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How the epitranscriptome is changing our world

Mary O'Connell

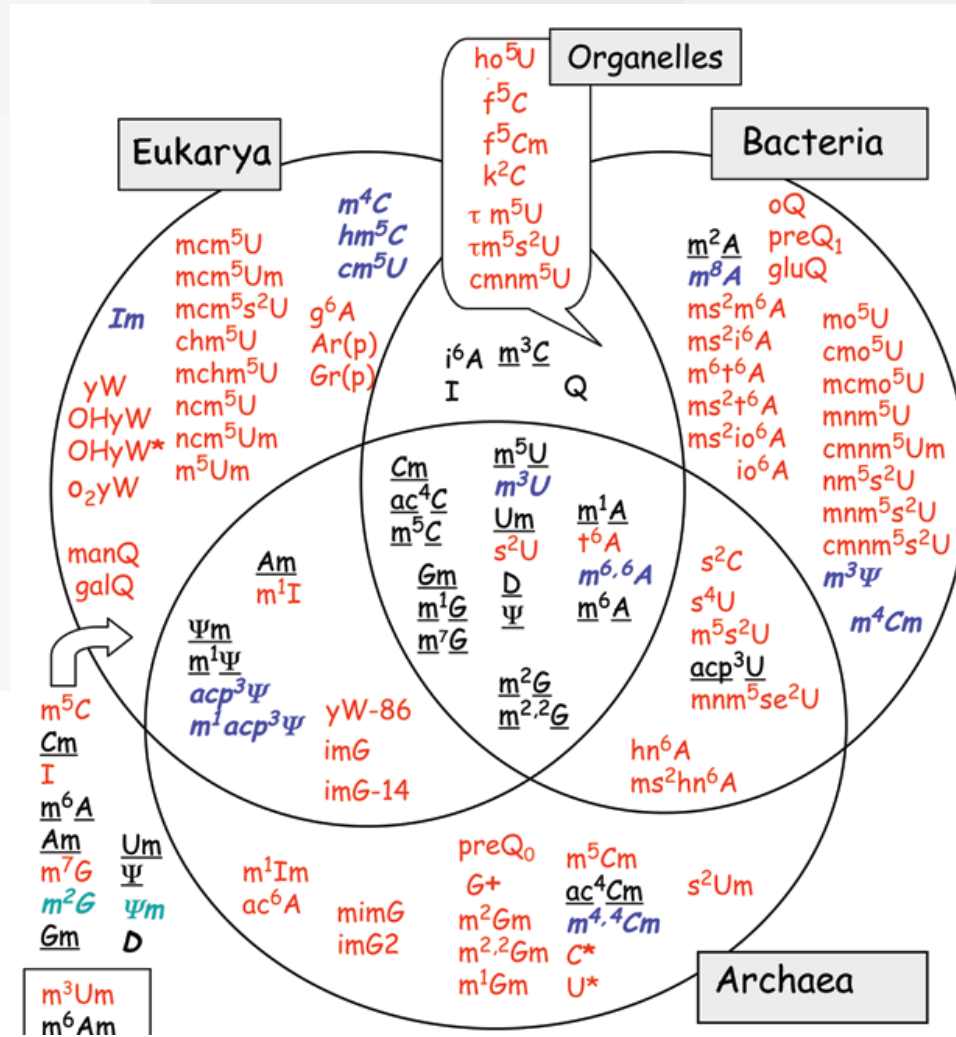


What is the epitranscriptome?

Guess!

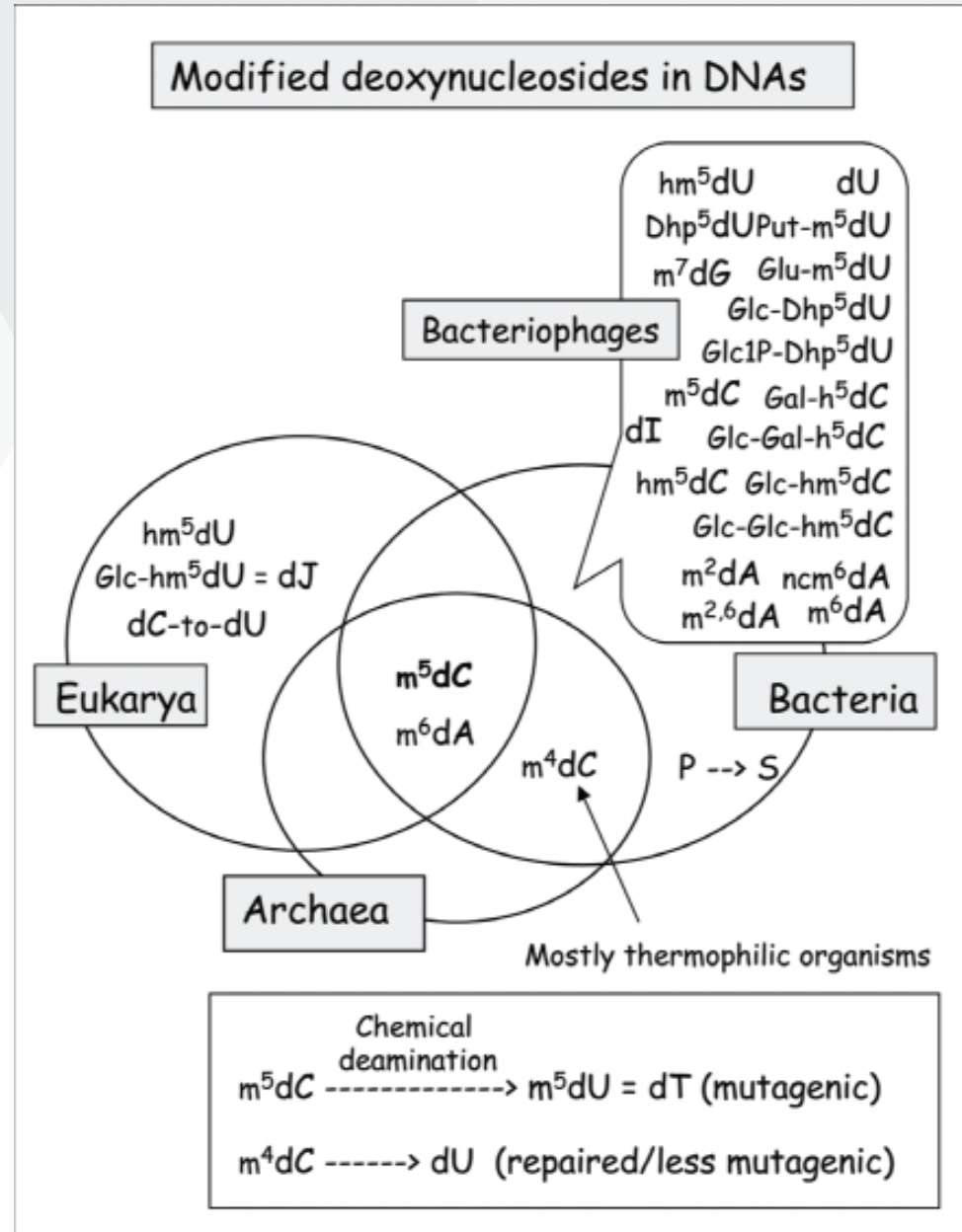
RNA modification

>170 RNA modifications



Henri Grosjean

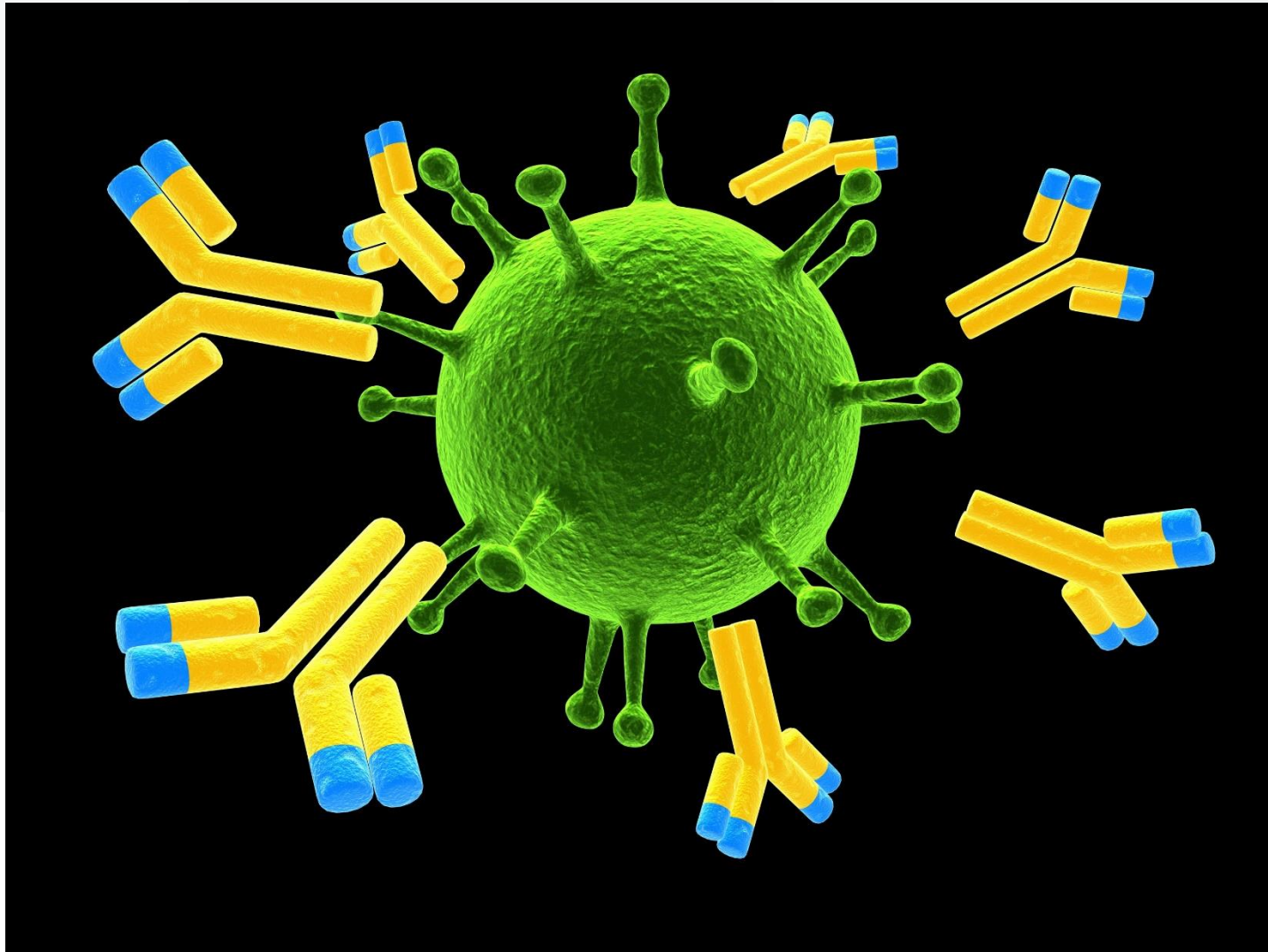
Modifications in DNA



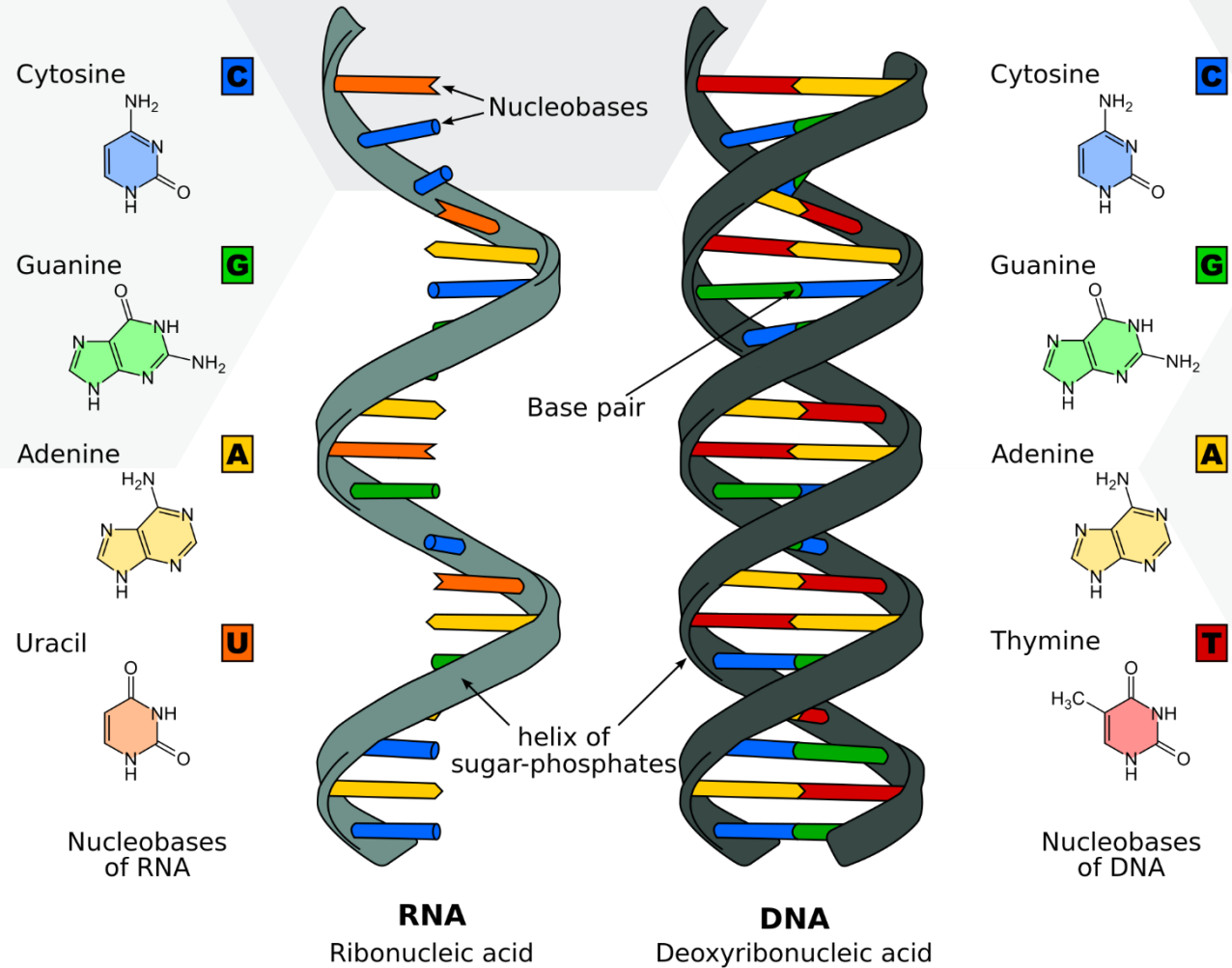
WHY?

Henri Grosjean

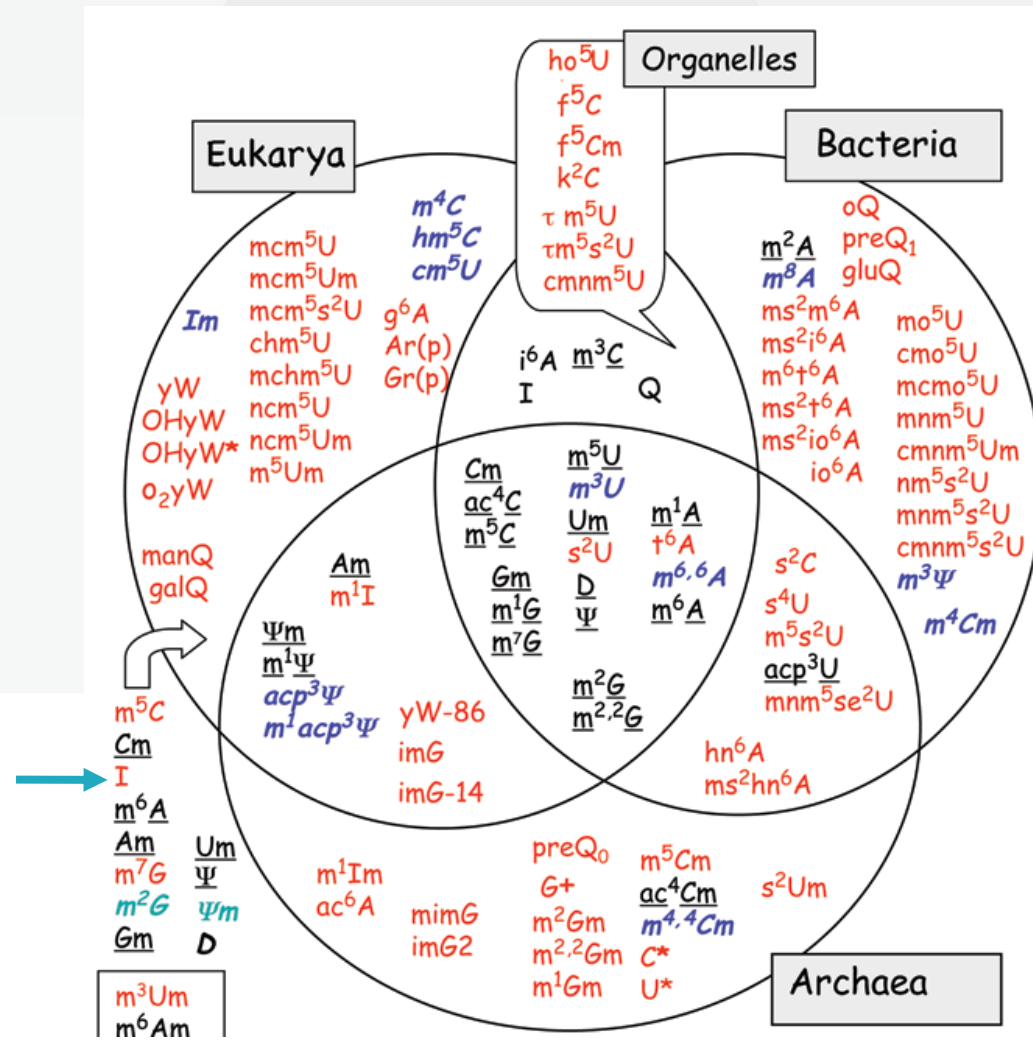
Antibody/Antigen interaction



Problem with nucleic acids



Approximately 150 RNA modifications



Henri Grosjean

RNA vaccines

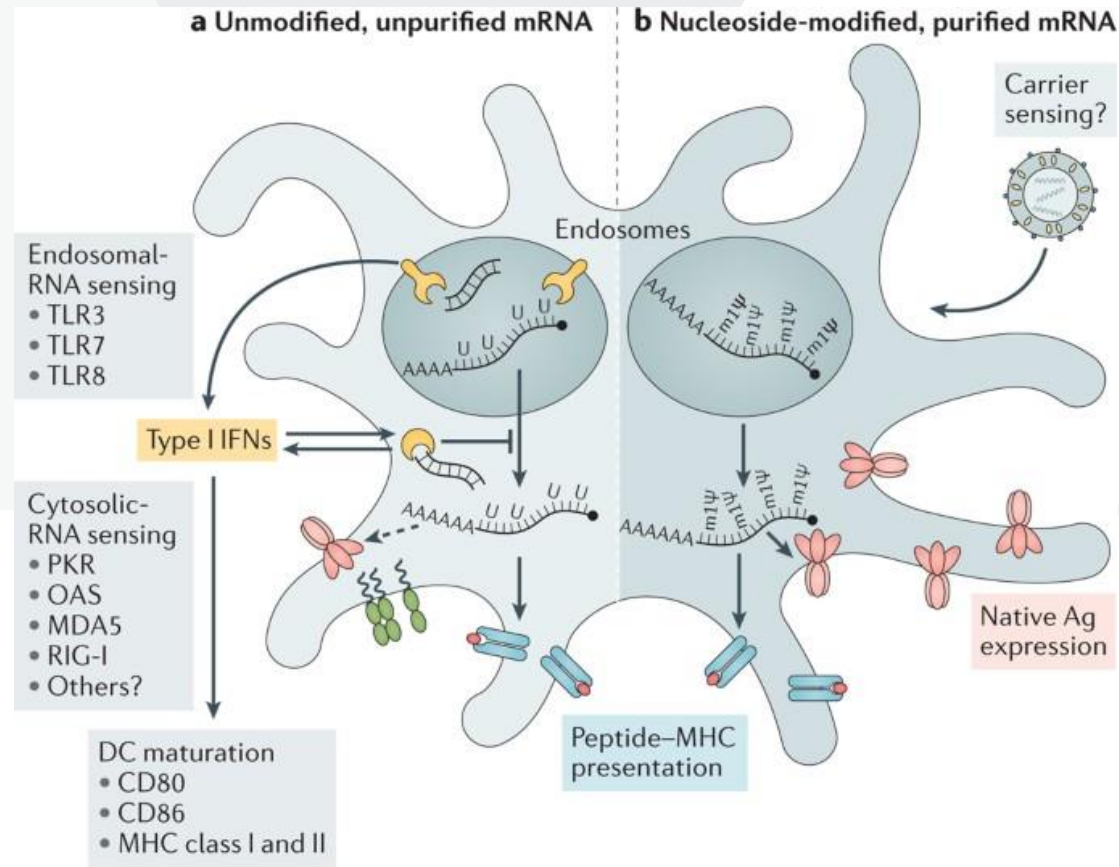


Katalin Kariko



Drew Weissman

Modified versus unmodified RNA



Fast vaccines

NEXT-GENERATION mRNA VACCINES



Computer-designed, a feature that makes them adaptable and rapidly scalable into millions of doses.



Rely on **genetic material called mRNA** to create an immune response against a specific antigen, or toxin—in this case, the coronavirus.



Production for the general population **can take just weeks** rather than the months required for conventional vaccines.

mRNA Vaccine

Components



mRNA (blueprint of protein)

Production



Faster because mRNA molecules are easier to produce

Process

Components are injected into the arm and serve as instructions for the body to make microbial protein

Traditional Vaccine

Components



Microbial protein or inactive microbe

Production



Slower and more difficult to produce the right type of protein

Process

Components are made in a lab and injected into the arm to stimulate immune response

R & D

Antigen determined for immune stimulation

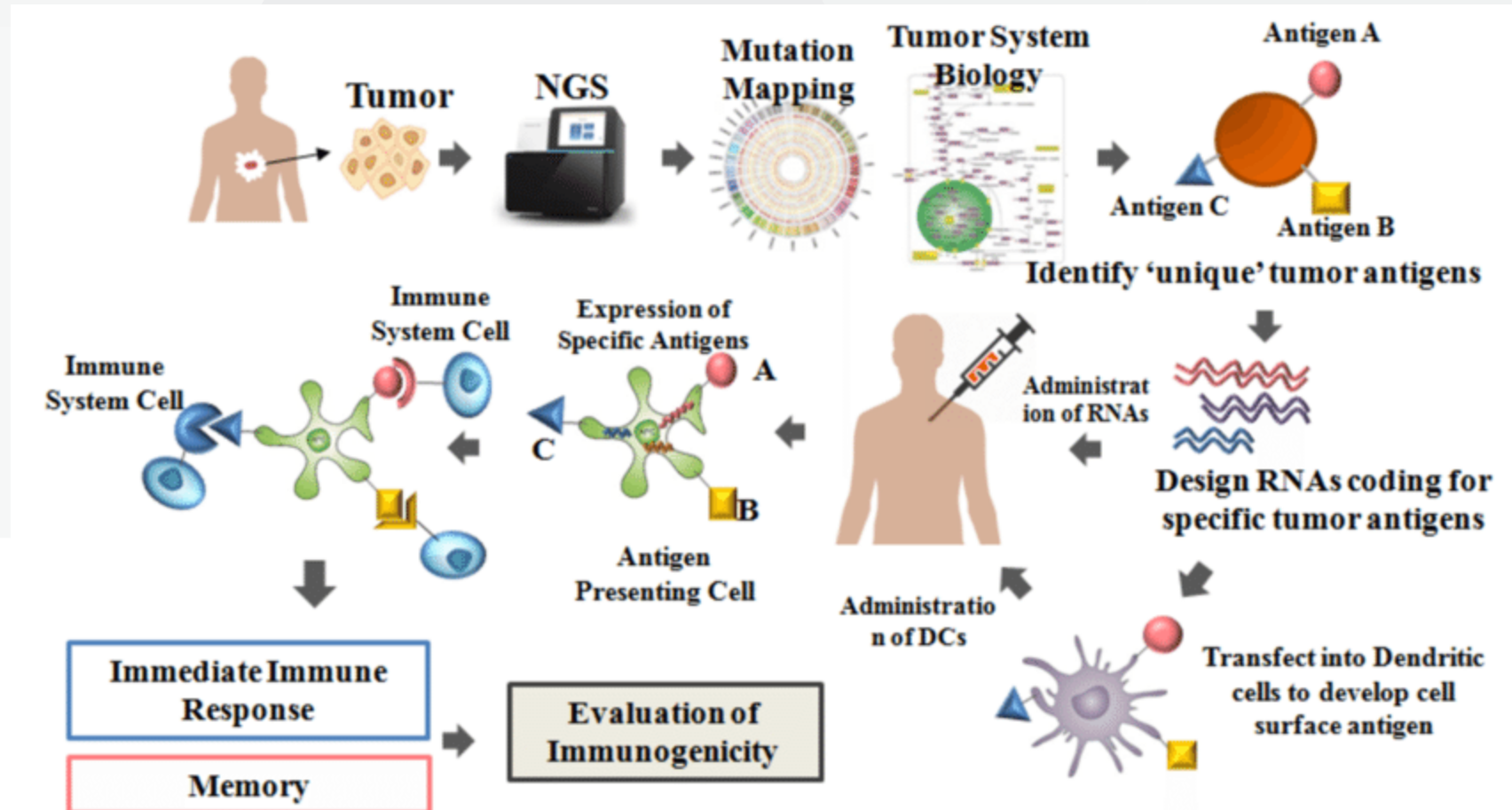


Result

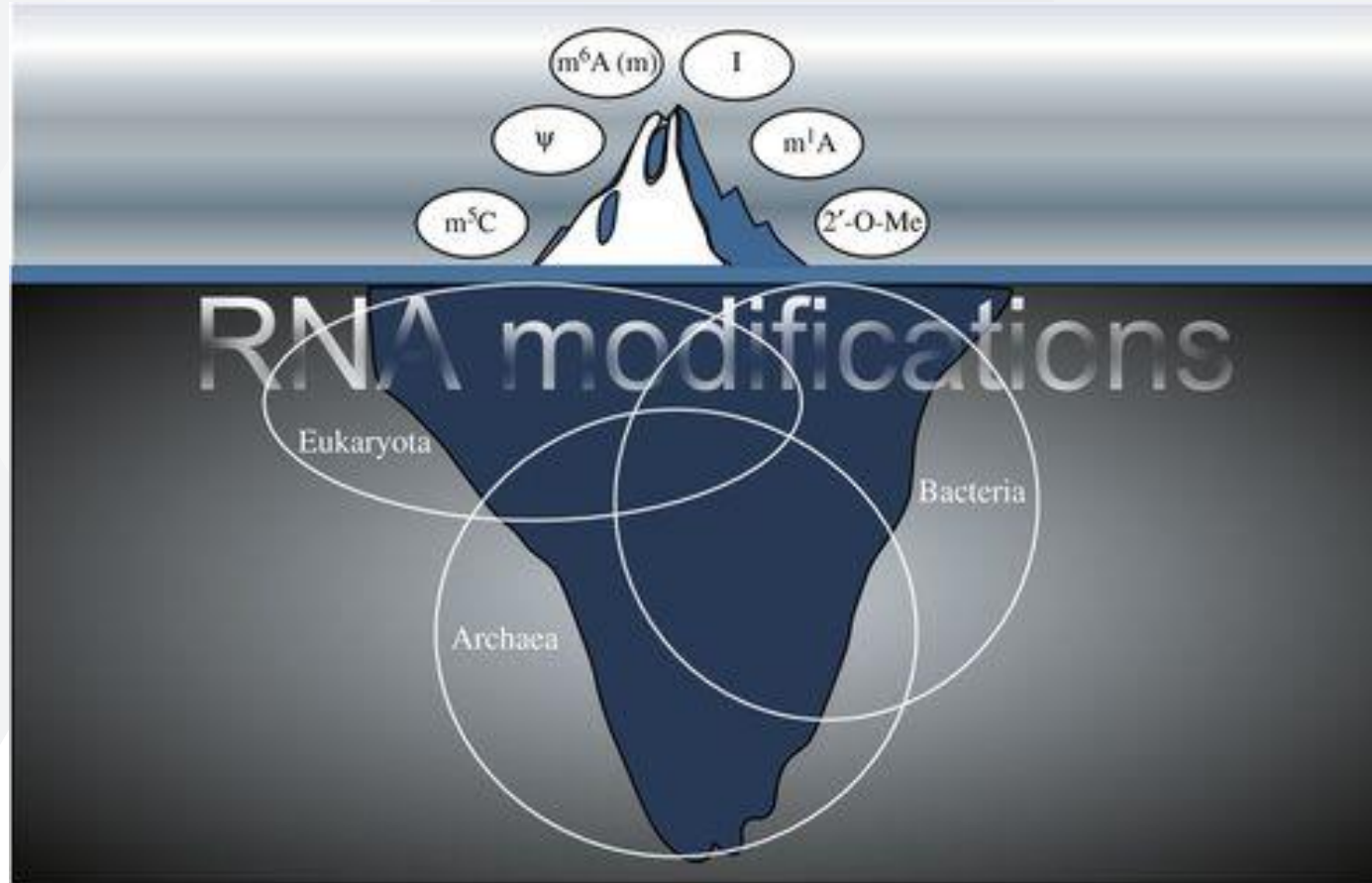
Teaches the body to protect itself against a microbe



Development of personalized RNA-based cancer vaccines

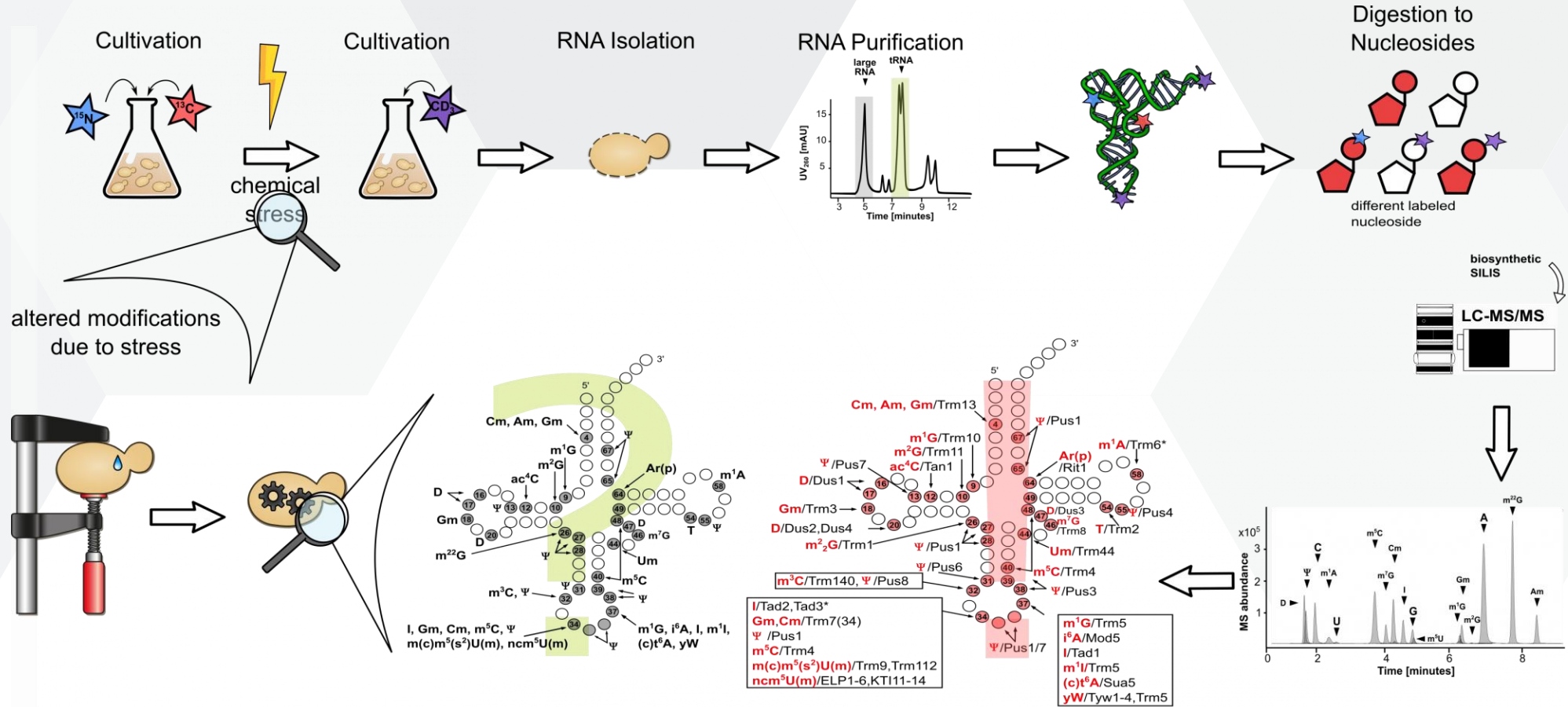


Challenges

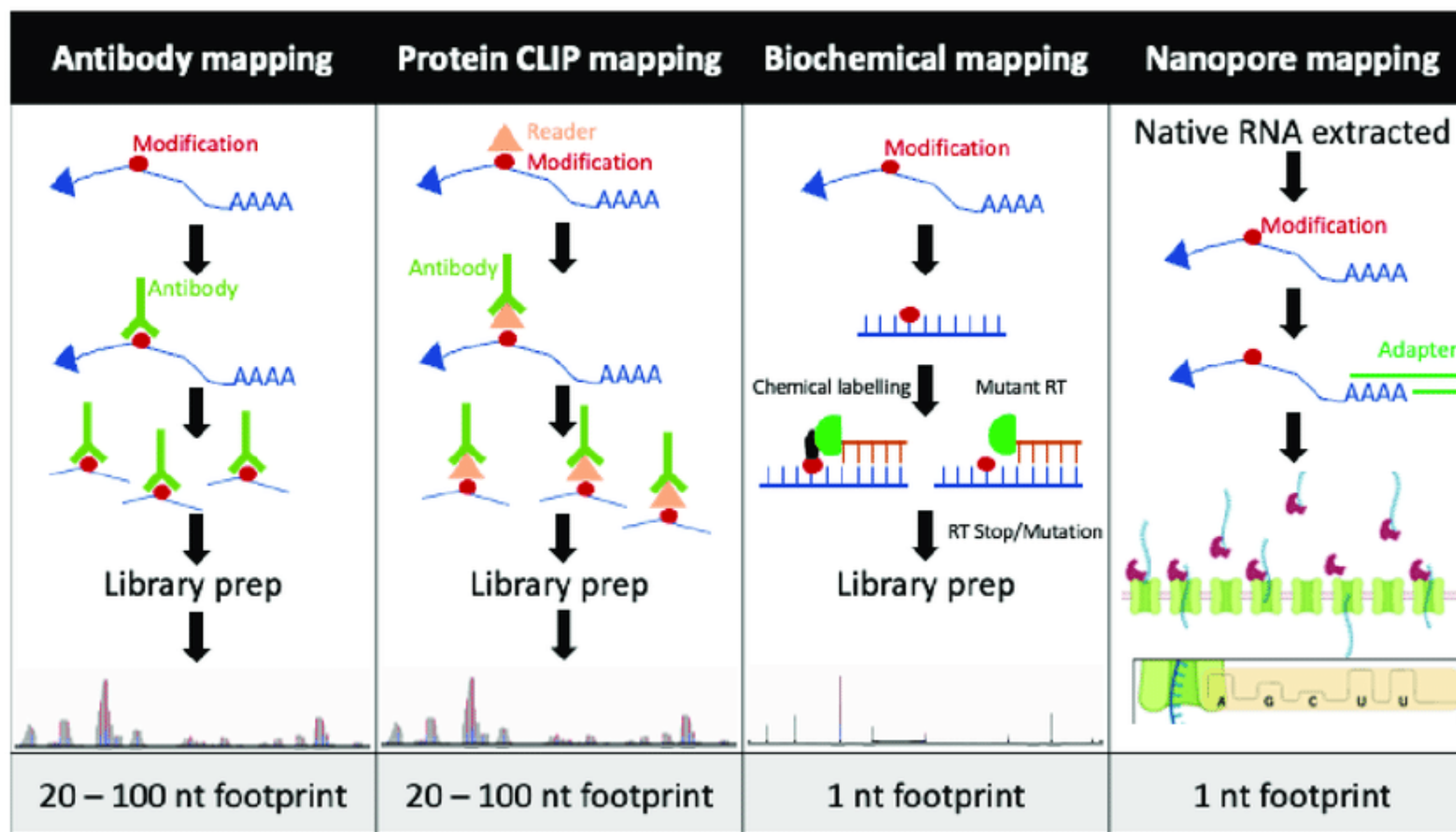


Problems with detection

Amount of starting material



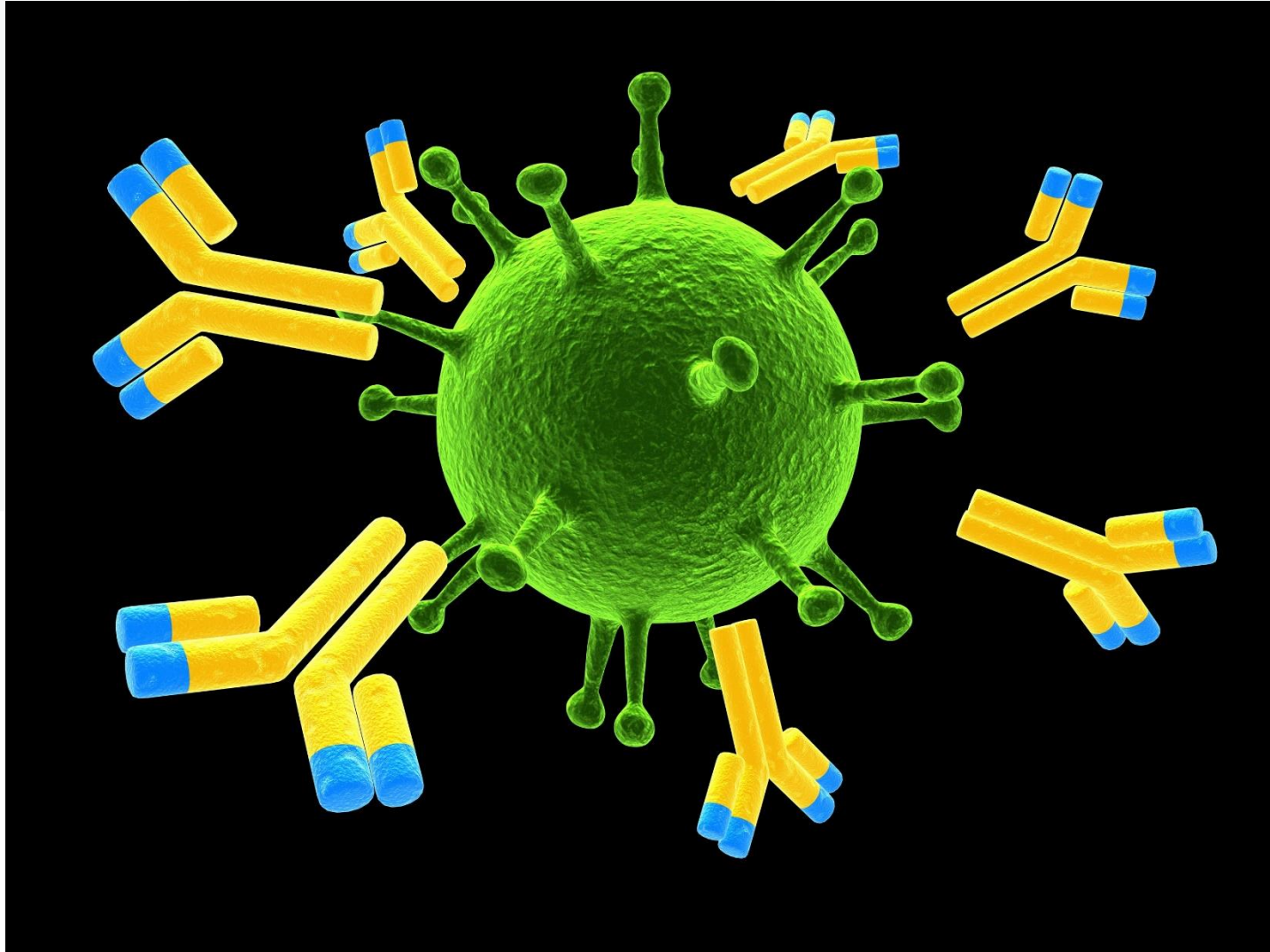
Current techniques



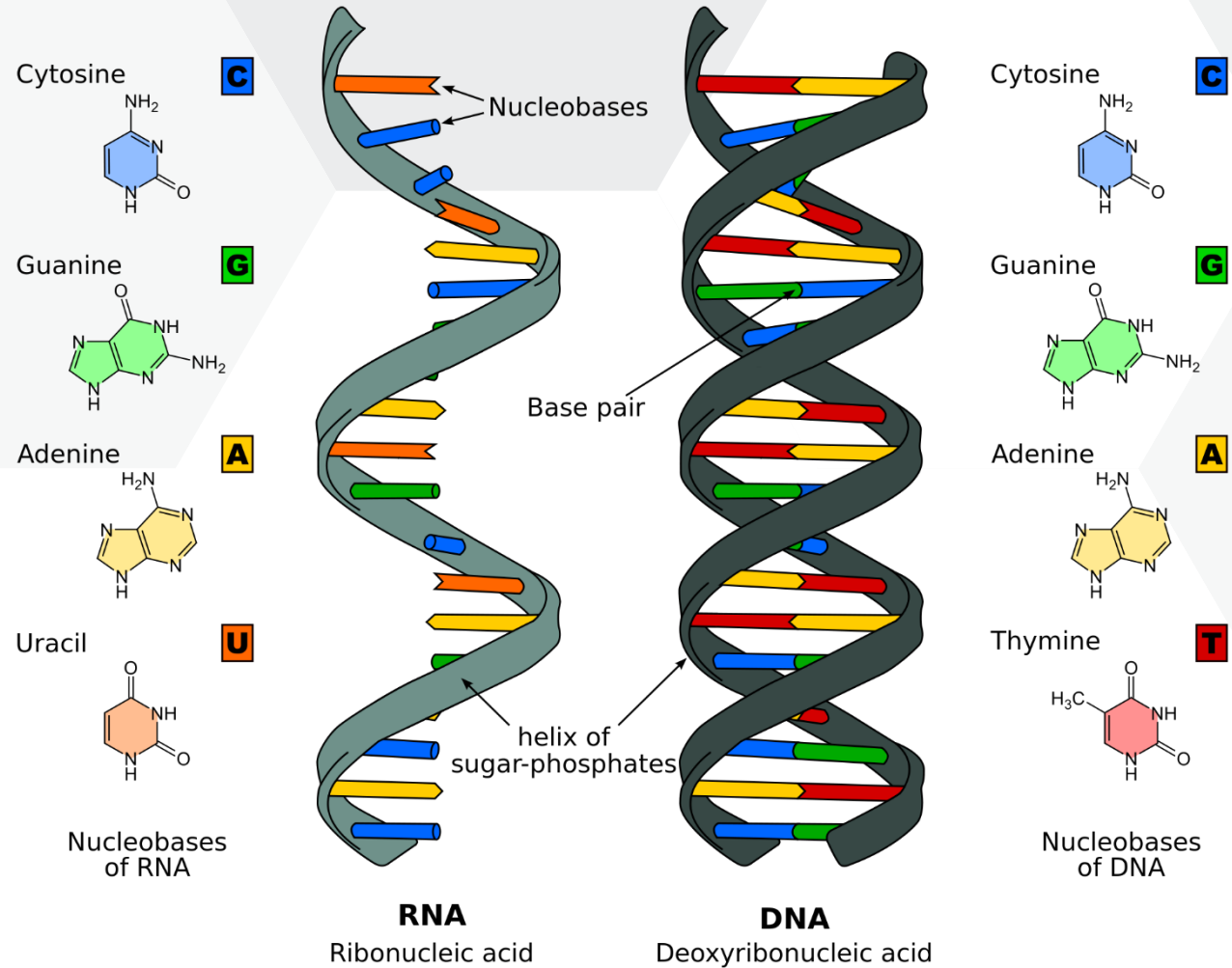
Problem with antibody specificity

What we do

Antibody/Antigen interaction



Problem with nucleic acids



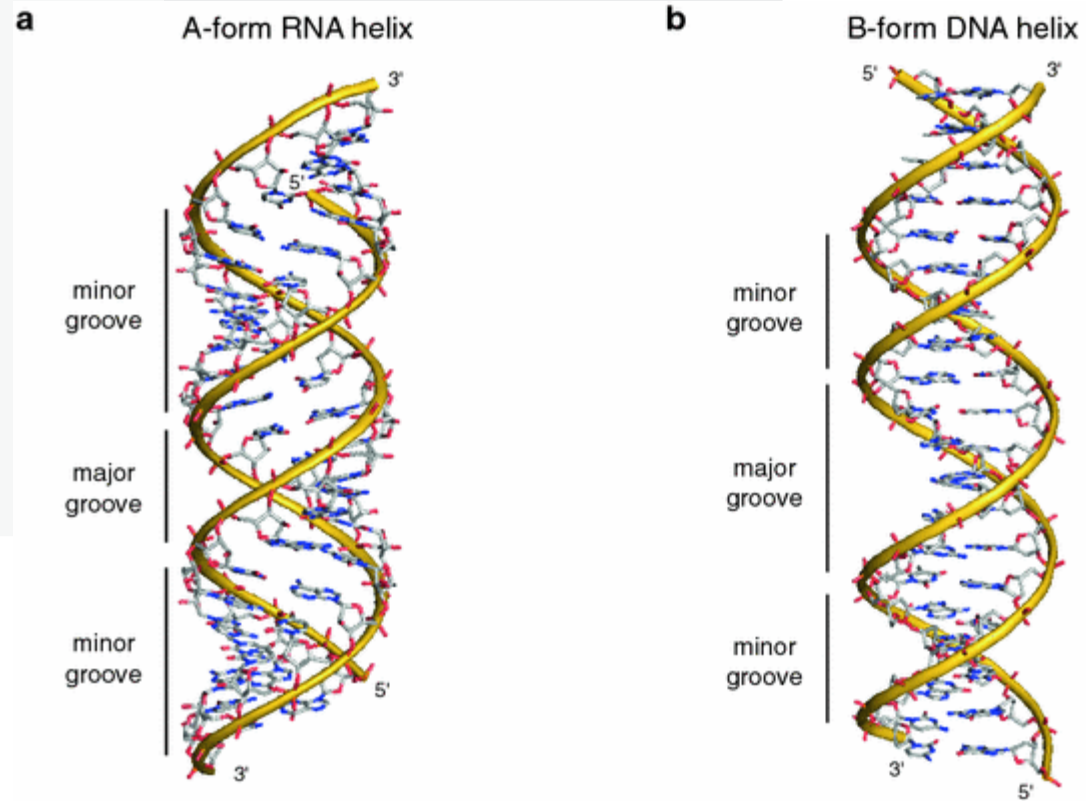
Innate immune pattern recognition receptors (PRRs) discriminate self from non-self



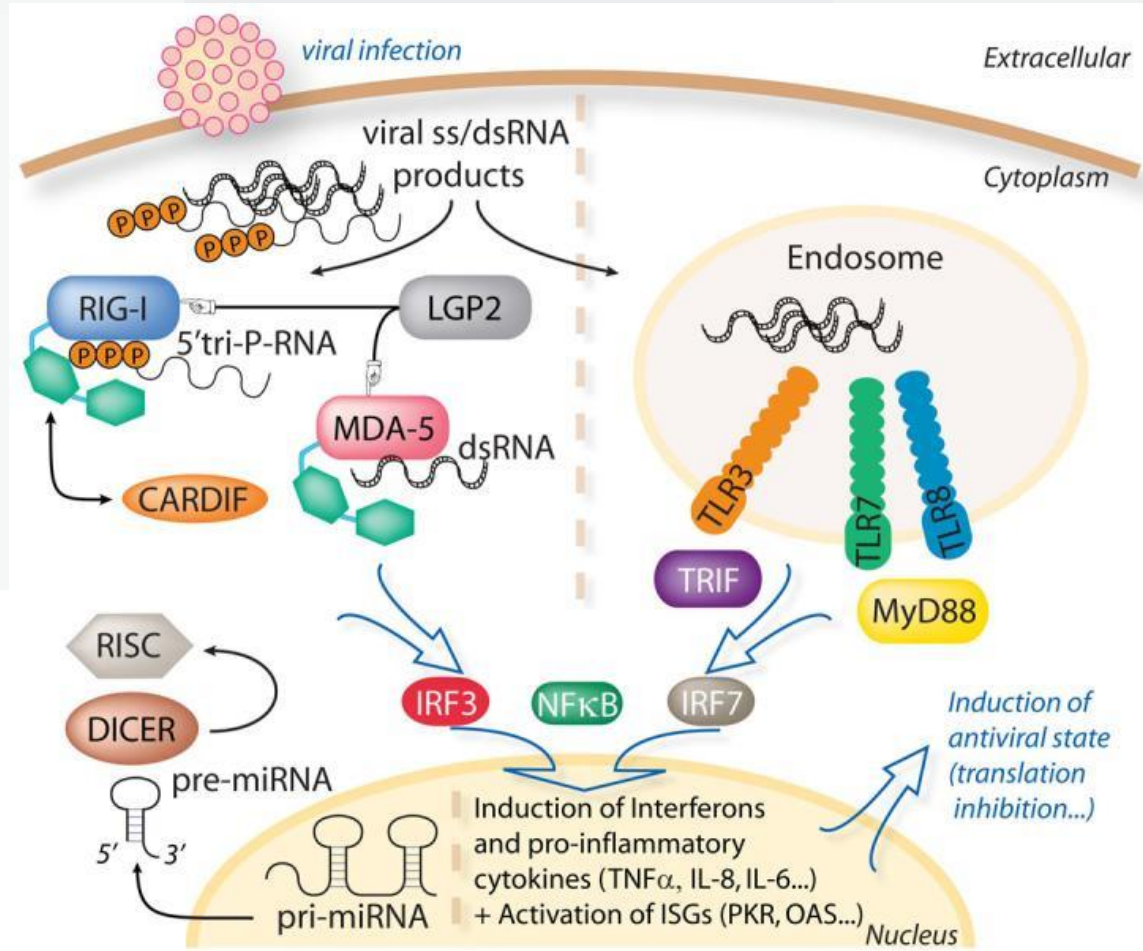
Charles Janeway

Polly Matzinger PRRs also react to damage or danger signals

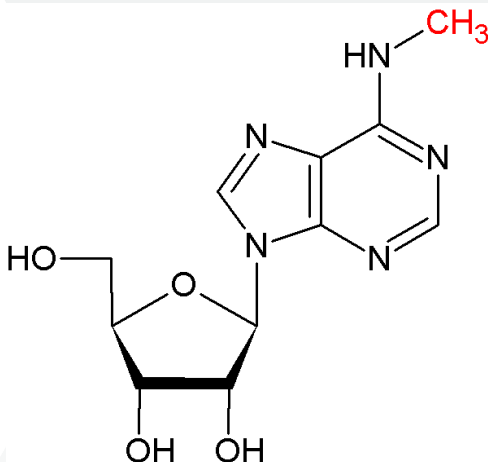
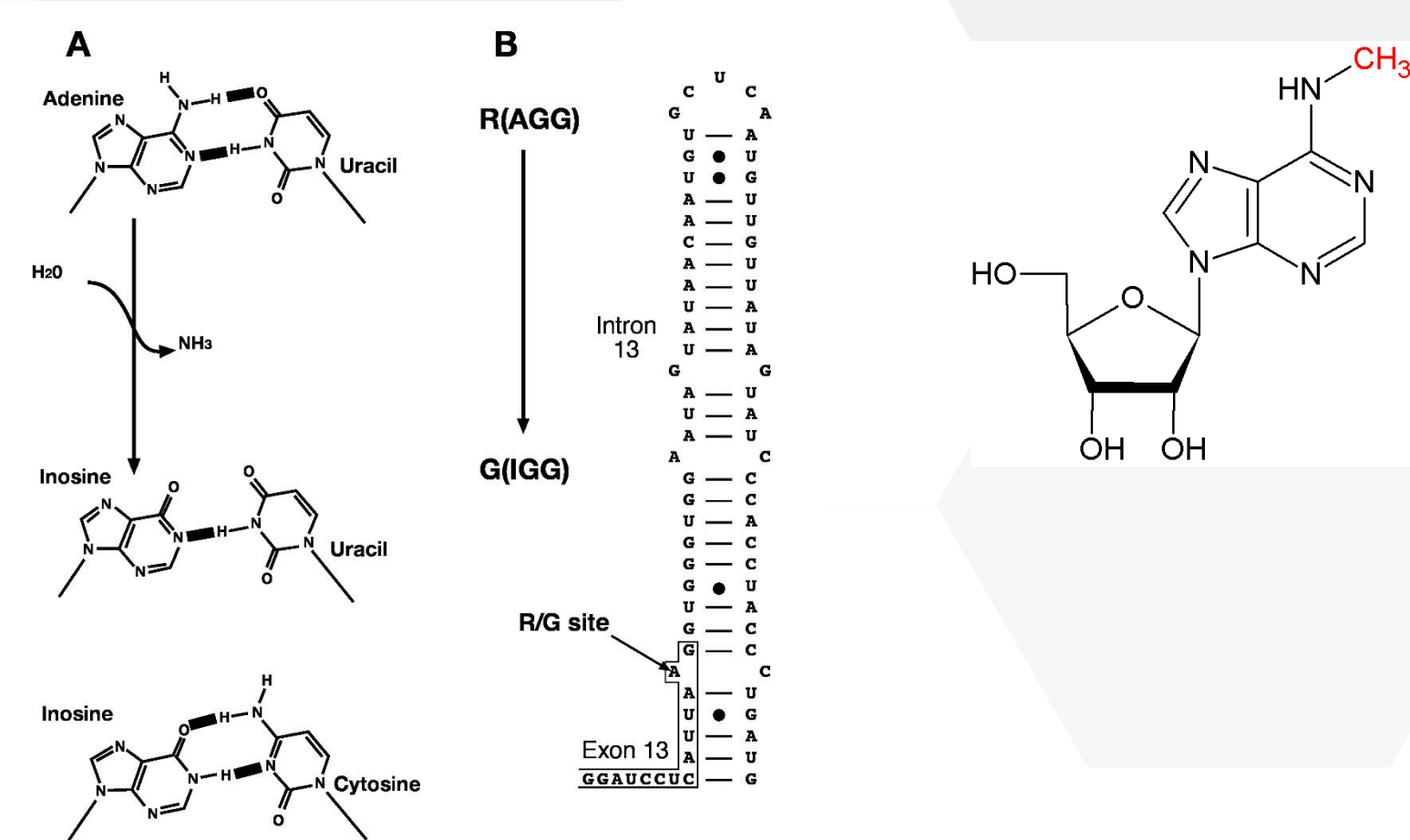
dsRNA versus DNA



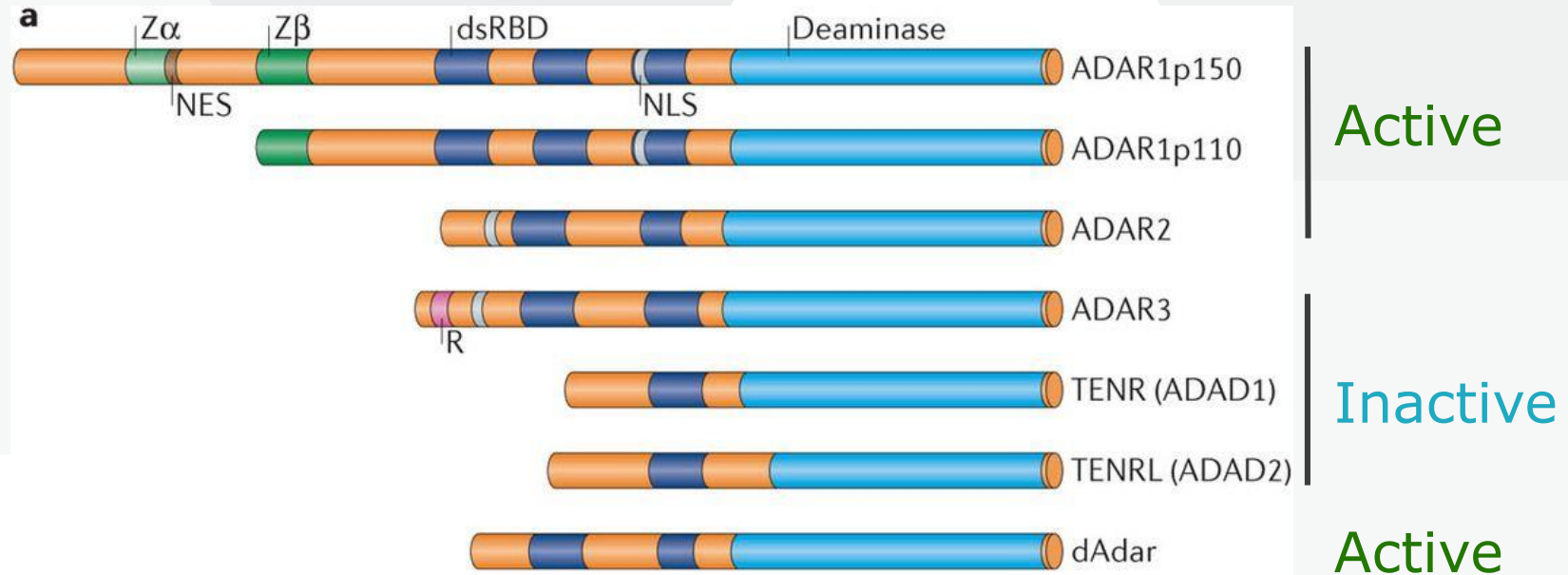
Cellular dsRNA is hazardous. Innate immune and RNAi trigger.



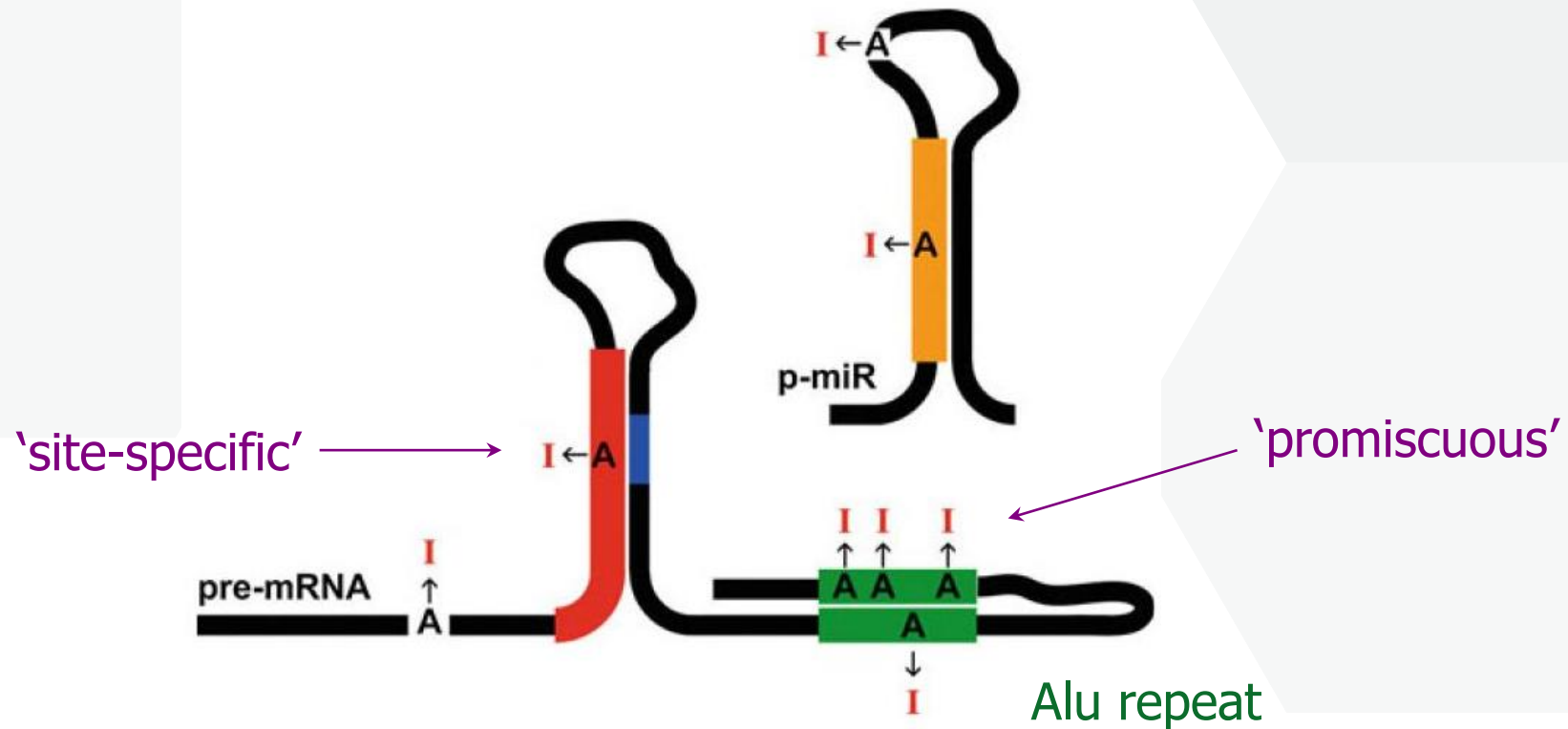
Conversion of adenosine to inosine change the encoded protein



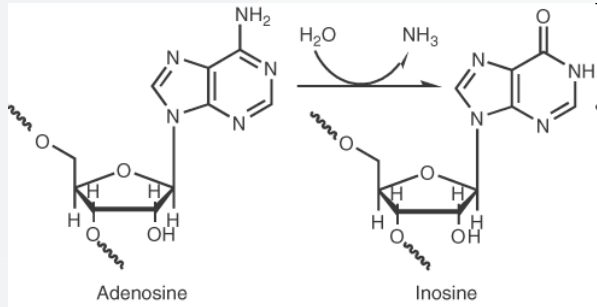
ADAR proteins in vertebrates and in *Drosophila*



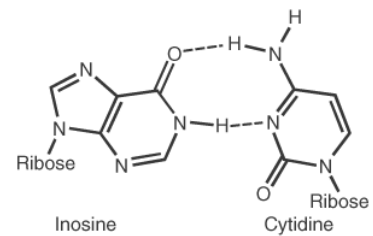
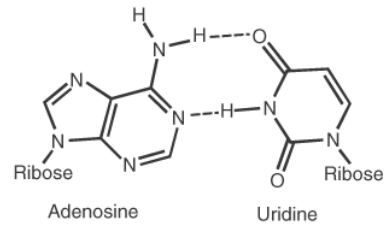
Adenosine Deaminases acting on RNA (ADARs) edit A-to-I in dsRNA



Prevalence of A-to-I editing

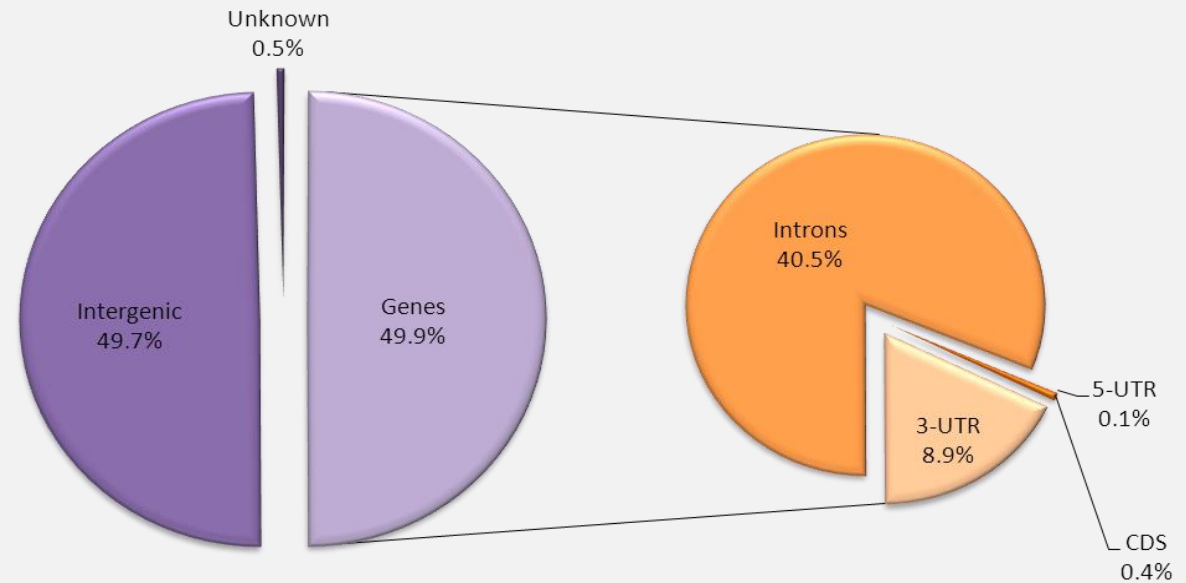


(a)



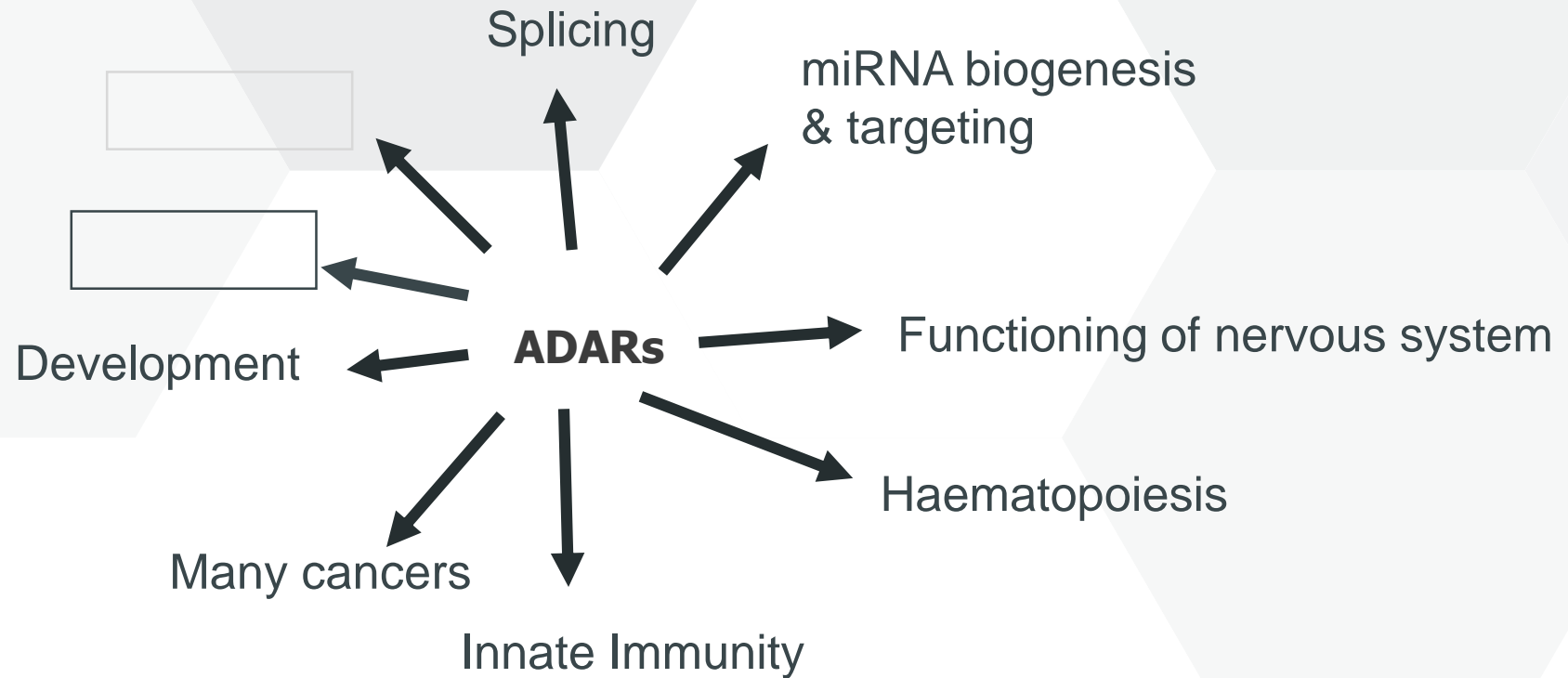
(b)

Human whole transcriptome A→G mismatches (~21,400)



Compiled from Peng et al. (2012) Nat Biotechnol 30, 253-60

Biological roles of ADARs



Editing dependent & independent roles of ADAR

Summary of ADAR mutant phenotypes in vertebrates and in *Drosophila*



Mutated gene *dAdar*^{-/-}

Adar1^{-/-}

Adar2^{-/-}

Tissue CNS

Liver

Brain

Transcript ?

Alus elements

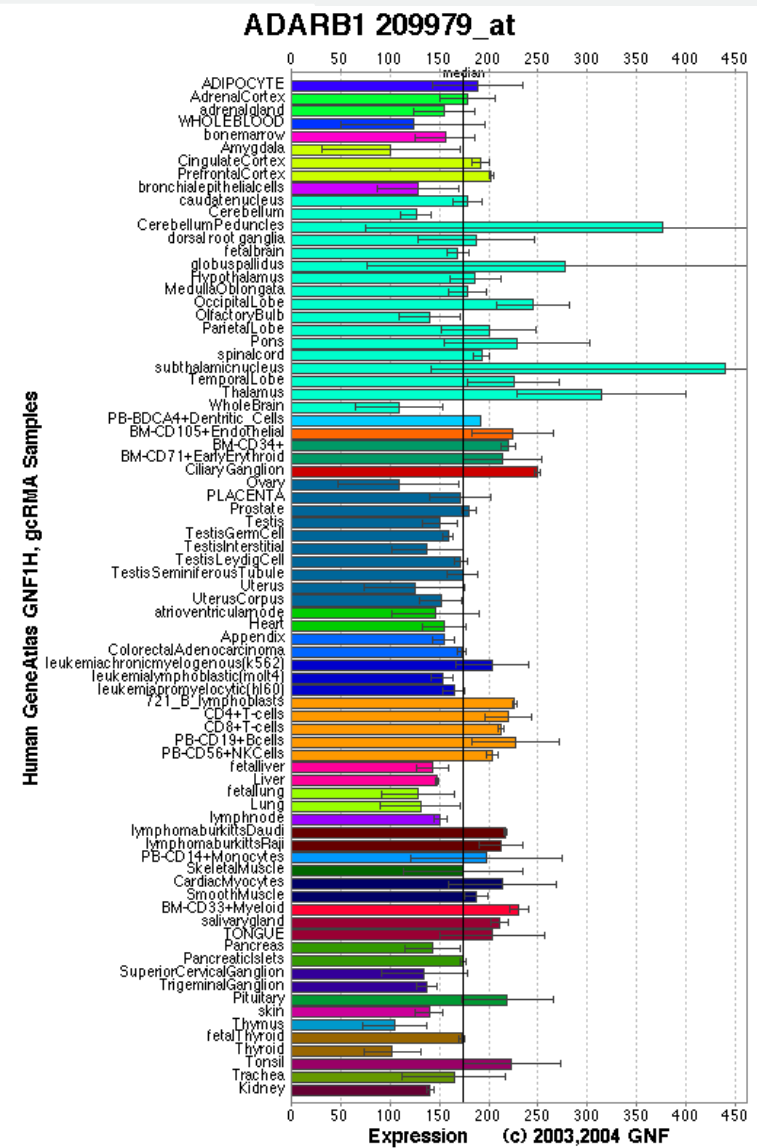
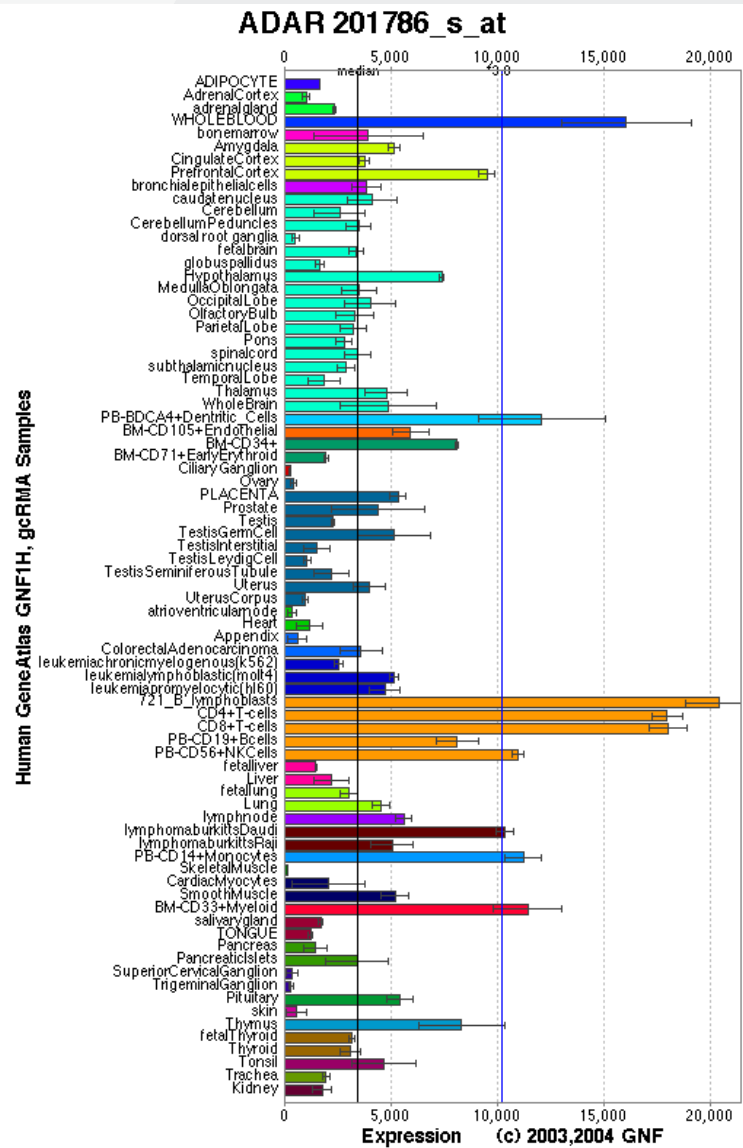
Gria2 Q/R (GluR2)

Phenotype Behavior
Locomotion
Neurodegeneration

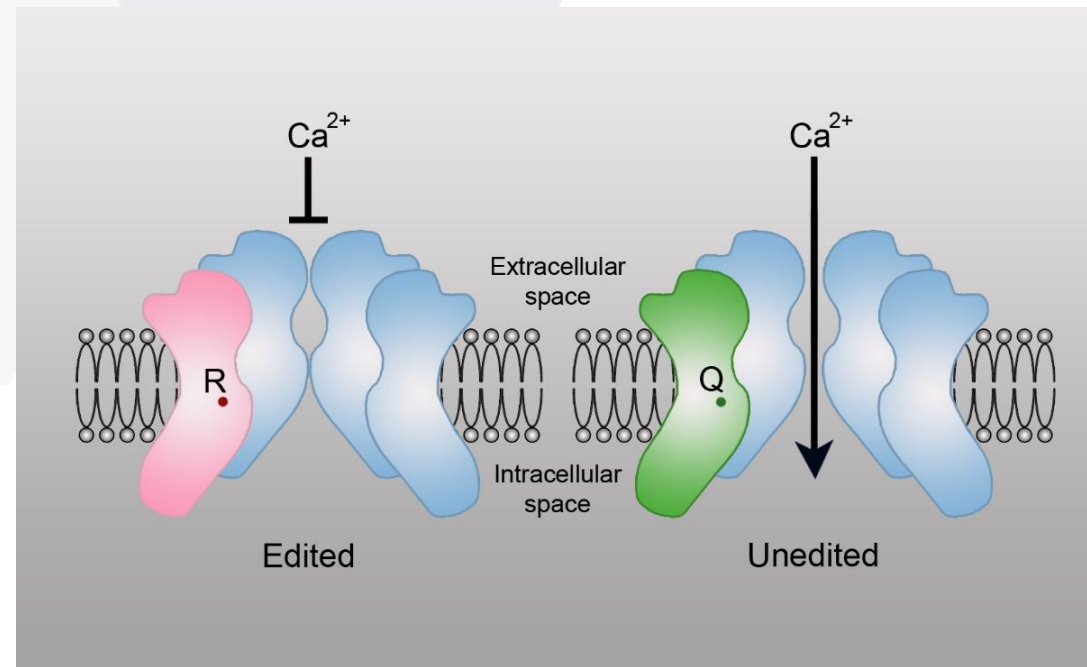
Hematopoiesis
Innate immunity
Death by E12.5

Seizures, death by P20

hADAR1 versus hADAR2 tissue expression



RNA editing of transcript encoding the GluR2 subunit of vertebrate glutamate AMPA receptors



ADARB1 mutations in childhood seizures EIMFS (epilepsy of infancy with migrating focal seizures)

- We have established a collaboration with Dr. Tiong Tan from the University of Melbourne and Dr. Mark Fitzgerald from Children's Hospital of Philadelphia, Dr. Riki Sukenik Halew, Meir Medical Center Israel,.
- h*ADAR2* variants were found in four patients (US patient now deceased) suffering with **microcephaly, severe intellectual disability and seizures**
- Australian patient: biallelic mutation in *ADAR2*; Lys367Asn, Thr498Ala
- US patient: homozygous mutation in *ADAR2*; Lys127Glu
- Israeli patient : homozygous mutation in *ADAR2*; Arg603Glu
- Iranian patients: homozygous mutation in *ADAR2*; Arg630Glu
 homozygous mutation in *ADAR2*; Ala722Val

Mutation in *KCNT1* also result in EIMSF

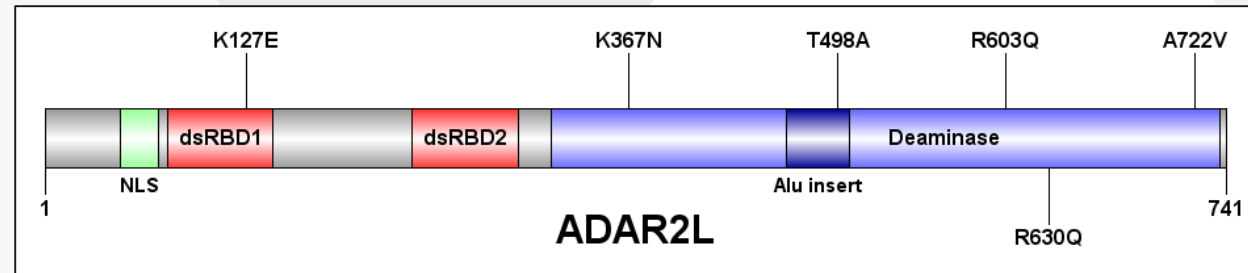
Patients with *ADARB1* variants



Tan *et al.* Am J Hum Genet. 2020,

Maroofian *et al.* J Med Genet. 2021

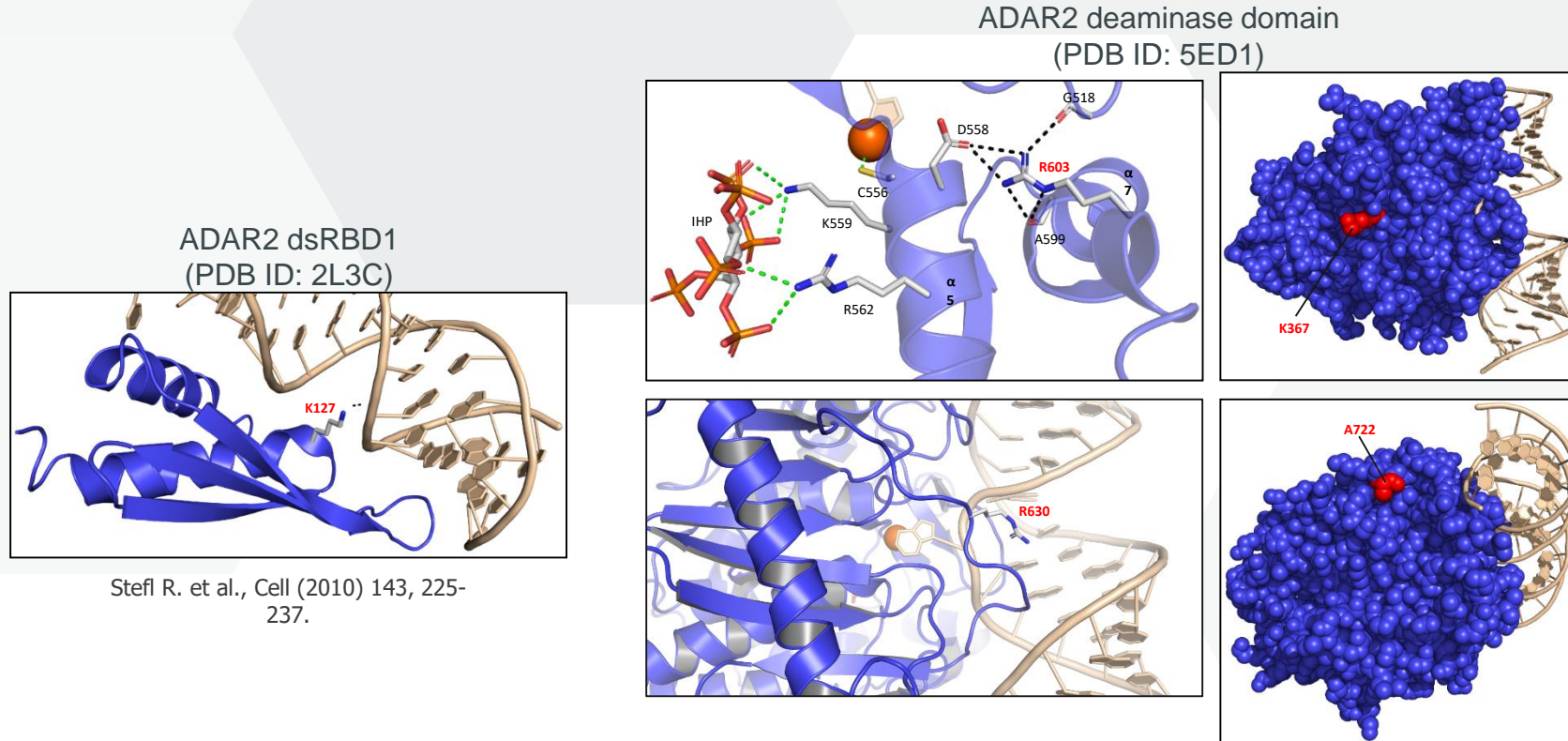
Locations of ADAR2 mutations in ADAR2 Variants



Tan *et al.* Am J Hum Genet. 2020,

Maroofian *et al.* J Med Genet. 2021

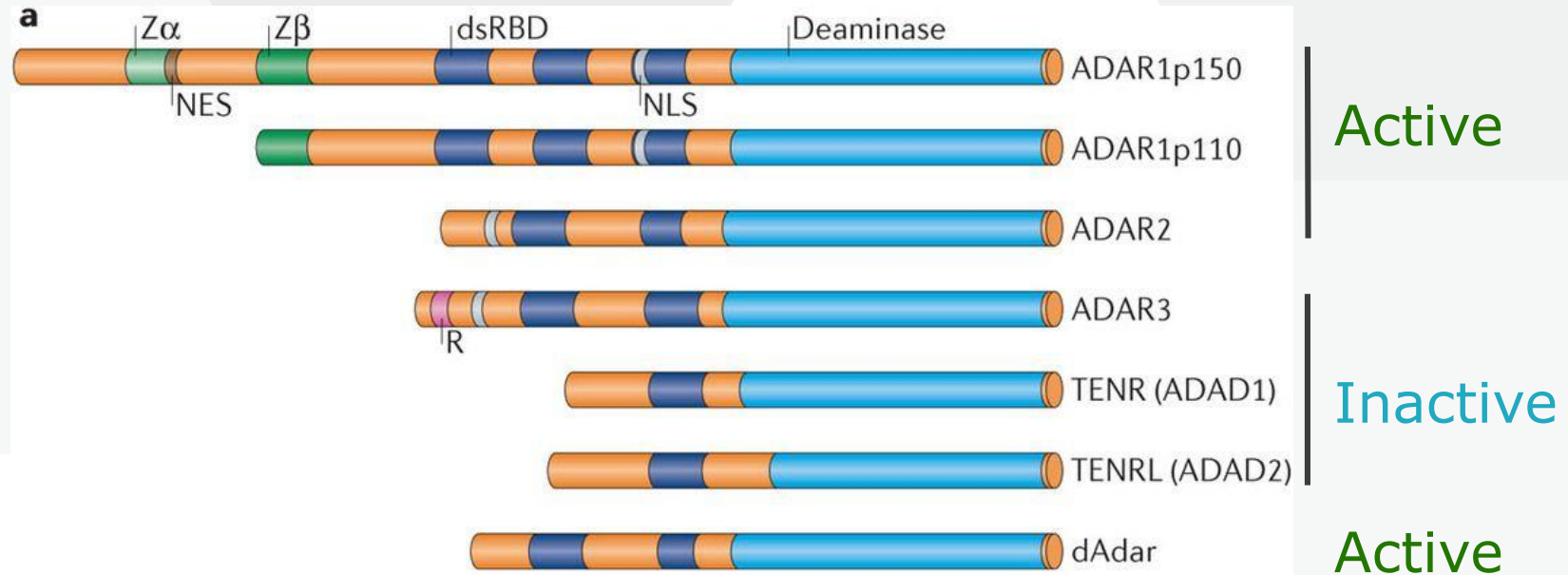
Location of Variants



- dsRBD1 or deaminase domain in **blue**
- dsRNA substrate in **wheat**, Zn²⁺ in **orange**

Tan *et al.* Am J Hum Genet. 2020,

ADAR proteins in vertebrates and in *Drosophila*



Mutations in *ADAR1* cause Aicardi-Goutières syndrome associated with a type I interferon signature

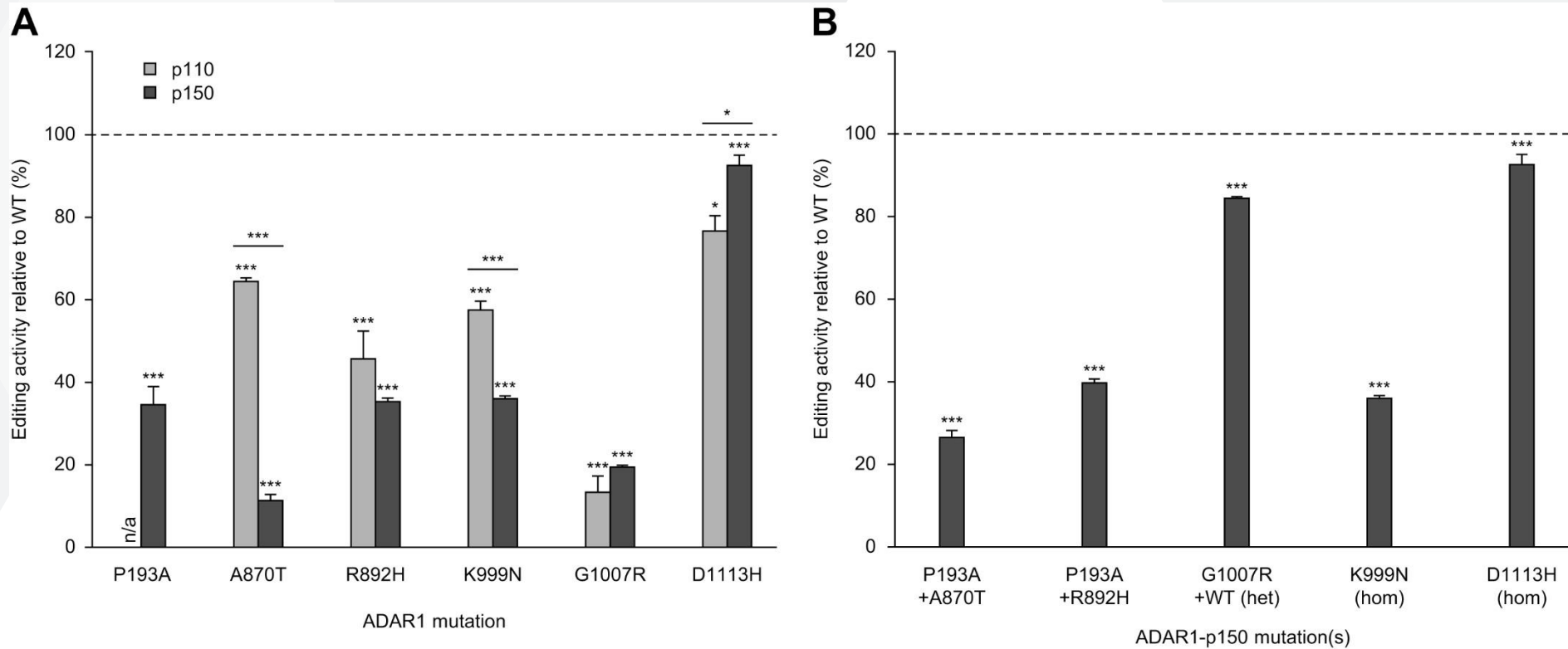
Gillian I Rice¹, Paul R Kasher¹, Gabriella M A Forte¹, Niamh M Mannion², Sam M Greenwood², Marcin Szykiewicz¹, Jonathan E Dickerson¹, Sanjeev S Bhaskar¹, Massimiliano Zampini¹, Tracy A Briggs¹, Emma M Jenkinson¹, Carlos A Bacino³, Roberta Battini⁴, Enrico Bertini⁵, Paul A Brogan⁶, Louise A Brueton⁷, Marialuisa Carpanelli⁸, Corinne De Laet⁹, Pascale de Lonlay¹⁰, Mireia del Toro¹¹, Isabelle Desguerre¹², Elisa Fazzi¹³, Àngels Garcia-Cazorla^{14,15}, Arvid Heiberg¹⁶, Masakazu Kawaguchi¹⁷, Ram Kumar¹⁸, Jean-Pierre S-M Lin¹⁹, Charles M Lourenco²⁰, Alison M Male²¹, Wilson Marques Jr²⁰, Cyril Mignot²²⁻²⁴, Ivana Olivieri²⁵, Simona Orcesi²⁵, Prab Prabhakar²⁶, Magnhild Rasmussen²⁷, Robert A Robinson²⁶, Flore Rozenberg²⁸, Johanna L Schmidt²⁹, Katharina Steindl³⁰, Tiong Y Tan³¹, William G van der Merwe³², Adeline Vanderver²⁹, Grace Vassallo³³, Emma L Wakeling³⁴, Evangeline Wassmer³⁵, Elizabeth Whittaker³⁶, John H Livingston³⁷, Pierre Lebon²⁸, Tamio Suzuki¹⁷, Paul J McLaughlin³⁸, Liam P Keegan², Mary A O'Connell², Simon C Lovell³⁹ & Yanick J Crow¹

Adenosine deaminases acting on RNA (ADARs) catalyze the hydrolytic deamination of adenosine to inosine in double-stranded RNA (dsRNA) and thereby potentially alter the information content and structure of cellular RNAs. Notably, although the overwhelming majority of such editing events occur in transcripts derived from Alu repeat elements, the biological function of non-coding RNA editing remains uncertain. Here, we show that mutations in *ADAR1* (also known as *ADAR*) cause the autoimmune disorder Aicardi-Goutières syndrome (AGS). As in *Adar1*-null mice, the human disease state is associated with upregulation of interferon-stimulated genes, indicating a possible role for ADAR1 as a suppressor of type I interferon signaling. Considering recent insights derived from the study of other AGS-related proteins, we speculate that ADAR1 may limit the cytoplasmic accumulation of the dsRNA generated from genomic repetitive elements.

autosomal recessive trait⁷, rare examples of disease due to *de novo* dominant mutations in *TREX1* have been reported⁸⁻¹⁰.

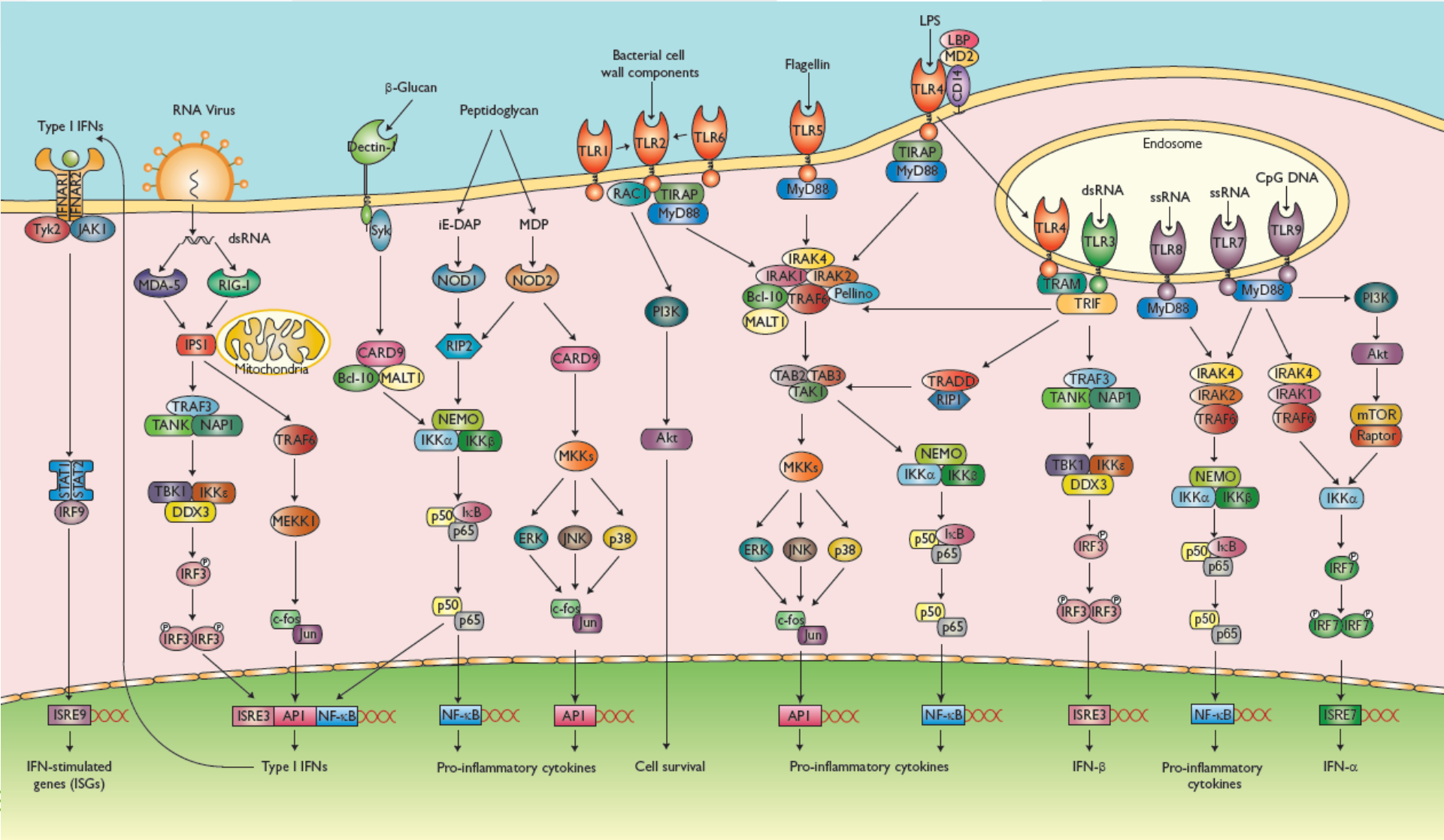
Studies of the function of *TREX1* have delineated a cell-intrinsic mechanism for the initiation of an autoimmune response by interferon (IFN)-stimulatory nucleic acid^{11,12}, begging the question of the source of nucleic acid inducing the type I IFN-mediated immune disturbance in AGS. In this regard, it has been shown that *TREX1* can metabolize reverse-transcribed DNA and that single-stranded DNA derived from endogenous retroelements accumulates in *TREX1*-deficient cells¹¹. On a related note, *TREX1* (ref. 13), *SAMHD1* (refs. 14-16) and RNase H2 (ref. 17) have been implicated in the metabolism of the (exogenous) retrovirus HIV-1. Perhaps most notably, a recent study showed rescue of the lethal inflammatory *TREX1*-null mouse phenotype by a combination of reverse transcriptase inhibitors (antiretroviral therapy as used to treat HIV-1)¹⁸, suggesting that the accumulation of cytosolic DNA in *TREX1*-null cells can be ameliorated by inhibiting endogenous retroelement cycling.

Combination of mutations in *ADAR1* as found in AGS patients.

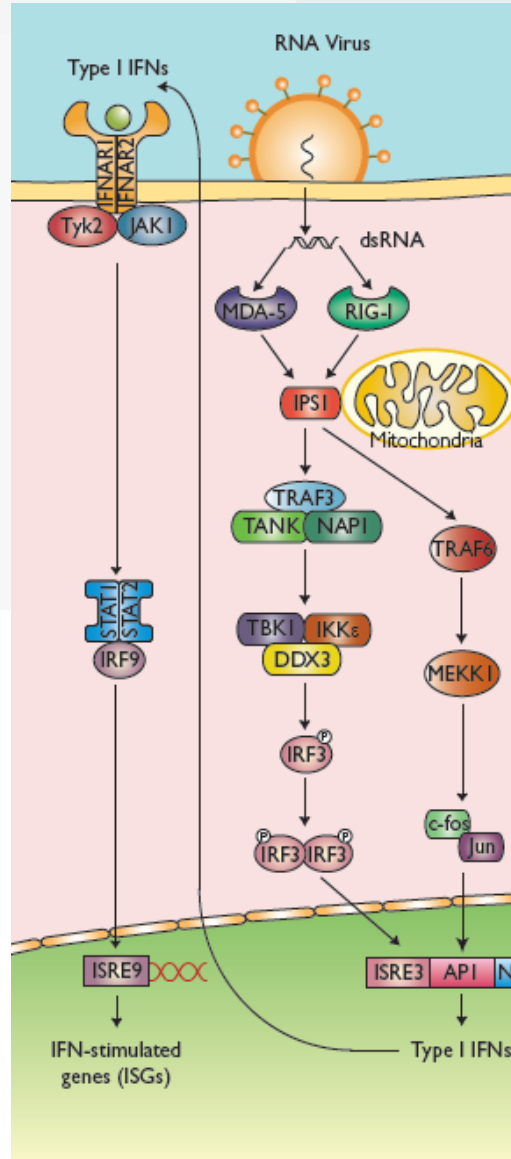


Innate immune system

Pattern Recognition Receptor signalling



Rescue of *Adar1* lethality by preventing innate immune sensing of intracellular dsRNA



Mavs/IPS-1 knockout prevents all known signalling in this pathway.

Mannion, 2014; complete null, death E12.5. Double mutant with *Mavs* dies at birth.

Liddicot 2015; inactive mutant, death by E14.5. Double mutant with *Mda5* has *no phenotype*.

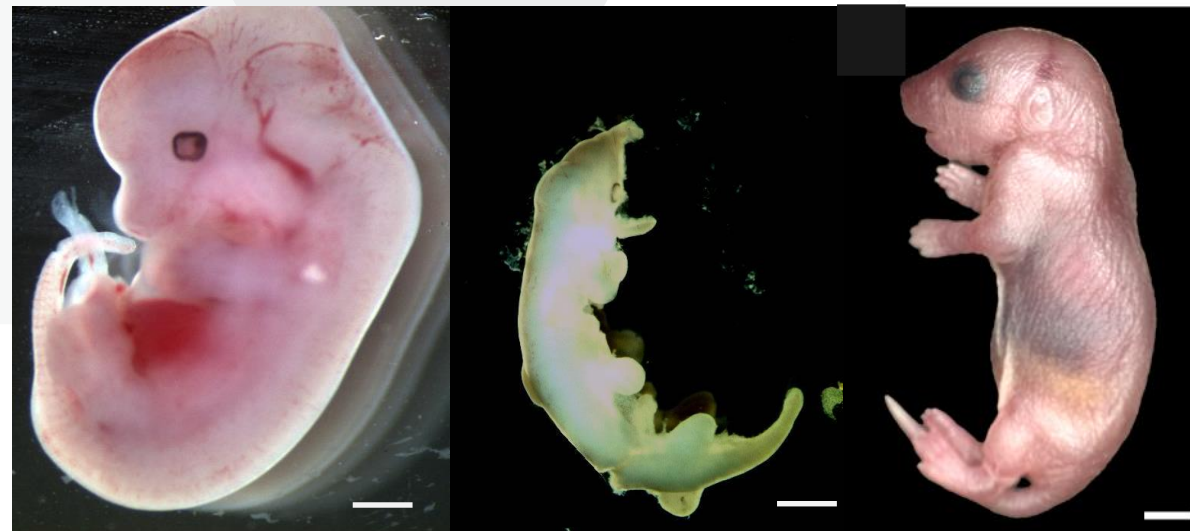
Sun Hur 2018 demonstrated that transcripts encoding inverted Alus activate MDA5

Mavs^{-/-} rescue of *Adar1*^{-/-} to birth

Adar1^{+/+}

Adar1^{-/-}

Adar1^{-/-}*Mavs*^{-/-}

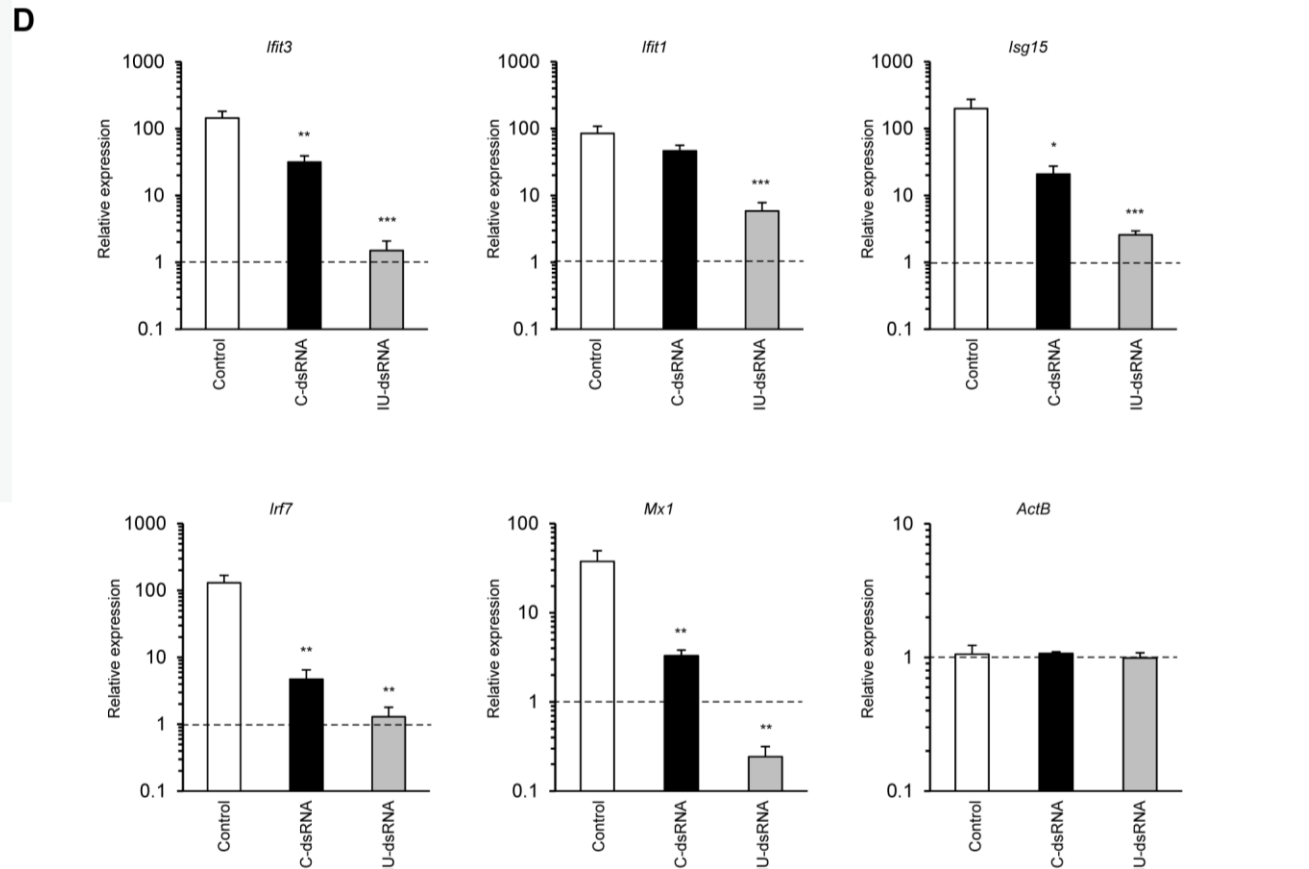


E12.5

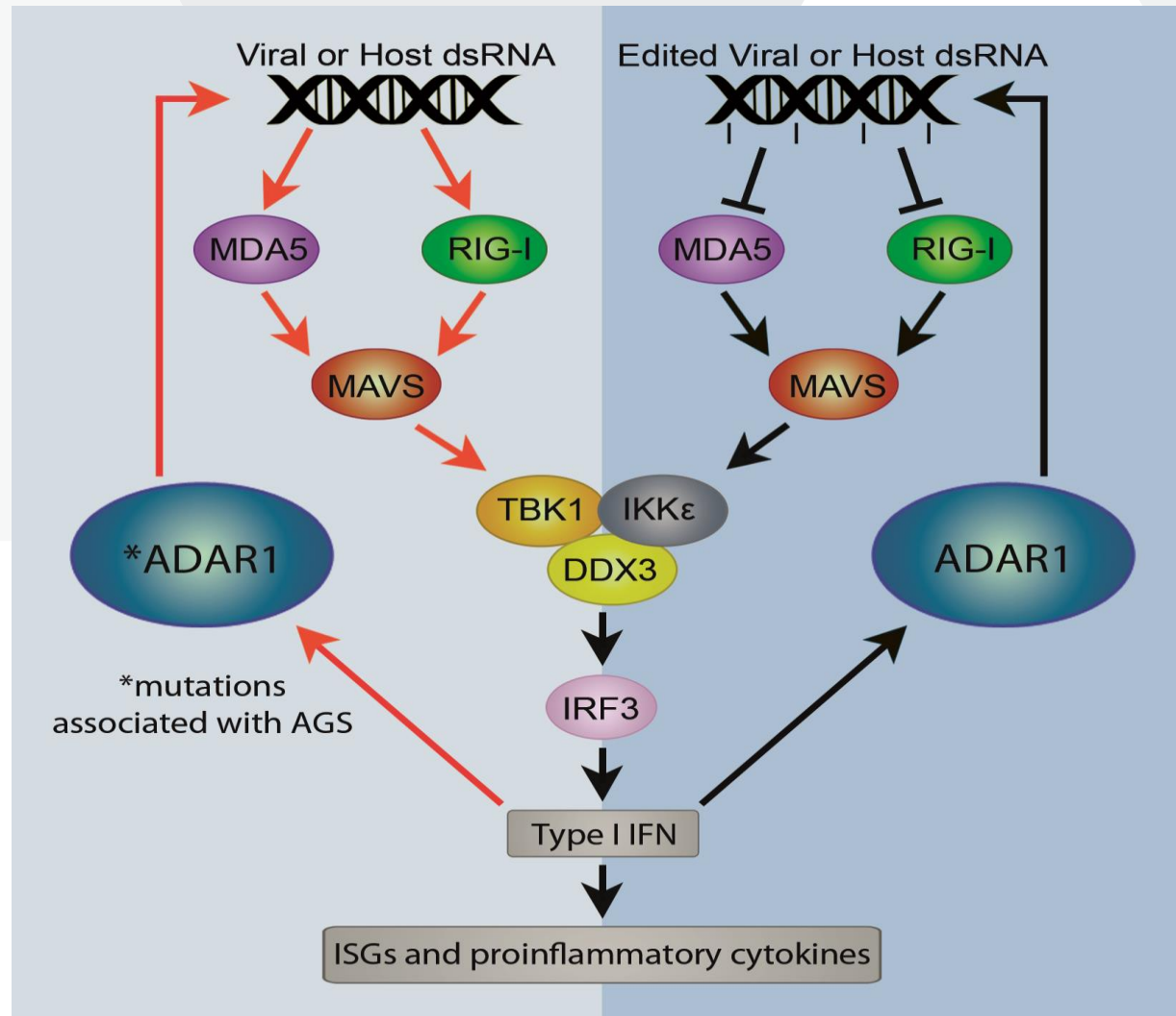
P0

Mannion *et al.* Cell Reports 2014

IU-dsRNA oligo reduces innate immune response in stressed *Adar1*; *p53* MEF



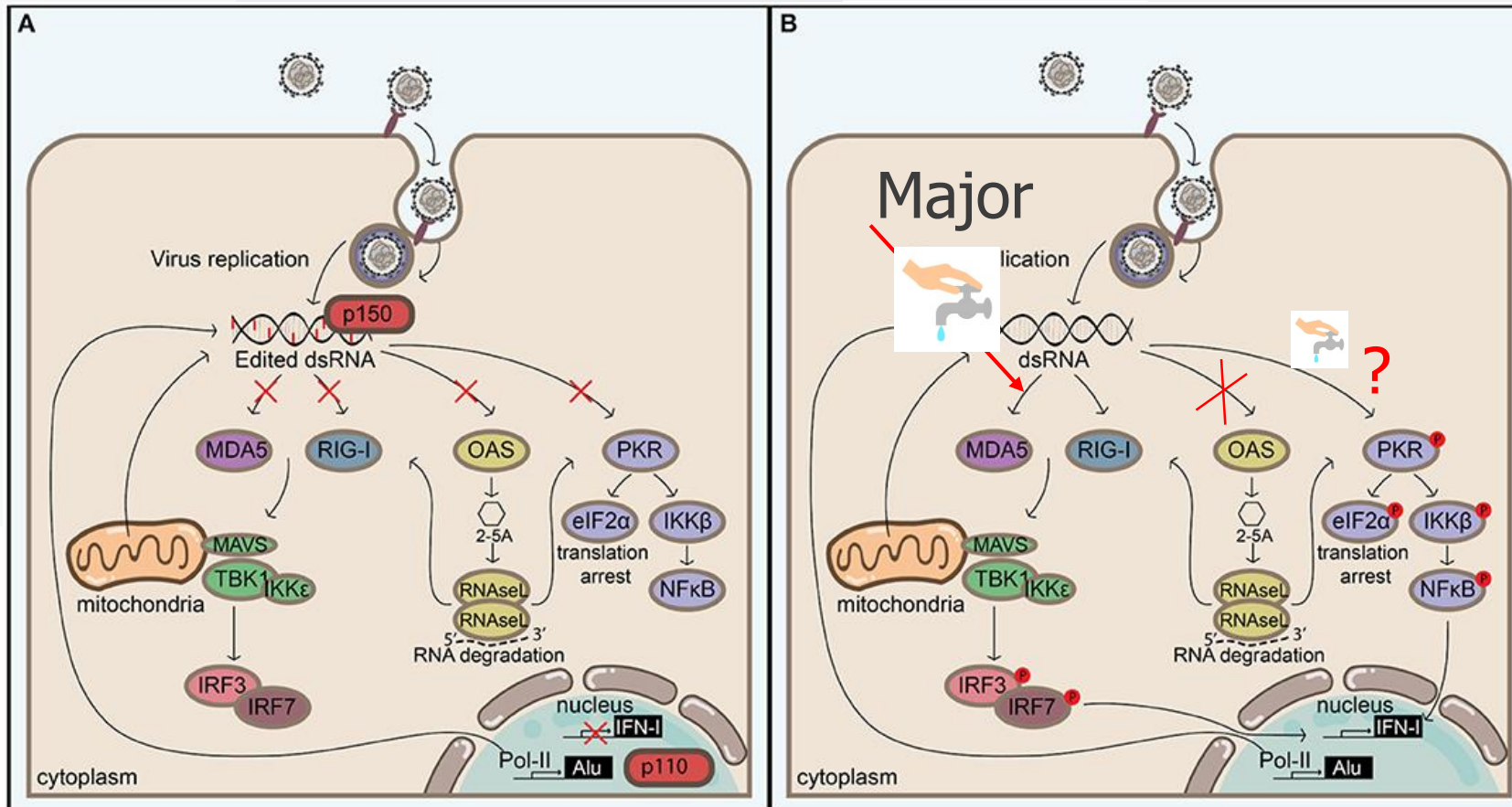
Inosine in RNA helps discriminate self from non-self



Adar1^{-/-} ; Mavs^{-/-} (complete null)

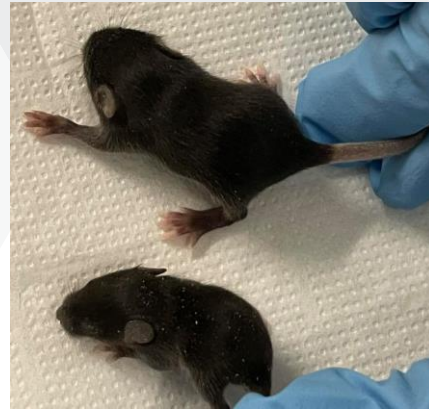


Pkr alone does not rescue *Adar*

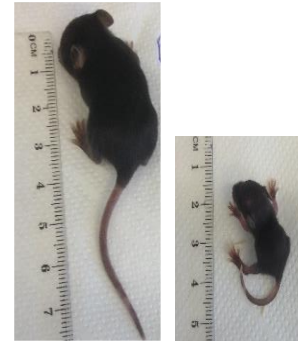


Adapted from Lamers *et al.* Front. Immunol., 25 July 2019

PKR deletion phenotype



Eif2ak2;Adar;Mavs 18days



Adar;Mavs 15d

***Eif2ak2* deletion in *Adar;Mavs* rescues even when it is heterozygous!**

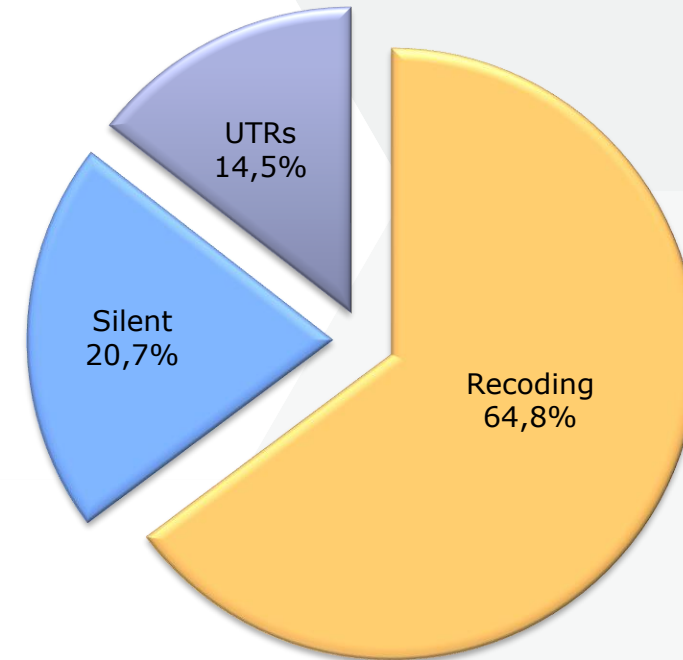
What is the inactive role of
ADARs?

A → I editing in *Drosophila melanogaster*

Poly(A)⁺ RNA-Seq: 972 sites in 597 transcripts

630 alter amino acids
201 silent
141 within UTRs

~20% of endogenous small RNAs bound to AGO2 are edited

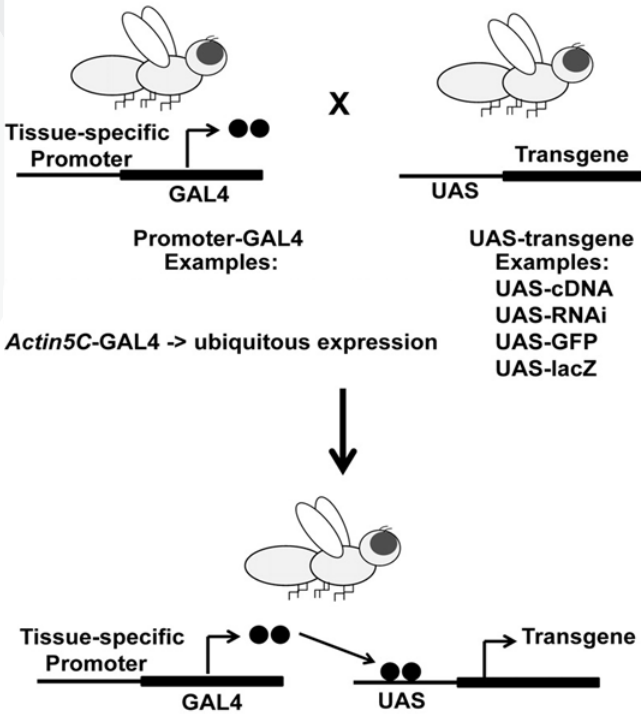


Graveley *et al.* Nature 2011

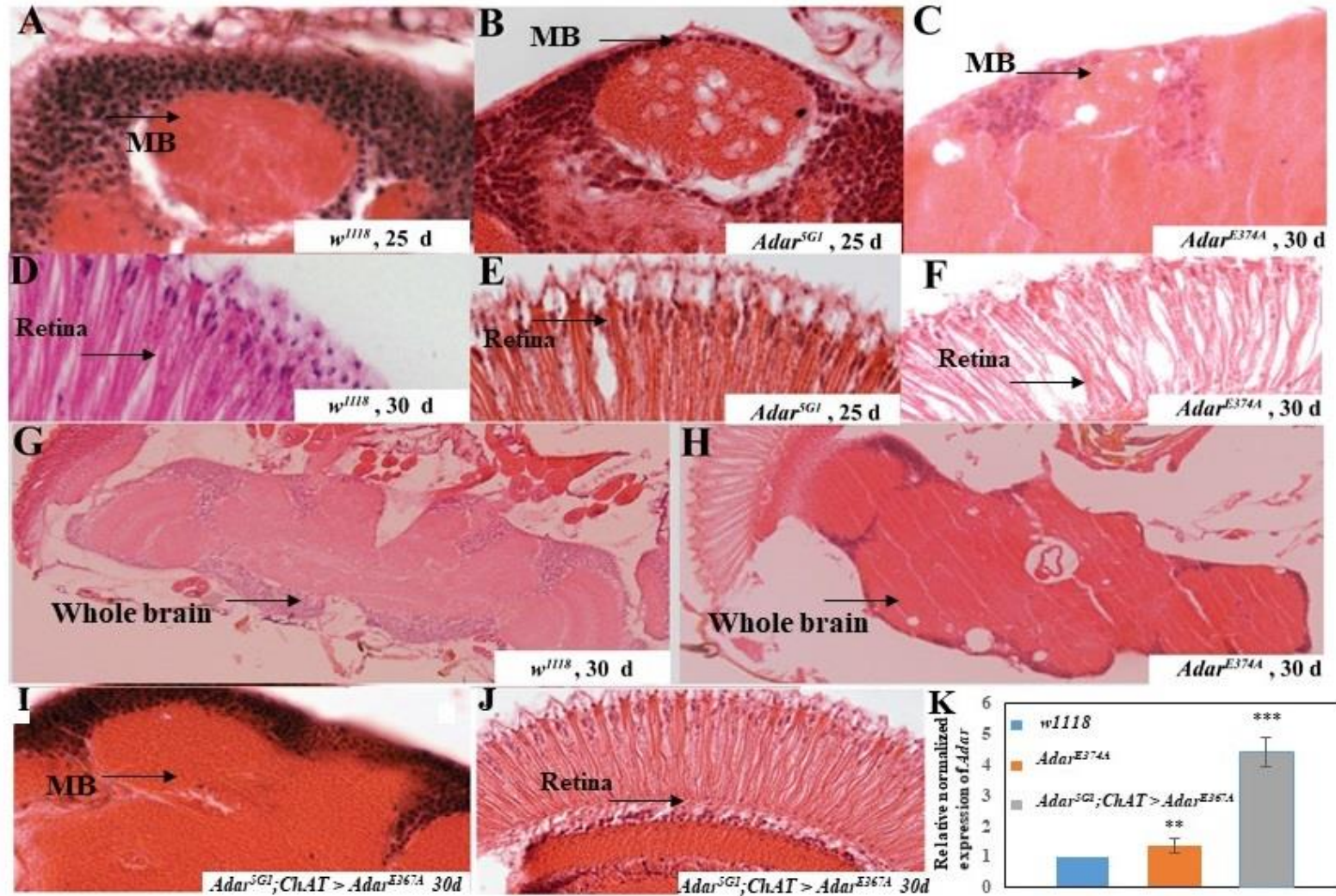
High RNA editing in neurotransmitter receptor subunits.

Name	Protein function	Number of sites	Functional consequence*	% editing at each site
<i>Voltage-gated ion channels</i>				
<i>DSCI</i>	Na ⁺ channel	1	+++	• 50%
Ca-alpha1T	Ca ²⁺ channel	1	+	• 30%
<i>DmCa1D</i>	Ca ²⁺ channel	5	++	• 30, 100, 95, 95, 100%
$\alpha_2\delta$	Ca ²⁺ channel	3	+	• 80, 50, 50%
	accessory subunit			
<i>Shaker (Sh)</i>	K ⁺ channel	6	+++	• 10, 10, 50, 50, 80, 50%
<i>ether-a-go-go (eag)</i>	K ⁺ channel	6	+++	• 50, 100, 20, 90, 10, 80%
<i>slowpoke (slo)</i>	K ⁺ channel	2	+	• 90, 90%
<i>Synaptic release machinery</i>				
<i>Synaptotagmin (syt)</i>	Ca ²⁺ sensor	4	+++	• 5, 50, 50, 100%
<i>Dunc-13</i>	SNARE binding	1	++	• 40%
<i>Stoned B (stnB)</i>	?	1	+	• 90%
<i>complexin (cpx)</i>	SNARE protein	3	?	• 50, 20, 30%
<i>lap</i>	Adaptor protein	1	?	• 10%
<i>Ligand-gated ion channels</i>				
<i>Dα5</i>	nAChR α subunit	7	+++	• 80, 80, 100, 50, 30, 60%
<i>ARD</i>	nAChR β subunit	4	?	• 50, 90, 50, 100%
<i>SBD</i>	nAChR β subunit	2	+++	• 30, 20%
<i>Resistance to dieldrin (Rdl)</i>	GABA-receptor	6	+++	• 15, 80, 90, 90, 20, 10%

GAL4-UAS system



Neurodegeneration is less severe in inactive *Adar* mutant and can be rescued by overexpression of inactive *Adar* protein



Conclusions

- *ADARs are important dsRNA binding protein so the expression level of the protein is critical.*
- *ADARs role in innate immune response is maintained despite the evolution of very different pathways.*



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Acknowledgements

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Grateful to the patients and their parents.



Thank you for your attention