

# Mutational signatures of redox stress in yeast and men

Natasha Degtyareva

Mutagenesis & DNA Repair Regulation Group, NIEHS

DNA repair interest group videoconference

May 18, 2021

# Redox stress is linked to human disease and ageing

## A hallmark of cancer



Luo, Solimini, Elledge, *Cell*, 2009

## Neurodegenerative diseases

- Alzheimer's, Parkinson's, Huntington's, Amyotrophic lateral sclerosis (ALS)
- Associated with cell loss/degeneration in high energy consuming cells



Lin and Beal, *Nature*, 2006

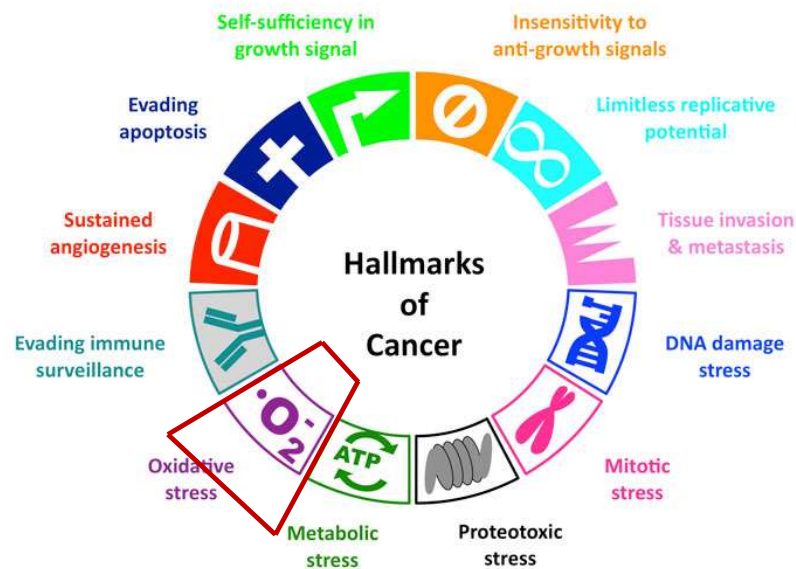
## Ageing

- Redox theory of ageing



Go and Jones, *Clin Sci*, 2017

# Redox stress is a hallmark of cancer



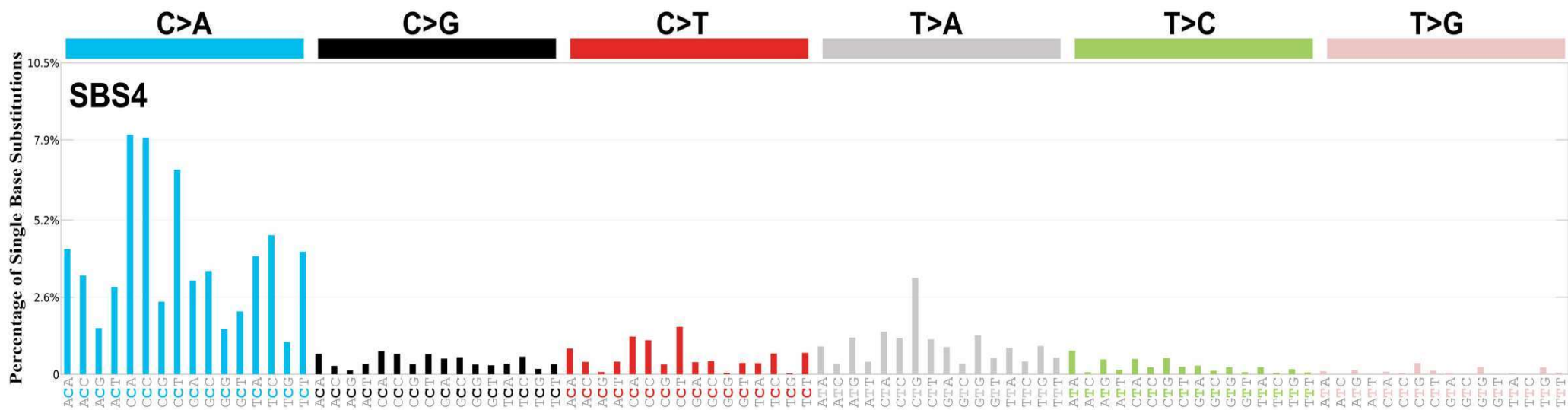
Redox status is deregulated in majority of cancers due to:

- altered metabolism;
- mitochondrial dysfunction;
- inflammation;
- malfunctioning of peroxisomes;
- over- or under-expression of ROS-producing and ROS-scavenging enzymes.

# How to assess the contribution of redox stress to disease and aging?

By analyzing the mutations accumulated following exposure to oxidizing agents  
and in tissues affected by a disease

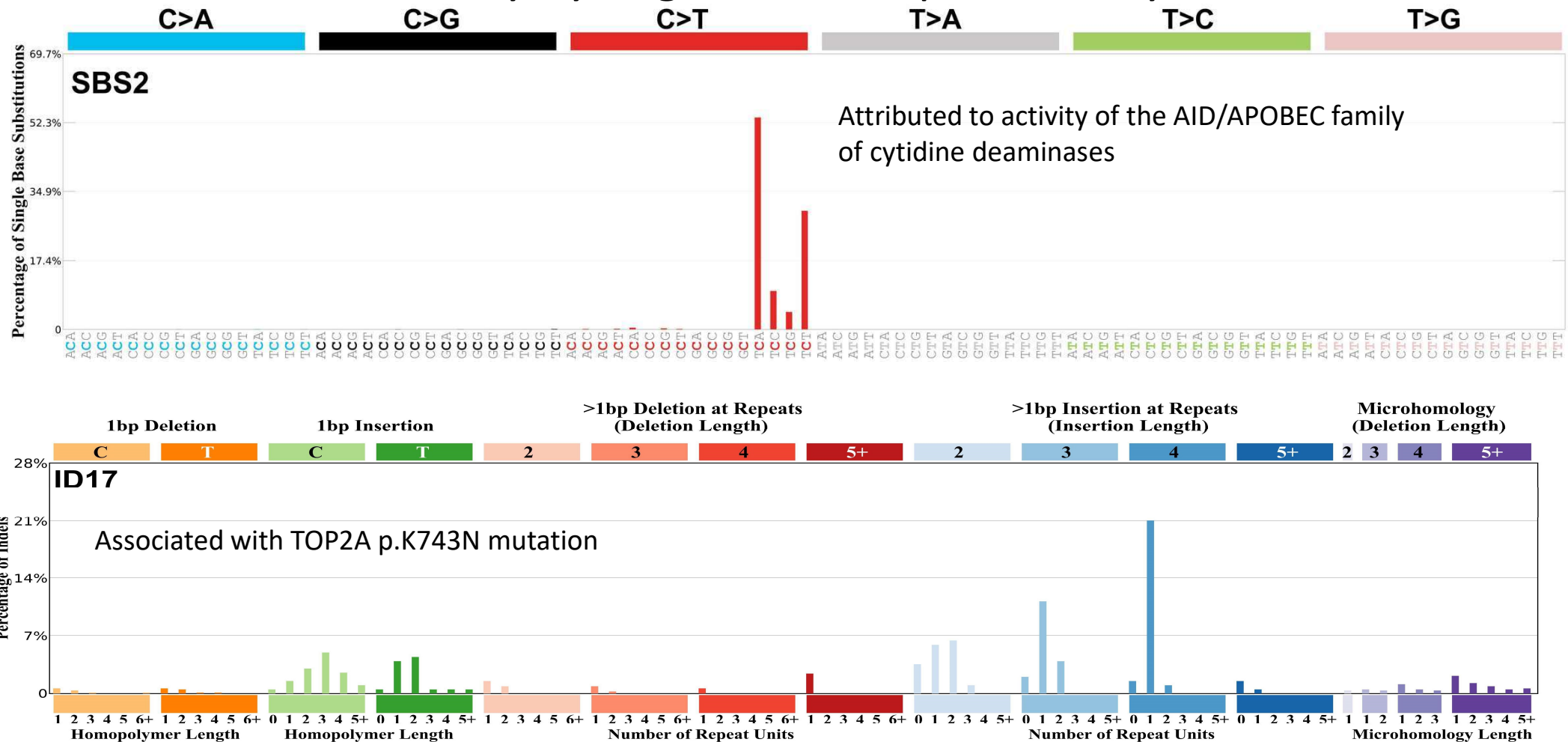
Mutational signature is a spectrum of mutations  
in the context of adjacent nucleotides



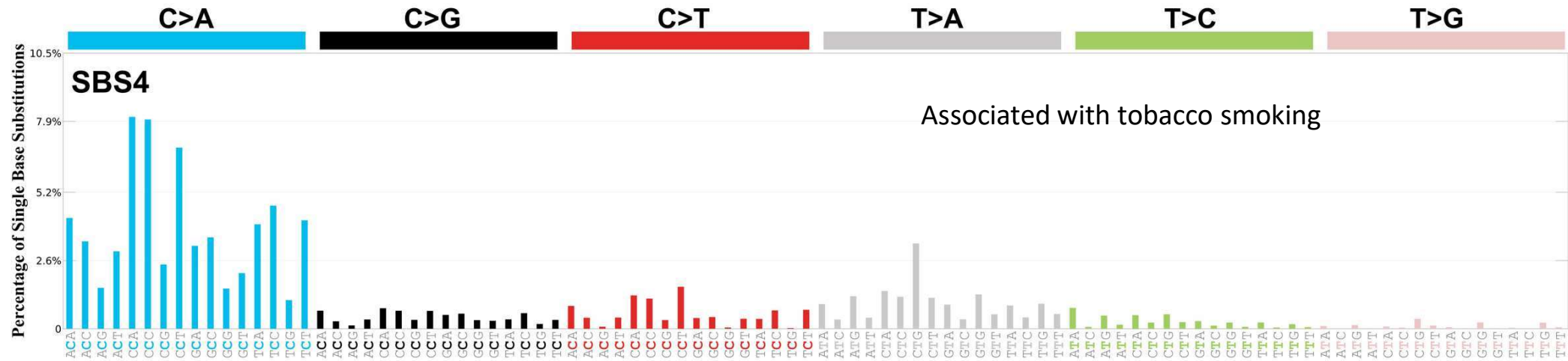


<https://cancer.sanger.ac.uk/signatures/sbs/>

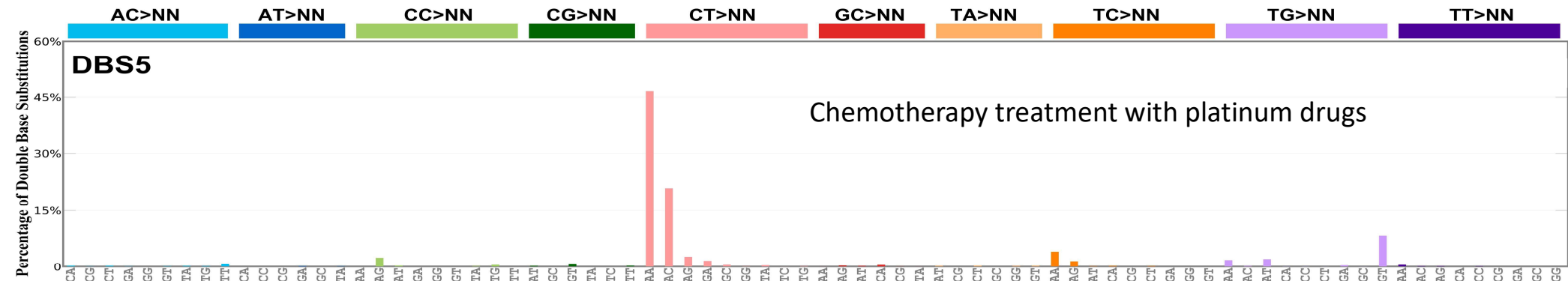
# Some of the mutational signatures reveal etiology of human cancers caused by dysregulation of specific enzymes



... or by environmental exposures

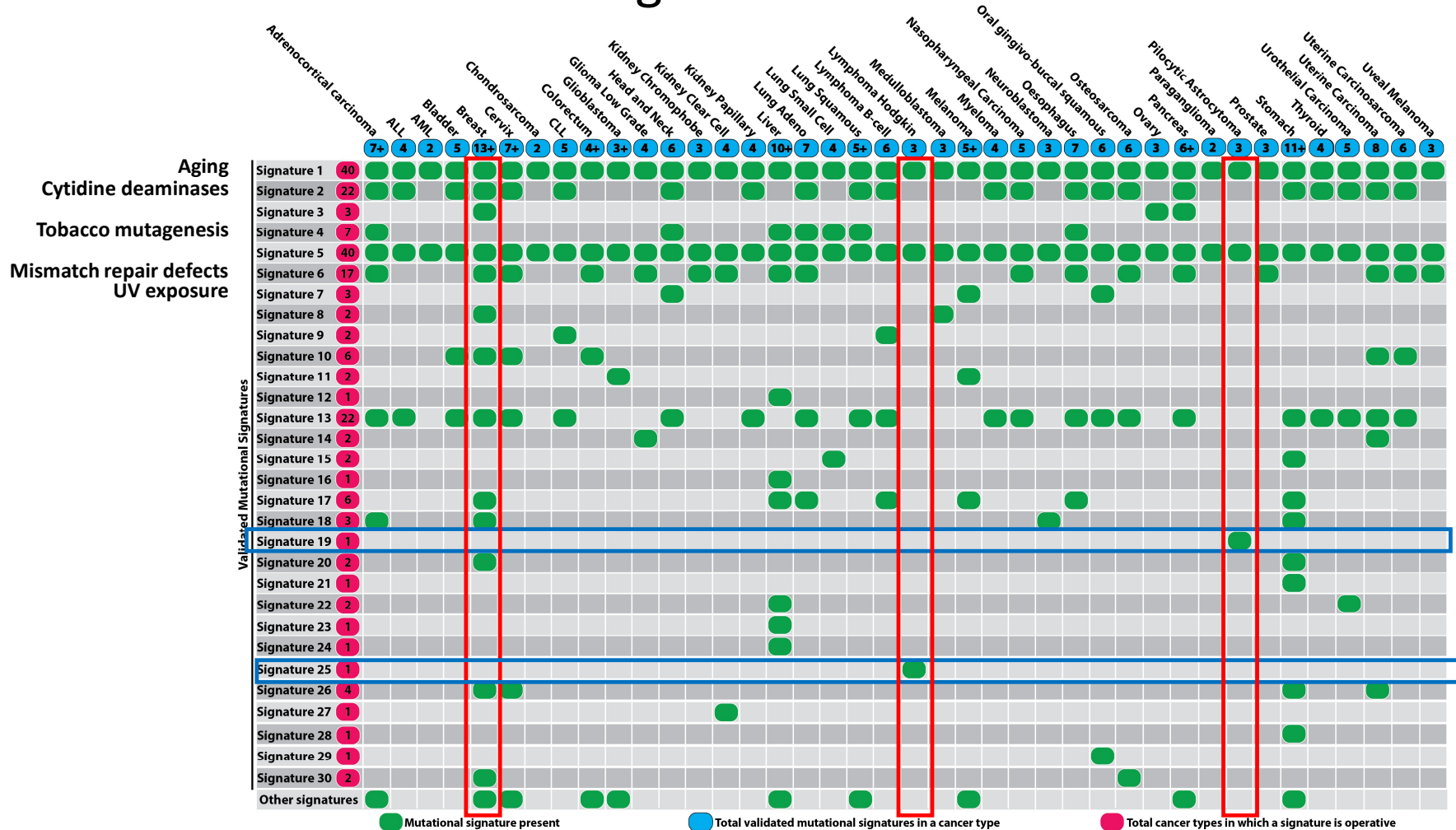


... or by chemotherapeutic interventions





# Is there a signature of redox stress?



Alexandrov et al., Nature, 2013

## **Approach:**

discerning the mutational signature of redox stress

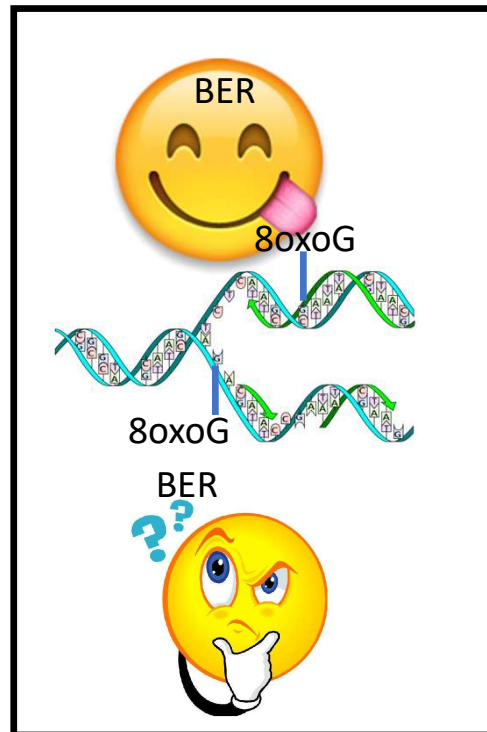
-in the model organism *Saccharomyces cerevisiae*;

-in single strand DNA;

- selecting clustered (closely-spaced) mutations.

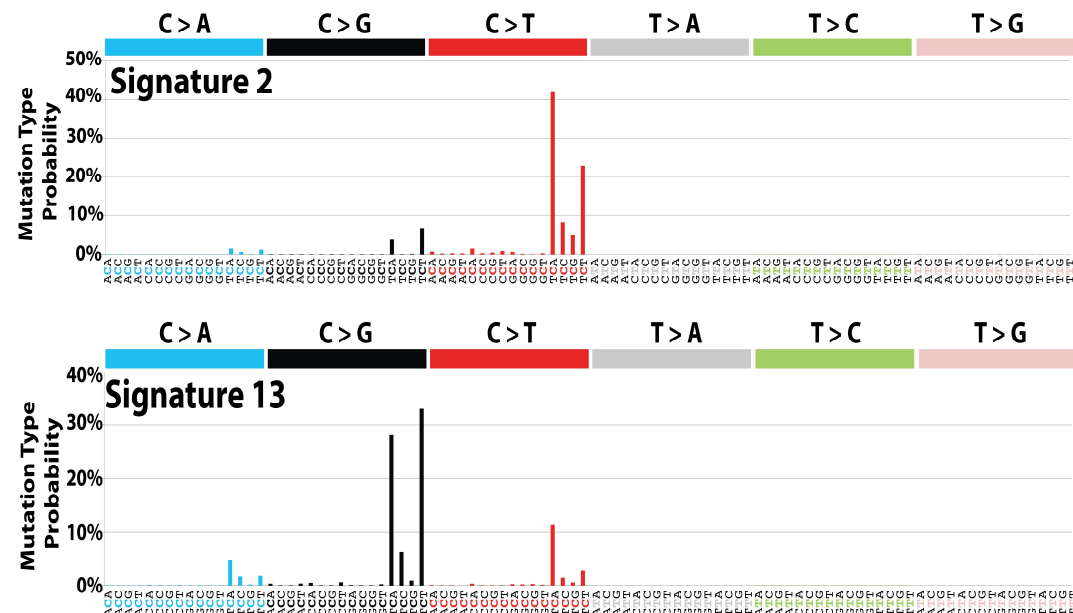
# Why did we look for a signature of oxidative damage in single strand DNA?

Base Excision Repair  
requires second  
DNA strand \*



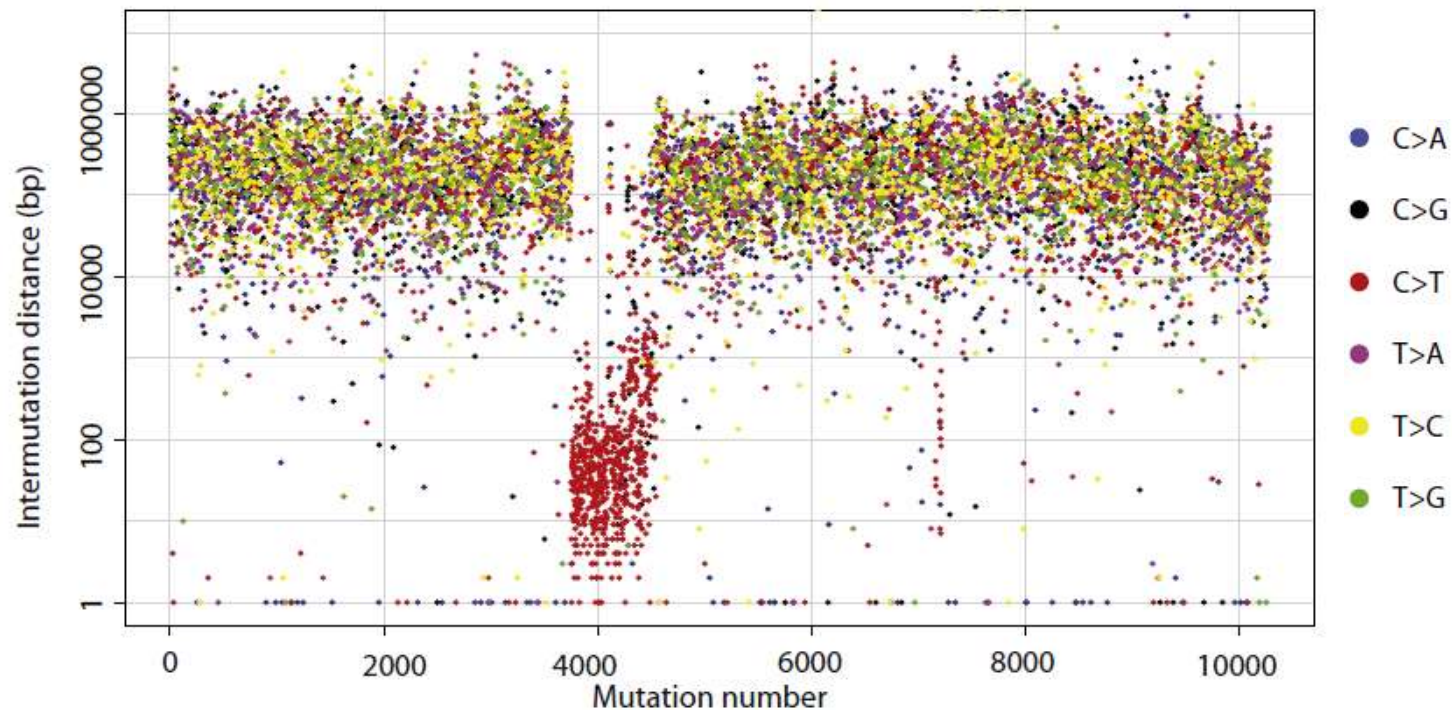
\* with some exceptions

Among published cancer signatures  
some are attributed to activity of enzymes that uses ssDNA  
as a substrate



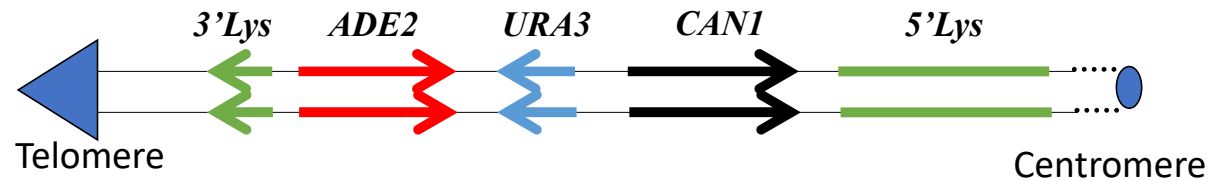
# Why did we look for a signature of oxidative damage in single strand DNA?

Localized hypermutability suggests persistent presence of ssDNA in cancer genomes

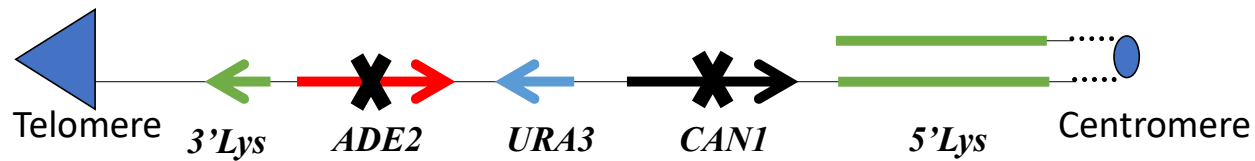


*Nik-Zainal et al., 2012*

The reporter system allows for the generation long stretches of ssDNA and selection for multiple mutations

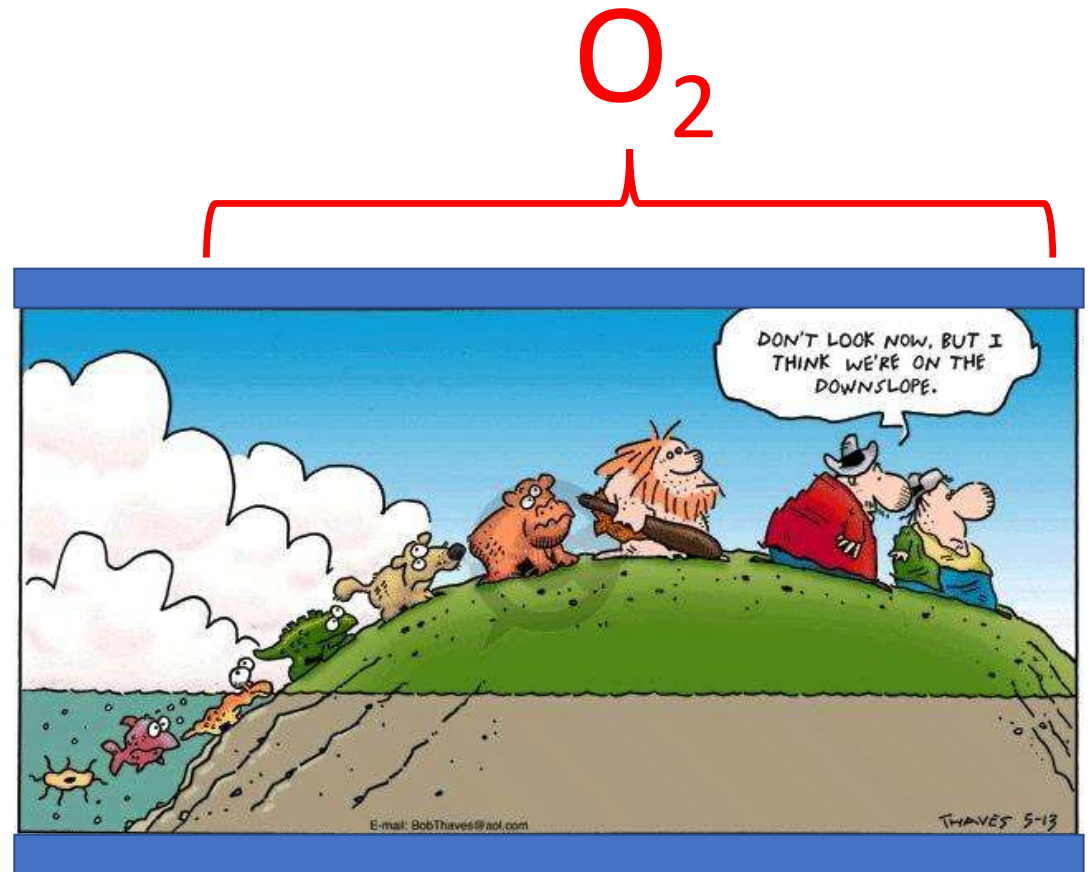
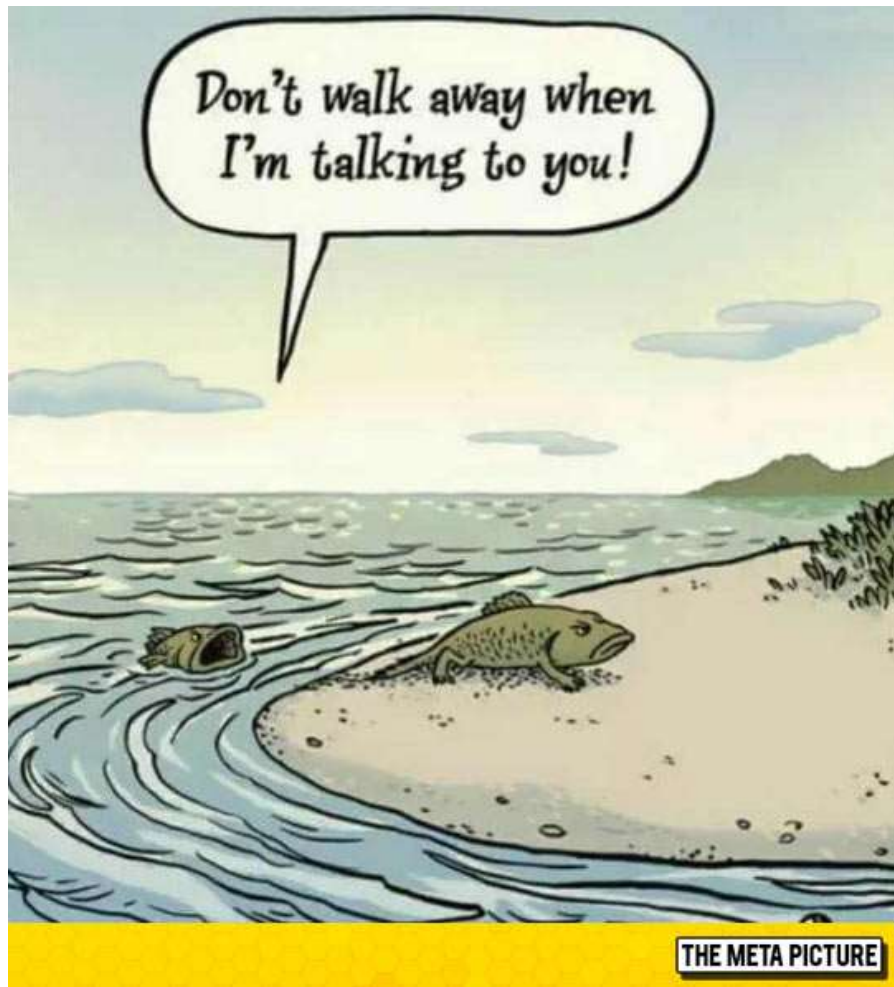


*cdc13-1ts* at 37°C



RED Can<sup>R</sup> clones = clustered mutations

Chan et al., 2012



# A Compendium of Mutational Signatures of Environmental Agents

Jill E. Kucab, Xueqing Zou, Sandro Morganella, Madeleine Joel, A. Scott Nanda, Eszter Nagy, Celine Gomez, Andrea Degasperi, Rebecca Harris, Stephen P. Jackson, Volker M. Arlt, David H. Phillips, Serena Nik-Zainal  
*Cell, 2019*

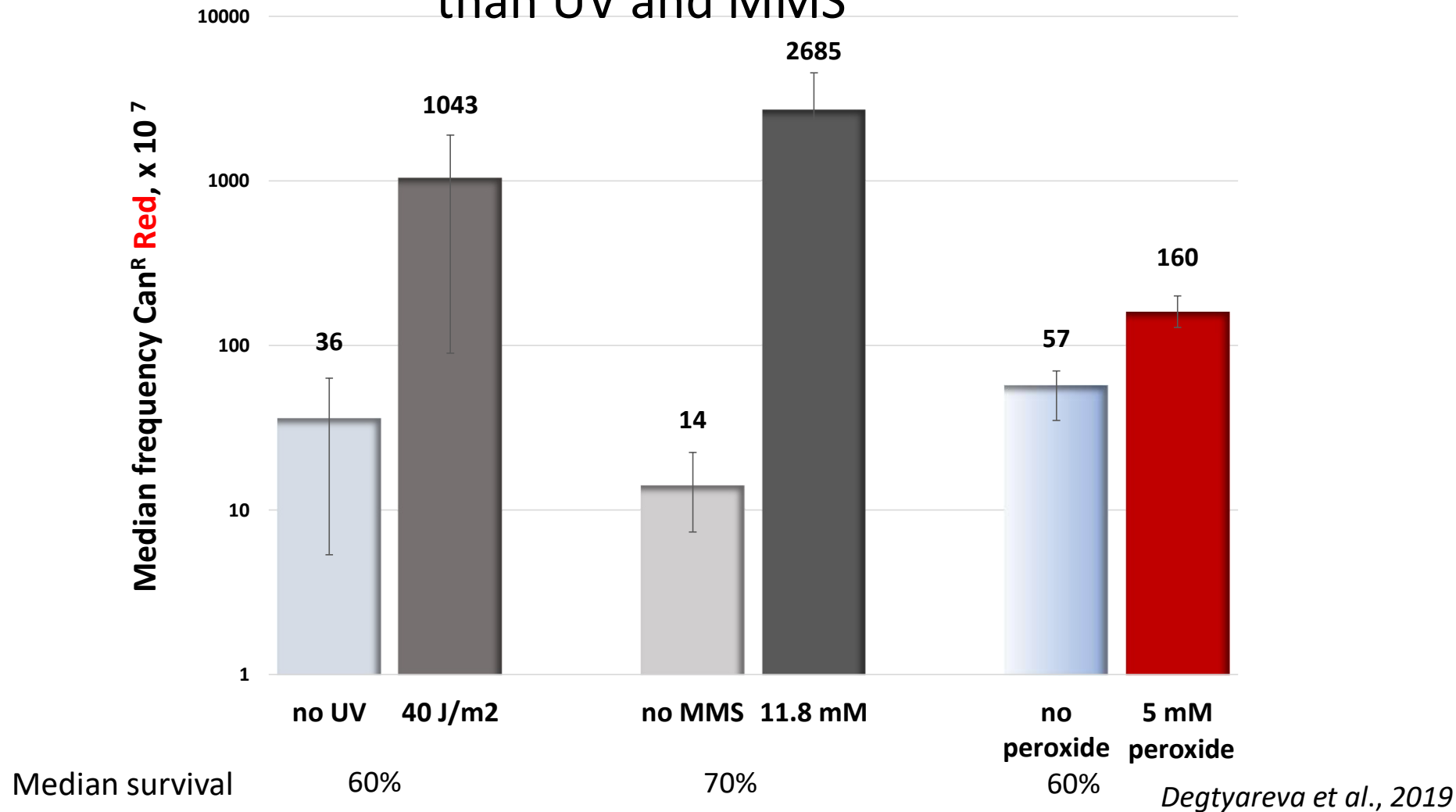
## Highlights

- 41 of 79 environmental agents yielded substitution signatures
- 6 agents produced double-substitution signatures and 8 produced indel signatures
  - Several signatures match or exhibit similarity with signatures found in human tumors
  - Topographical mutational asymmetries reveal mechanistic insights

“...hydrogen peroxide, anticipated to create ROS, and peroxynitrite, which generates reactive nitrogen (nitric oxide) species, did not yield clear mutation patterns”...



At an equitoxic dose hydrogen peroxide induces fewer mutations, than UV and MMS





Even if it is possible to discern a mutational signature of oxidative stress  
what to expect?

8oxoG  
mis-pairing with  
A

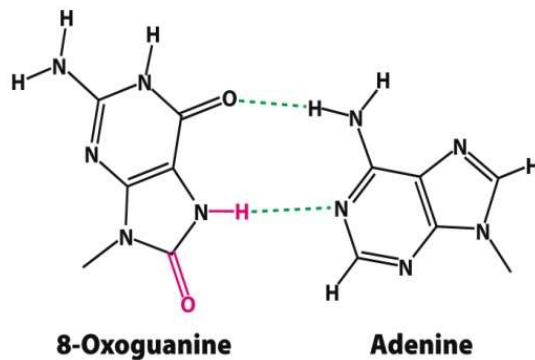


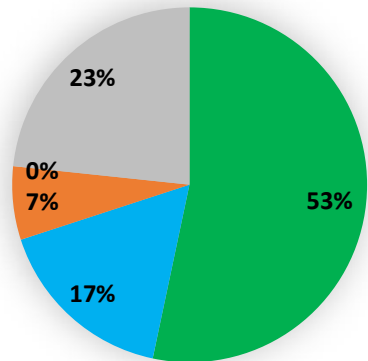
Figure 28.30  
Biochemistry, Seventh Edition  
© 2012 W. H. Freeman and Company

leads to  
**G** to **T**  
transversions

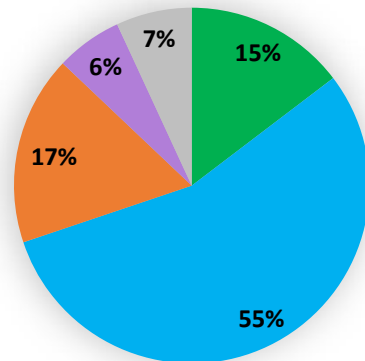


# Oxidative stress-induced mutagenesis in ssDNA occurs primarily at C

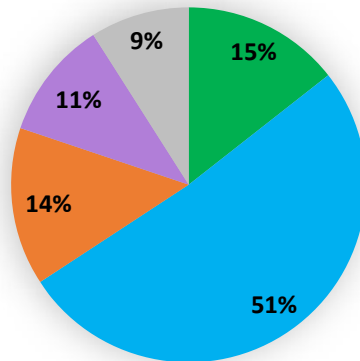
Wt spontaneous



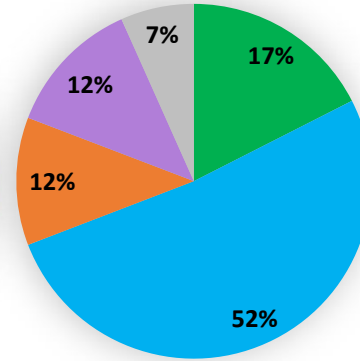
Hydrogen peroxide -induced



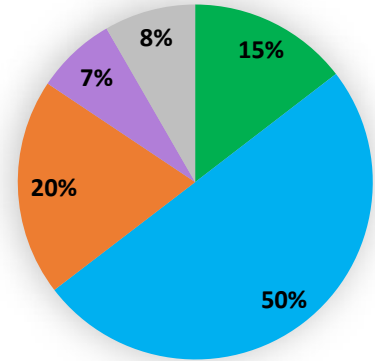
wt



*ogg1*



*rtt109*



*gcn5*

C

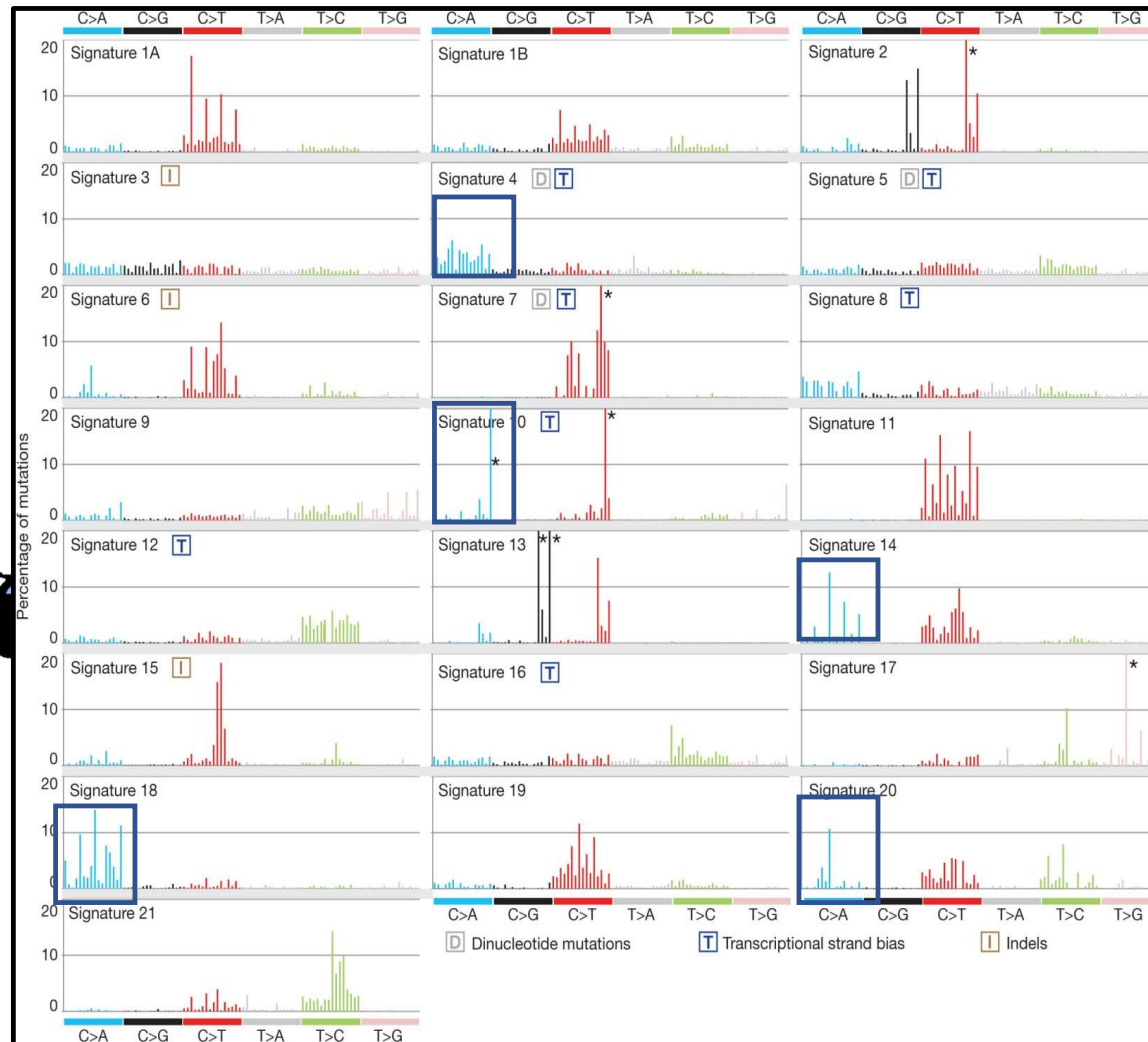
G

A

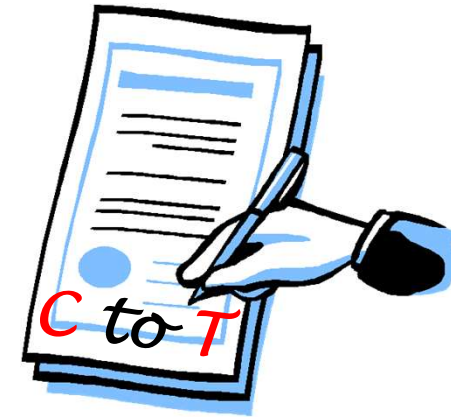
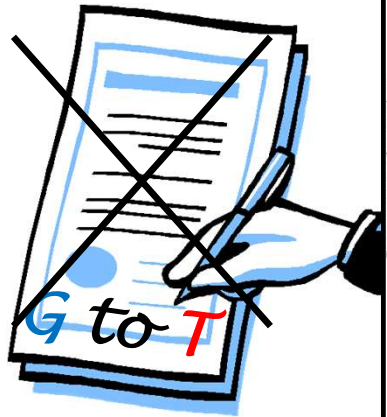
T

other

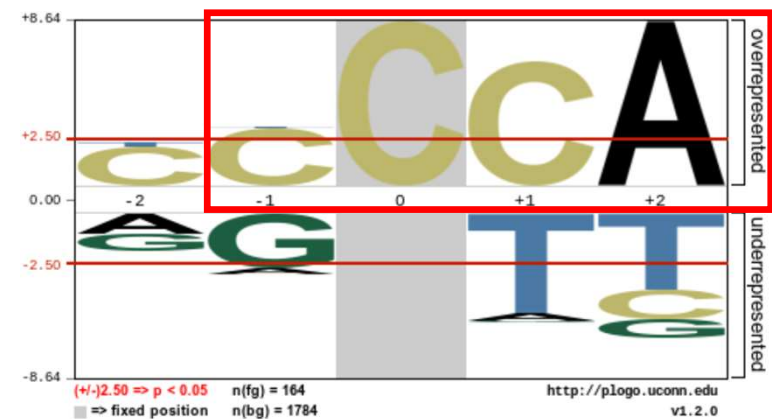
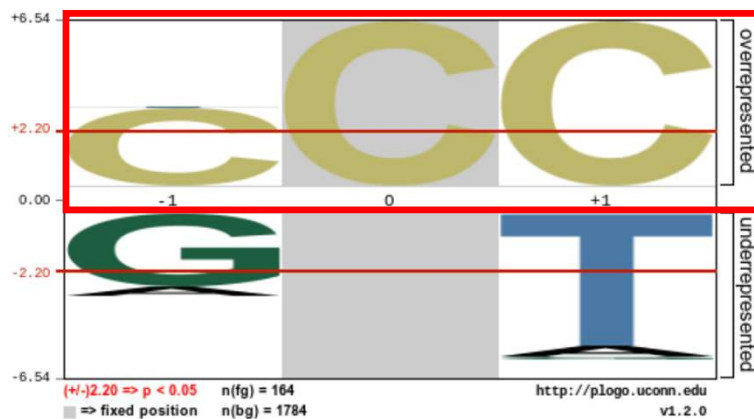
Expected: G to T enrichment in oxidative stress signature



# C to T enrichment in oxidative stress signature?



# Signature of hydrogen peroxide - induced oxidative stress



<https://plogo.uconn.edu/>

Significant enrichment in C at position +1 and -1

Significant enrichment in A at position +2

## Confirmed signature of hydrogen peroxide - induced oxidative stress

$$(E)nrichment = \frac{Mutations_{(cCc \rightarrow cTc)} \times Context_{(c)}}{Mutations_{(C \rightarrow T)} \times Context_{(ccc)}}$$

- Produces sample-specific P-values
- Not affected by “topography” preferences

*Roberts et al., 2012*

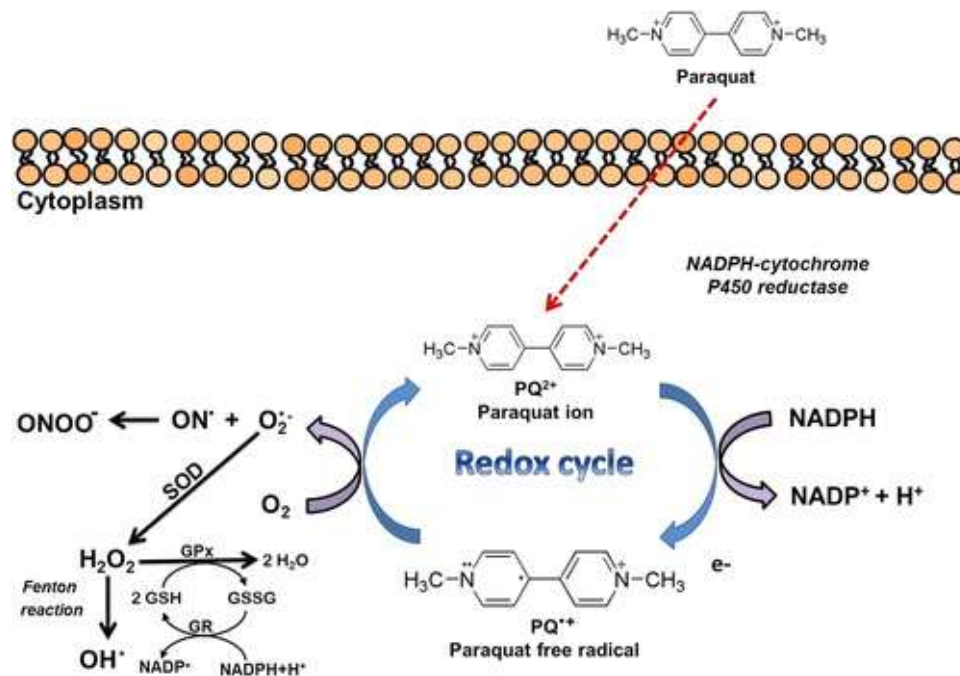
Motif	Fold enrichment	Mutational load	Bonferroni-corrected Fisher P value
Cca to Tca	3.00	21	1.95E-08
cCca to cTca	3.40	4	1.04E-02
Cca toTna	1.49	19	2.43E-04

Is this signature hydrogen peroxide-specific?

## Paraquat (PQ)



- Widely used herbicide
- Toxic to humans and animals
- Exposure has been linked to Parkinson's Disease



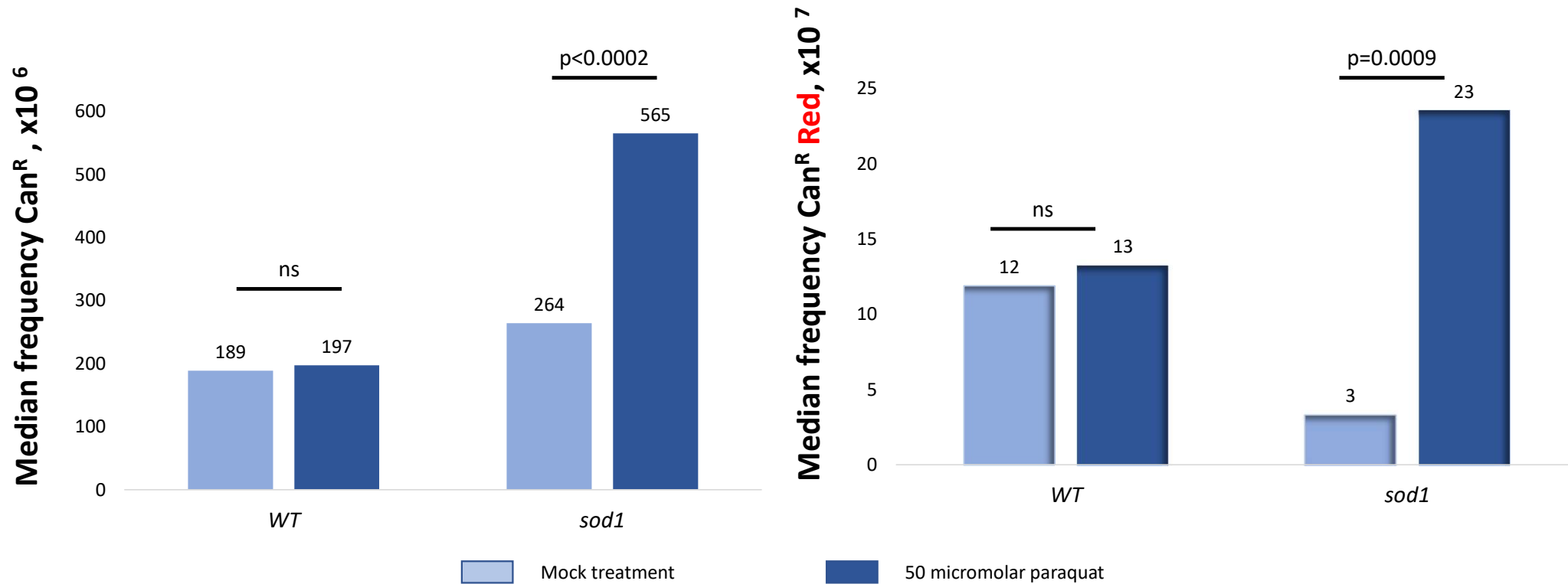
<https://emergency.cdc.gov/agent/paraquat/basics/facts.asp>

*Blanco-Ayala et. al., Free Radical Research, 2014*

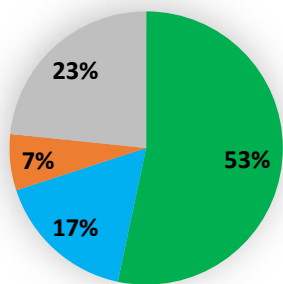
Is there a signature of paraquat – induced mutagenesis?



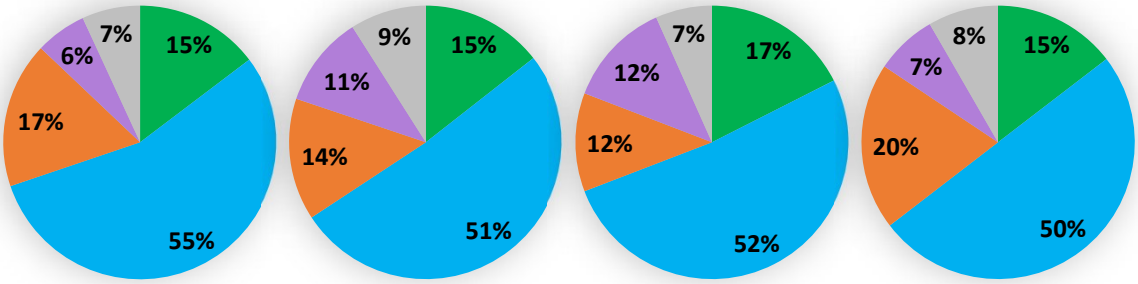
# Exposure to paraquat increases mutation frequencies in *sod1* mutants



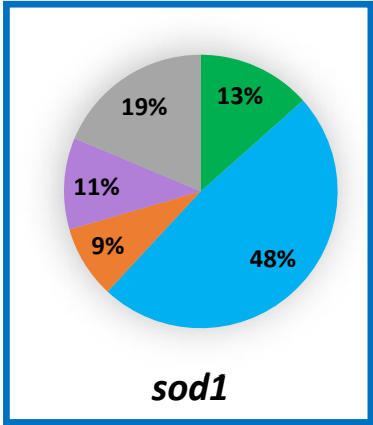
WT spontaneous



Hydrogen peroxide -induced



Paraquat -induced



WT

*ogg1*

*rtt109*

*gcn5*

*sod1*

C

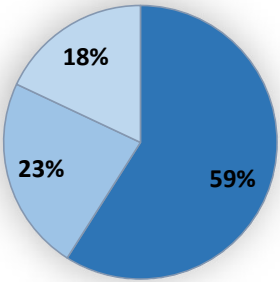
G

A

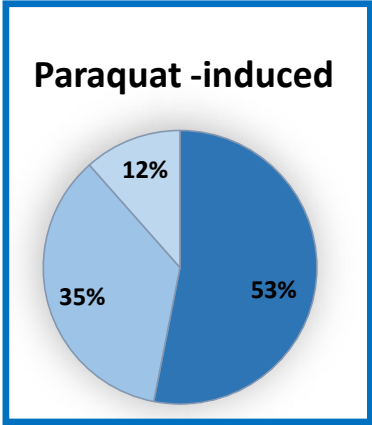
T

other

Hydrogen peroxide -induced



Paraquat -induced



C-T

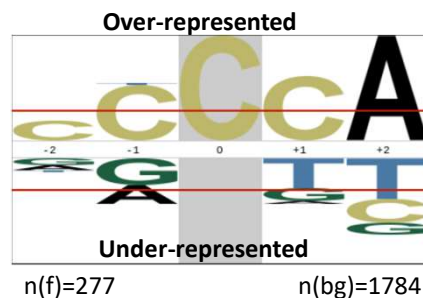
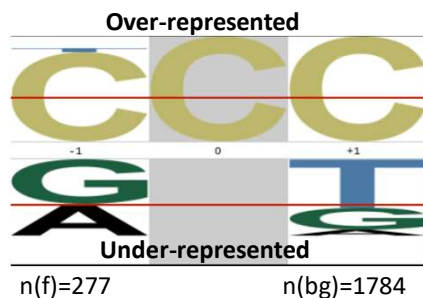
C-G

C-A

# Mutational signatures of H<sub>2</sub>O<sub>2</sub> and paraquat in ssDNA are similar

Hydrogen peroxide

*WT*

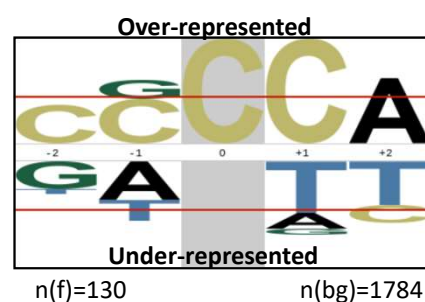
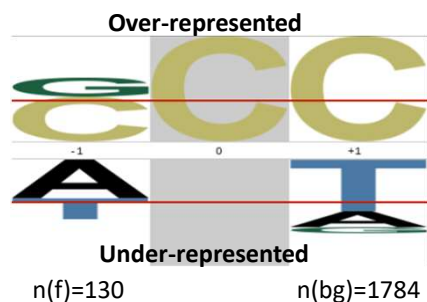


H<sub>2</sub>O<sub>2</sub>

cCca to cTca

Paraquat

*sod1*



O<sub>2</sub><sup>•-</sup>

Cca to Tca

Two different oxidizing agents, hydrogen peroxide and paraquat, generate similar mutational signatures in ssDNA in yeast

From yeast to men:

Is oxidative stress a major contributor to aging-related somatic mutations ?

Where to look for redox stress – related signatures in human DNA?



Mitochondria accumulate somatic mutations with age,  
but they are not oxidative stress-induced (?)

OPEN  ACCESS Freely available online

 **PLOS** | GENETICS

## Ultra-Sensitive Sequencing Reveals an Age-Related Increase in Somatic Mitochondrial Mutations That Are Inconsistent with Oxidative Damage

Scott R. Kennedy<sup>1</sup>, Jesse J. Salk<sup>1,2</sup>, Michael W. Schmitt<sup>1,2</sup>, Lawrence A. Loeb<sup>1,3\*</sup>

PLOS Genetics | www.plosgenetics.org

1

September 2013 | Volume 9 | Issue 9 | e1003794

OPEN  ACCESS Freely available online

 **PLOS** | GENETICS

## Oxidative Stress Is Not a Major Contributor to Somatic Mitochondrial DNA Mutations

Leslie S. Itsara<sup>1,2</sup>, Scott R. Kennedy<sup>3</sup>, Edward J. Fox<sup>3</sup>, Selina Yu<sup>1</sup>, Joshua J. Hewitt<sup>1,4</sup>,

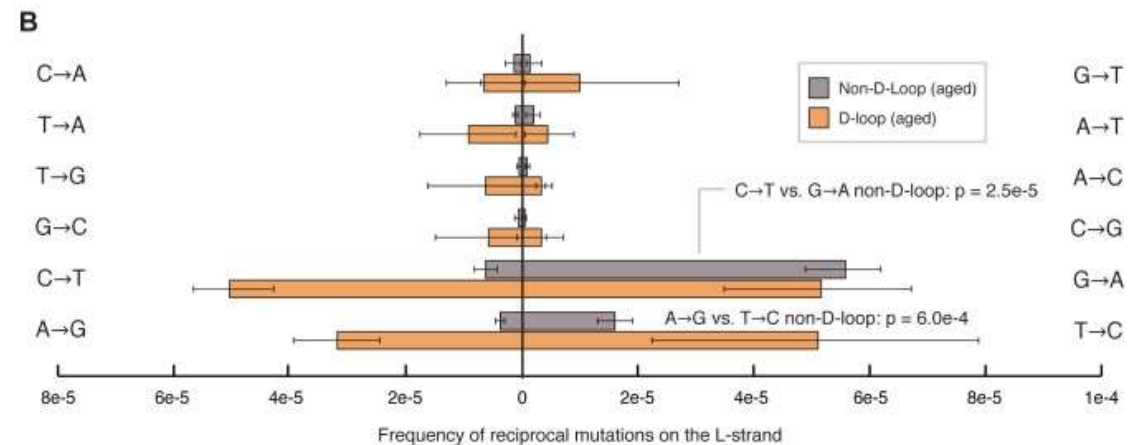
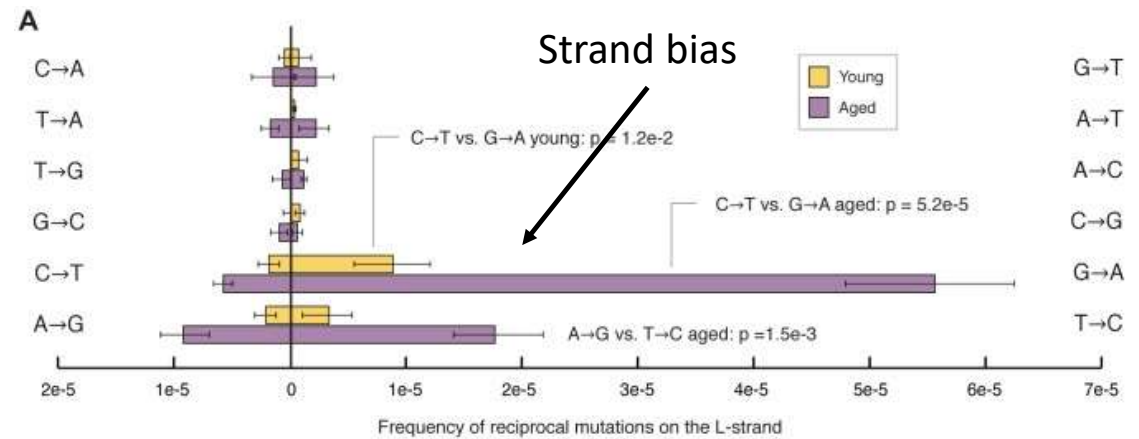
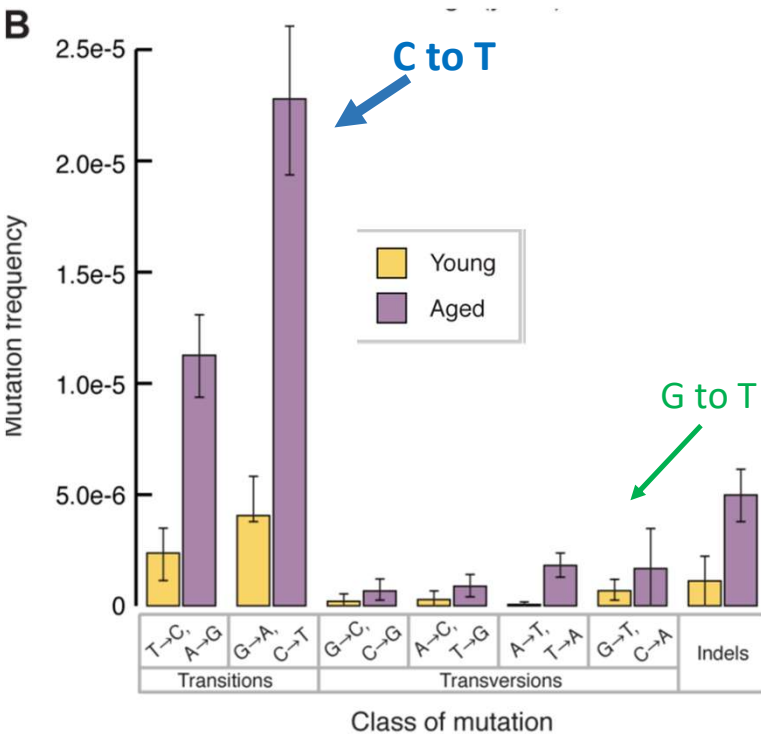
Monica Sanchez-Contreras<sup>5</sup>, Fernando Cardozo-Pelaez<sup>6</sup>, Leo J. Pallanck<sup>1\*</sup>

PLOS Genetics | www.plosgenetics.org

1

February 2014 | Volume 10 | Issue 2 | e1003974

# The major type of somatic mutations in aging mitochondria is C to T changes



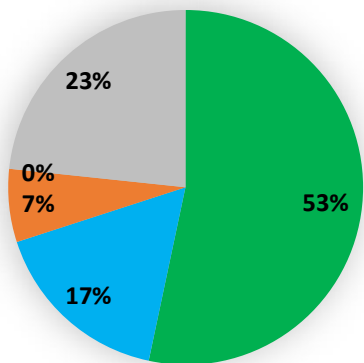
“Surprisingly, comparison of the mutation spectra of the young and old samples reveals a **notable absence of the mutational signature of oxidative damage**... “We failed to find either a preponderance of **GtoT/CtoA** substitutions or a proportionally greater increase with age in this type of mutation relative to other types, despite a span of 80 years between our sequenced sample groups”

*Kennedy et al., 2013*

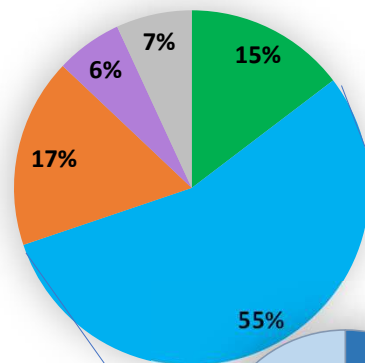
# Mutational spectra of redox stress in yeast ssDNA and of aging in human mitochondrial DNA share a common feature

Yeast

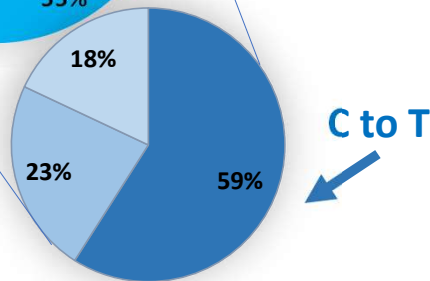
Wt spontaneous



Hydrogen peroxide -induced

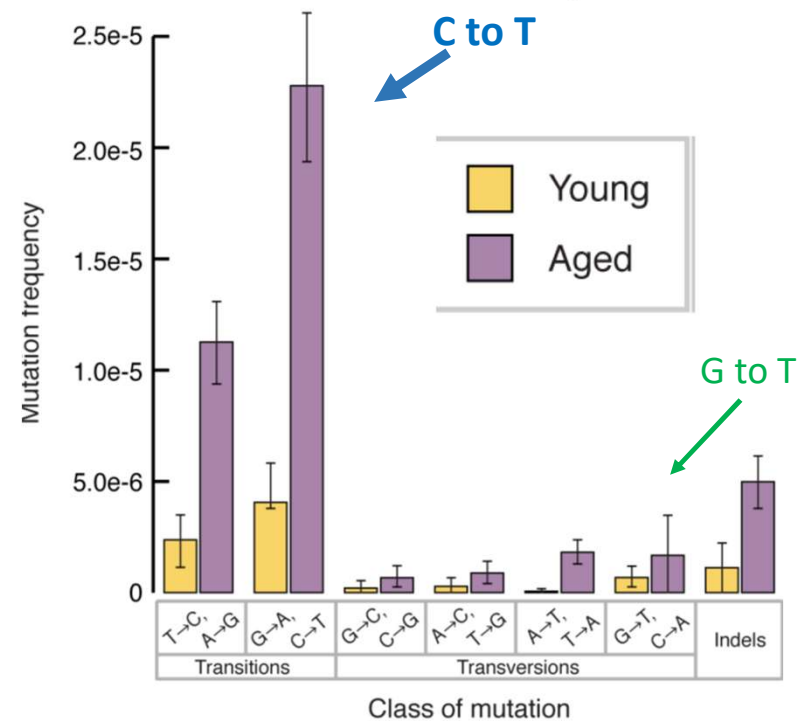


C G A T other



C to T

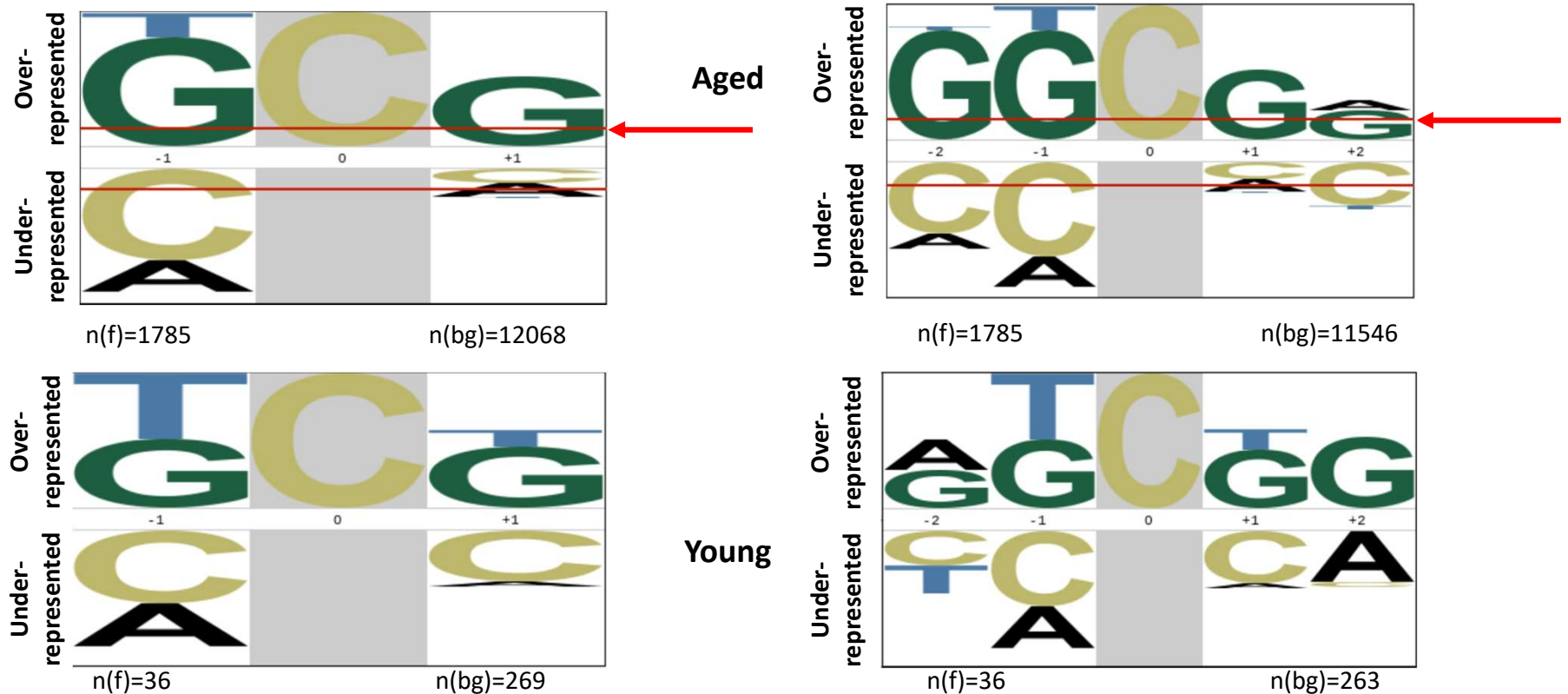
Men



Kennedy et al., 2013

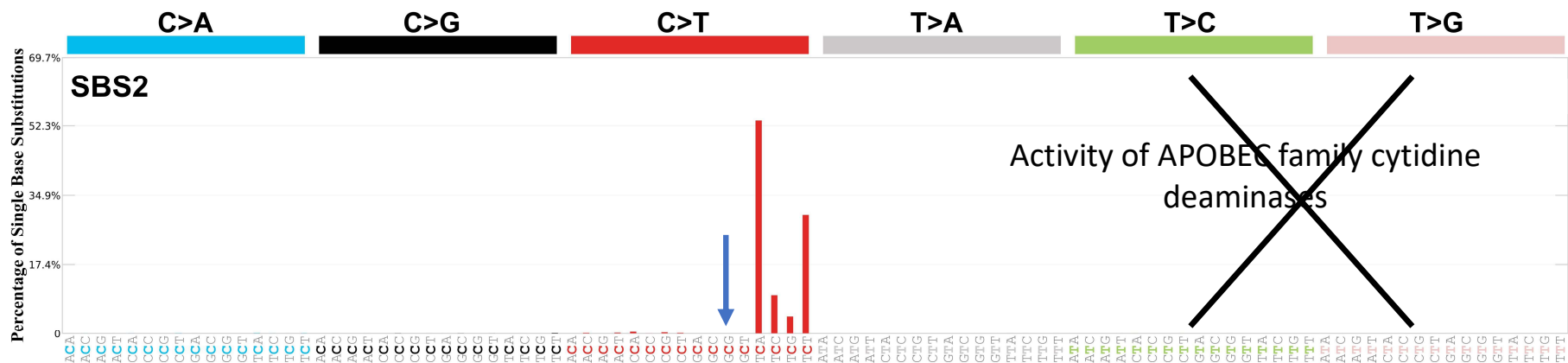
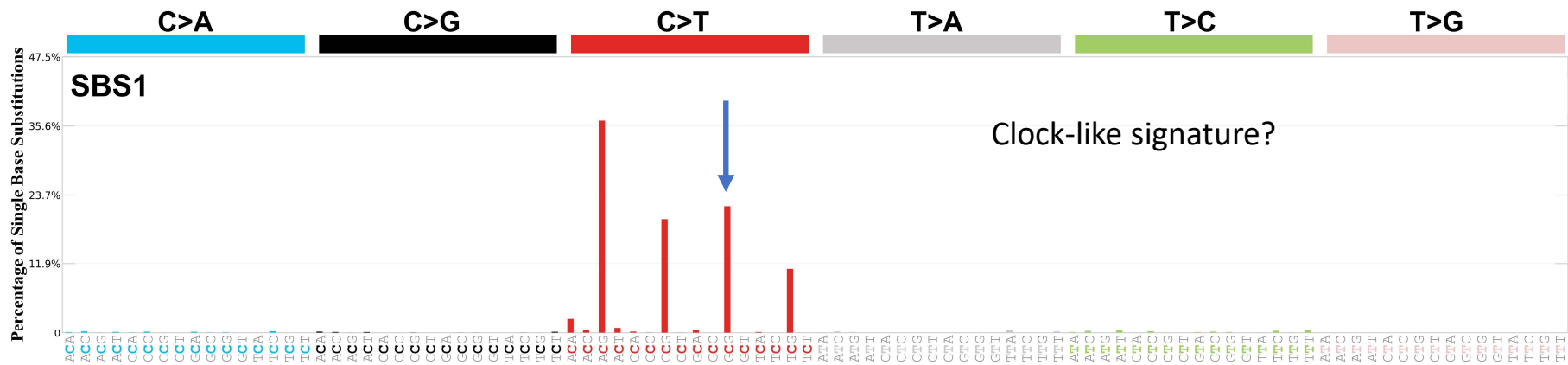


# Is there a distinct signature of aging in mtDNA?

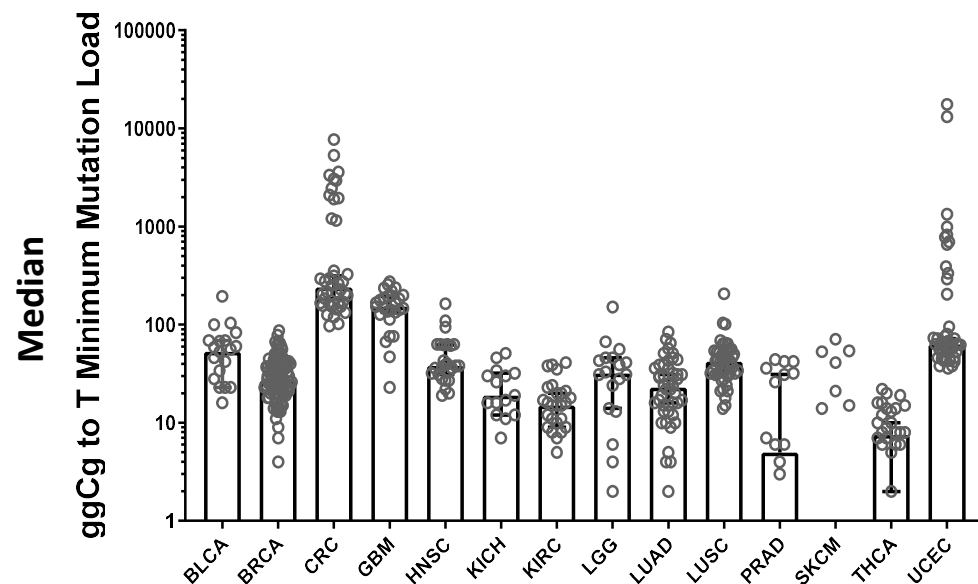
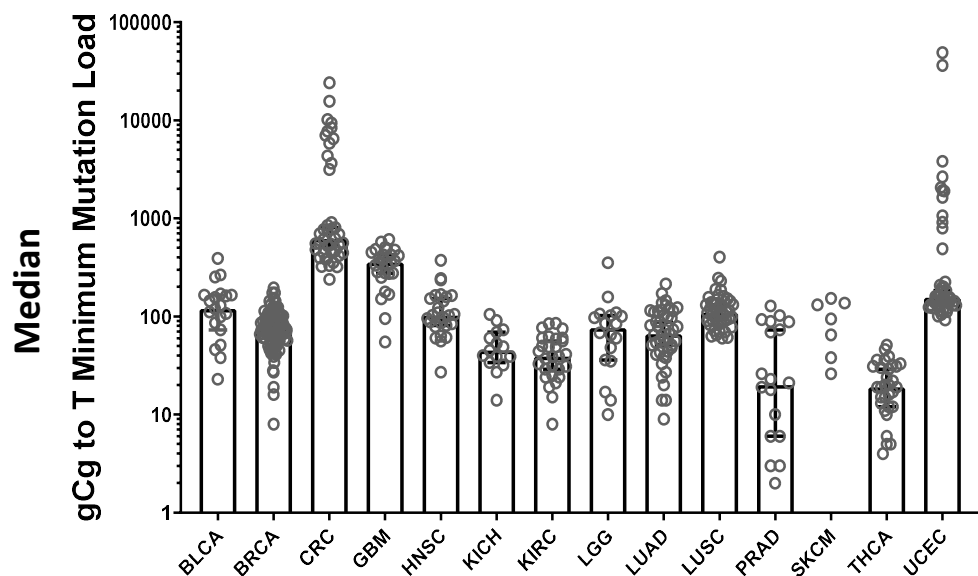


Motif	Fold enrichment	Mutational load	Bonferroni-corrected Fisher P value
gCg to gTg	2.44	25	3.26E-06
ggC to ggT	2.04	36	2.89E-07
ggCg to ggTg	3.10	15	6.50E-05

ggCg to ggTg signature of mitochondria aging:  
(methyl)cytosine deamination?



# Evidence for redox-stress related signature in human cancers



# Conclusions

- The majority of hydrogen peroxide- and paraquat-induced mutations in ssDNA occurs at C;
- C to T is the main type of redox stress-induced substitutions in ssDNA in yeast;
- 8oxoG is not the major redox stress-induced mutagenic lesion in ssDNA;
- The mutational signatures of hydrogen peroxide and paraquat in ssDNA in yeast are similar;
- Mutational signatures of redox stress in yeast ssDNA and of aging in human mitochondrial DNA share a common feature
- Many cancer genomes are enriched for redox stress-related signatures.

# Questions

- What is the underlying DNA lesion at C?
- What are the molecular mechanisms of protection of ssDNA from oxidative damage?
- Do antioxidants prevent oxidative damage in ssDNA?
- Is it possible to find mutational signatures of oxidative stress in cancers by utilizing the new, broadened databases?
- Is there a mutational signature of redox stress in cells exposed to chronic inflammation?
- Are there other mutational signatures attributable to oxidative stress?

# Acknowledgements

## NIEHS

Paul Doetsch

Dmitry Gordenin

Les Klimczak

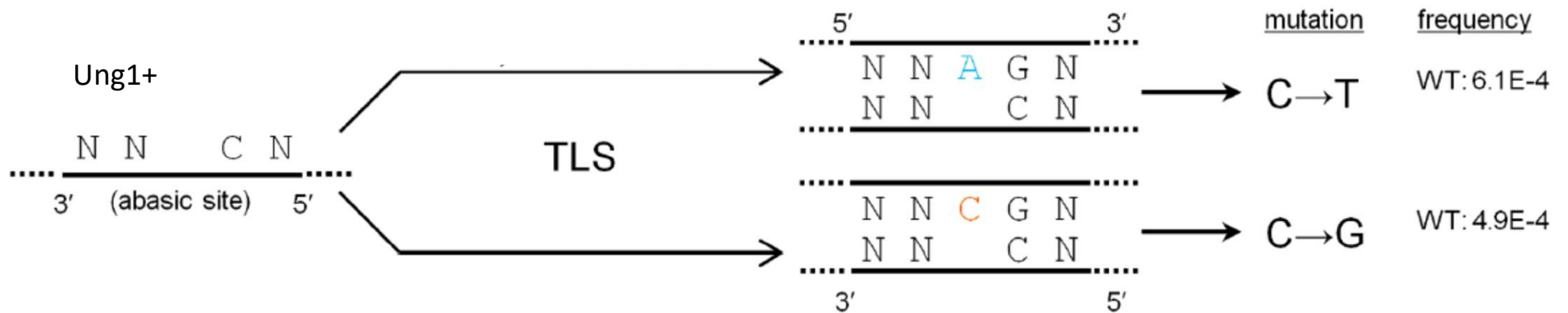
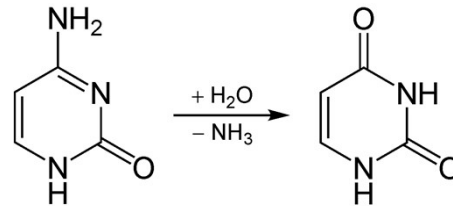
Joan Sterling

Tori Placentra

Natalie Saini (MUSC)

Scott Kennedy (University of Washington)

# Cytosine deamination?



Frequency C to T = Frequency C to G

*Chen et al., 2013*