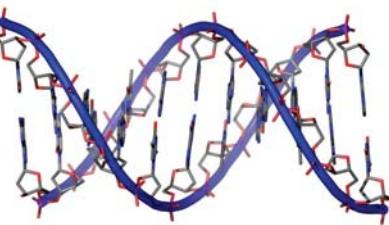


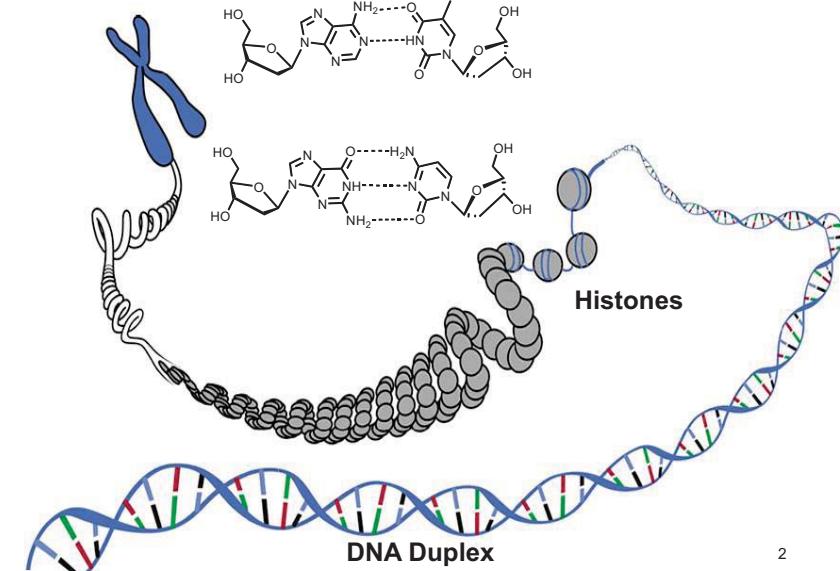
## Ischia 2014, Summerschool



Thomas Carell  
Department für Chemie  
Ludwig-Maximilians Universität (LMU)  
München  
  
How Chemistry  
defines Stem Cell Development  
[www.carellgroup.de](http://www.carellgroup.de)

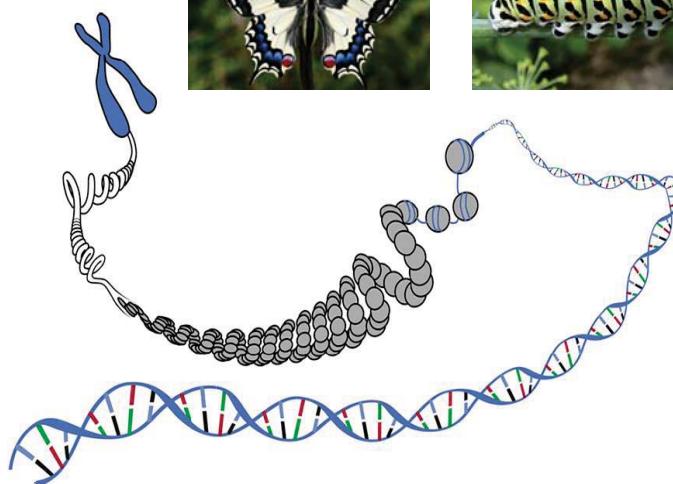
1

## The complexity of DNA

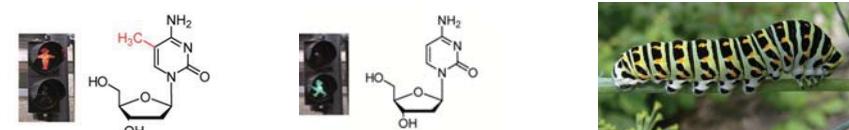


2

## Epigenetic control of transcription



3



**GC TTAGTTAGCCGATGCAAATTGCCCTGTGCACTGCTGTCAGCAATTGCTACTACGCATGGATTGCTATCG**

Gene 2  
↓  
Protein 2

Gene 3  
↓  
Protein 3



**on** **off** **on** **off** **on**

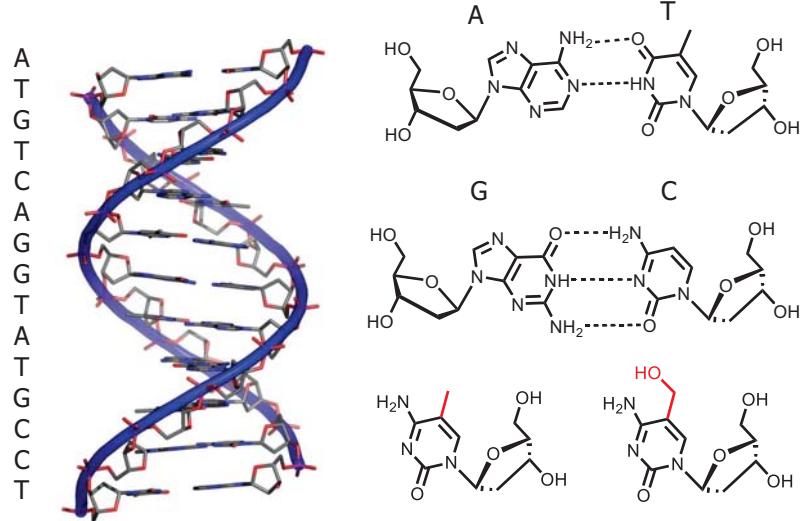
Gene 1  
↓  
Protein 1

Gene 3  
↓  
Protein 3

Gene 5  
↓  
Protein 5

4

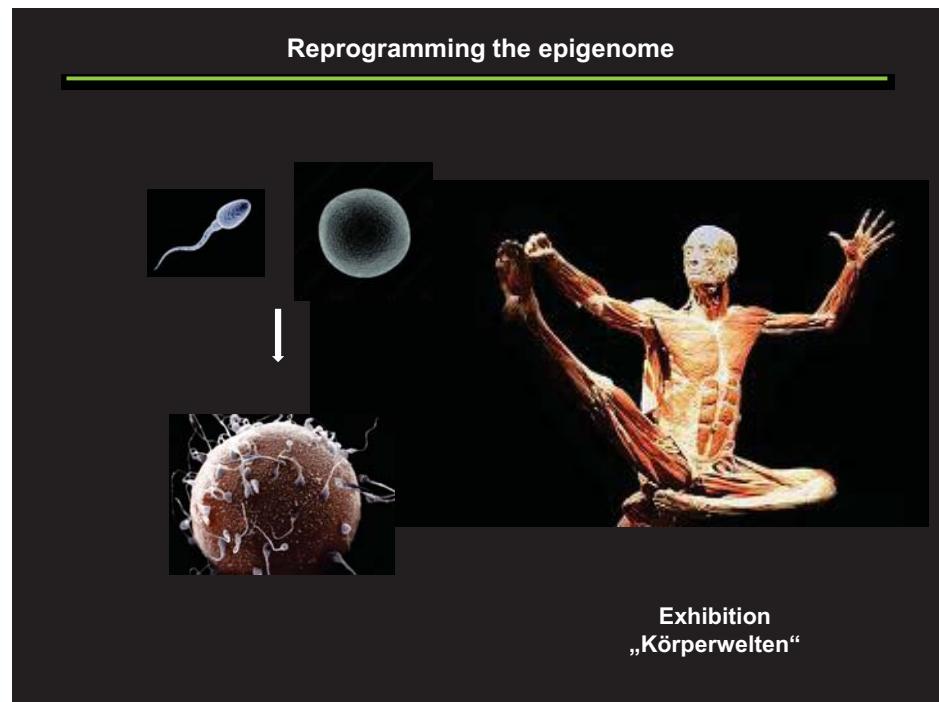
## The molecules of the genetic code



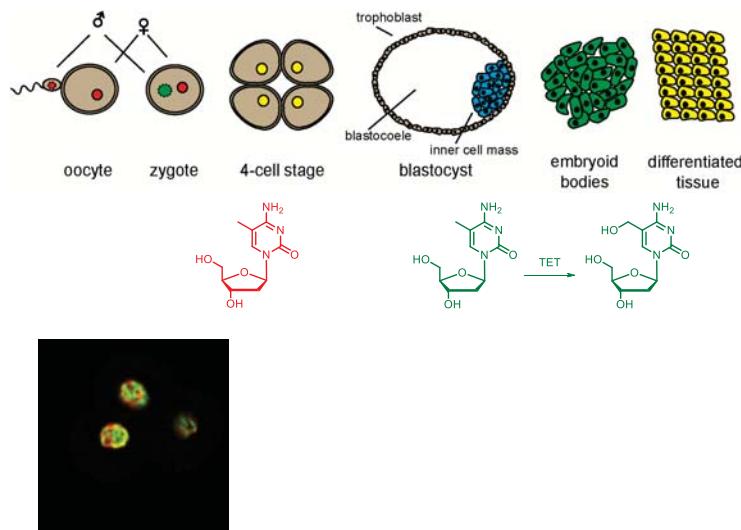
S. Kriaucionis, et al. *Science*, 2009, 324, 929. M. Tahiliani et al. *Science*, 2009, 324, 930

5

## Reprogramming the epigenome

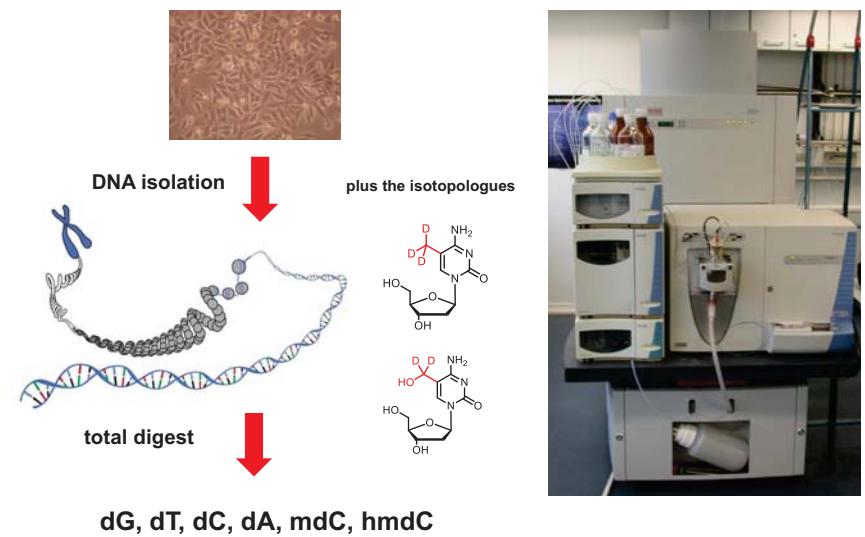


## Chemistry in zygotes



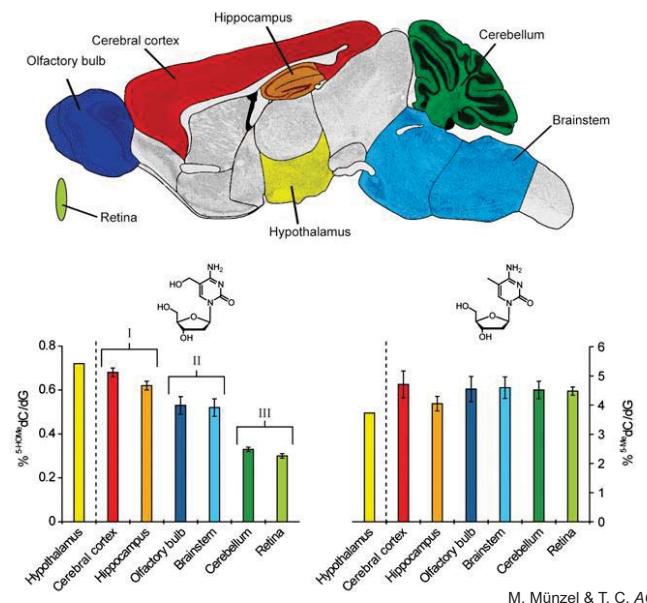
Images: PNAS 2011, 3642 – 3647.

## The isotope standardized LC-MS quantification method (*is*-LC-MS)

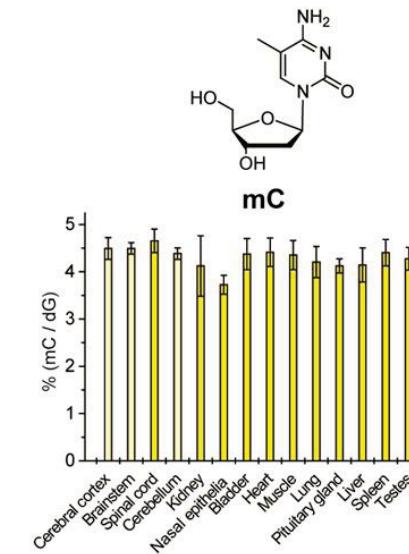


8

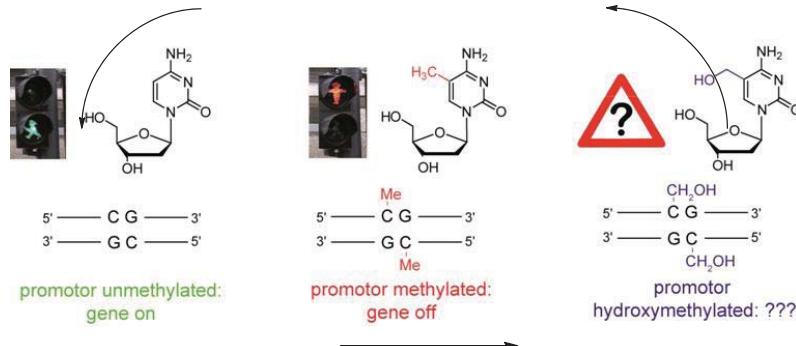
## Quantification in brain tissues



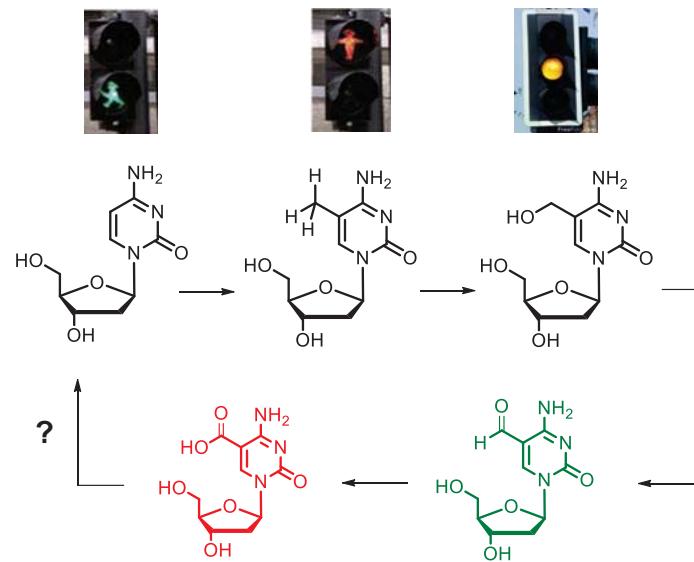
## Distribution of mC and hmC in various tissues



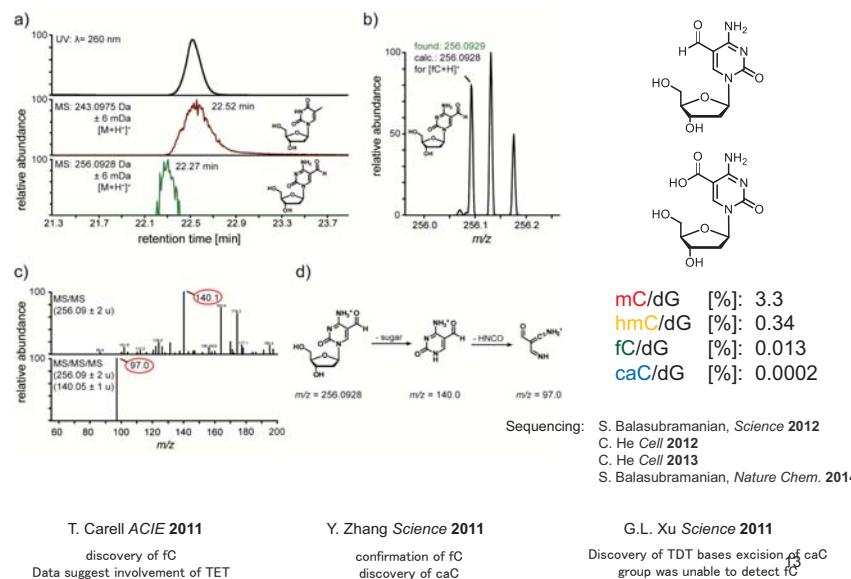
## The function of hmC, fC and caC



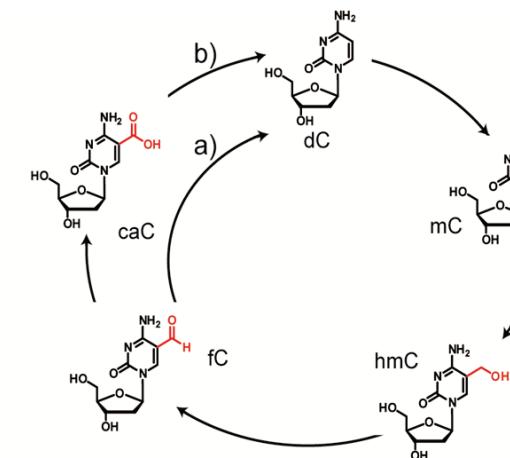
11



## Discovery of fC and caC in ESC



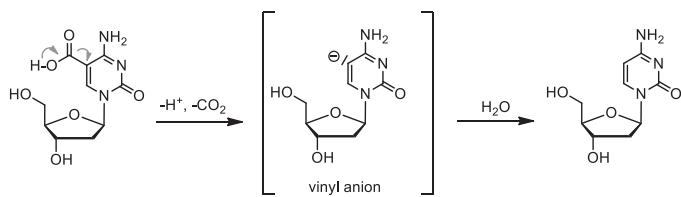
## Potential chemistry of active demethylation



T. Carell *ACIE* 2011, Y. Zhang *Science* 2011 and G.L. Xu *Science* 2011

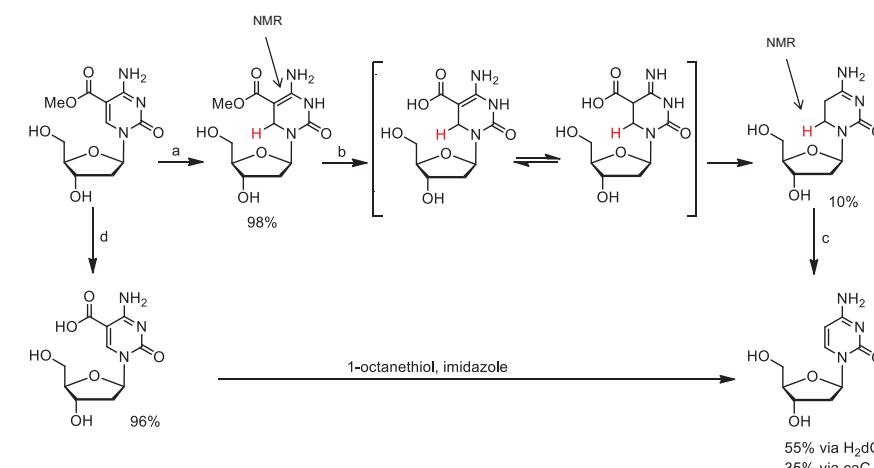
14

## Chemical mechanism of caC decarboxylation

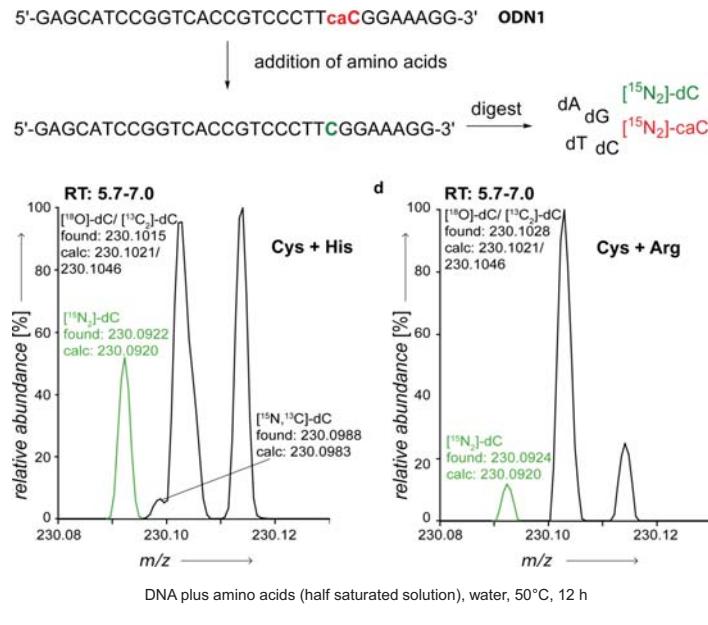


S.Schiess & T.C. *JACS* 2013, 51, 6516-6520  
15

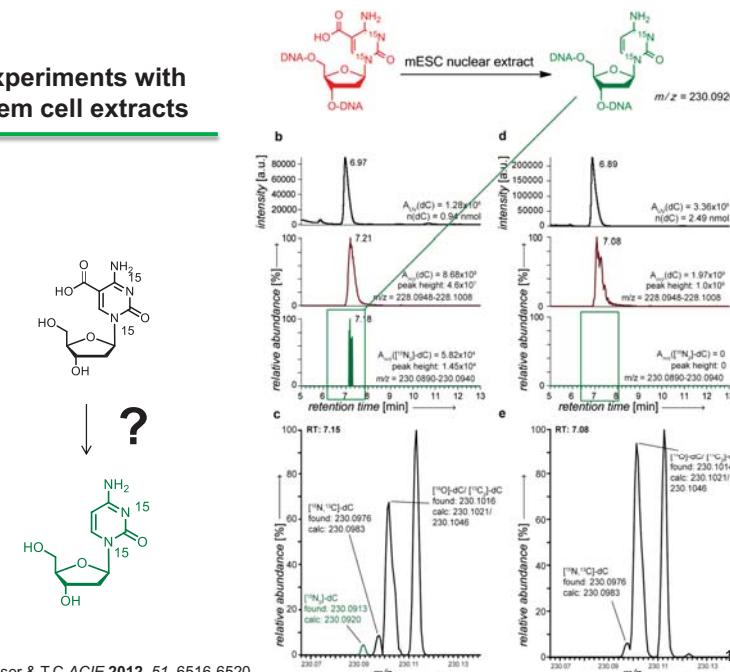
## Chemical mechanism of caC decarboxylation



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## Experiments with stem cell extracts

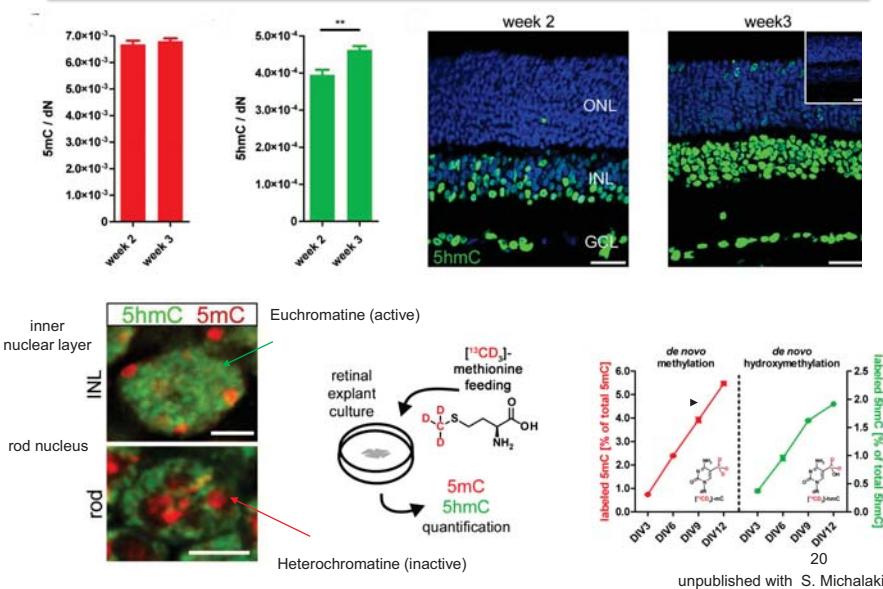


## Analysis of the function of hmC

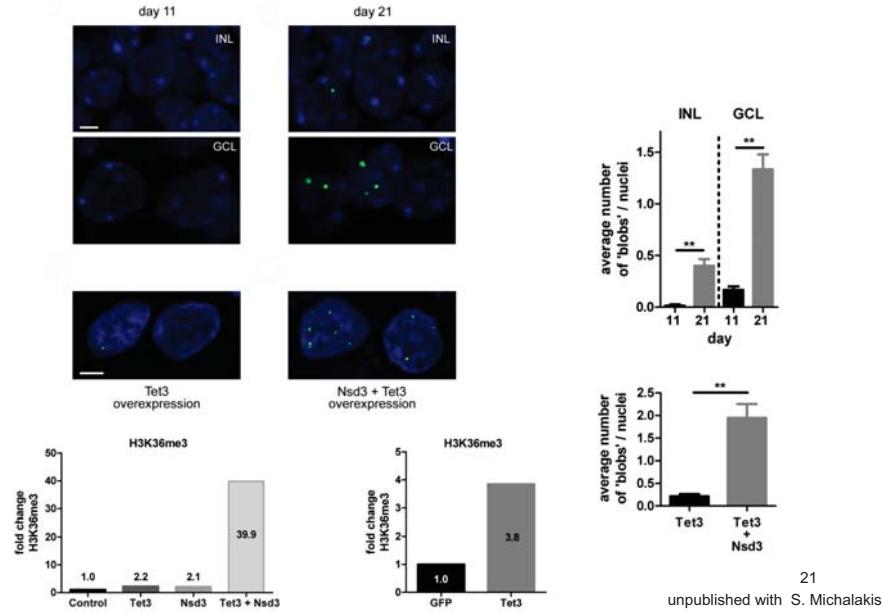
eye opening in mice occurs between week 2 and 3 and goes in hand with large genomic changes in the neuroreceptor cells of the retina



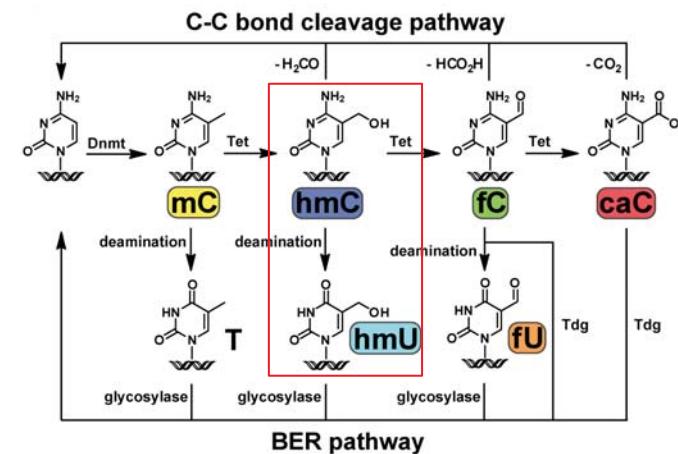
## Studies of brain development shows that hmC is associated with active transcription



## Upregulation of expression leads to increased H3K36me<sup>3</sup>



## Discussed mechanisms of active demethylation



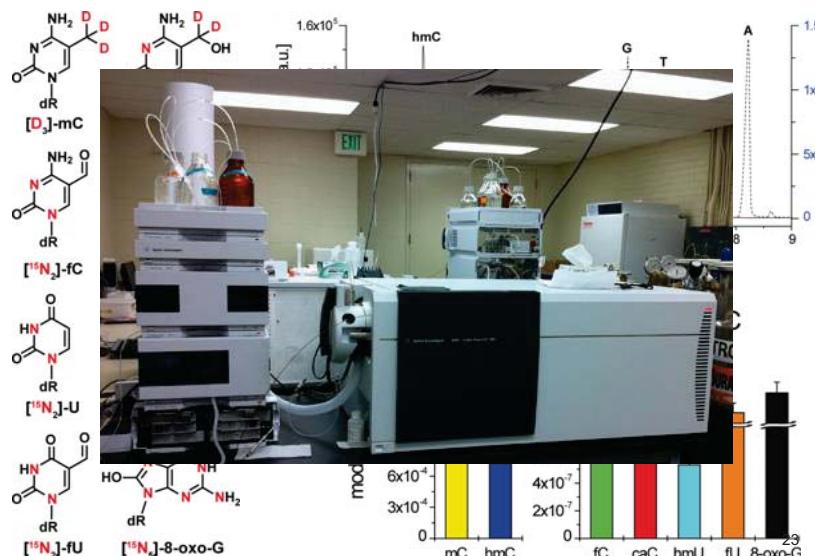
D. Globisch & T. C. PLoS One, 2010, 5, e15367

H. Song, Cell, 2011, 145(3), 423-434.

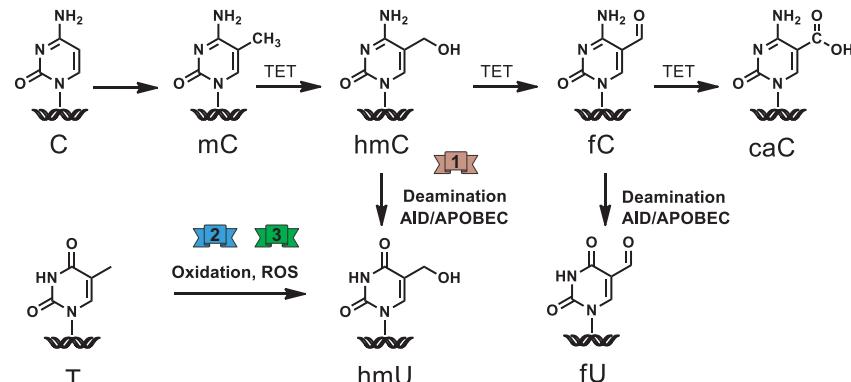
R. M. Kohli, Nature Chem. Biol. 2012, DOI: 10.1038/NCHEMBIO.1042

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## Parallel quantification shows presence of hmU, fU and 8-oxo-G



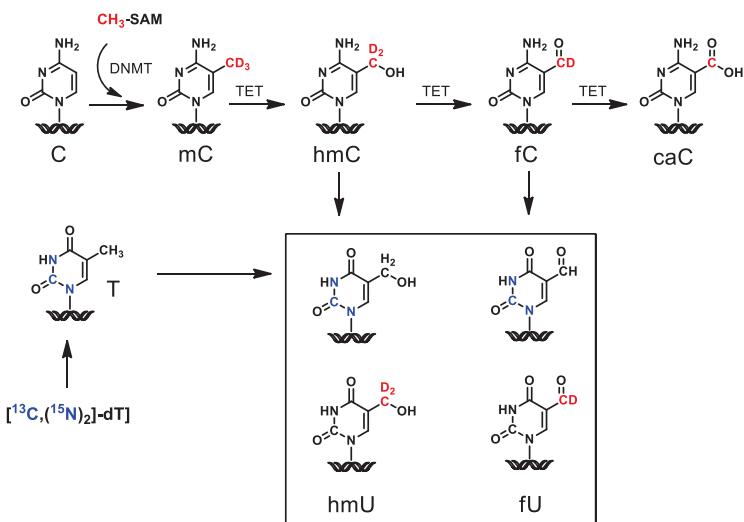
## What is the origin of hmU ?



T. Pfaffeneder, T. C. Nature Chem. Biol. 2014

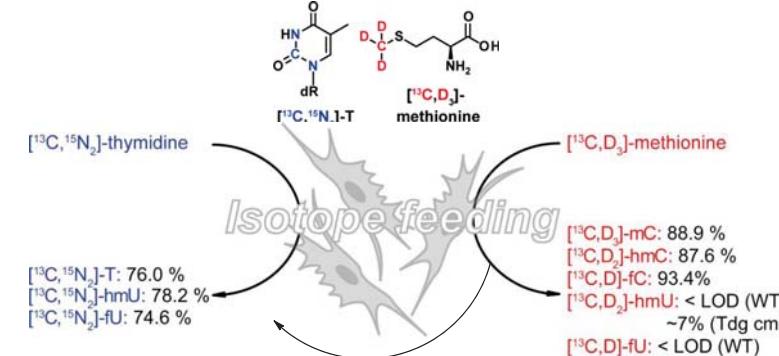
24

### Labelling schemes



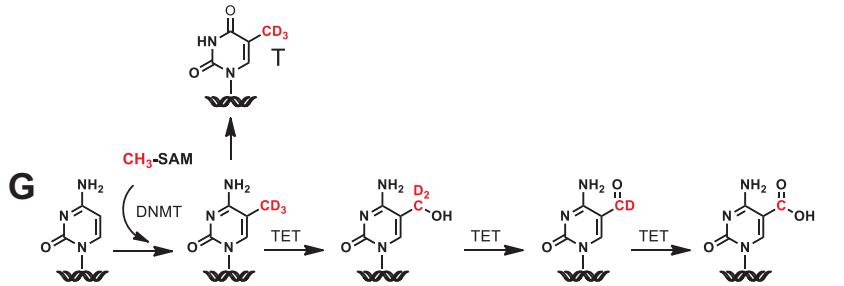
25

### Isotope tracing shows hmU is a dT derivative

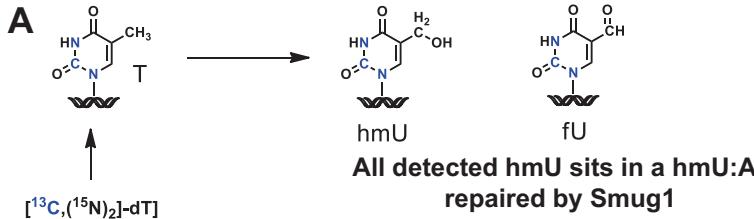


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### Isotope tracing reveals the origin of the new bases

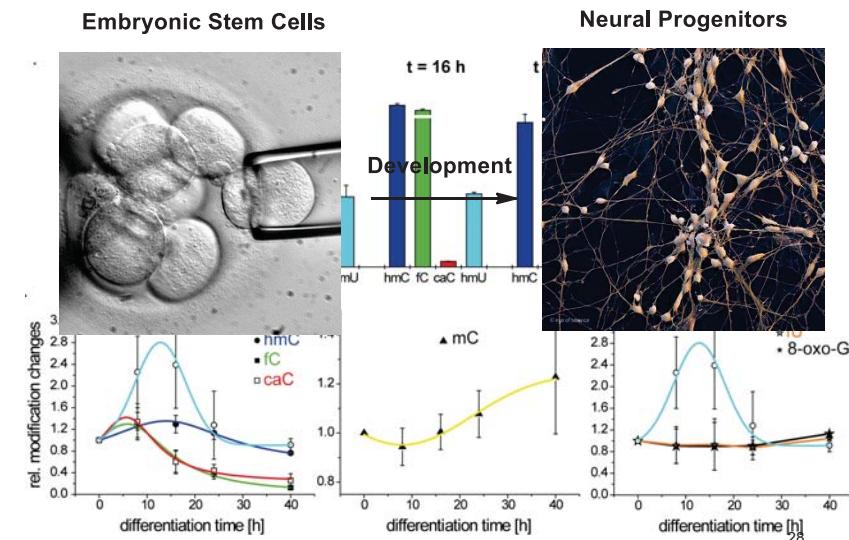


**Who or what oxidizes T to hmU ?**



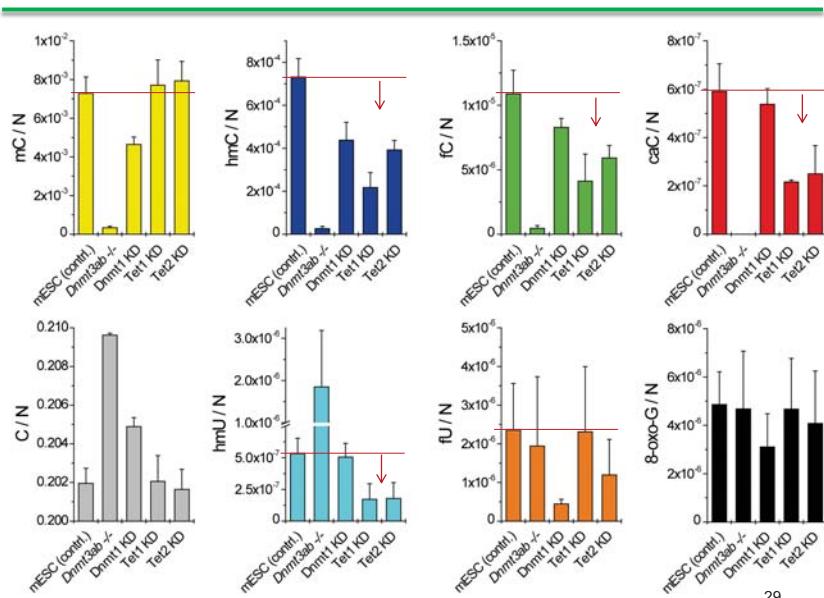
27

### Kinetic analysis shows that hmU is a new base

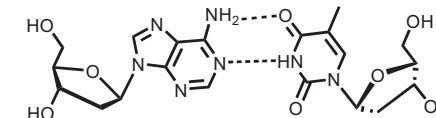


28

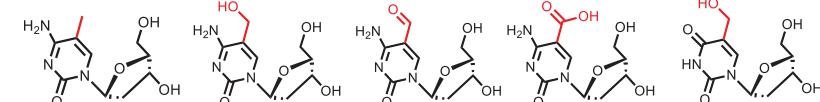
## Tet enzymes oxidize T to hmU



29



## What are the functions of hmC, fC, caC and hmU?



Sequencing:

S. Balasubramanian, *Science* 2012

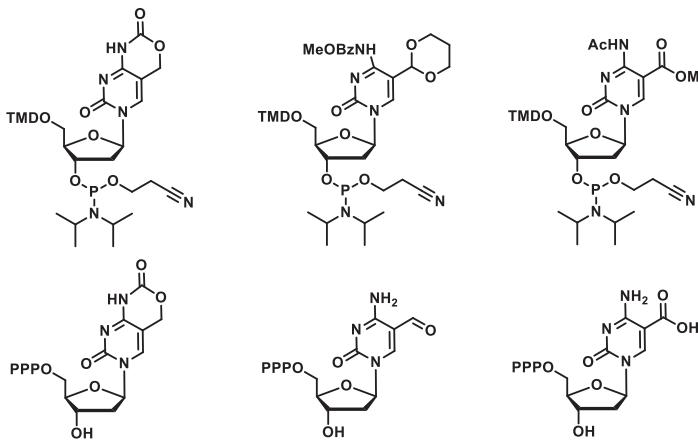
C. He *Cell* 2012

C. He *Cell* 2013

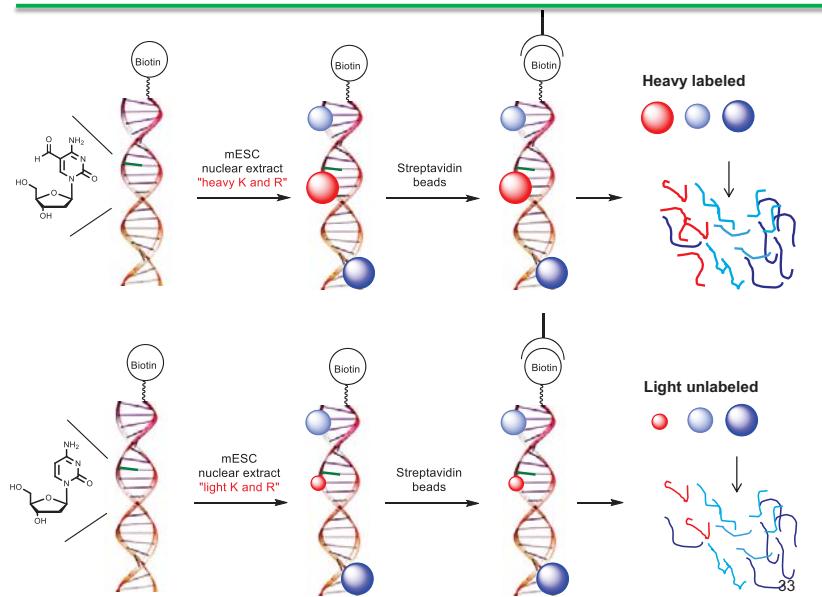
S. Balasubramanian, *Nature Chem.* 2014

30

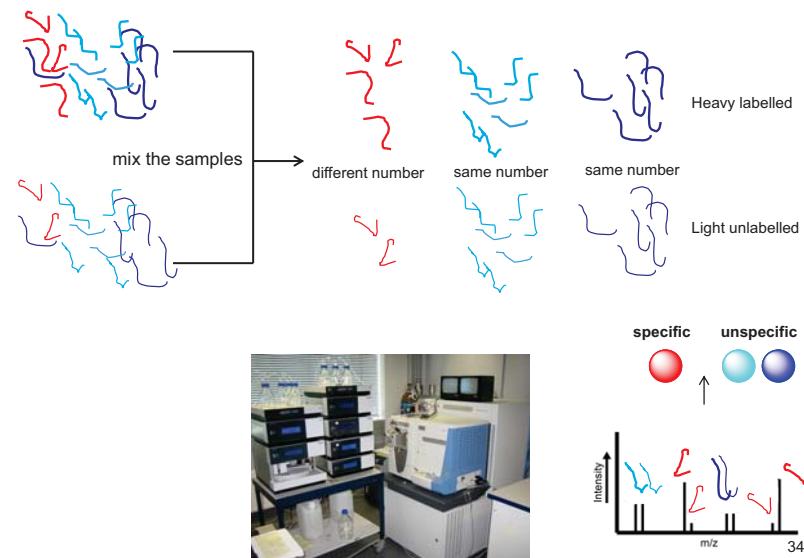
## Building blocks for the synthesis of hmC, fC and caC containing oligonucleotides and PCR products



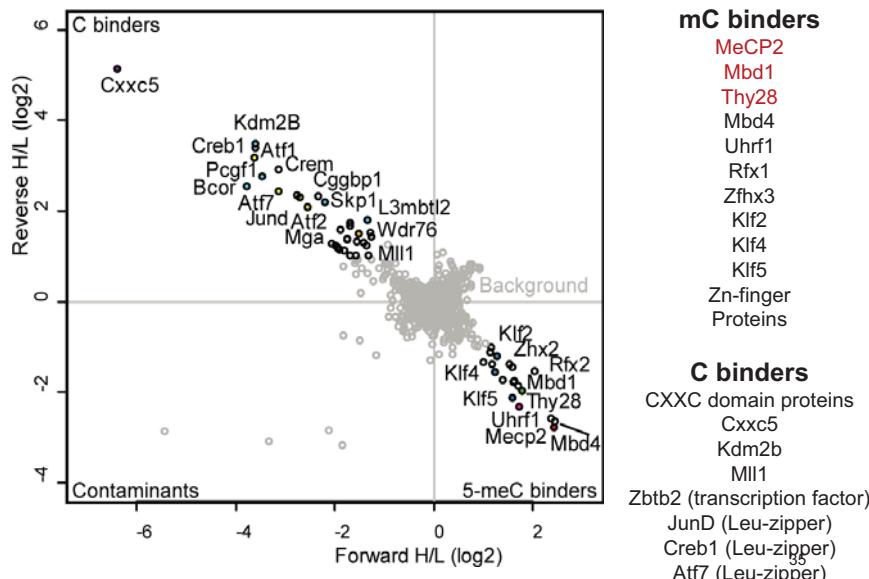
## Which proteins bind to hmC, fC and caC *in vivo* ?



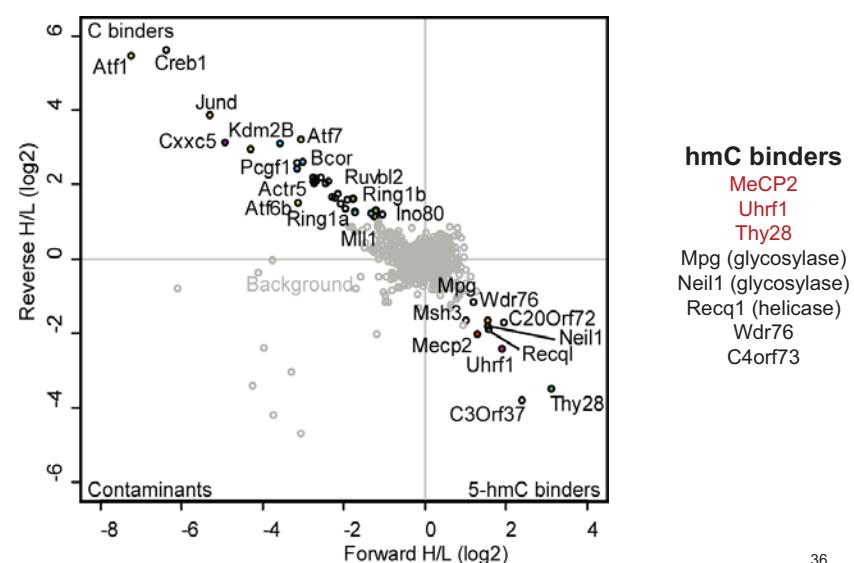
## Which proteins bind to hmC, fC and caC *in vivo* ?



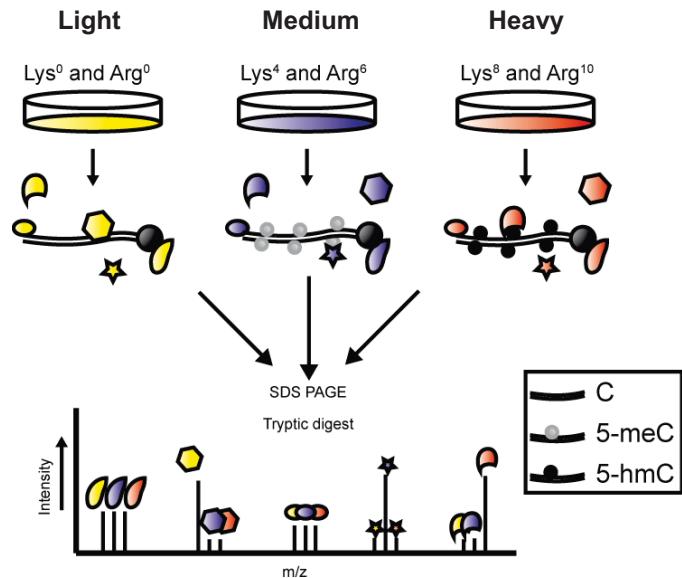
## SILAC derived 5-mC binders



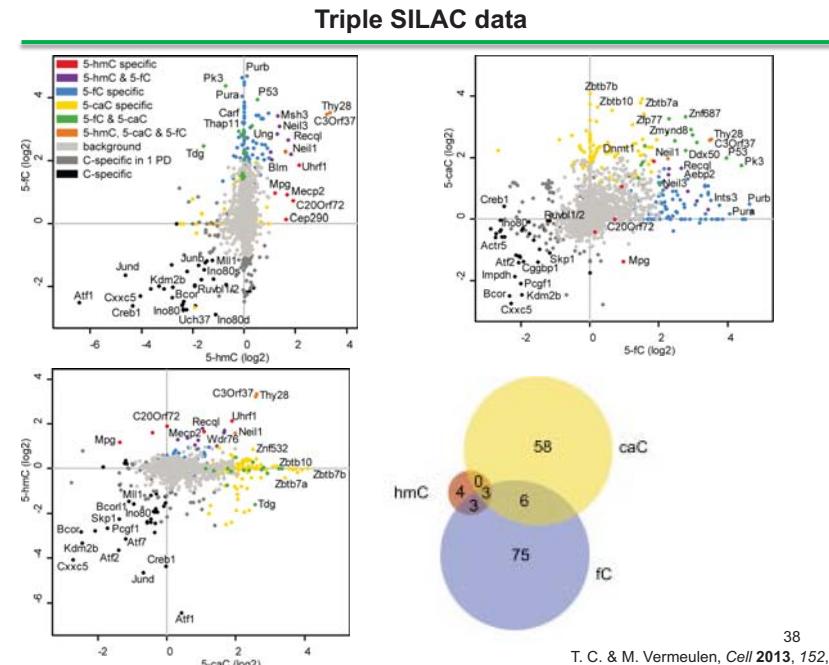
## SILAC derived 5-hmC binders



## Triple-SILAC (M. Mann, M. Vermeulen)

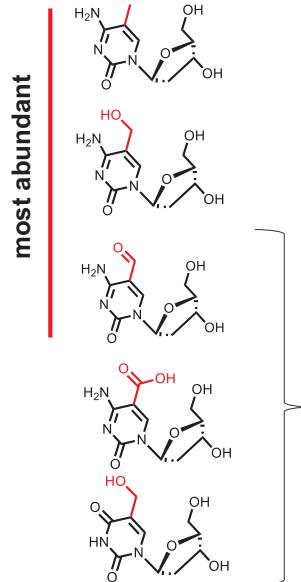


T. C. & M. Vermeulen, *Cell* 2013, 152, 1-14



T. C. & M. Vermeulen, *Cell* 2013, 152, 1-14.

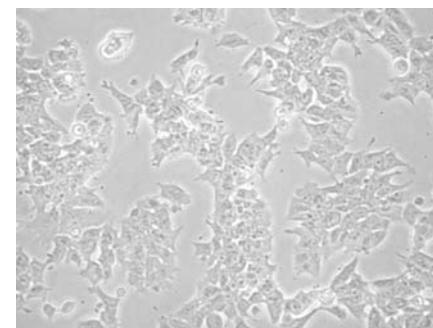
## **Proposed functions of the new bases**



**Creates transcriptionally silenced genes**

**Creates transcriptionally very active genes.  
It triggers H3K36me<sup>3</sup> and  
general histone remodeling**

Required for active demethylation  
and potentially re-methylation.  
These bases attract repair proteins



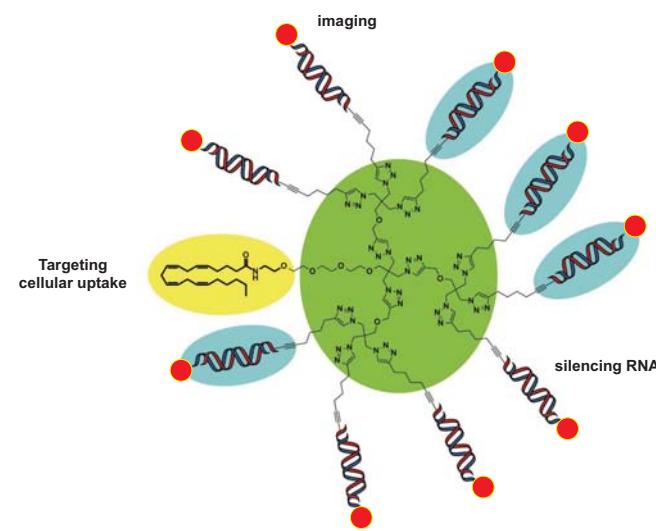
## Axolotl



### Nanog, Oct4, Sox2, Ssea1 (cMyc, KLF4)

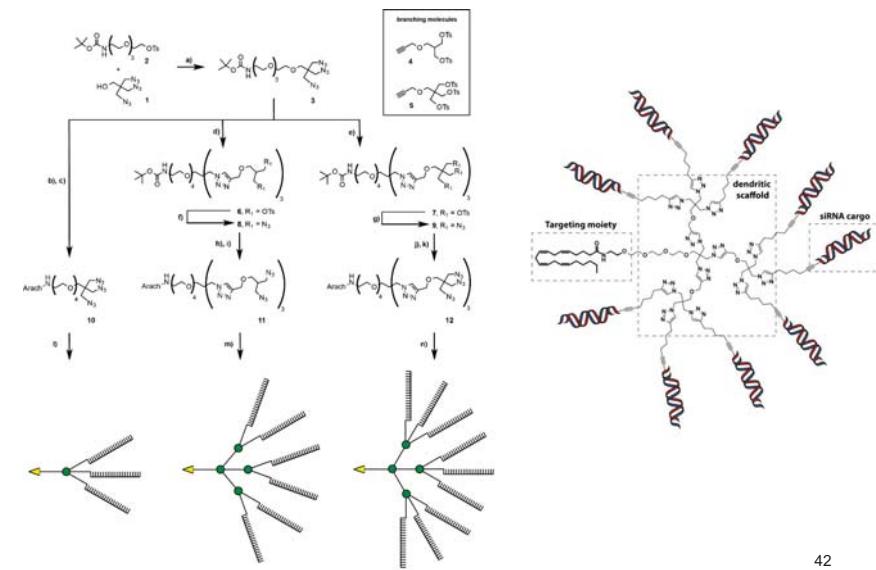
**mmu-miRNA-302a,b,c,d**  
**mmu-mirRNA-367**

## siRNA dendrimers to control stem cell fate assembled by „Click-Chemistry“



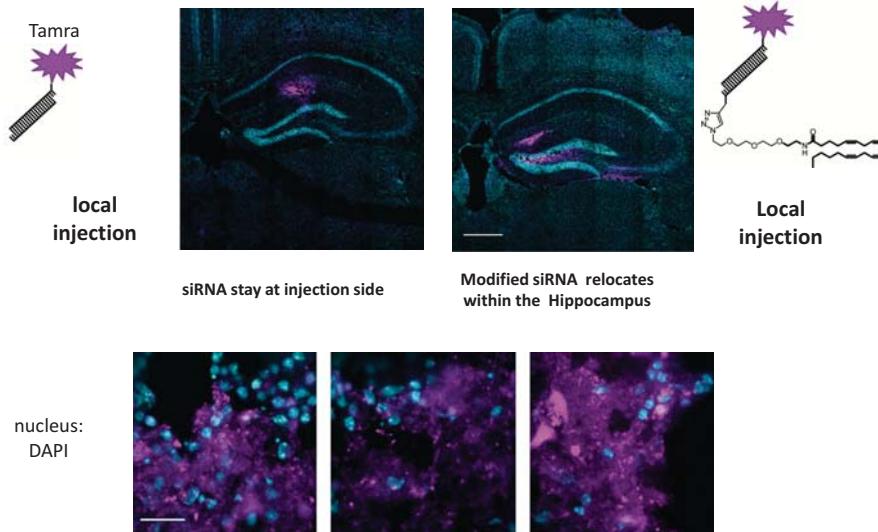
41

## siRNA dendrimers created by „Click Chemistry“



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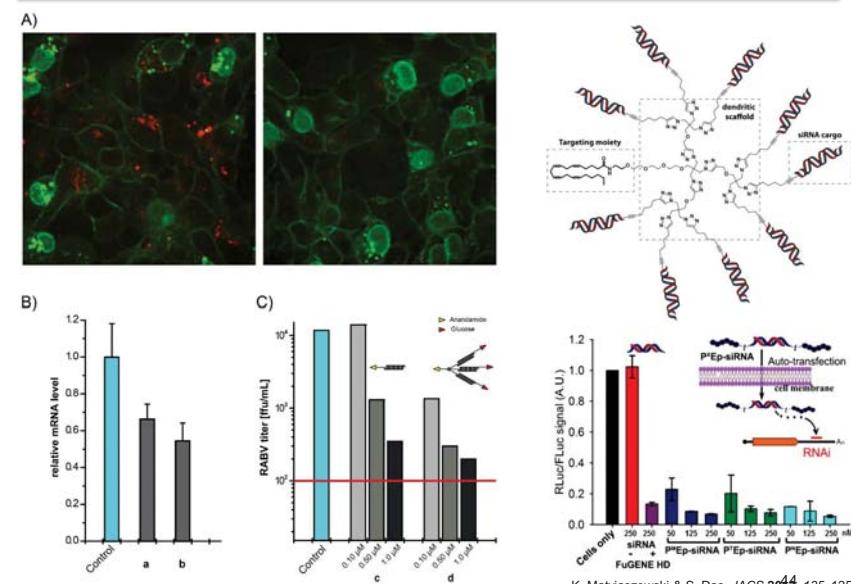
## Arachidonic acid modified siRNA



60-time enlargement: Z-scan demonstrates cellular uptake of (AEA)-siRNA into brain cells

43

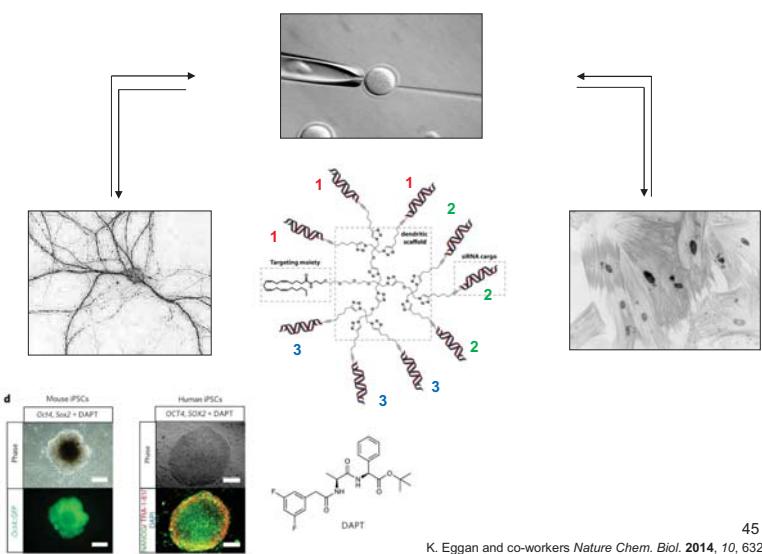
## The molecules of the genetic code



J. Willibald, T. C. JACS 2012, 134, 1230-3.

K. Matyjaszewski & S. Das, JACS 2013, 135, 12508  
K. Matyjaszewski & S. Das, ACIE 2014, in press

## The molecules of the genetic code



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Dr. Markus Müller  
Dr. Fabio Spada  
Dr. David Eisen  
Dr. Gengo Kashiwasaki  
Dr. Maria Gamasa Tomas

Michael Ehrlich  
Benjamin Hackner  
Olesea Kosmatchev  
Olga Kotlarova  
Silvia Laube  
Toni Pfaffeneder  
Stephan Schießer  
Arne Schröder  
Barbara Steigenberger  
Jessica Steinebacher  
Mirko Wagner  
Thomas Wildenhofer

Korbinian Brunner  
Johannes Harder  
Nada Raddouj  
Julian Willibald

T. BEIN, LMU  
C. BRÄUCHLE, LMU

H. LEONHARDT, LMU  
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S. Michalakis, LMU  
M. VERMEULEN, Utrecht  
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M. TRUSS, Charite  
P. SCHÄR BASEL

