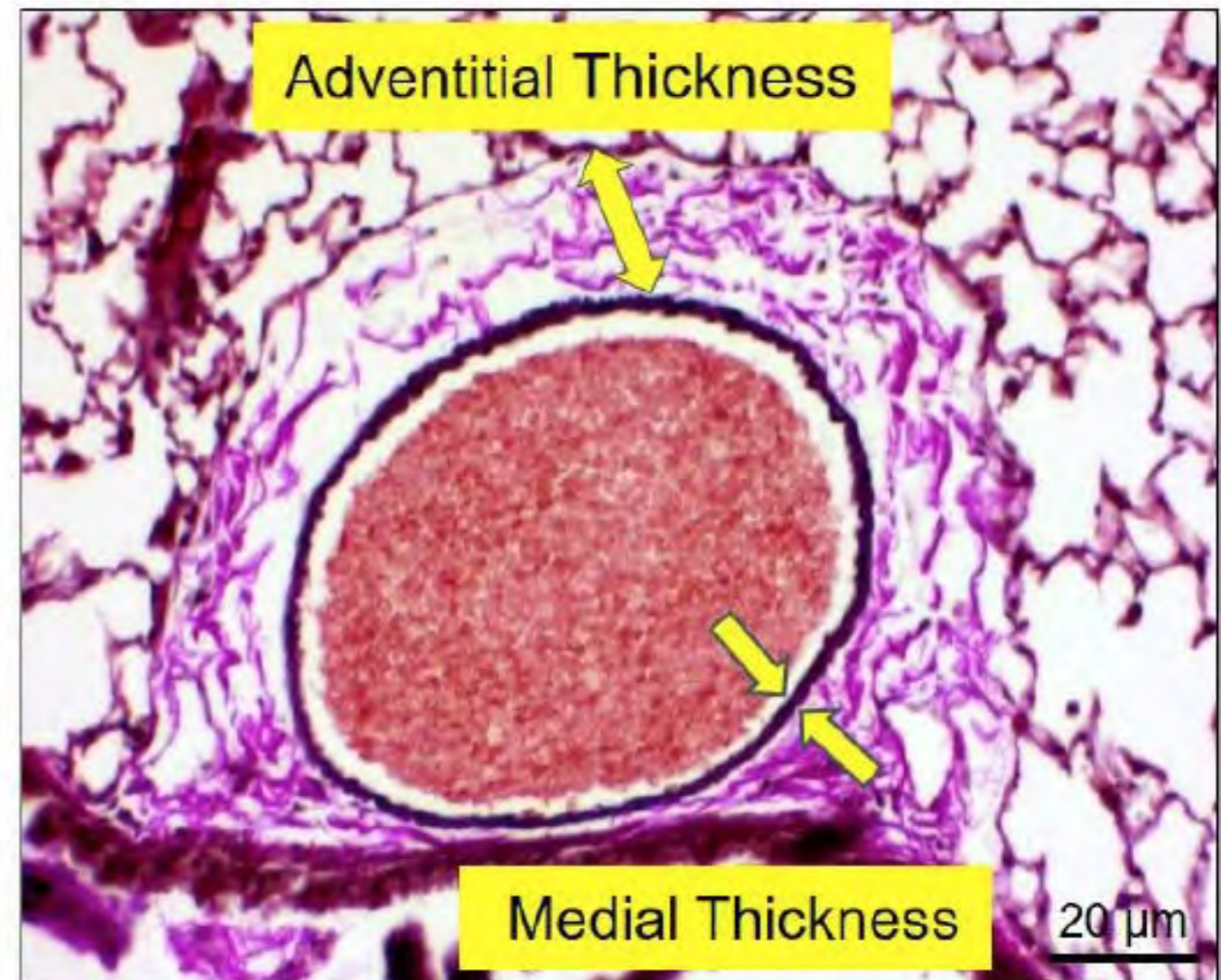


miR-200b^{-/-} lungs have thicker vessel walls

WT

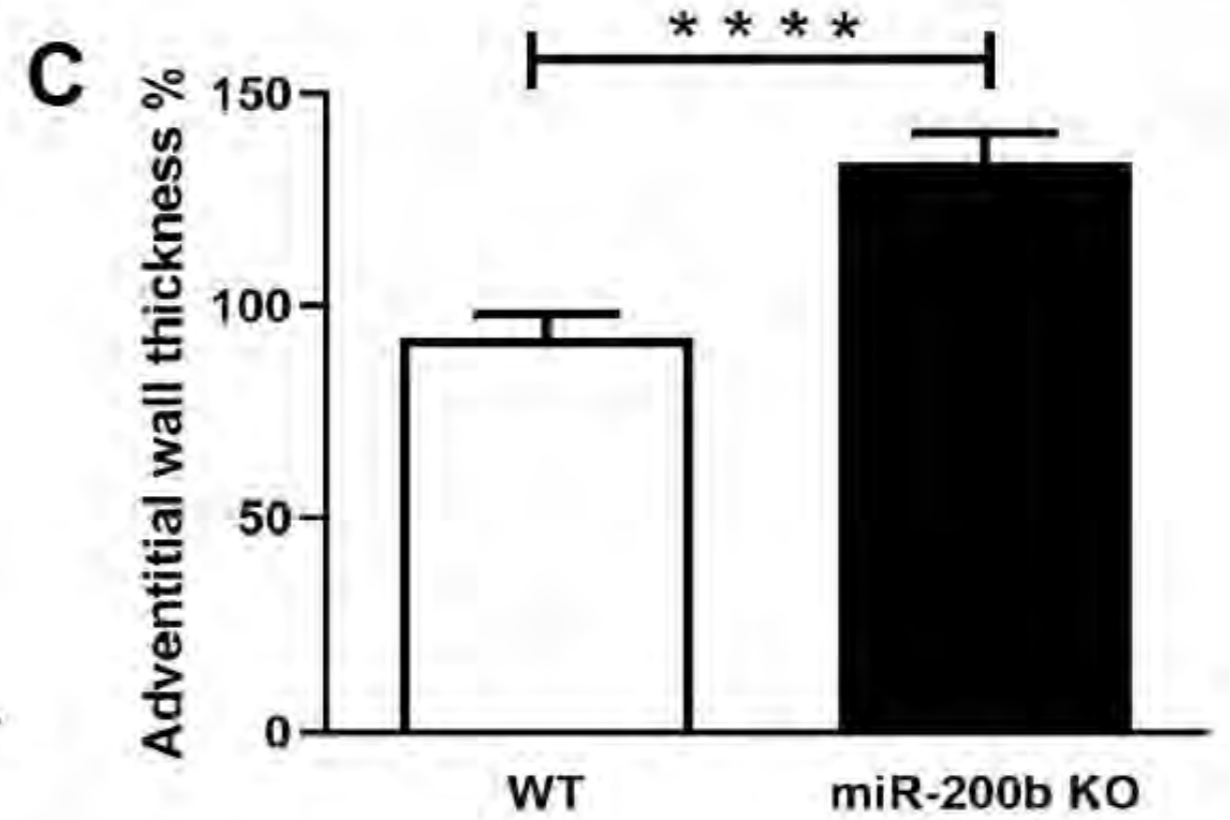
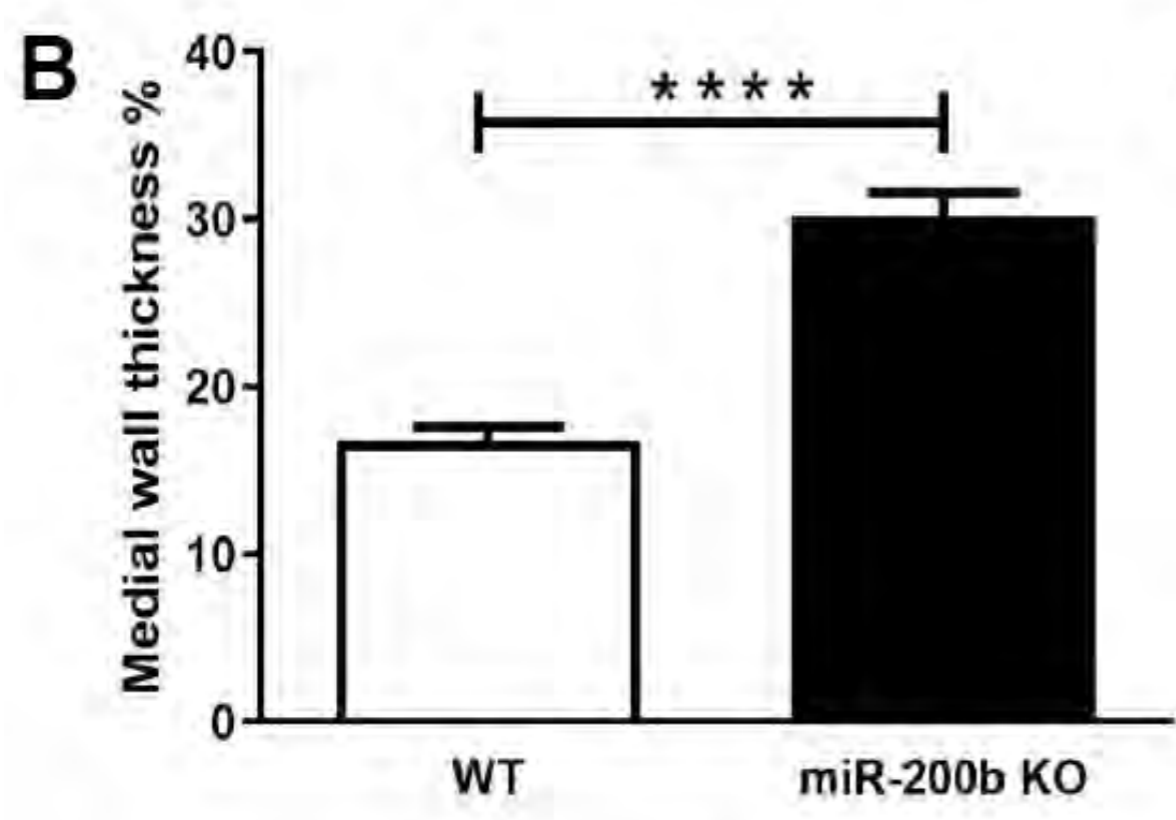
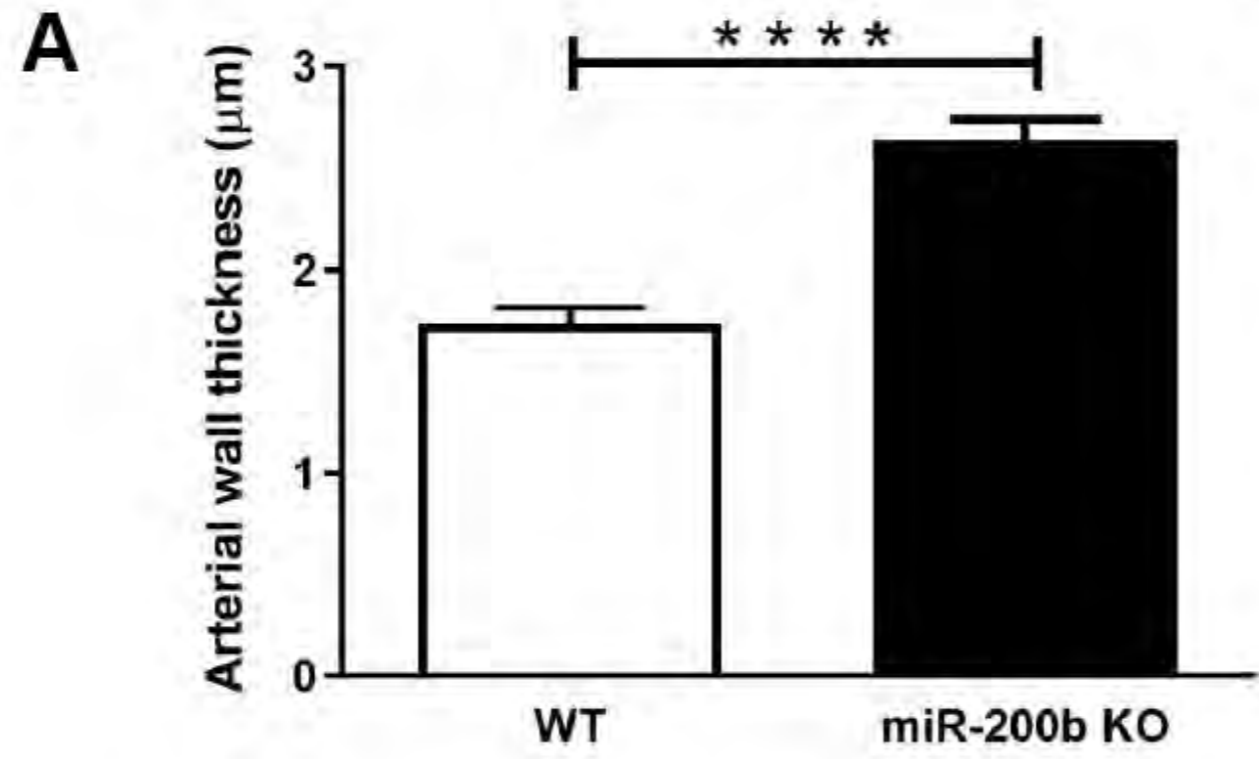


miR-200b KO



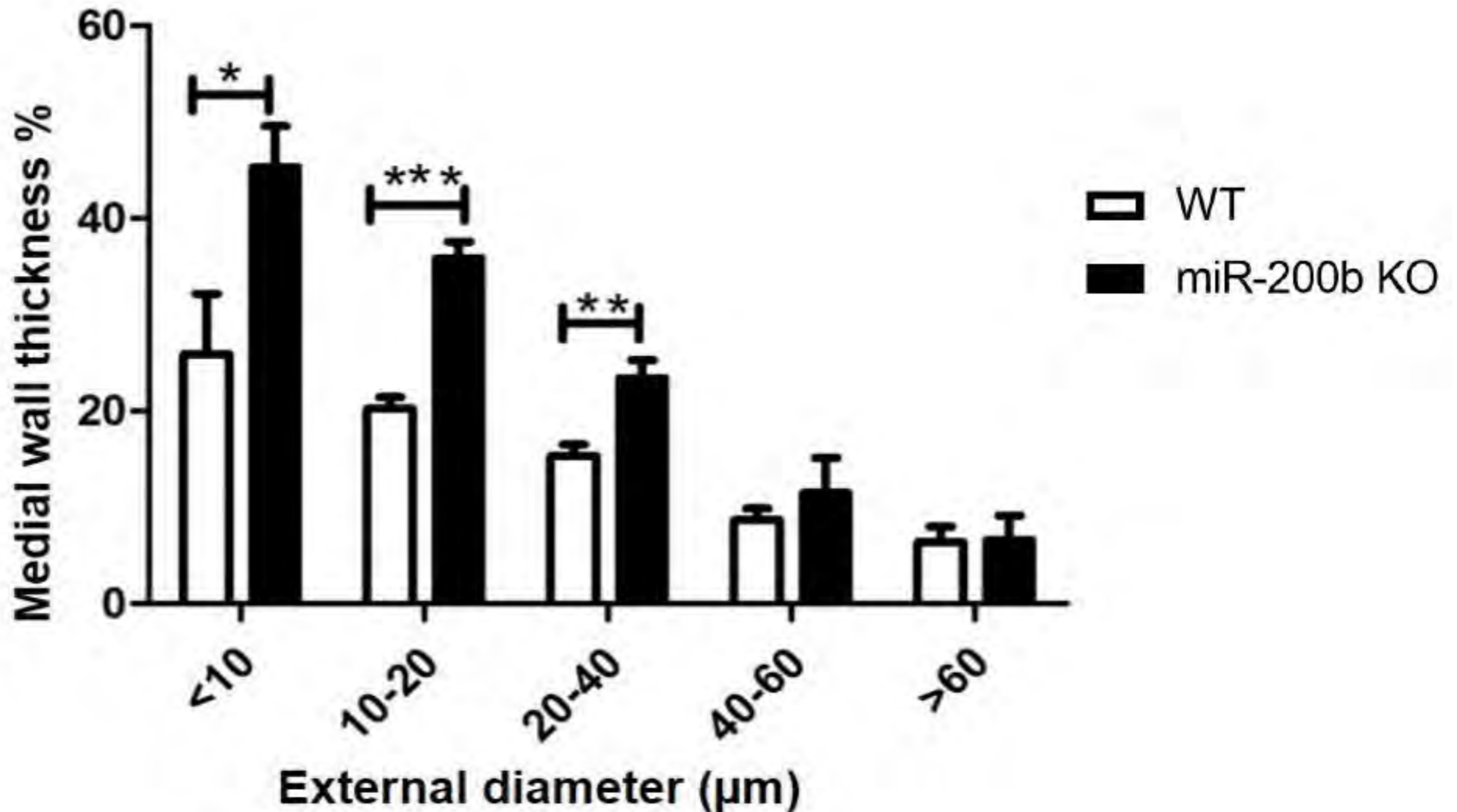
miR-200b^{-/-} lungs have thicker vessel walls

**** $P < 0.0001$

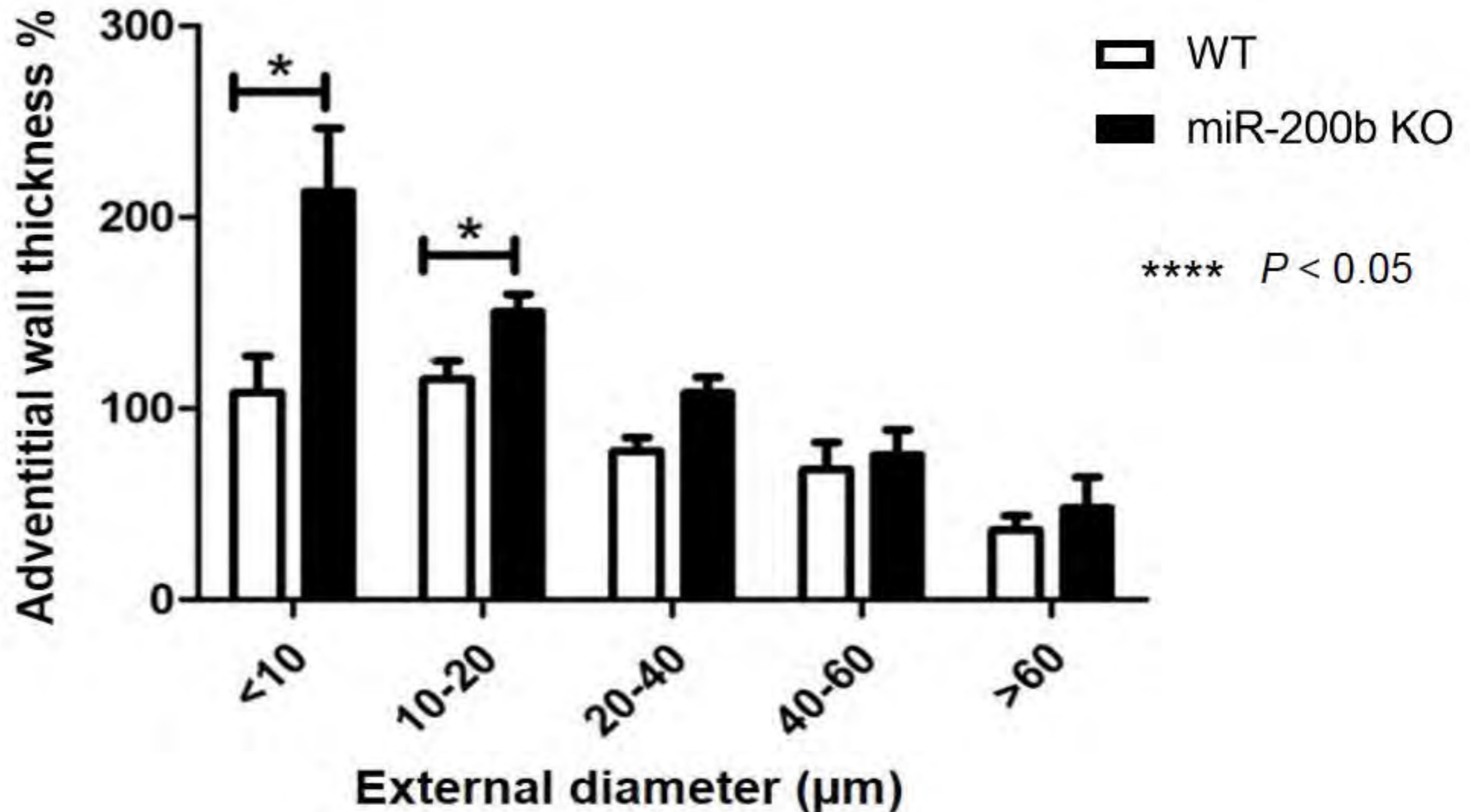


Unpublished results

miR-200b^{-/-} lungs have thicker vessel walls



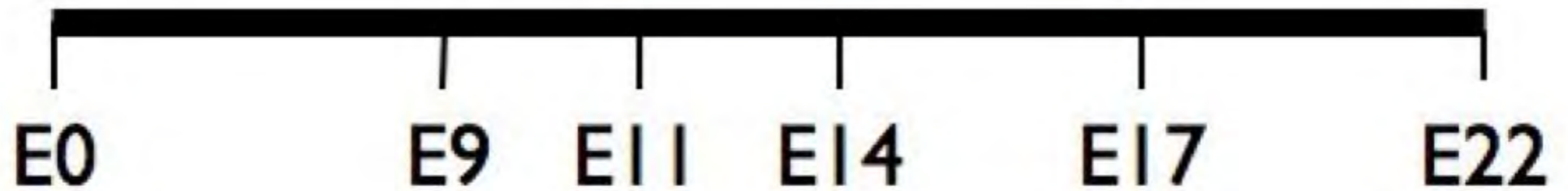
miR-200b^{-/-} lungs have thicker vessel walls



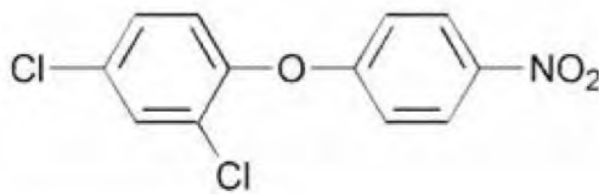
NITROFEN MODEL OF CDH



Lung Diaphragm Fetal breathing Birth



NITROFEN



80% CDH
100% PH



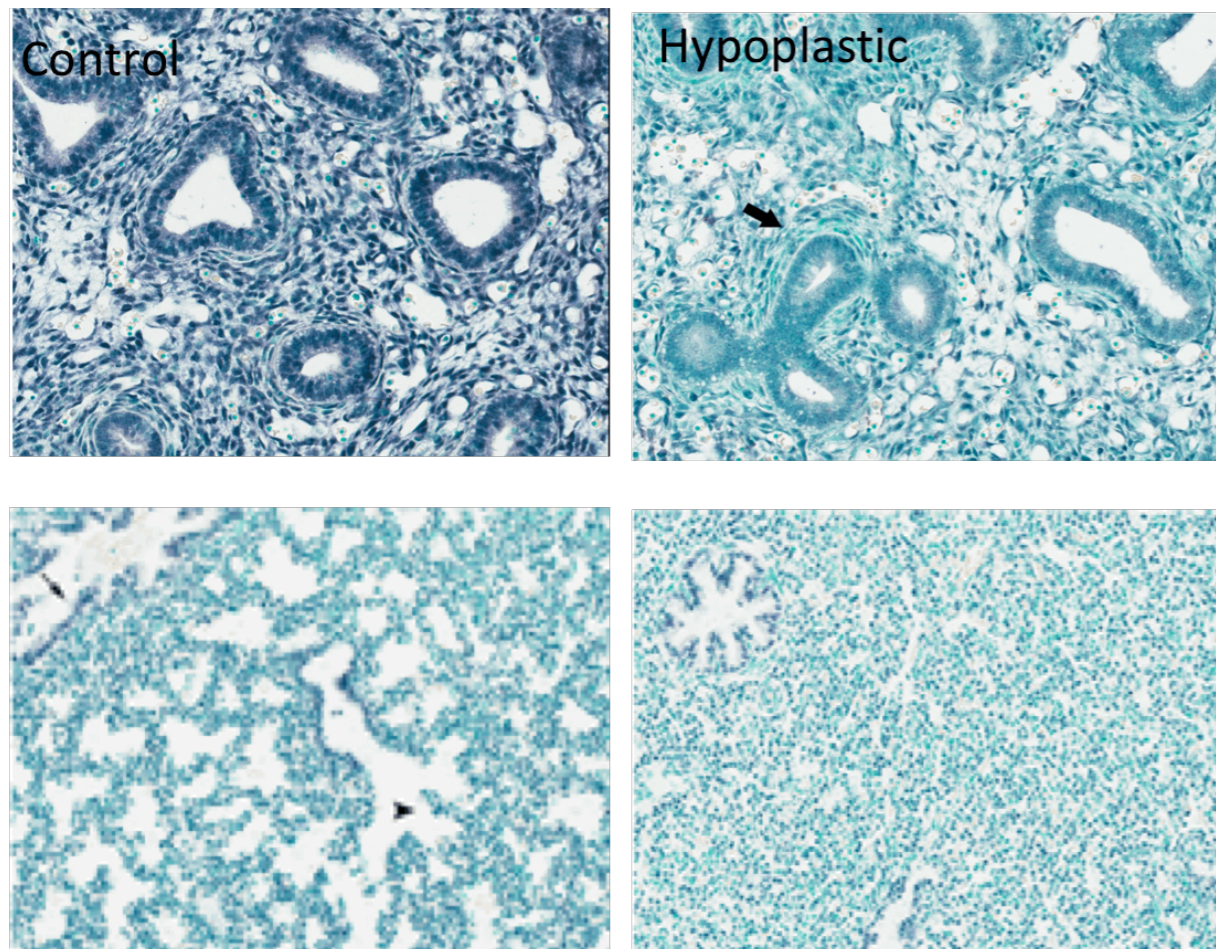
Prenatal microRNA miR-200b Therapy Improves Nitrofen-induced Pulmonary Hypoplasia Associated With Congenital Diaphragmatic Hernia

Naghmeleh Khoshgoo, MSc, Ramin Kholdebarin, MD, MSc,* Patricia Pereira-Terra, PhD,*†
Thomas H. Mahood, MSc,* Landon Falk, BSc,* Chelsea A. Day, BSc,* Barbara M. Iwaszow, MSc,*
Fuqin Zhu, BSc,* Drew Mulhall, BSc,* Carly Fraser, BSc,* Jorge Correia-Pinto, MD, PhD,†‡
and Richard Keijzer, MD, PhD, MSc, FACS**

In Situ Hybridization

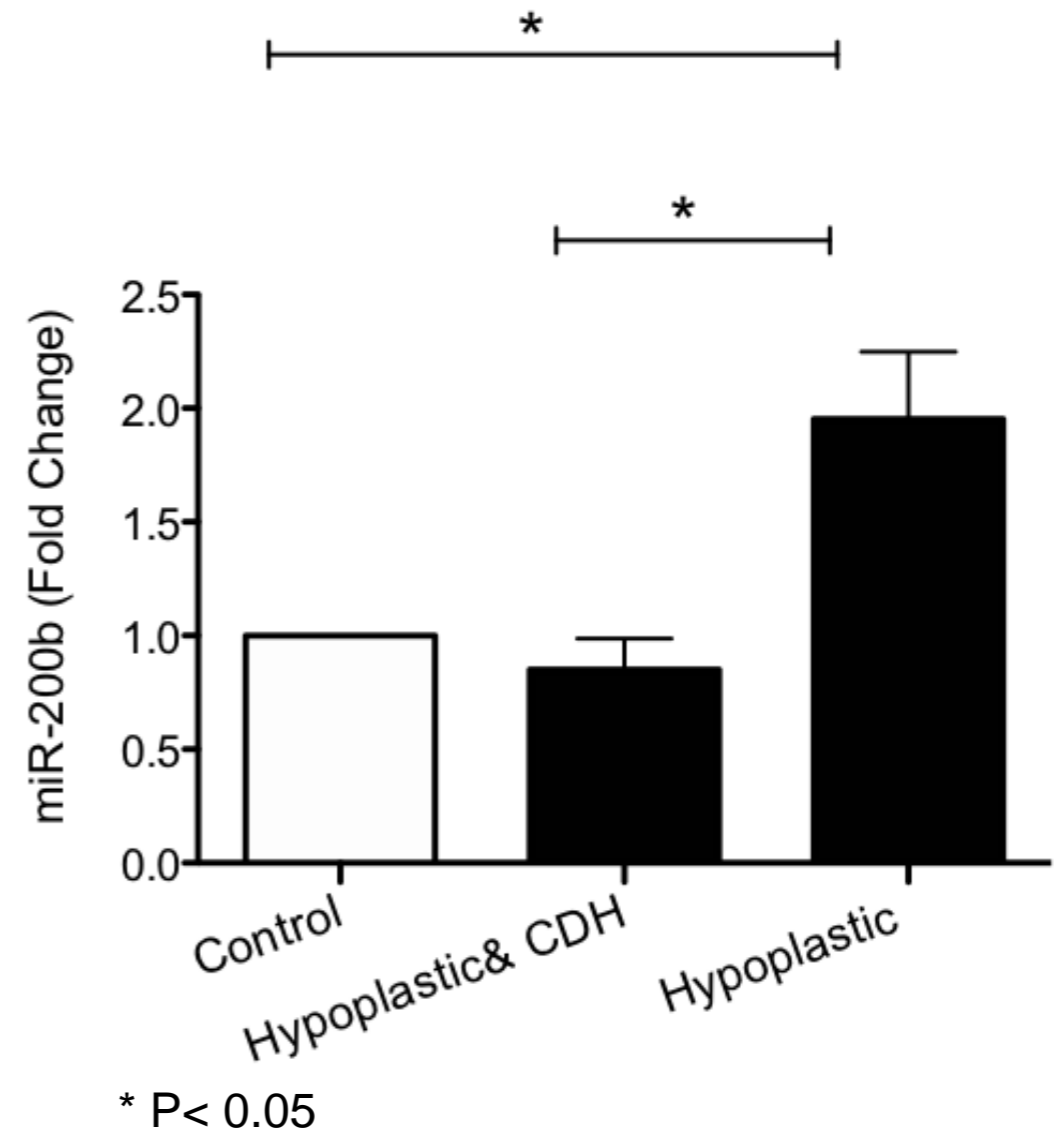
Control

Hypoplastic+ CDH

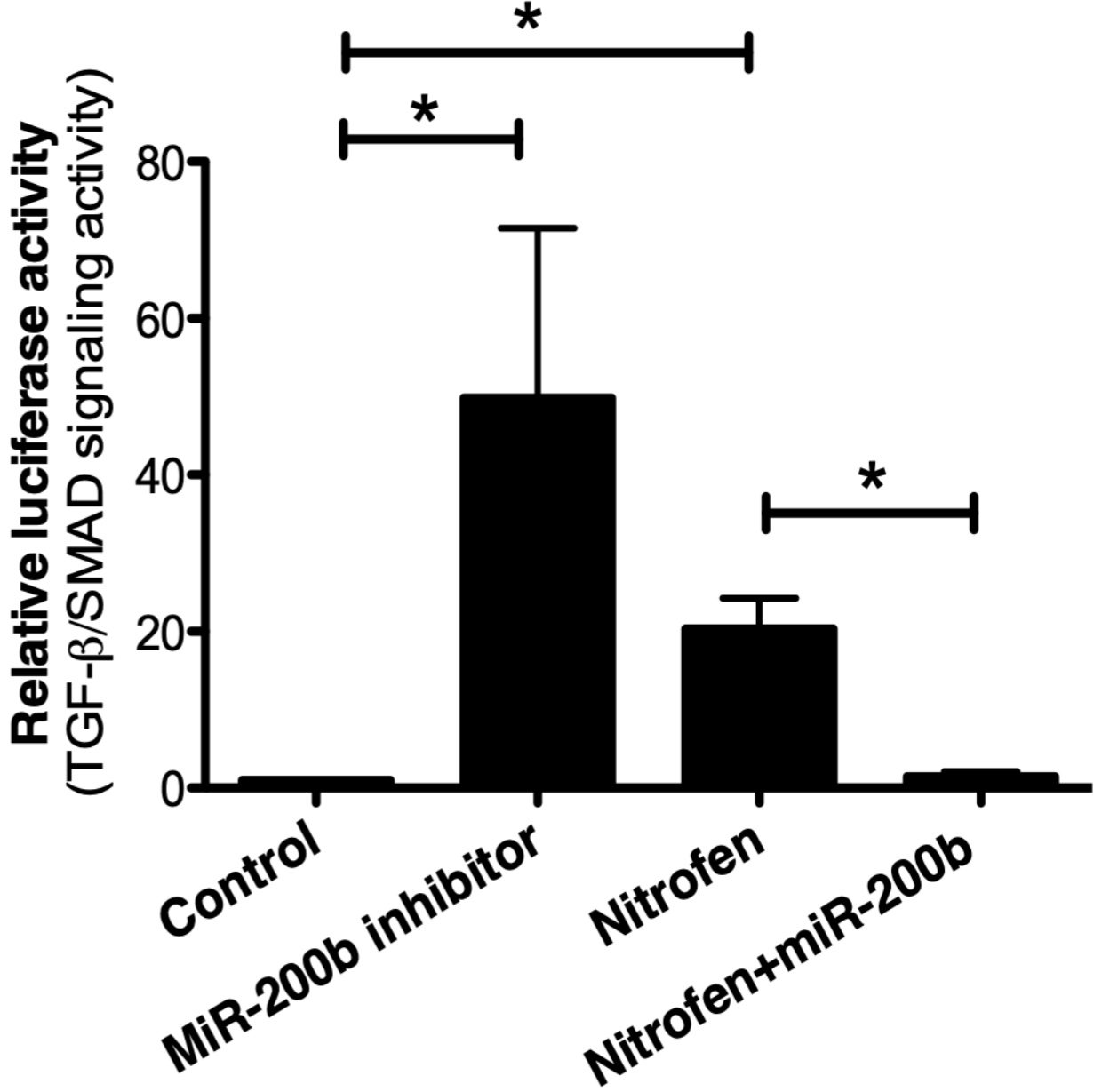
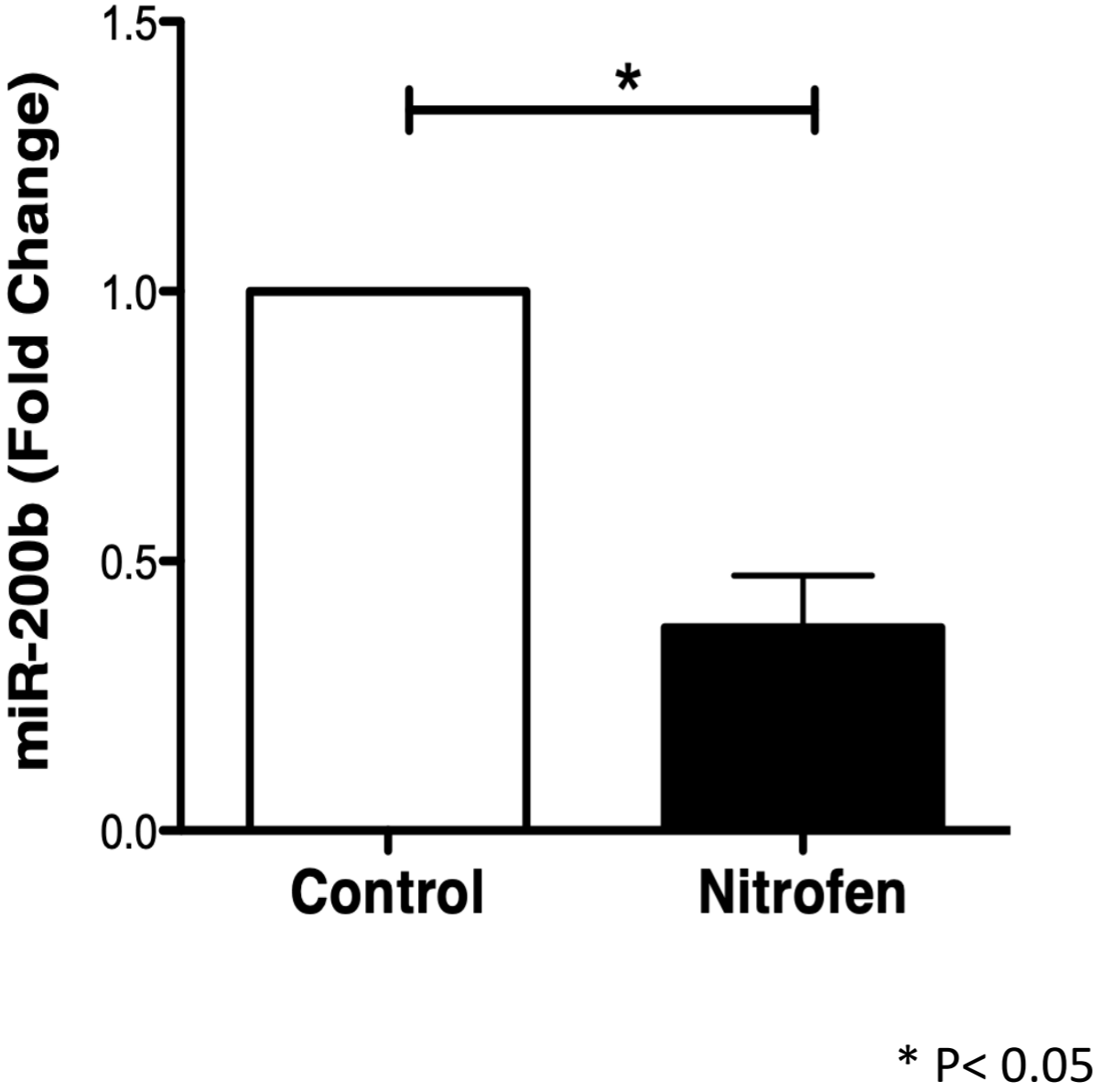


Blue staining: miR-200b

PCR (E21)



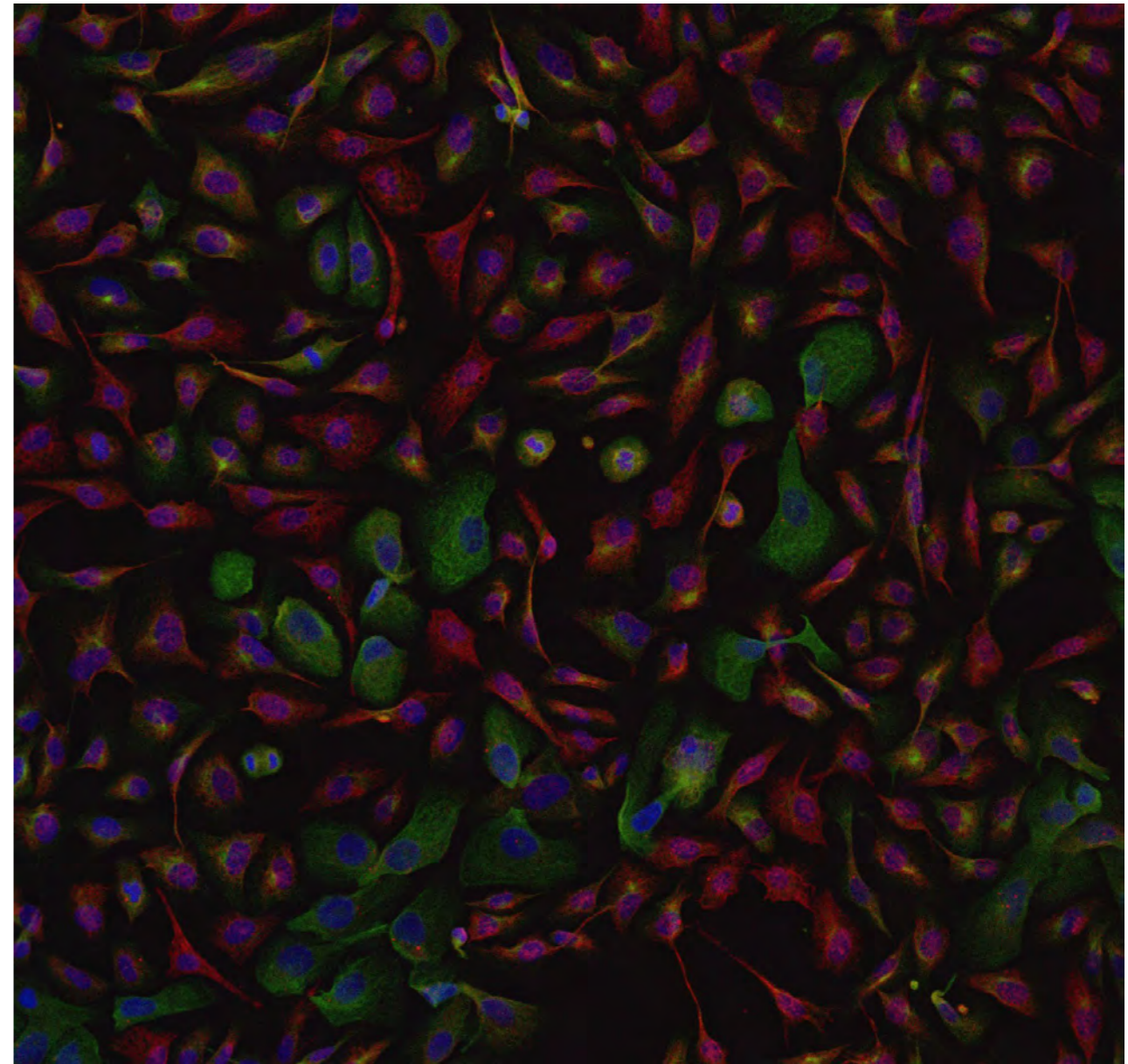
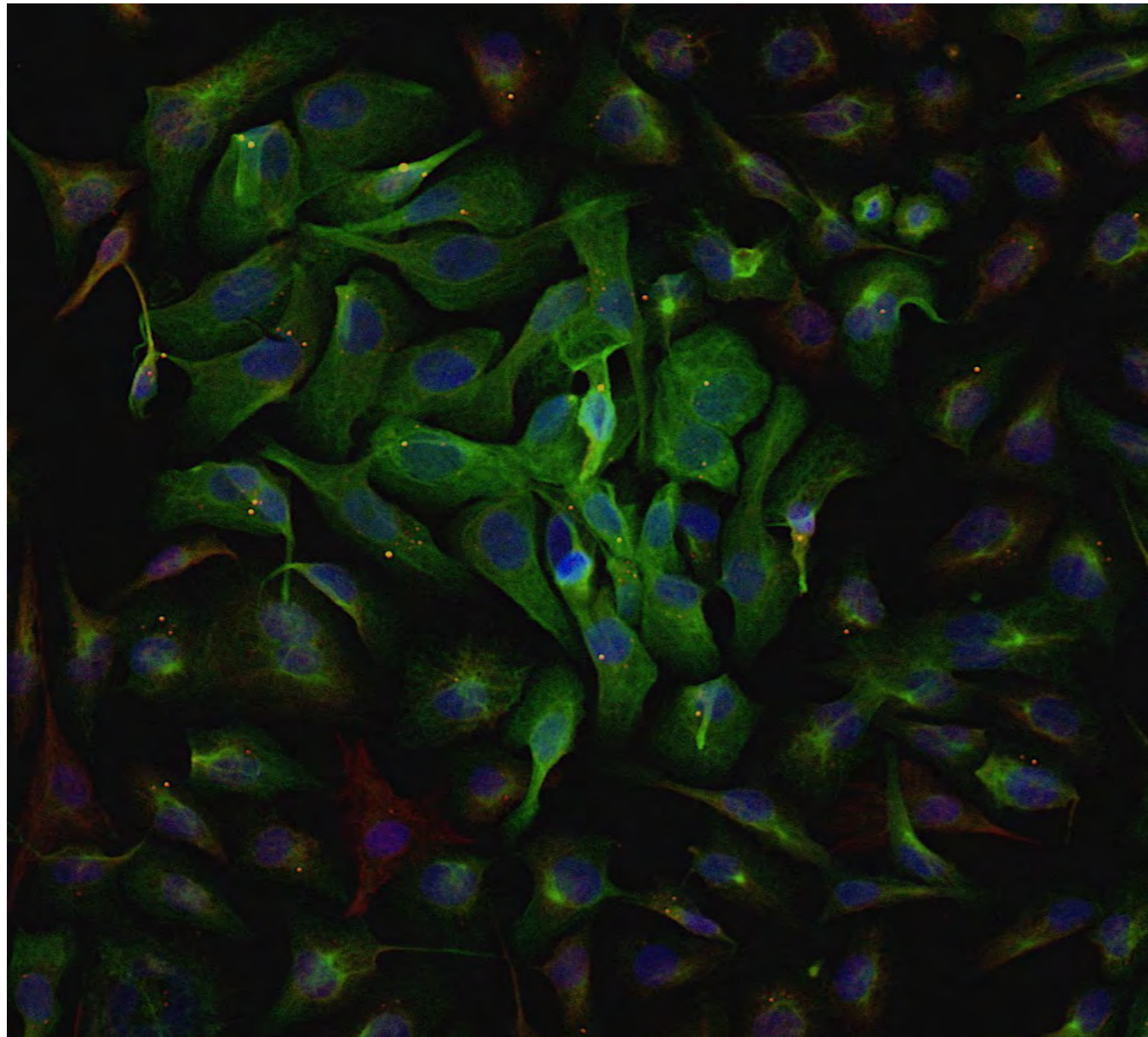
MiR-200b & Human Bronchial Epithelial Cells



miR-200b maintained epithelial cell phenotype in bronchial epithelial cells

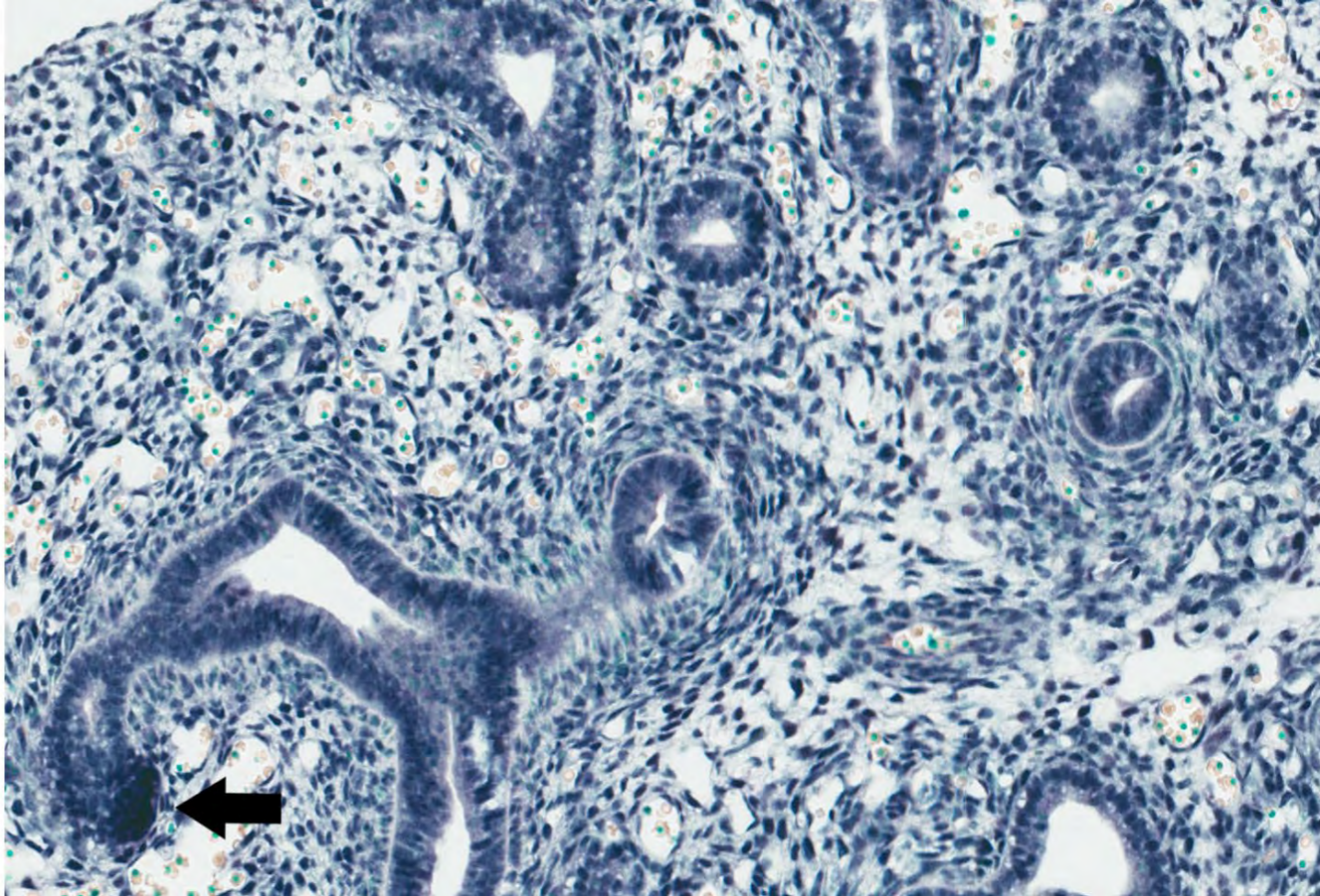
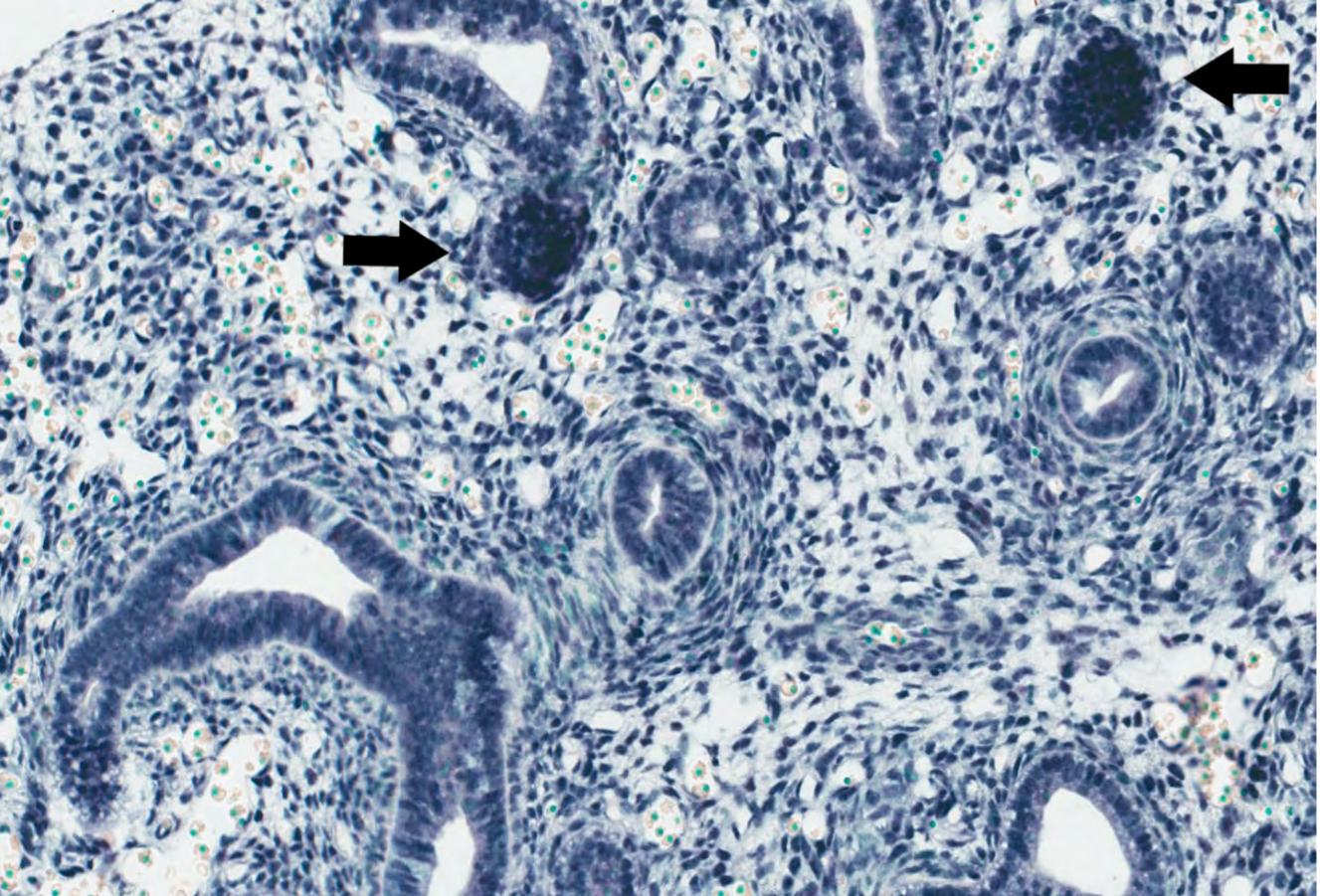
Control

MiR-200b inhibitor



Green : Epithelial Marker (cytokeratin)
Red: Mesenchymal Marker (Vimentin)

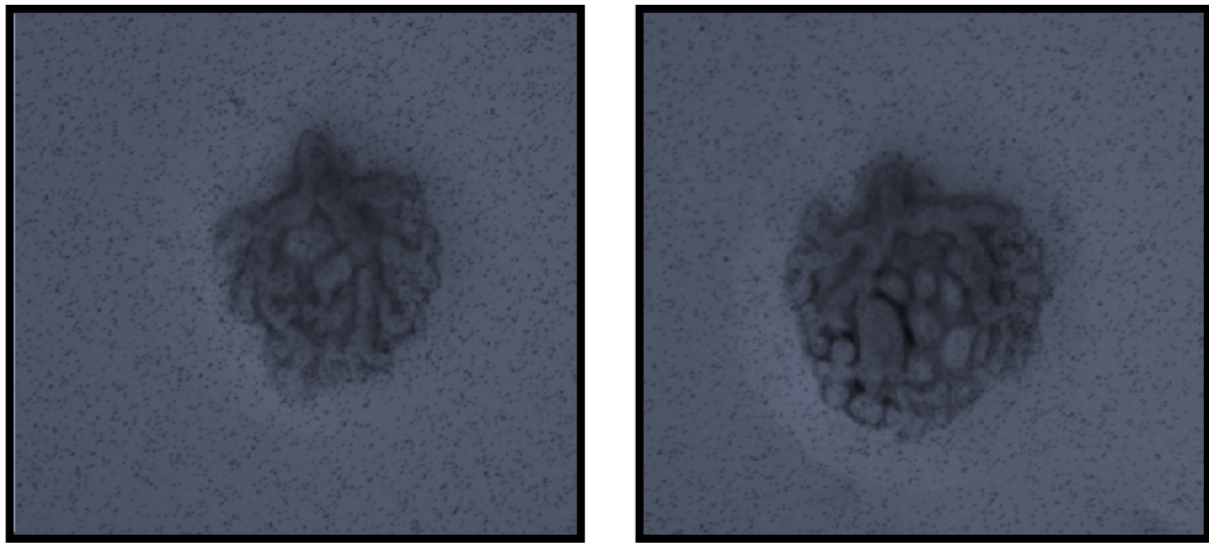
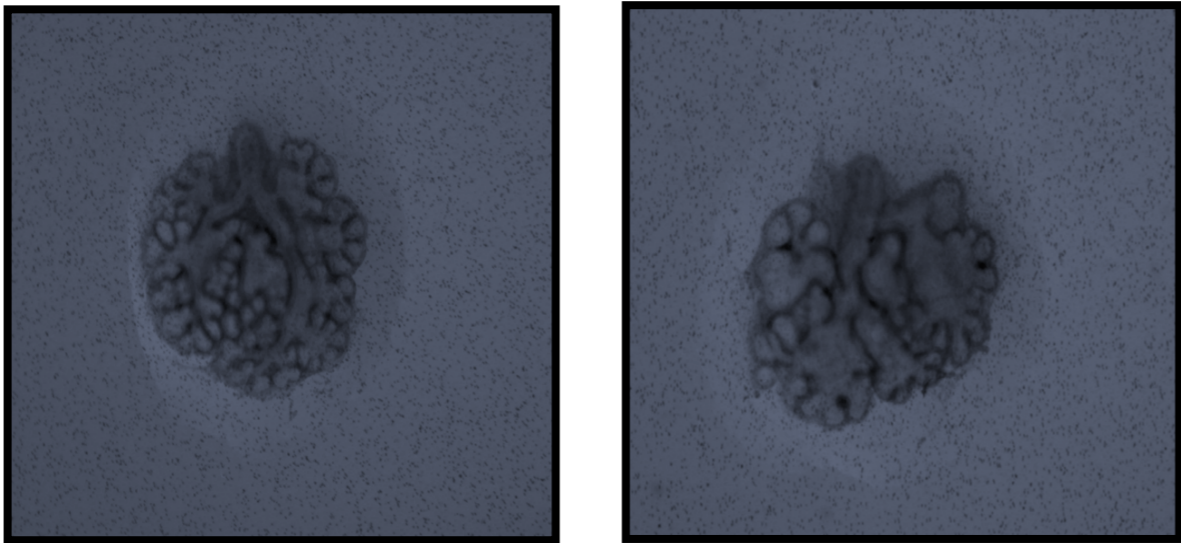
miR-200b expression is high at branching lung tips



miR-200b improves branching hypoplastic lungs

Normal lungs

Hypoplastic lungs

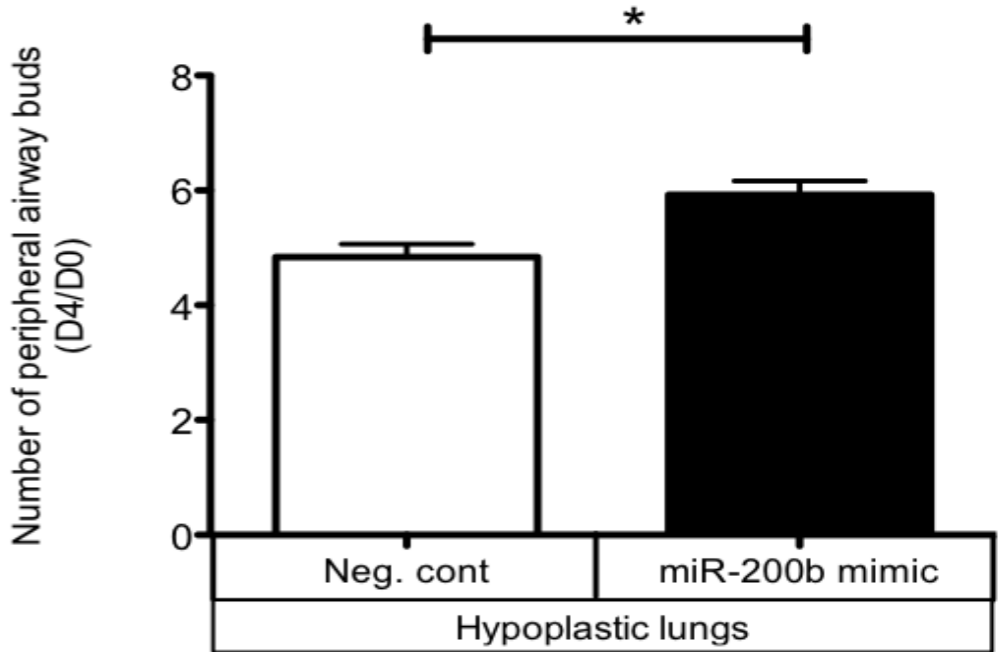
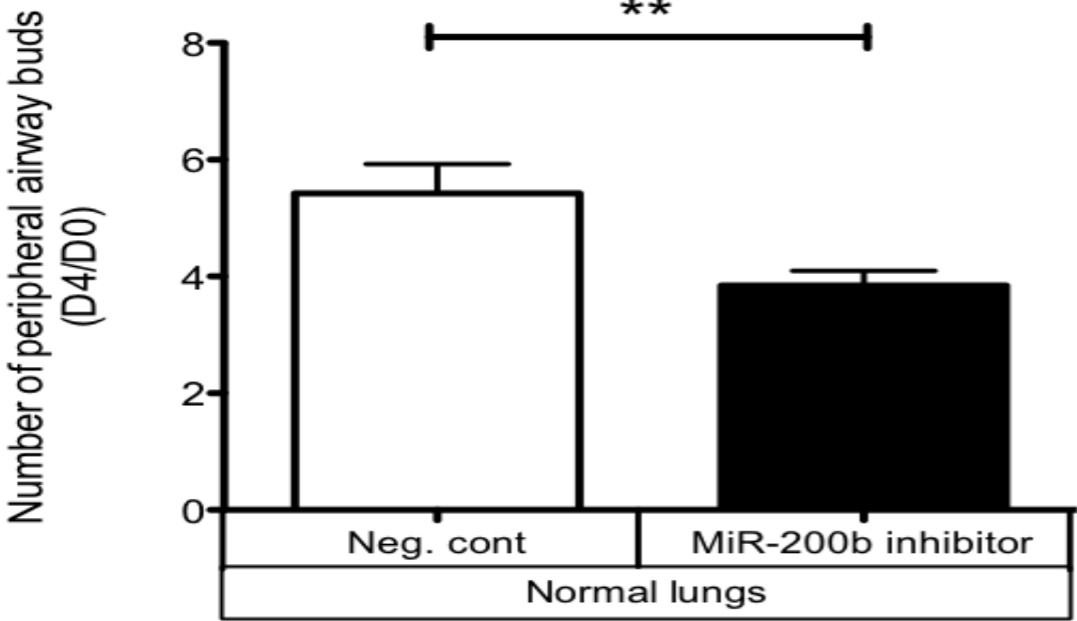


Control

200b inhibitor

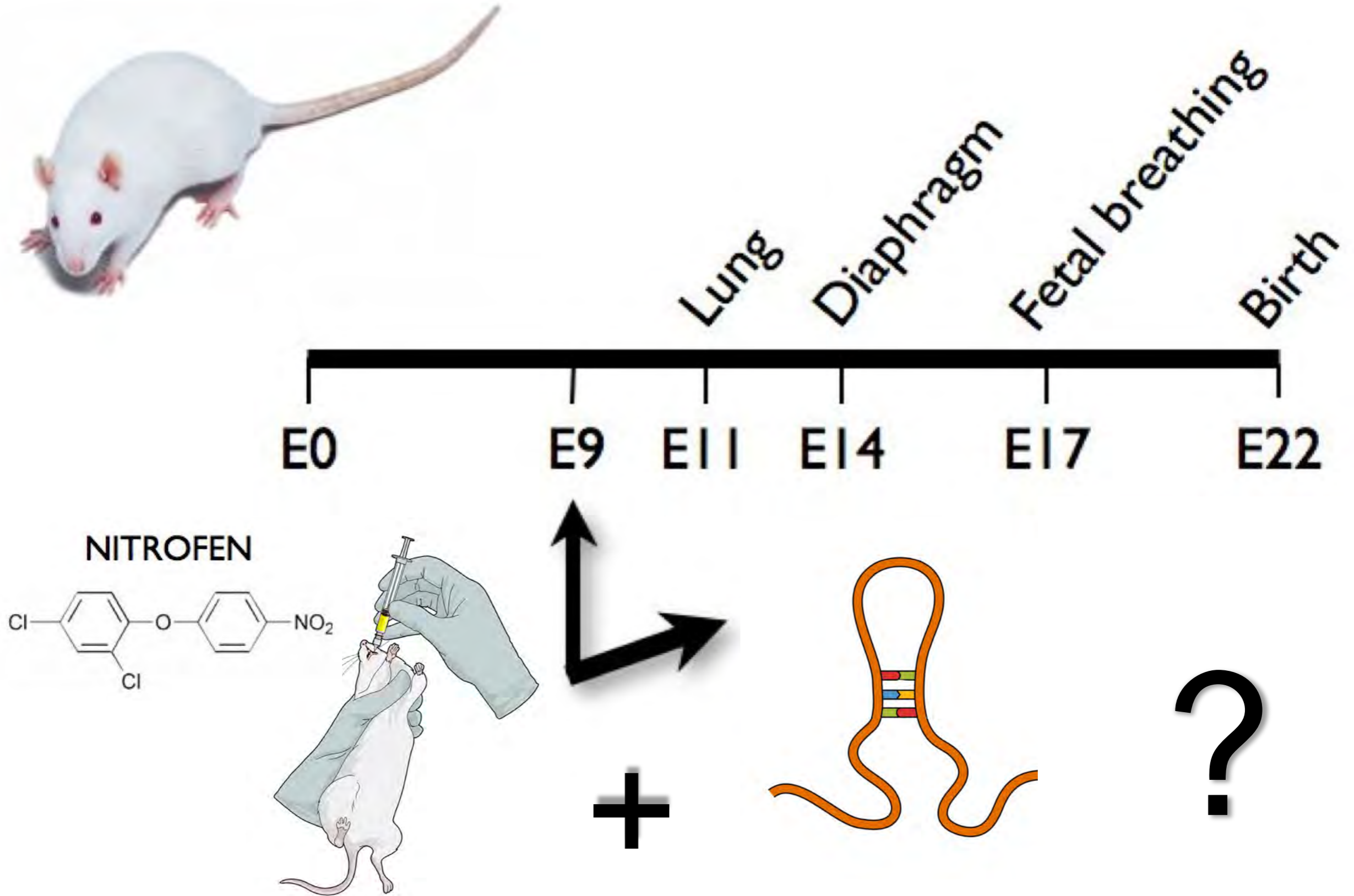
Hypoplastic

Hypoplastic+ miR-200b



* P < 0.05, ** P < 0.01

NITROFEN MODEL OF CDH



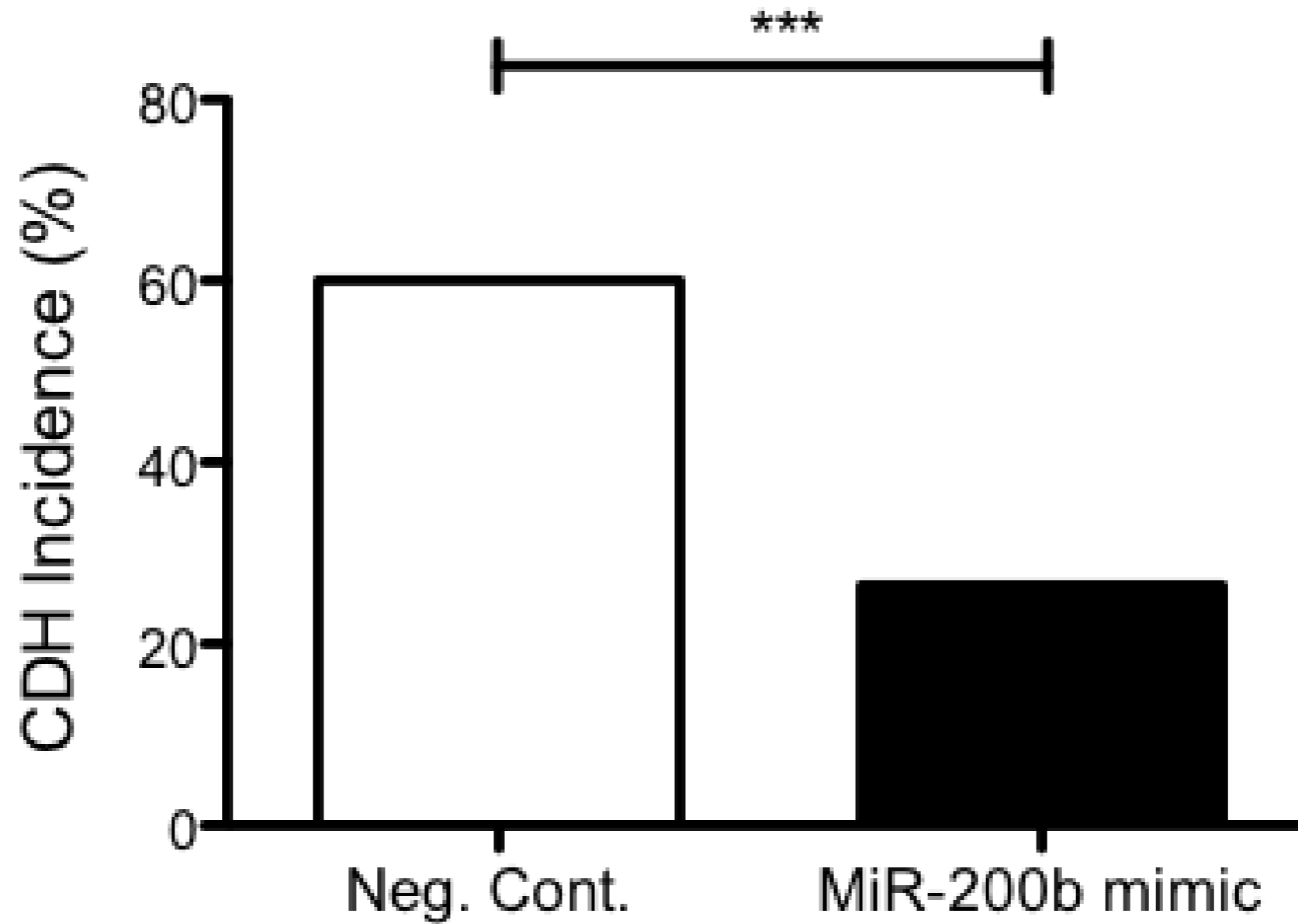
Prenatal miR-200b improves lung hypoplasia



hypoplastic
(80% CDH) ←

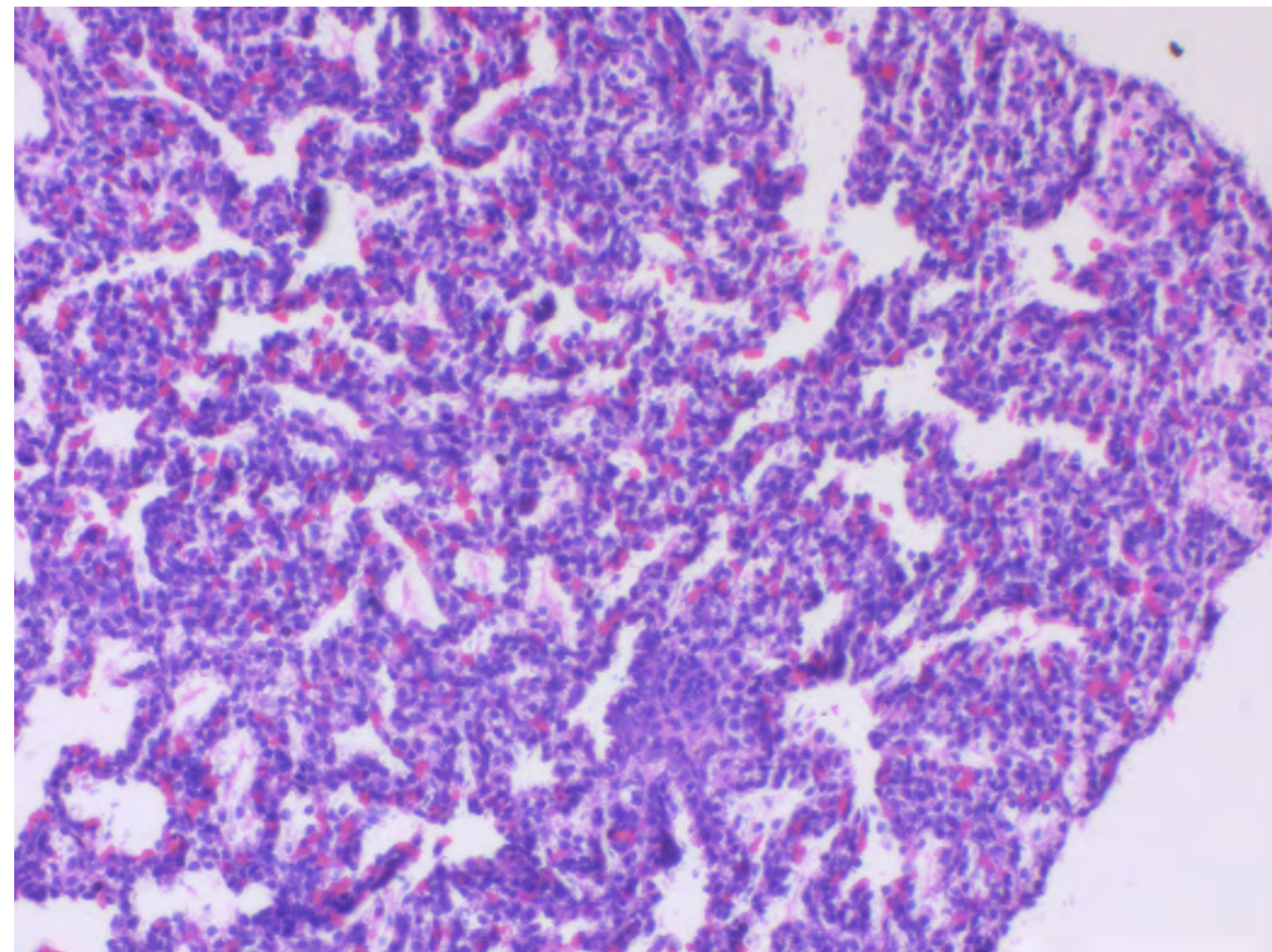
hypoplastic +
miR-200b
(15% CDH)

Prenatal miR-200b reduces CDH incidence

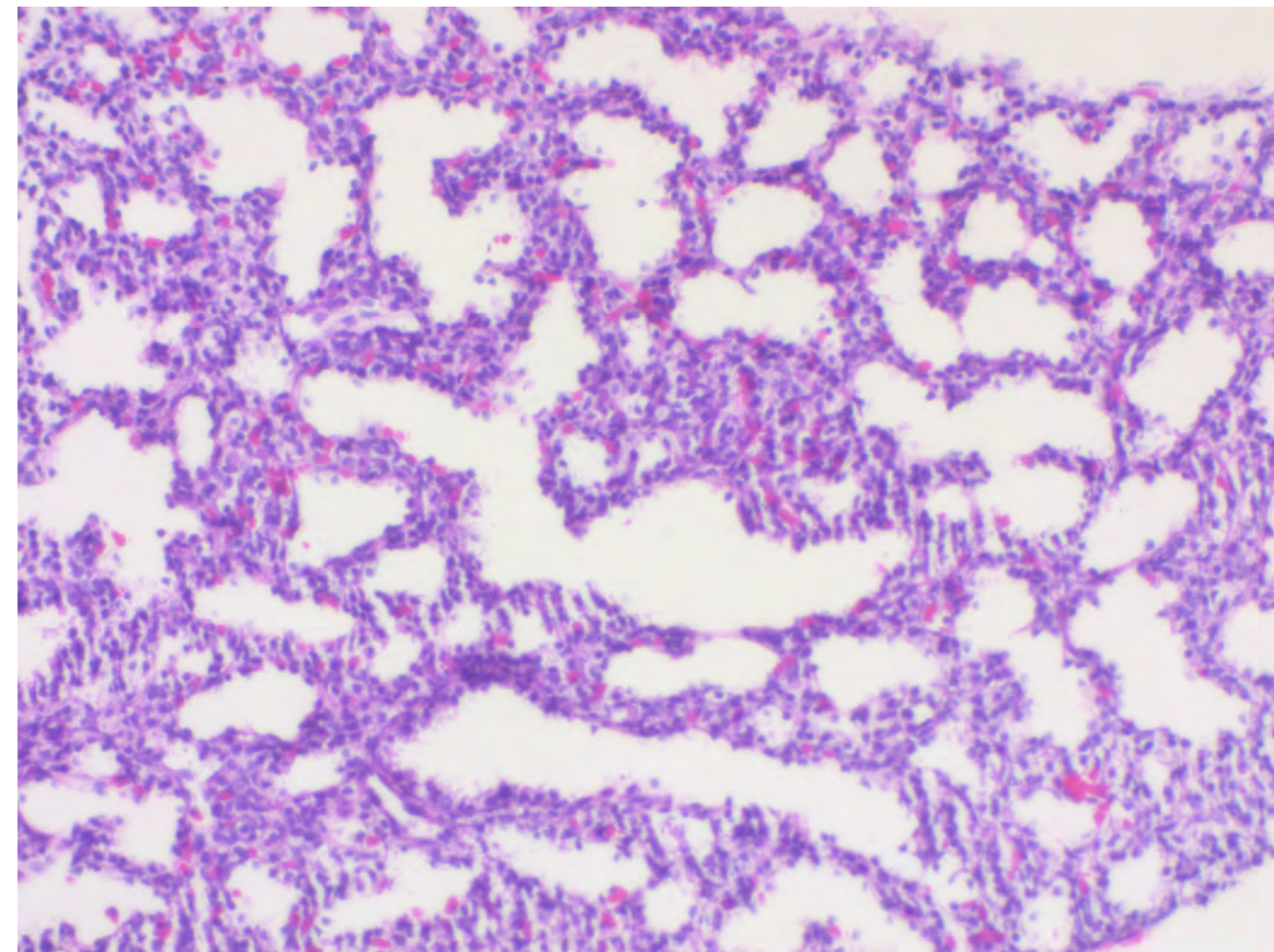


Prenatal miR-200b improves lung hypoplasia

Negative Control

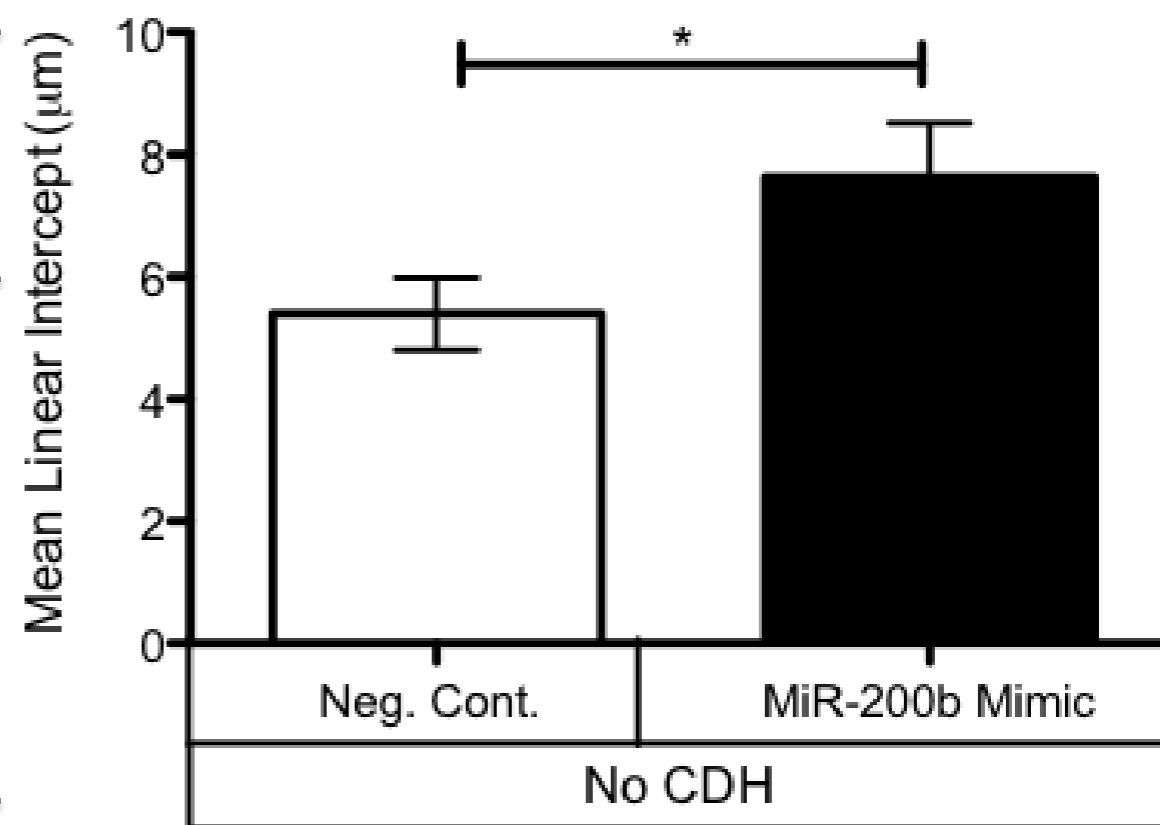
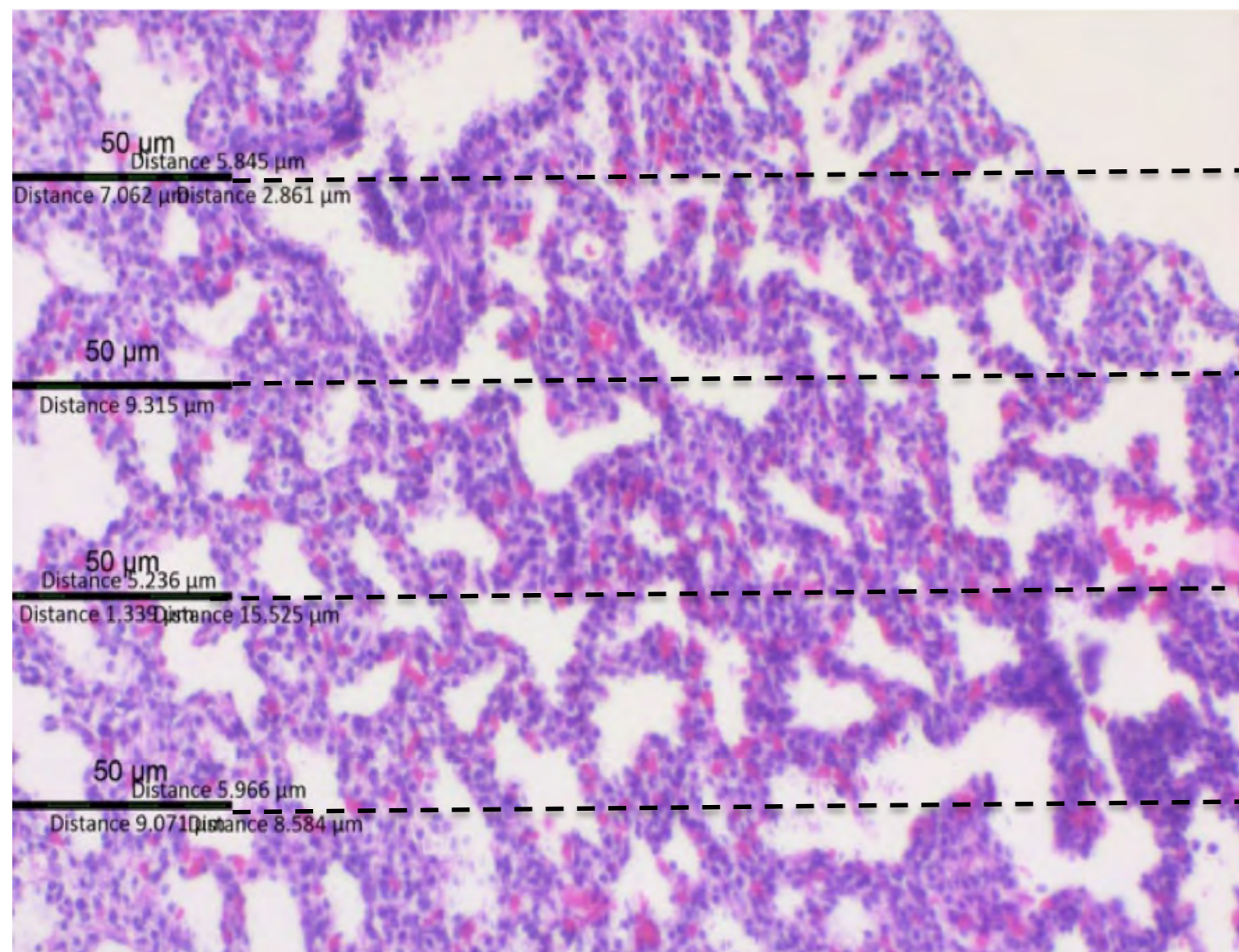


MiR-200b



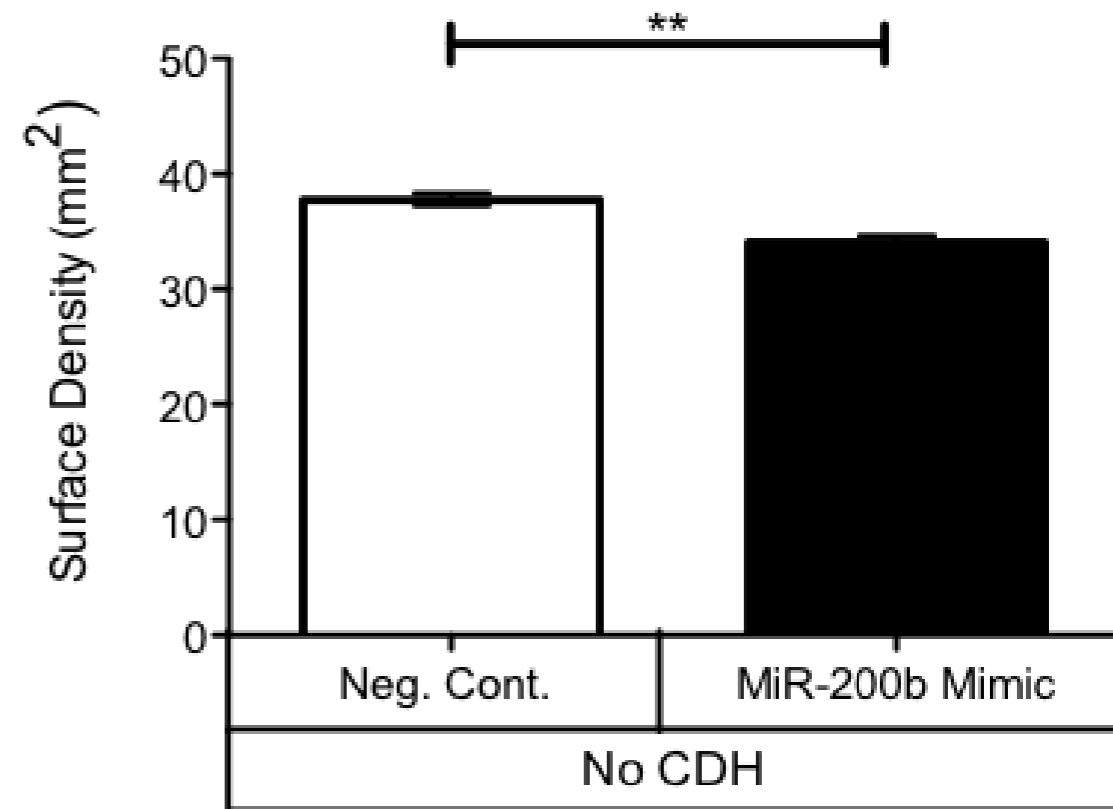
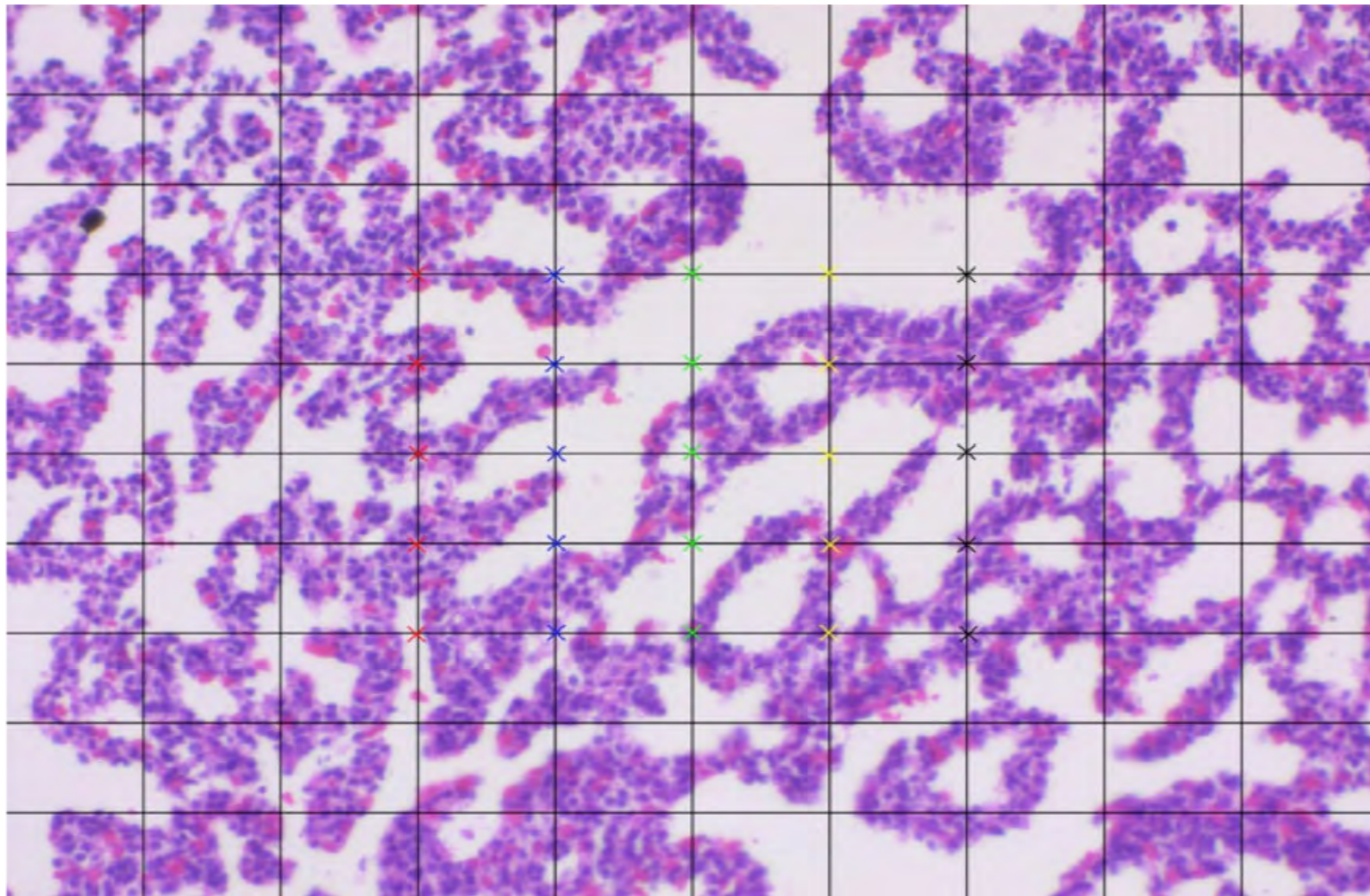
Prenatal miR-200b improves lung hypoplasia

Mean Linear Intercept



Prenatal miR-200b improves lung hypoplasia

Surface Density



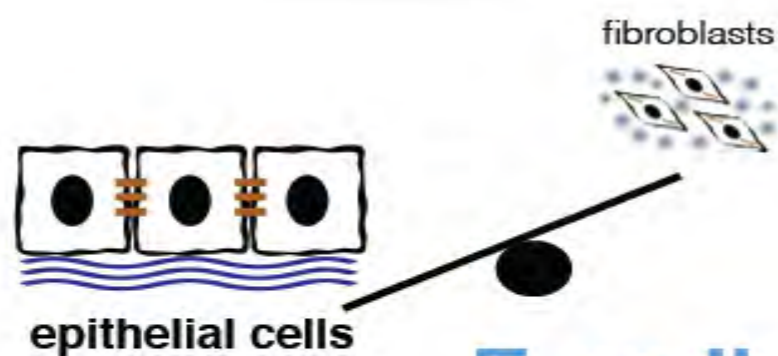
control lung



CDH lung

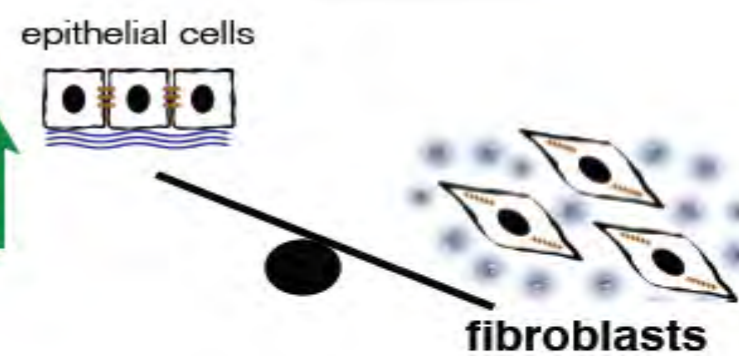


miR-200b



E-cadherin ↑

TGF-beta ↑



fibroblasts

ZEB

miR-200b ↑



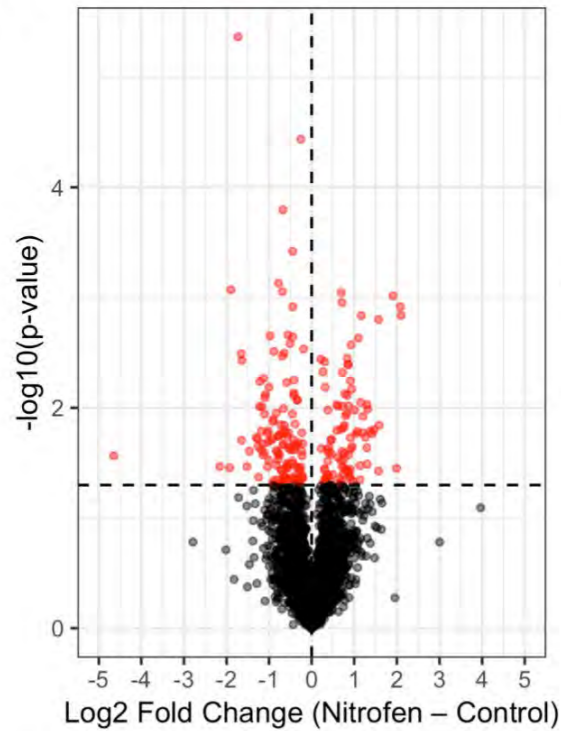
normal lung development



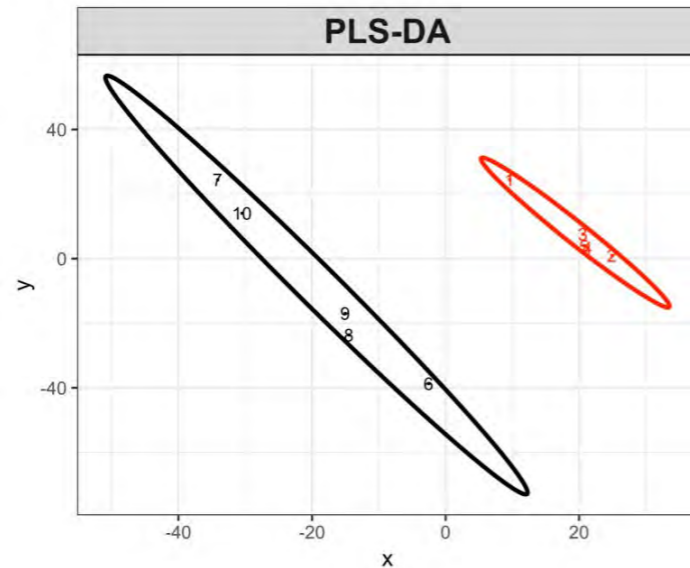
lung hypoplasia

Proteome suggests that nitrofen-induced abnormal lung development is an immune disease?

A



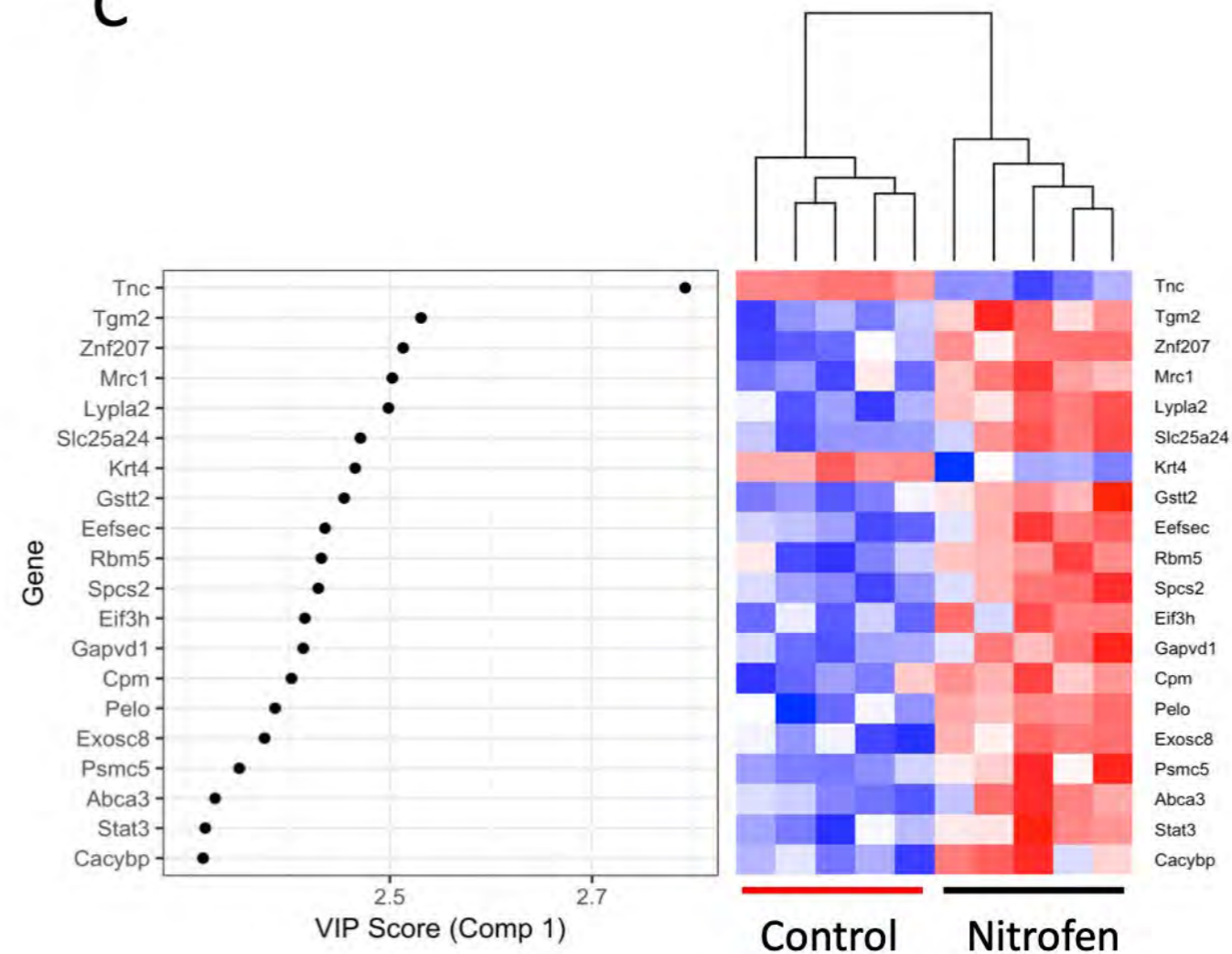
B



Legend

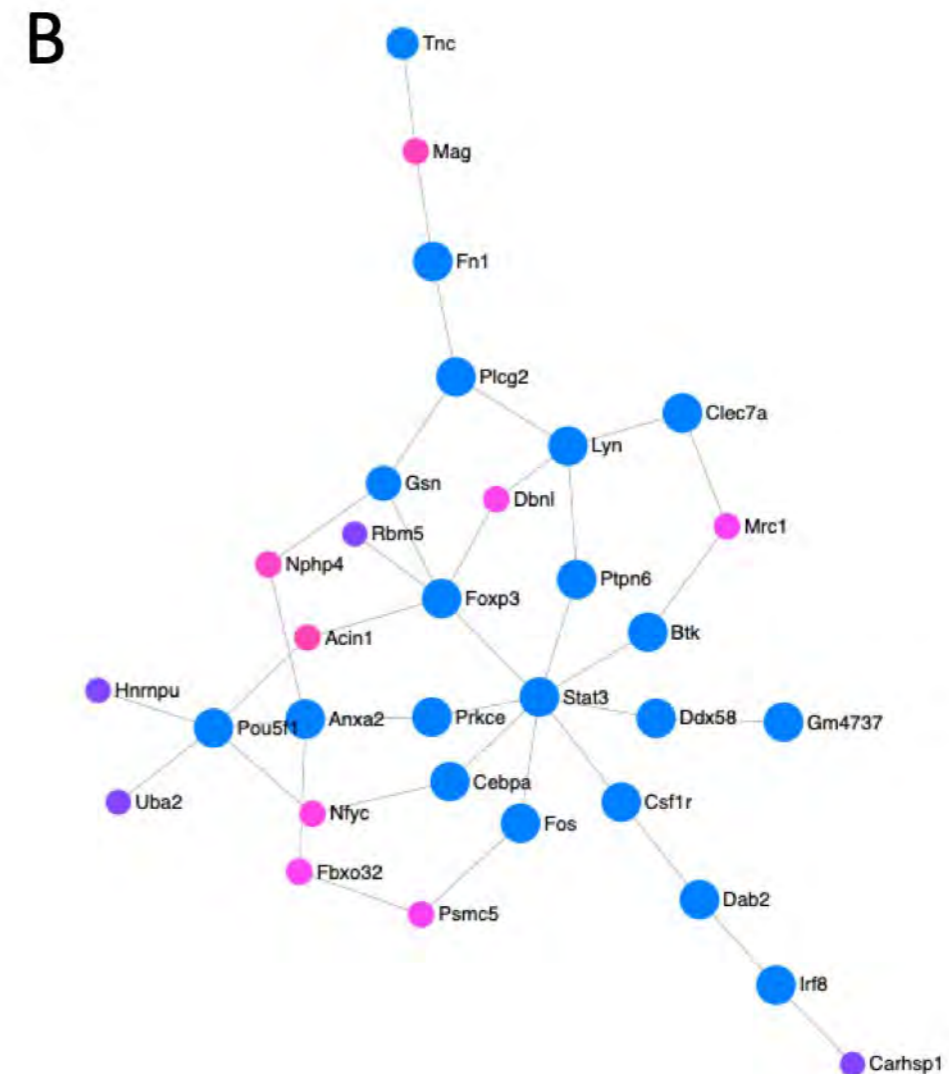
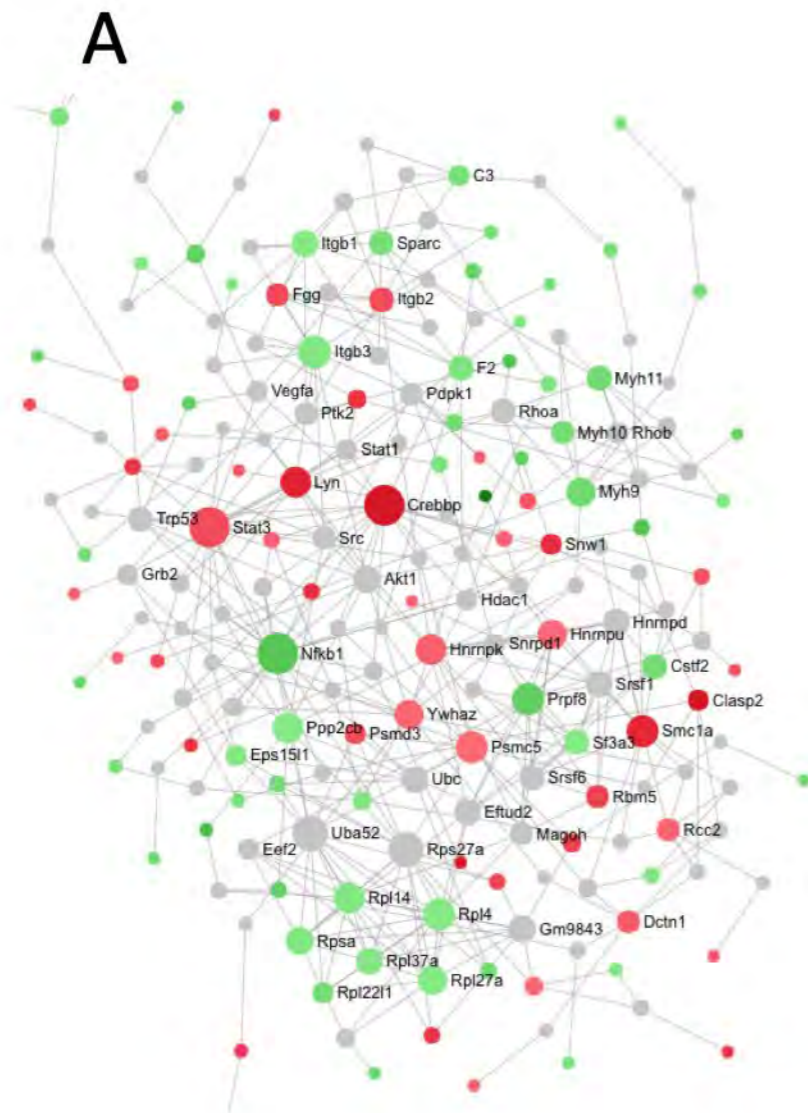
- Control
- Nitrofen

C



Unpublished results

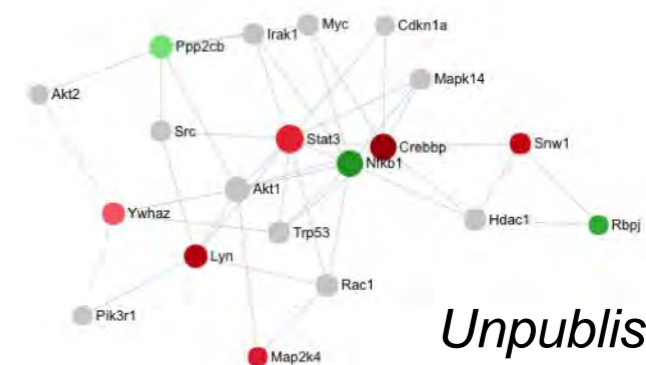
Proteome suggests that nitrofen-induced abnormal lung development is an immune disease?



C

Pathway	Total	Expected	Hits	P.Value	FDR
Immune System	1110	27.3	62	4.25E-11	3.33E-08
Signaling by Interleukins	118	2.91	19	5.09E-11	3.33E-08
Cytokine Signaling in Immune system	270	6.66	26	1.62E-09	7.09E-07

D



Unpublished results

Upregulated nodes KEGG

Pathway	Total	Expected	Hits	P.Value	FDR
Epstein-Barr virus infection	114	0.458	4	0.000973	0.207
Cell cycle	127	0.51	3	0.0136	0.75
Phagosome	49	0.197	2	0.0161	0.75
Notch signaling pathway	49	0.197	2	0.0161	0.75
Wnt signaling pathway	147	0.59	3	0.02	0.75
Tuberculosis	170	0.682	3	0.0293	0.75
Salmonella infection	68	0.273	2	0.0299	0.75
Long-term potentiation	69	0.277	2	0.0307	0.75
Phenylalanine, tyrosine and tryptophan biosynthesis	8	0.0321	1	0.0317	0.75
Fc epsilon RI signaling pathway	75	0.301	2	0.0358	0.762
ErbB signaling pathway	87	0.349	2	0.0469	0.829
Malaria	12	0.0482	1	0.0472	0.829

Does EBV cause CDH?????

JOURNAL OF VIROLOGY, Oct. 2010, p. 10329–10343
0022-538X/10/\$12.00 doi:10.1128/JVI.00923-10
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Vol. 84, No. 19

Cellular MicroRNAs 200b and 429 Regulate the Epstein-Barr Virus Switch between Latency and Lytic Replication[∇]

Amy L. Ellis-Connell,[†] Tawin Iempridee, Iris Xu, and Janet E. Mertz^{*}

Tumor and Stem Cell Biology

Cancer
Research

Downregulation of MicroRNA-200 in EBV-Associated Gastric Carcinoma

Aya Shinozaki¹, Takashi Sakatani¹, Tetsuo Ushiku¹, Rumi Hino¹, Maya Isogai¹, Shunpei Ishikawa¹, Hiroshi Uozaki¹, Kenzo Takada², and Masashi Fukayama¹

JOURNAL OF VIROLOGY, Aug. 2010, p. 7892–7897
0022-538X/10/\$12.00 doi:10.1128/JVI.00379-10
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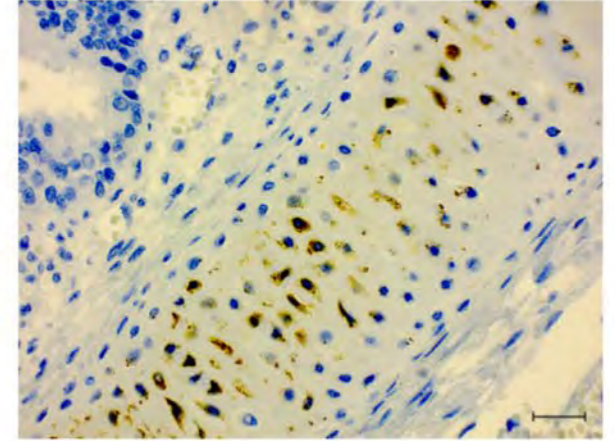
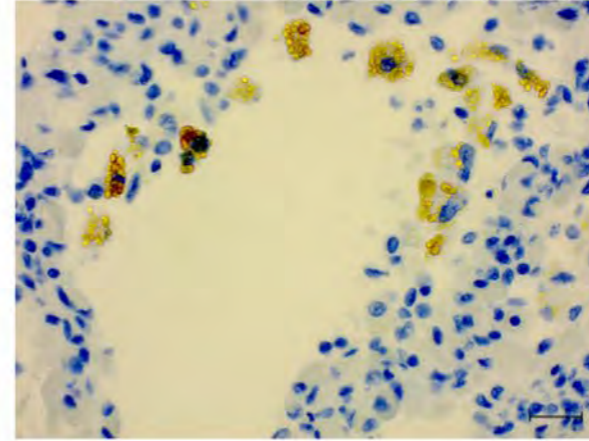
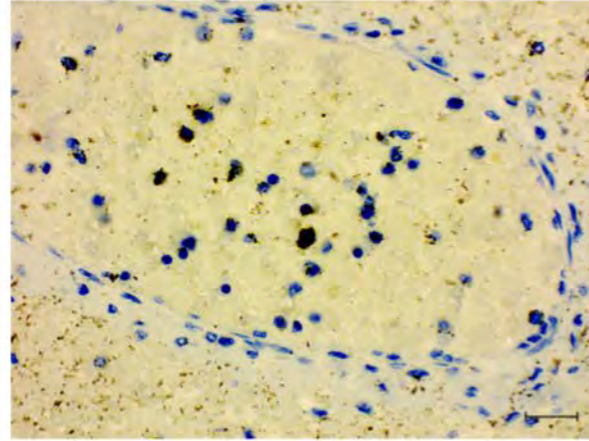
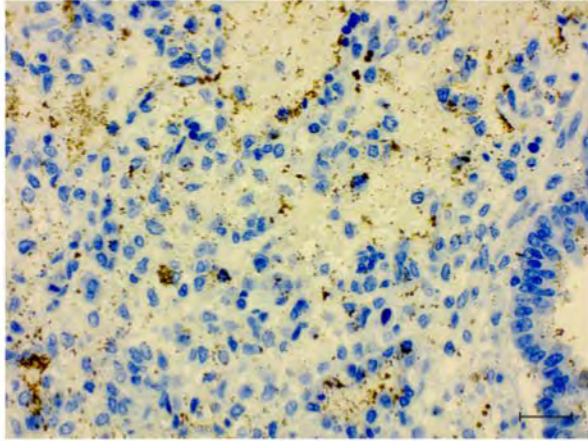
Vol. 84, No. 15

Differential Expression of the miR-200 Family MicroRNAs in Epithelial and B Cells and Regulation of Epstein-Barr Virus Reactivation by the miR-200 Family Member miR-429[∇]

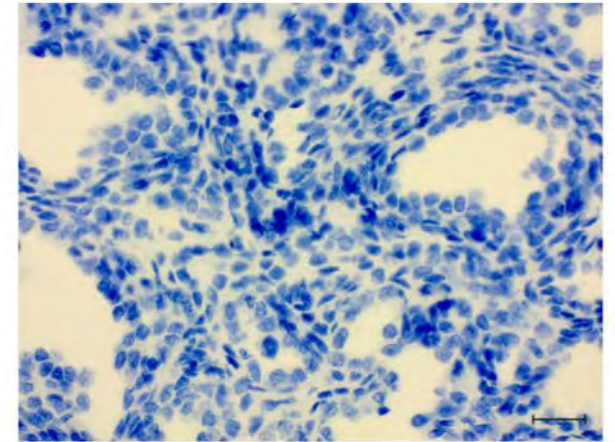
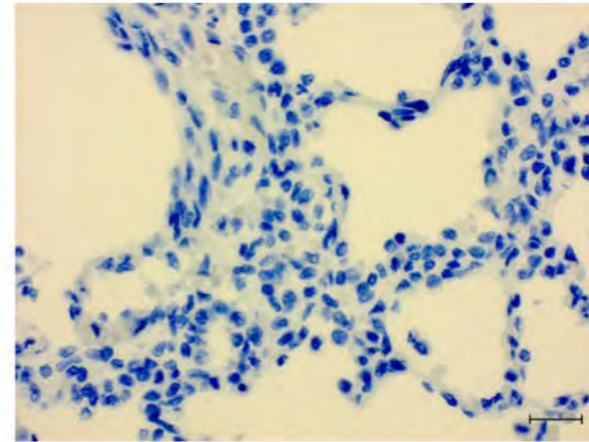
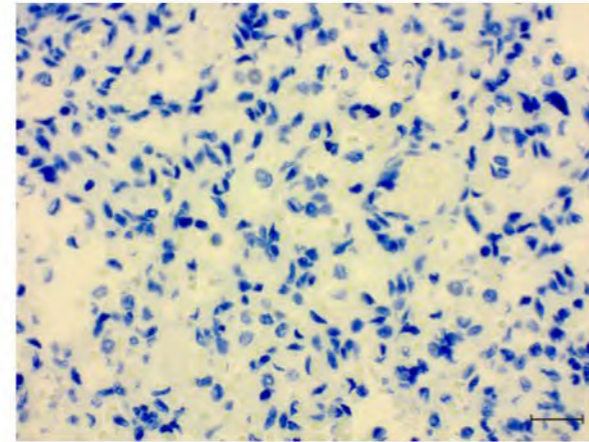
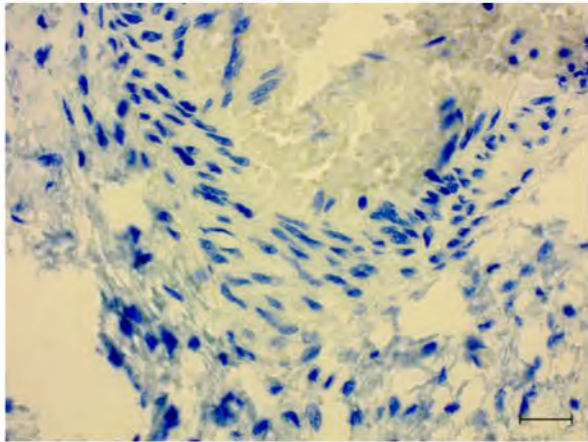
Zhen Lin, Xia Wang, Claire Fewell, Jennifer Cameron, Qinyan Yin, and Erik K. Flemington^{*}

Does EBV cause CDH?????

CDH



Control



WiSDOM: WINNIPEG'S SURGICAL DATABASE & OUTCOMES MANAGEMENT

DEVOTION Legacy Project

Dr. Richard Kuzner, Suyeon Lum Min, Melonie Morris, Anni Shawyer, Chelsea Ruth, Marni Brownell, Francesca Ryan, and Robert Bralshaw



ABOUT WiSDOM:

WiSDOM is all about children with congenital anomalies. A congenital anomaly is a birth defect. Some babies are born with a congenital anomaly that requires surgery immediately after birth. Until a few decades ago, most of these babies died. Recent improved surgical and intensive care techniques have resulted in better survival. Currently, we do not know how these babies do later in life when they grow up. To find out, we created a database of almost 800 surgical congenital anomaly patients, recording birth and surgery details. We plan to link our surgical database to population data managed by the Manitoba Centre for Health Policy (MCHP). MCHP databases contain unique information about healthcare, education, and social service utilization for all Manitobans. This linkage will allow us to answer the following questions:

1. **Who** is at greatest risk of giving birth to a baby with a congenital anomaly requiring surgery?
2. **How** do children with surgical congenital anomalies do at school and in life compared to children without a defect?
3. **Why** do some children with surgical congenital anomalies do better than other children with surgical congenital anomalies?

The answers to these questions will help us understand the causes and long-term outcomes of surgical birth defects and improve the care of babies born with surgical congenital anomalies.



OBJECTIVES:

1. WHO is at risk: Identify maternal factors associated with surgical congenital anomalies: Maternal demographic information will be identified using MCHP databases to determine the populations with highest risk of having a baby with surgical congenital anomalies.

2. HOW are children doing: Compare the long-term outcomes of patients with surgical congenital anomalies to those of age-matched controls: Virtual long-term follow-up of each patient in WiSDOM using MCHP linkages to determine their medical, educational, psychosocial, socioeconomic outcomes and comparing them with an age-matched control from MCHP.

3. WHY do some children do better: Determine patient and maternal factors that affect long-term patient outcomes: After defining the maternal risk factors and long-term outcomes for the WiSDOM patient-cohort, subgroups who have the highest risk of poor outcomes will be identified, as well as the demographic, medical, educational, psychosocial and socioeconomic determinants of favourable long-term outcomes.



HYPOTHESIS:

Demographic, medical, educational, psychosocial and socioeconomic differences exist between the mothers of, and children with surgical congenital anomalies and the general population of Manitoba.



TEAM:

The WiSDOM team includes expertise in:

- pediatric surgery
- neonatology
- nursing
- allied health care
- pediatric medicine
- child health outcomes
- epidemiology
- biostatistics
- data analysis



WiSDOM CLINIC:

WiSDOM has established a multidisciplinary long-term follow-up clinic at the Children's Hospital for patients with surgical congenital anomalies. Children attending the clinic receive evidence-based follow-up for early identification and intervention for patients and families. Other advantages of the WiSDOM clinic include improved and coordinated care, ongoing education, increased quality of life, and ease of access to health care professionals.



DEVOTION:

1. Provides support for prospective data collection and entry for babies born with surgical congenital anomalies after 2016
2. Facilitates consent and enrollment of participants into the WiSDOM study and collection of research data in the WiSDOM Clinic
3. Helps to distribute information to high-risk populations, healthcare providers and stakeholders using research findings from the WiSDOM database.
4. Aids in developing policies to direct risk mitigation strategies
5. Assists in developing a plan to establish a multidisciplinary, long-term follow-up clinic for WiSDOM children born in Manitoba within the next 5 years
6. DEVOTION provided initial funding for the retrospective data collection for all babies born in Manitoba from 1991 - 2016 in addition to the funding for prospective data collection and linkages to MCHP.



SIGNIFICANCE:

Surviving congenital abnormalities has drastically improved, therefore, we need to refocus our attention on optimizing long-term medical, educational, psychosocial and socioeconomic outcomes. By linking our surgical congenital anomalies patient cohort with the comprehensive information in the MCHP databases we will be able to begin this optimization. Linking databases will provide the opportunity for virtual long-term follow-up for the first time in this patient population. This virtual follow-up will identify the risk factors and outcomes of surgical congenital anomalies. This information will guide the development of preventative strategies to provide better care to babies born with surgical congenital anomalies.

Does EBV cause CDH?????_WiSDOM

Infectious Mononucleosis (ICD-9:075)

Odds ratio = 0.49 for mothers with CDH baby compared to controls from the general population

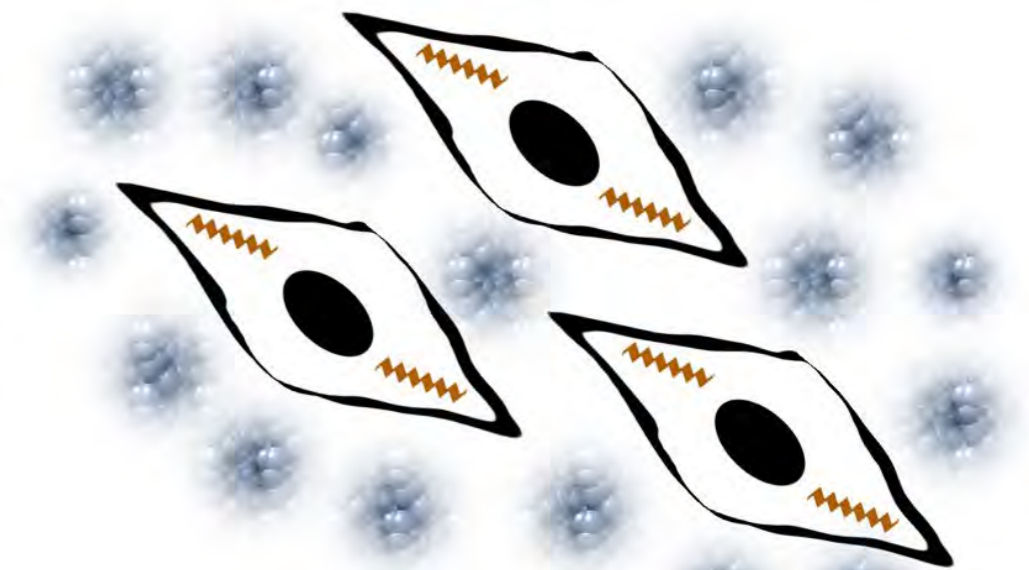
95% CI = 0.12-1.36

P-value = 0.2371

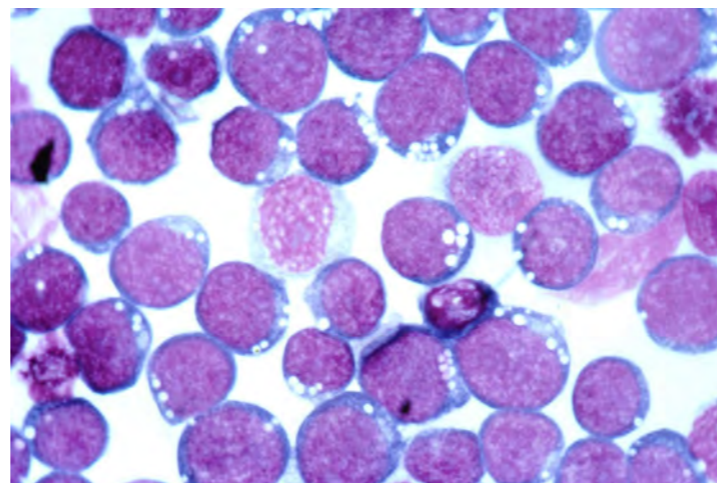
Epithelial-to-Mesenchymal Transition, CDH and EBV



Epithelial cells



Mesenchymal cells



?

Prenatal treatment

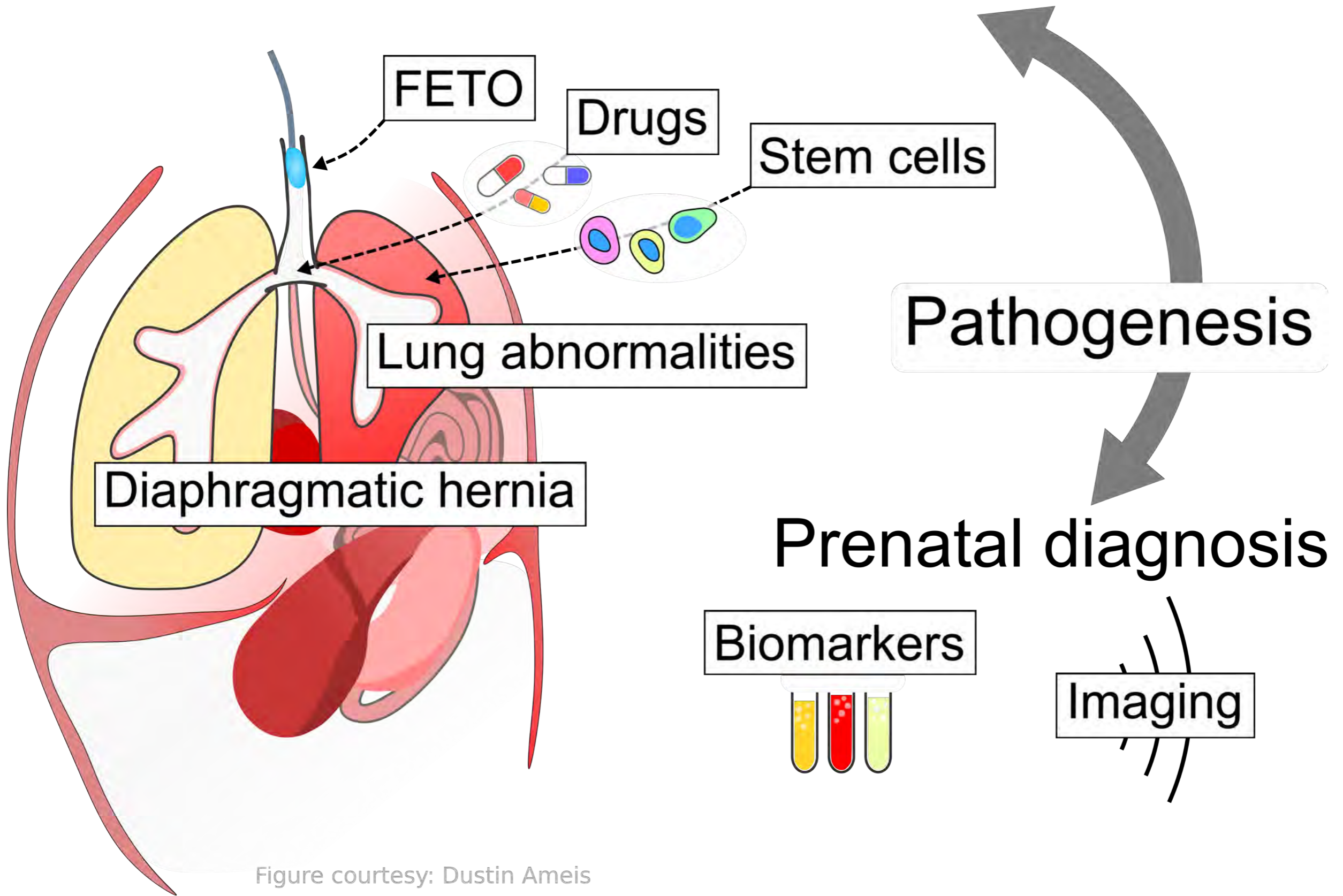
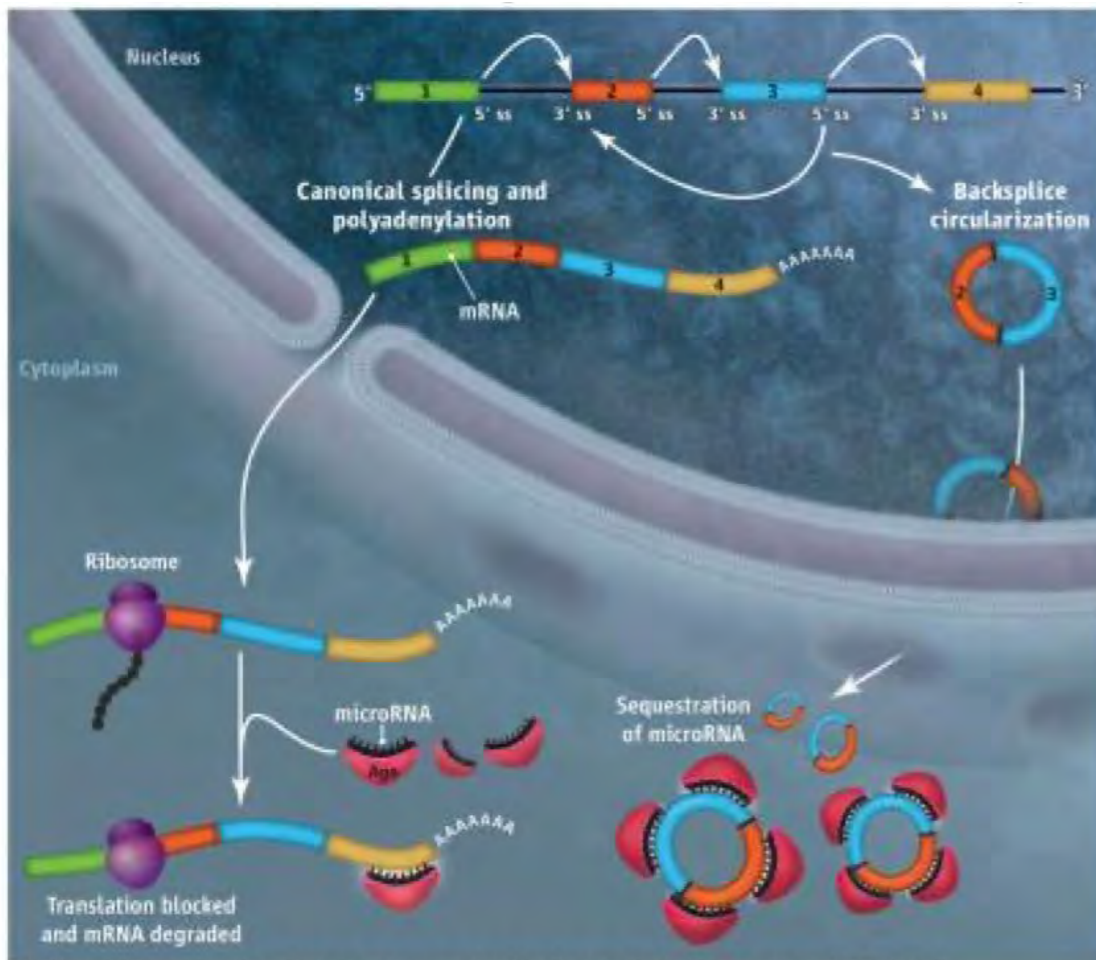


Figure courtesy: Dustin Ameis

CircularRNA

- More stable, more abundant and specific than linear RNAs.
- Regulate gene expression at transcriptional and post-transcriptional level by serving as **microRNA sponges** and interacting with mRNA and proteins.



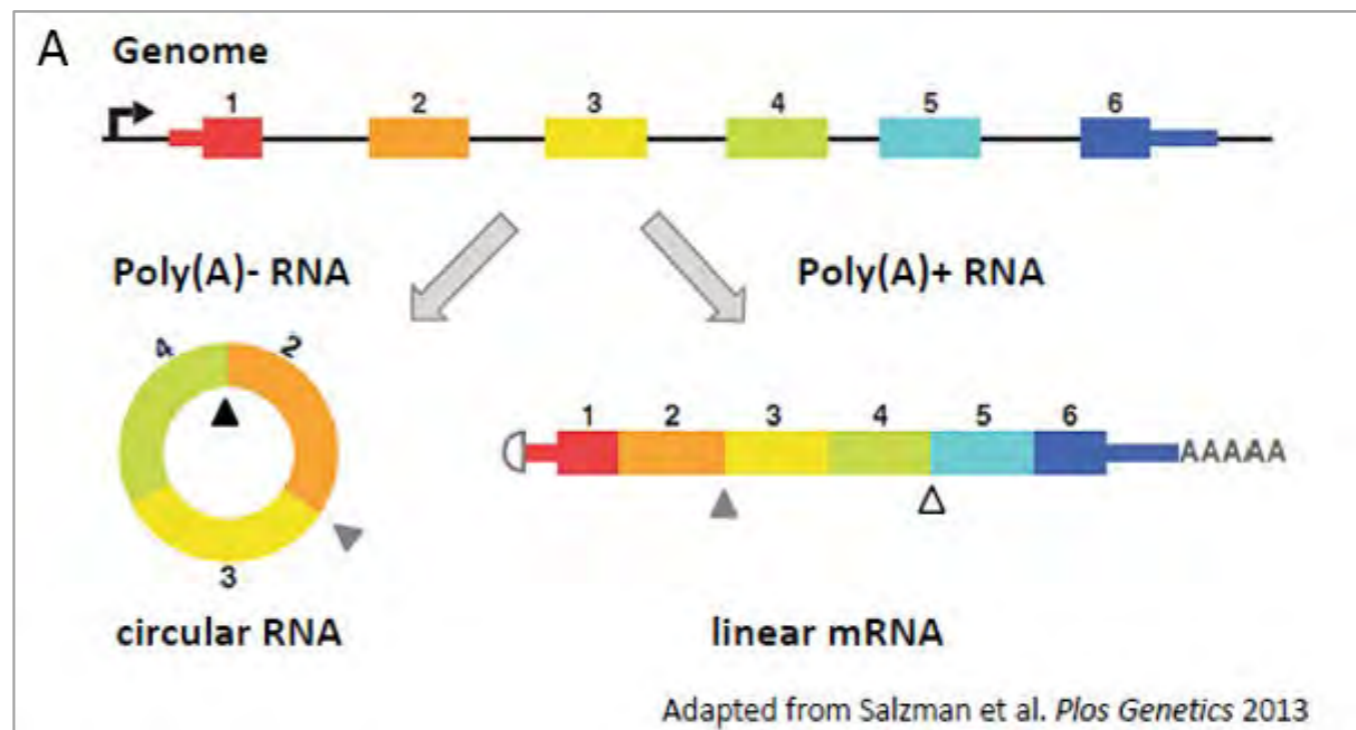
<http://www.med.upenn.edu/wiluszlab/research.html>



<http://www.epibeat.com/developmental-biology-stem-cells/circular-rnas-as-molecular-sponges-for-mirnas/518/>

Circular RNAs

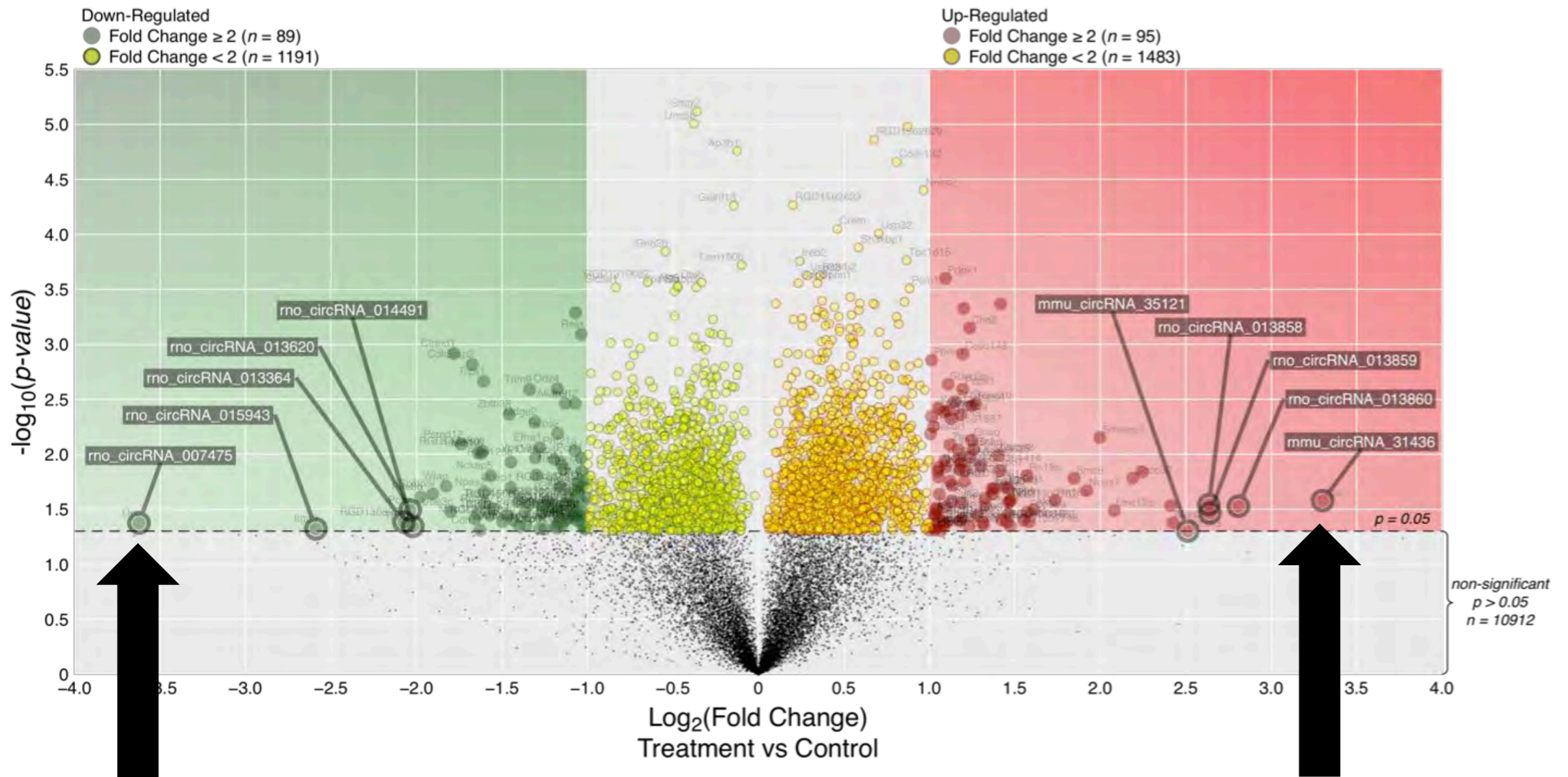
- Upstream regulators of miRNAs (epigenetic regulation)
- “Head-to-tail” splicing



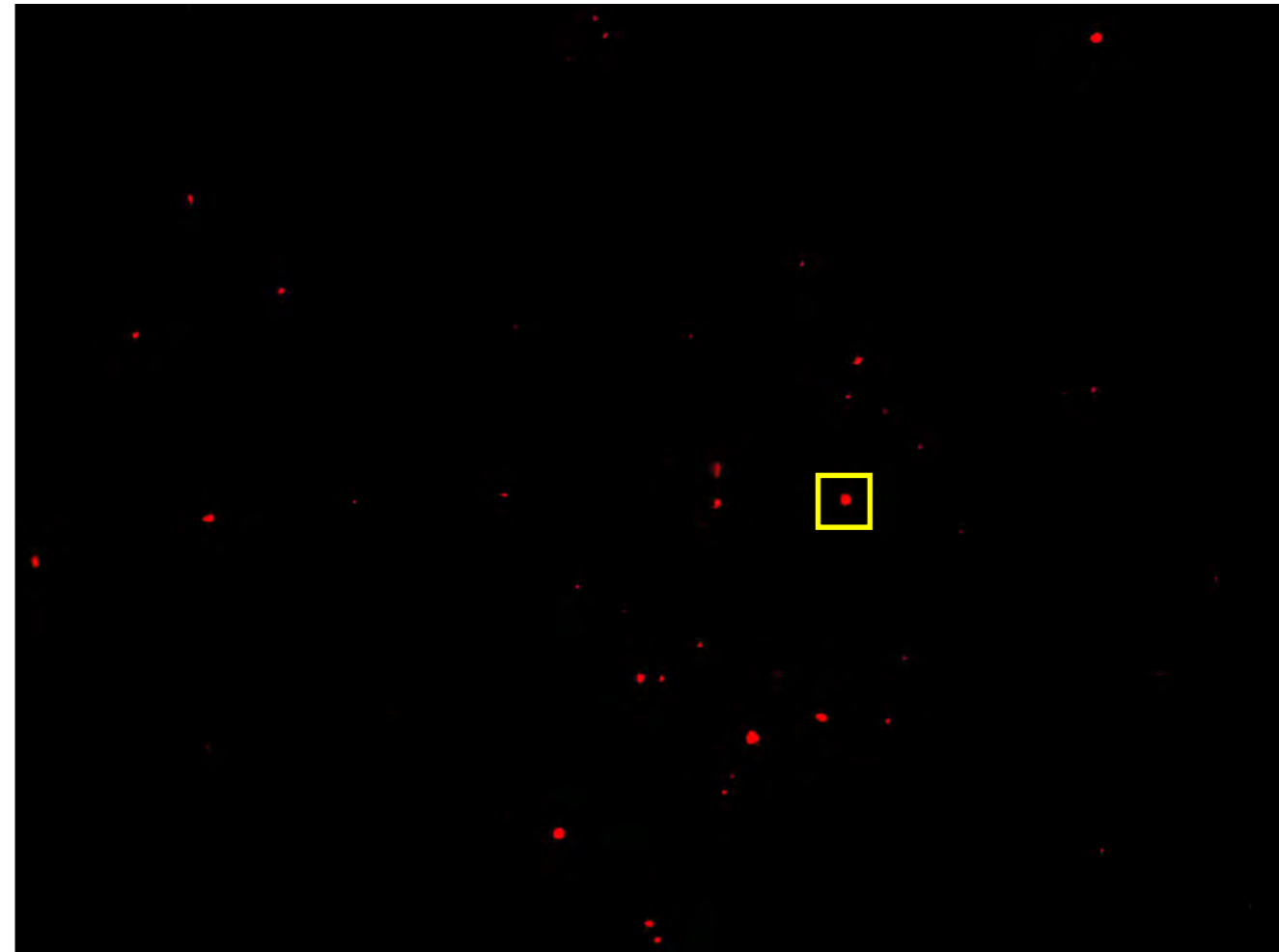
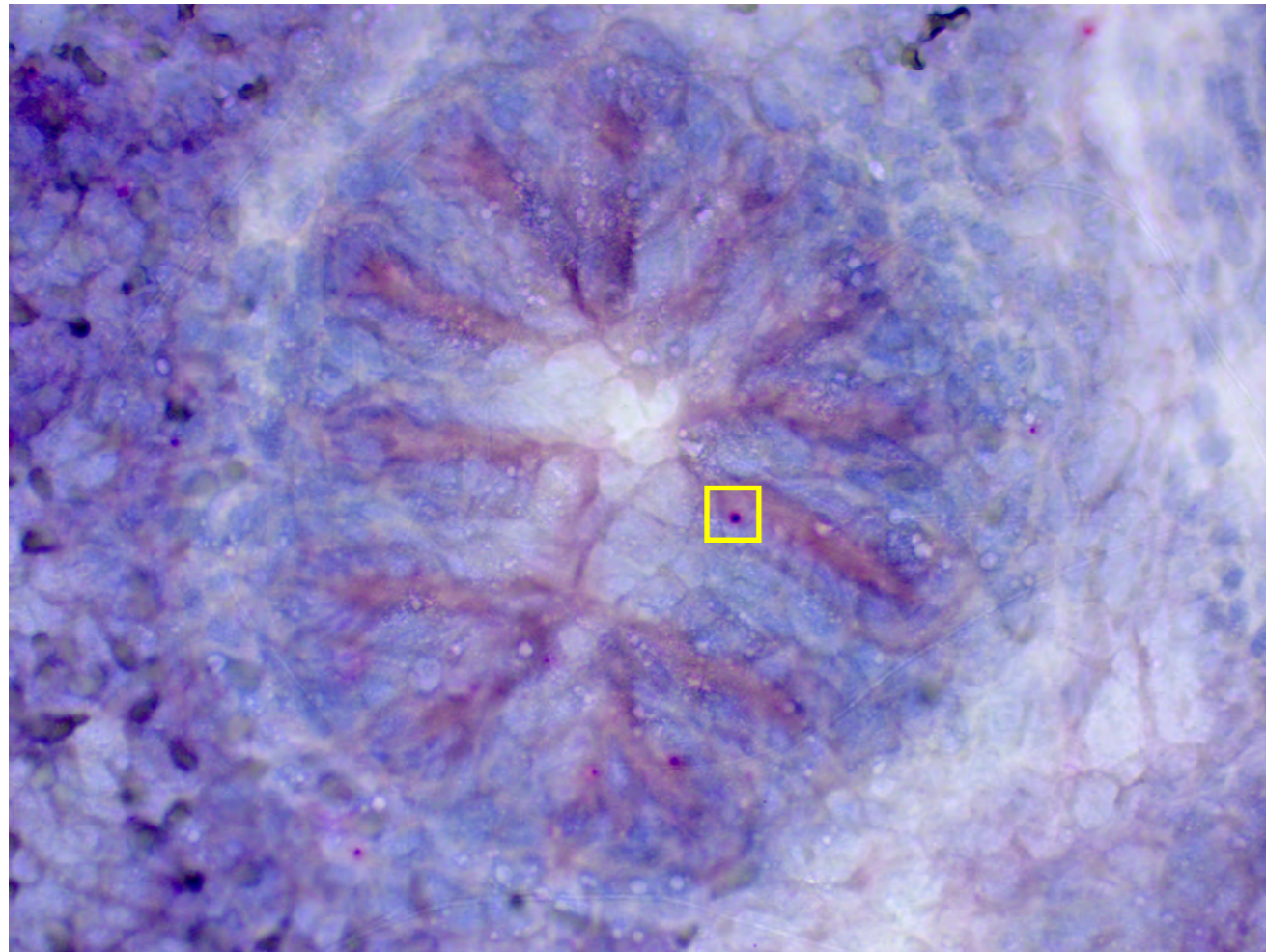
Ideal Biomarkers:

- Stable
- Conserved
- Tissue- and development stage-specific expression
- High abundance in extracellular compartments

Circular RNA profile is dysregulated in E21 nitrofen-induced hypoplastic lungs



BaseScope™ E21 Rat Lung



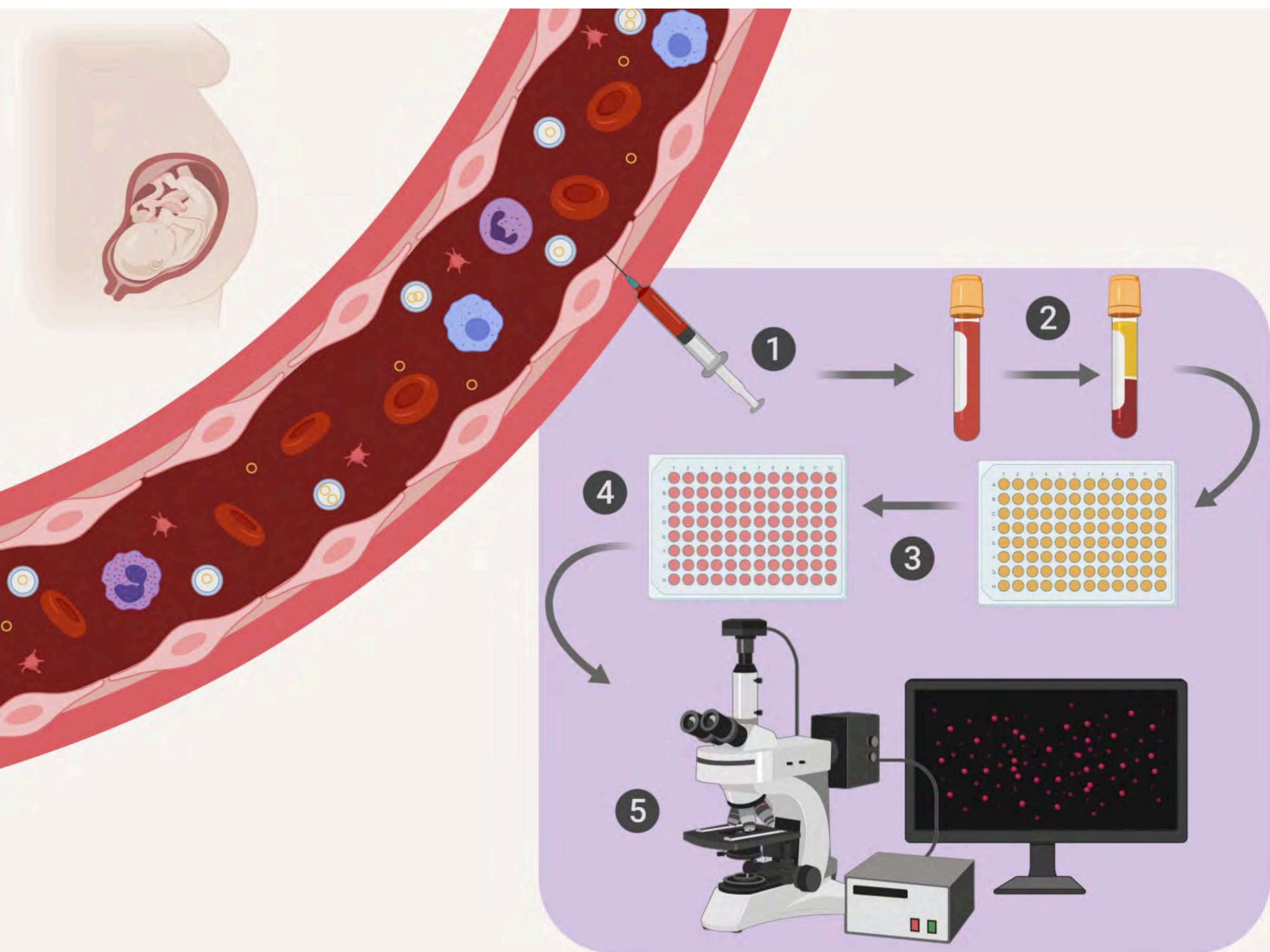
(rno_circRNA_007475)

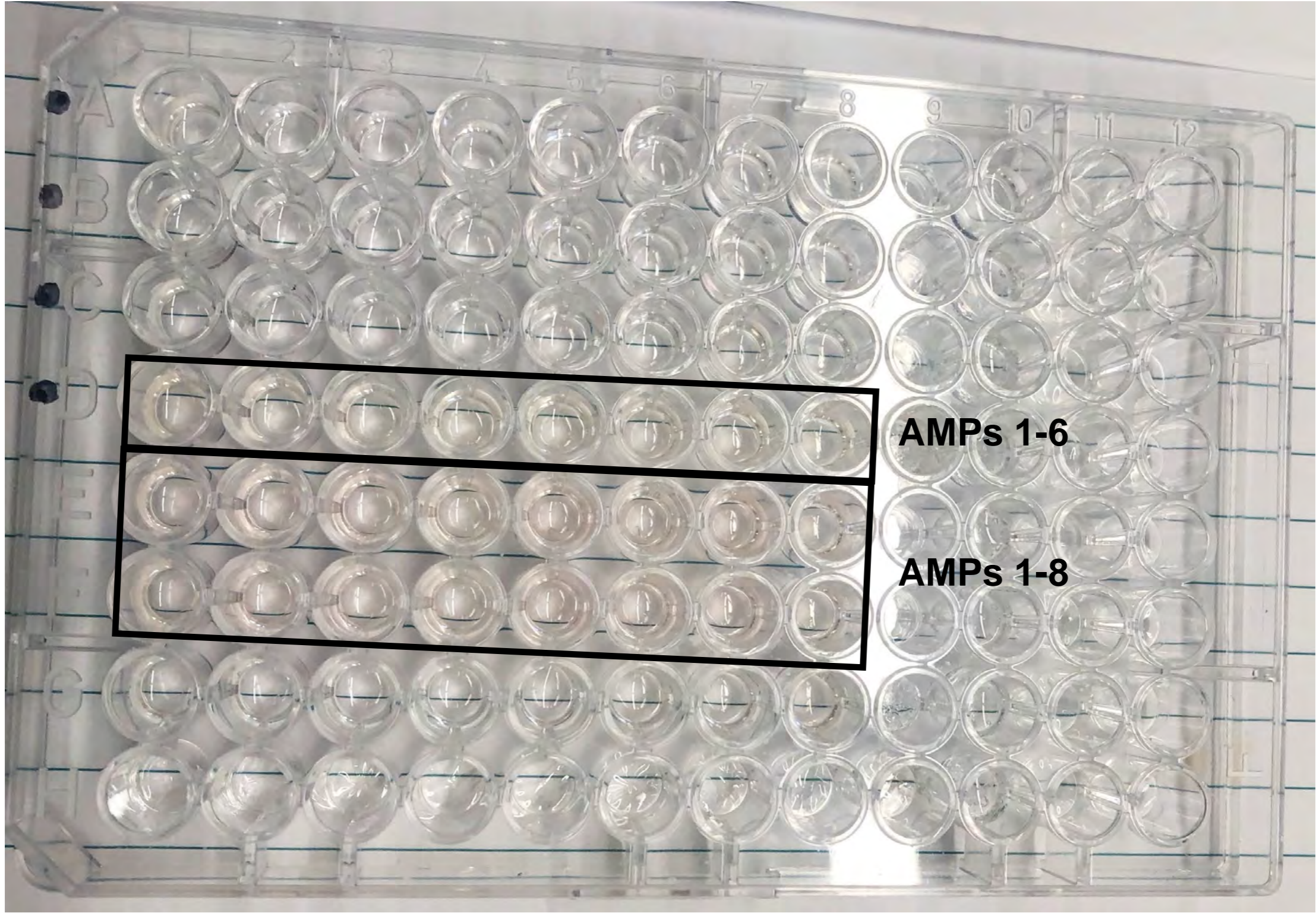
Pediatric Surgery International
<https://doi.org/10.1007/s00383-019-04558-2>

ORIGINAL ARTICLE

First steps in the development of a liquid biopsy in situ hybridization protocol to determine circular RNA biomarkers in rat biofluids



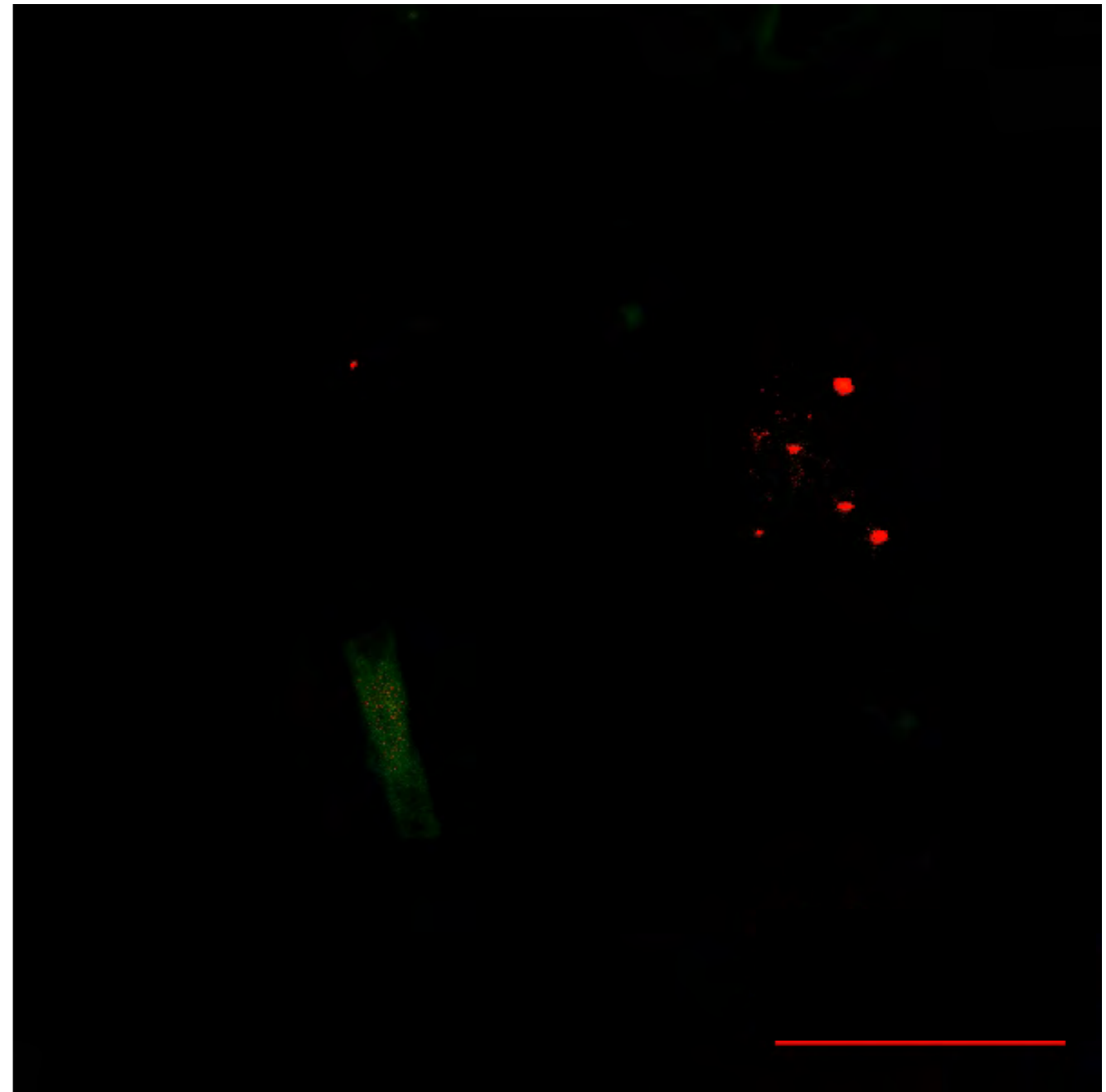
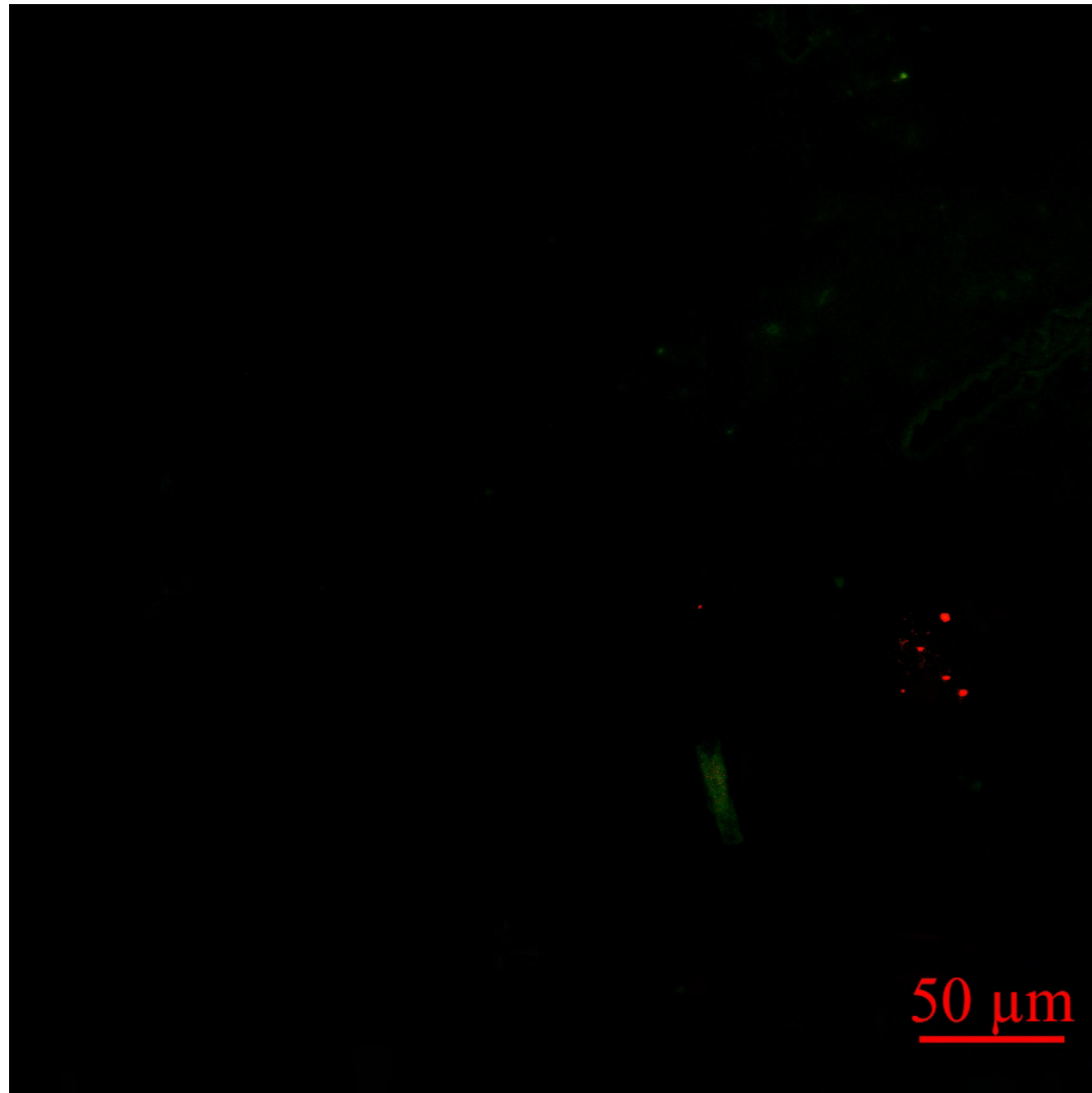




AMPs 1-6

AMPs 1-8

Control Adult Rat Serum: rno_circRNA_007475



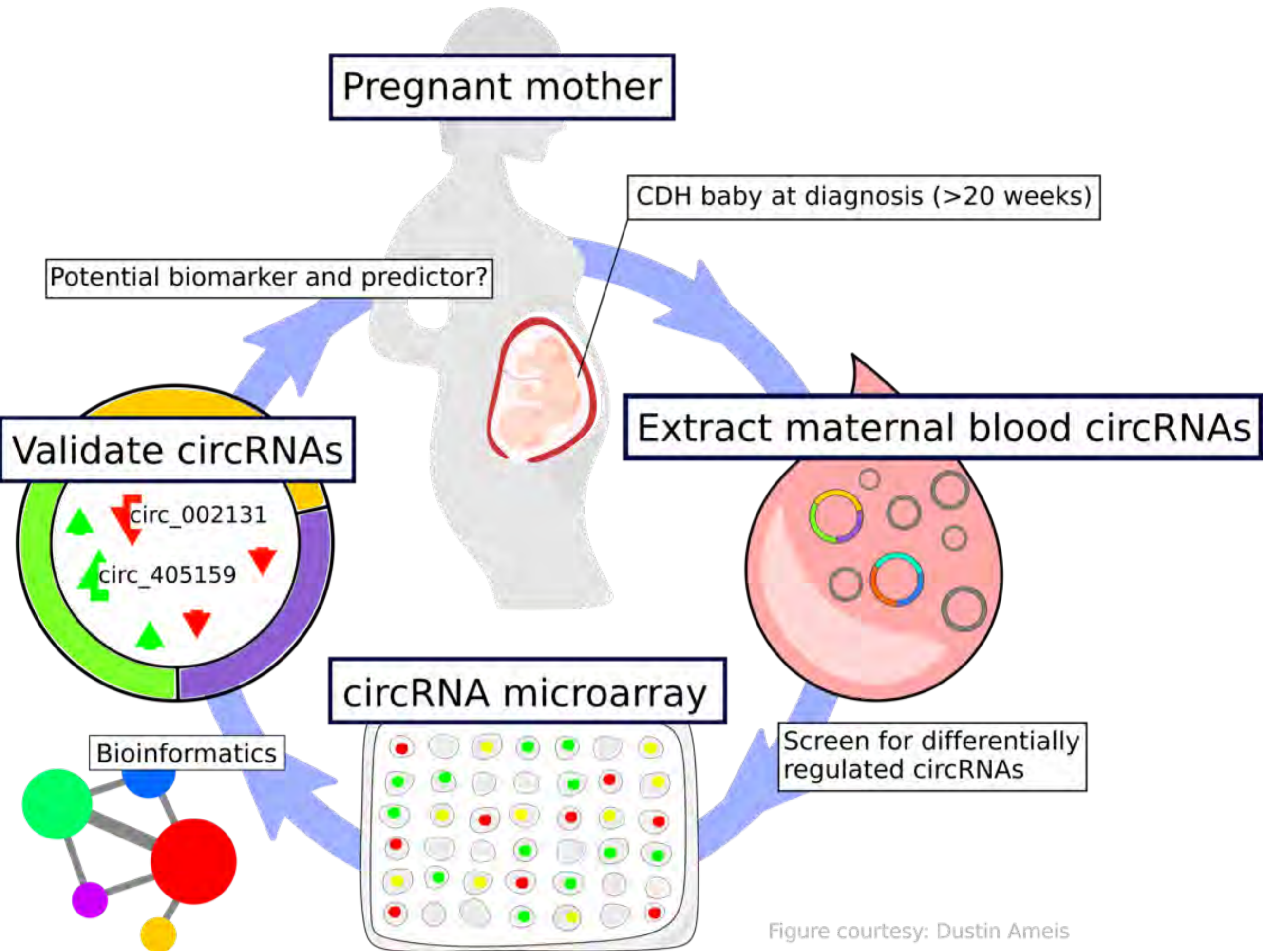
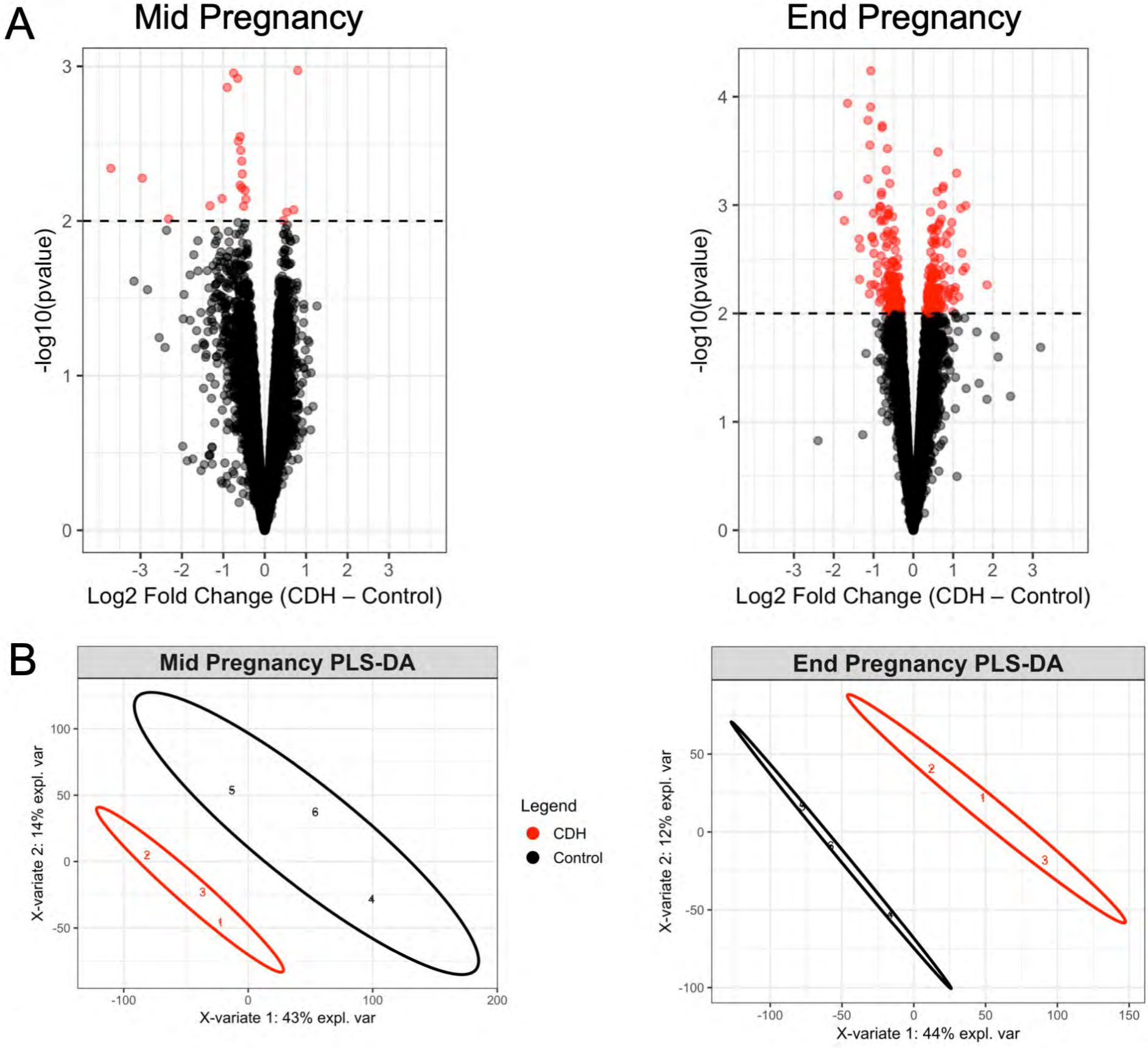
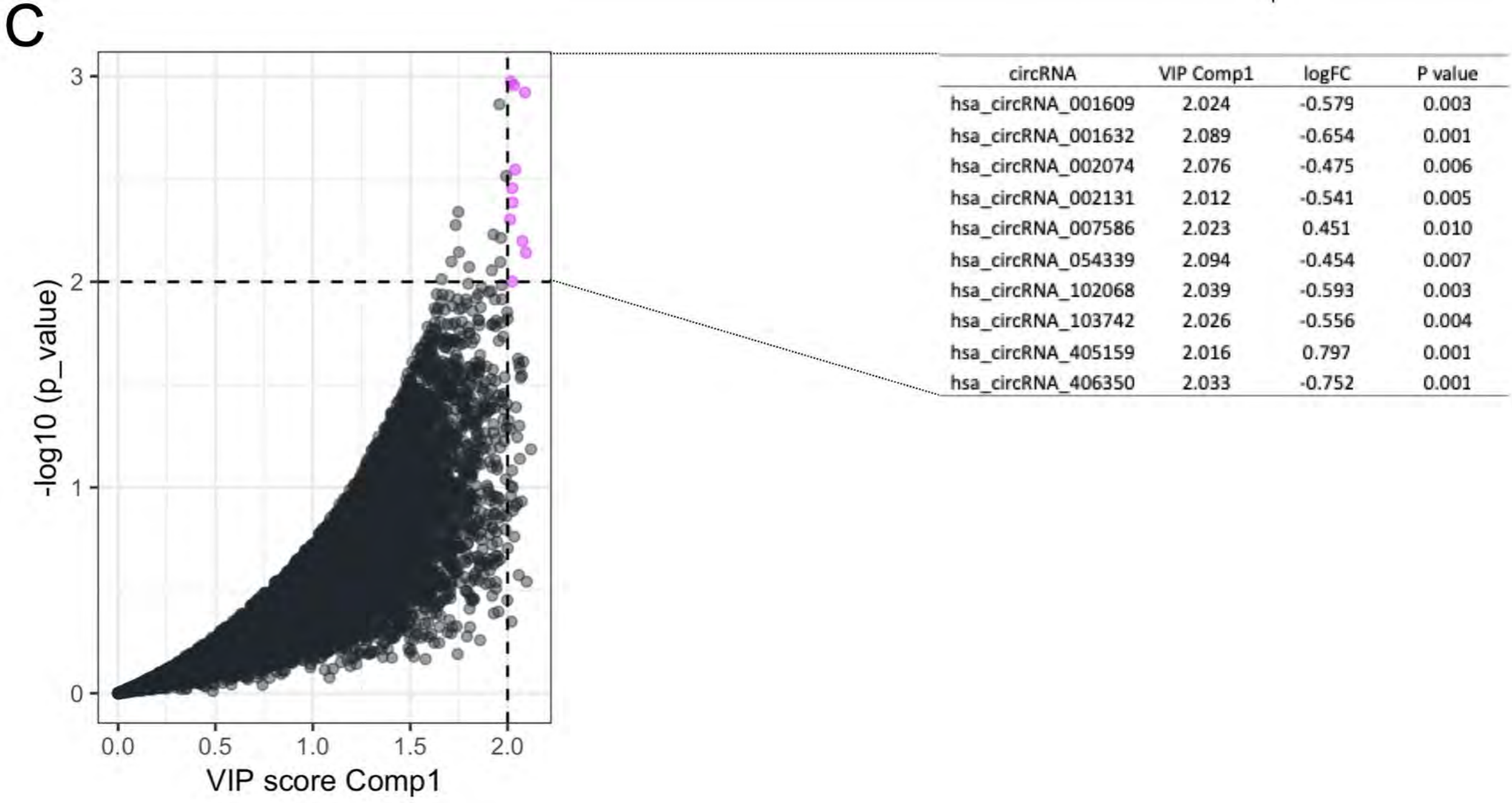


Figure courtesy: Dustin Ameis

CircRNA profile distinguishes CDH lung from control



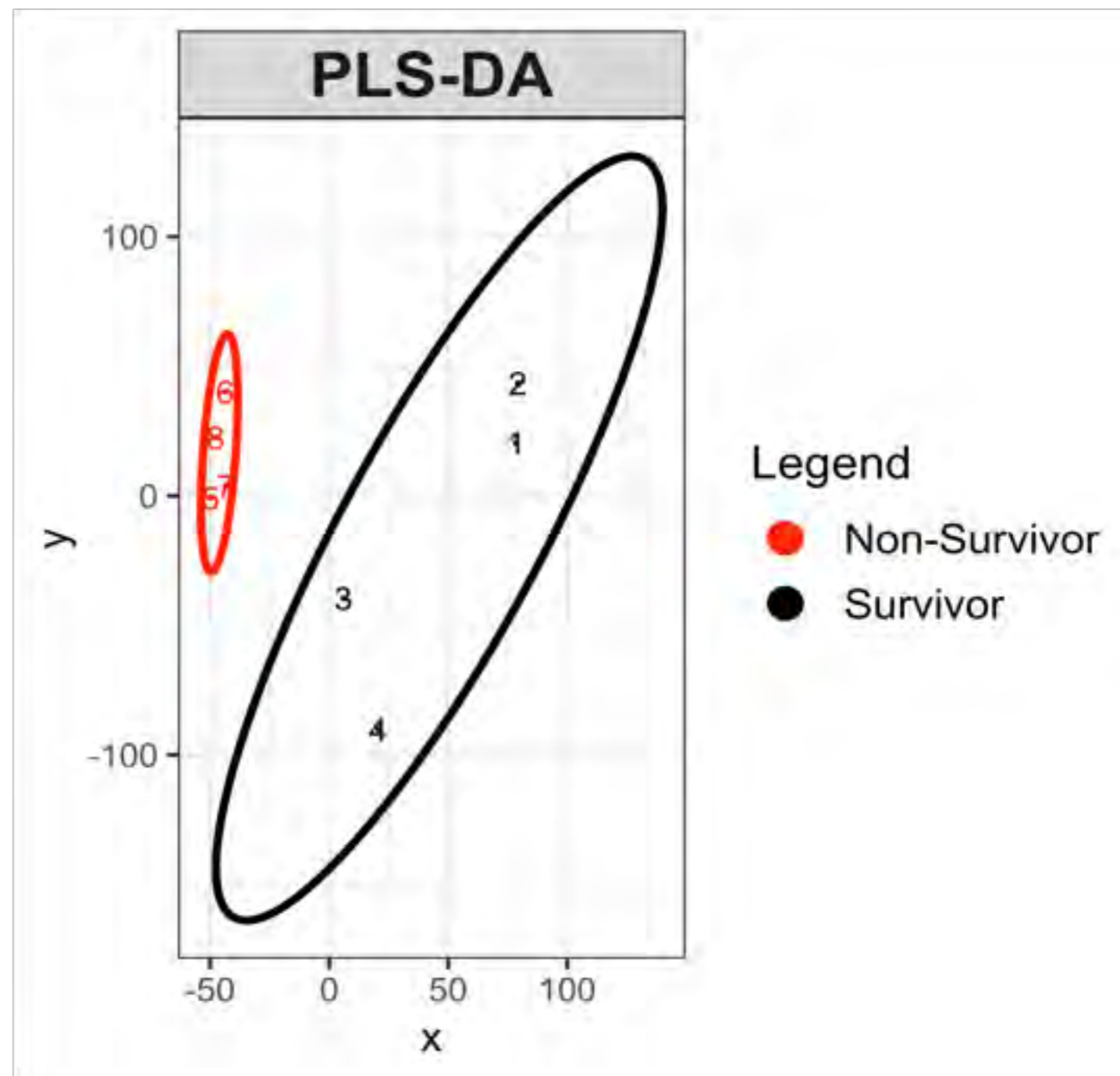
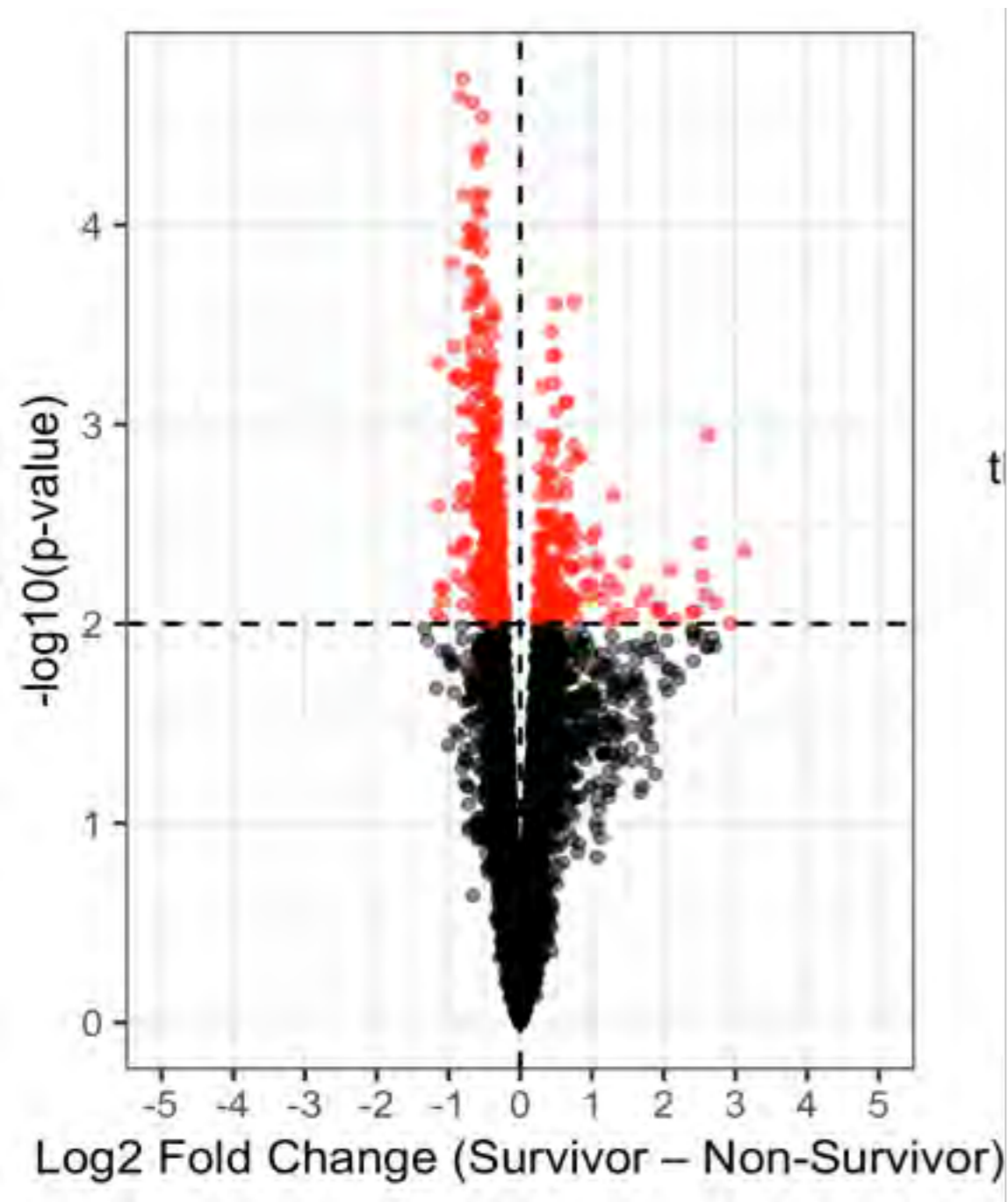
CircRNA profile distinguishes CDH lung from control



CircRNA profile can distinguish FETO survivors from non-survivors

	Survivors	Non-Survivors	P-value
Gestational Age at Plug (weeks)	28.1 (28.7 - 27.1)	27.9 (29.1 - 27.0)	0.84
Observed/ Expected Lung to Head Ratio (%)	22.5 (23.6 - 17.5)	21.6 (24.0 - 15.9)	0.87
Liver herniated	10 (91%)	9 (100%)	1.00
Fetal gender	7 male/ 4 female	4 male/ 5 female	0.65
Birth weight (g)	2780 (3180 - 2160)	3195 (3278 - 2650)	0.3

CircRNA profile can distinguish FETO survivors from non-survivors



Prenatal treatment

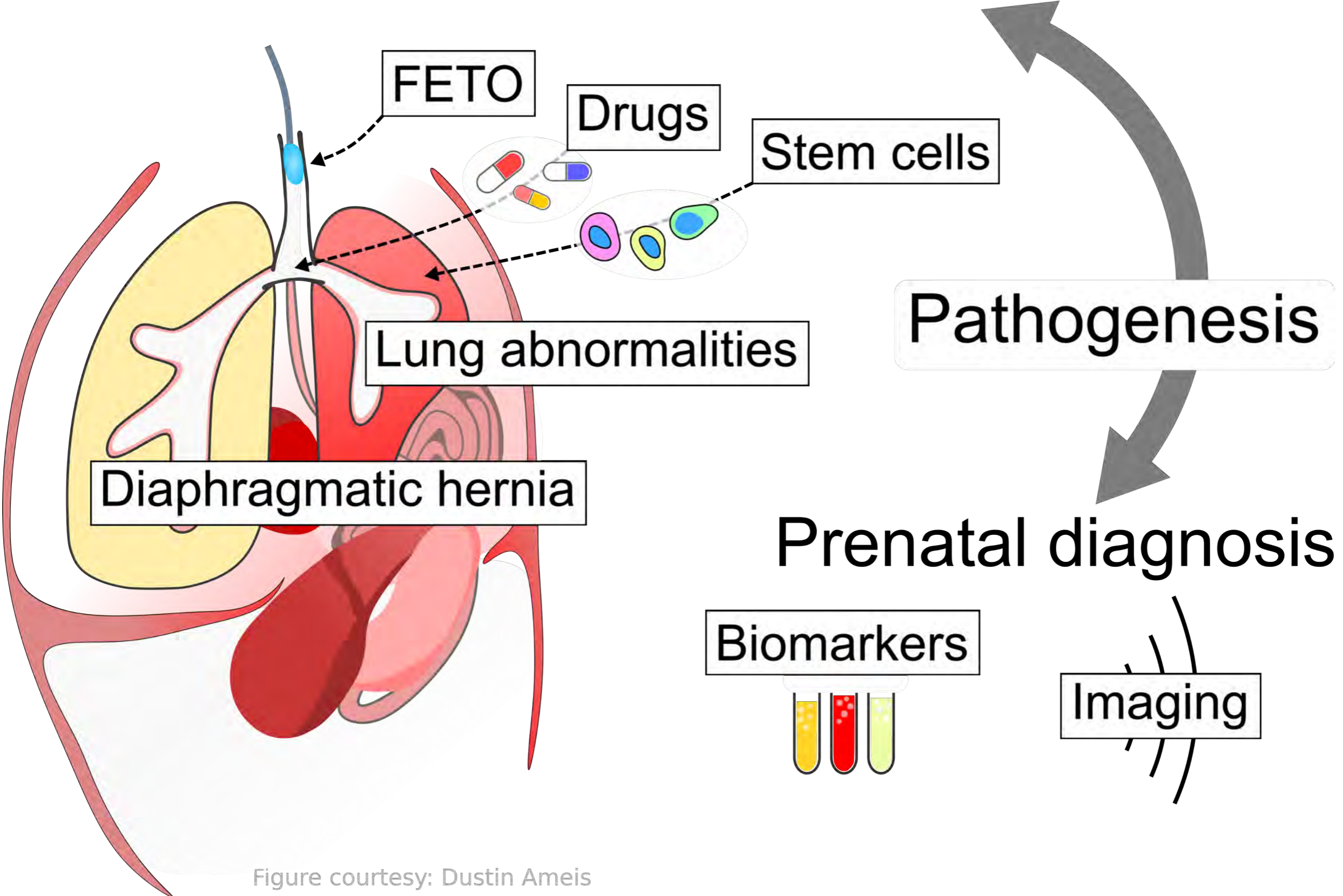


Figure courtesy: Dustin Ameis



<http://store.winnipegfreepress.com/photostore/details/139975/>

Thank you MIRACLA Lab



Acknowledgements

Laboratory members

Nolan Deleon

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Daywin Patel

Pediatric Surgery

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Suyin Lum Min

BJ Hancock

Nathan Wiseman

Cindy Holland

Previous Laboratory members

Dustin Ameis

Lojine Ayoub

Carly Fraser

Barbara Iwasiew

Shana Kahnamoui Zadeh

Ramin Kholdebarin

Naghme Khoshgoo

Eimear Kirby

Thomas Mahood

Samira Seif

Phillip Snarr

Robin Visser

Fuqin Zhu

UofM collaborators

Geoff Hicks's group

Andrew Halayko's group

Malcolm Xing's group

Neeloffer Mookherjee's group

Other Collaborators

Robbert Rottier

Dick Tibboel

Martin Post

Jan Deprest

Patrice Eastwood

Francesca Russo

Martin Lacher

Richard Wagner

Jorge Correia-Pinto

Patricia Pereira-Terra

Patients and families

GFT Group Academic Surgeons Winnipeg



Molly Towell
PERINATAL RESEARCH
FOUNDATION



Canada Foundation
for Innovation

Fondation canadienne
pour l'innovation

CCHCSP
PCCCSE



Research
Manitoba

EXIQON



**The Children's
Hospital
Research
Institute
of Manitoba**



CIHR IRSC
Canadian Institutes of Health Research
Instituts de recherche en santé du Canada

Thorlakson Chair
in Surgical Research



ADVENTURER TRAILBLAZER CHALLENGER DEFENDER VISIONARY INNOVATOR
TRAILBLAZER CHALLENGER DEFENDER VISIONARY INNOVATOR EXPLORER TRAILBLAZER CHALLENGER DEFENDER VISIONARY INNOVATOR EXPLORER

Word cloud featuring the phrase "thank you" in various languages and scripts, including: danke, 謝謝, ngiyabonga, tesekkür ederim, tapadh leat, gracias, thank you, obrigado, dziękuje, hvala, maururu, kōsōnōm, bedankt, nanni, nandri, kiitos, dankie, dhanyavad, bayarlalaa, gracie, faafetai lava, mersi, kia ora, barka, welalin, tack, spas, vinaka, blagodaram, dank je, misaotra, matondo, paldies, grazzi, mahalo, хвала, asante, manana, obrigada, tenki, chokrane, murakoze, mochchakkeram, djiere dieuf, tau, дякую, mamnun, go raibh maith agat, sulpáy, taiku, sukriya, kop khun krap, arigatō, takk, dakujem, trugarez, mèsì, didi madloba, sagolun, najis tuke, kam sah hamnida, rahmat, terima kasih, tanemirt, rahmet, dhanyavadagalu, shukriya, merce, мерси, তোমাকে ধন্যবাদ, 감사합니다, xiexie, ευχαριστώ, diolch, dyauf, and many others.

