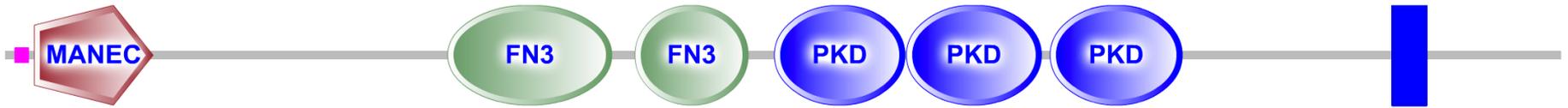


Domains, Motifs, and Pfam/SMART Analysis



Sara Grange and Katie Ness



Protein Domains/ DNA motifs

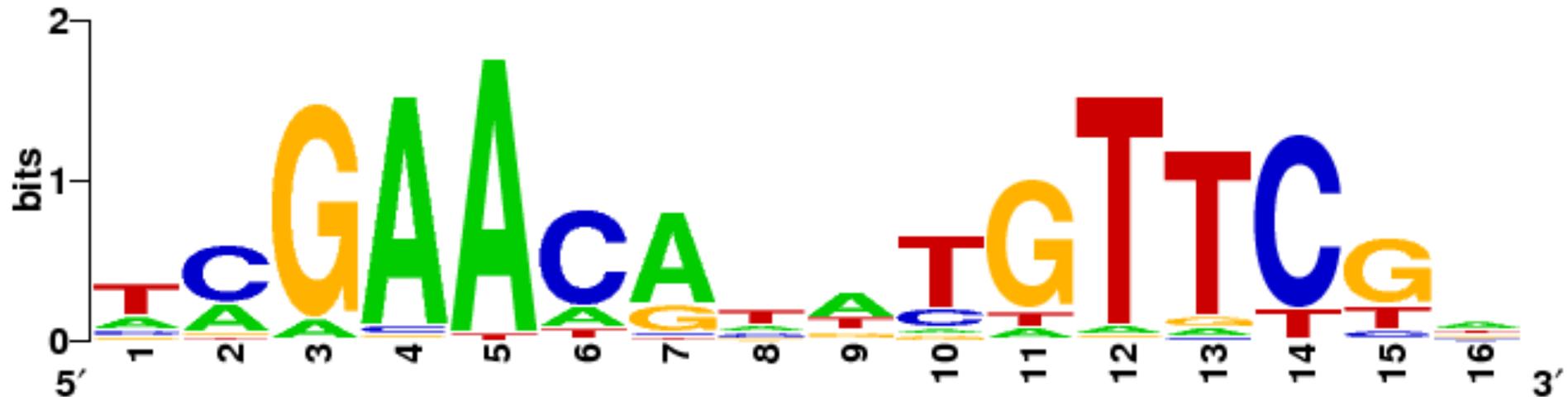
Why are they important?

How are they built?

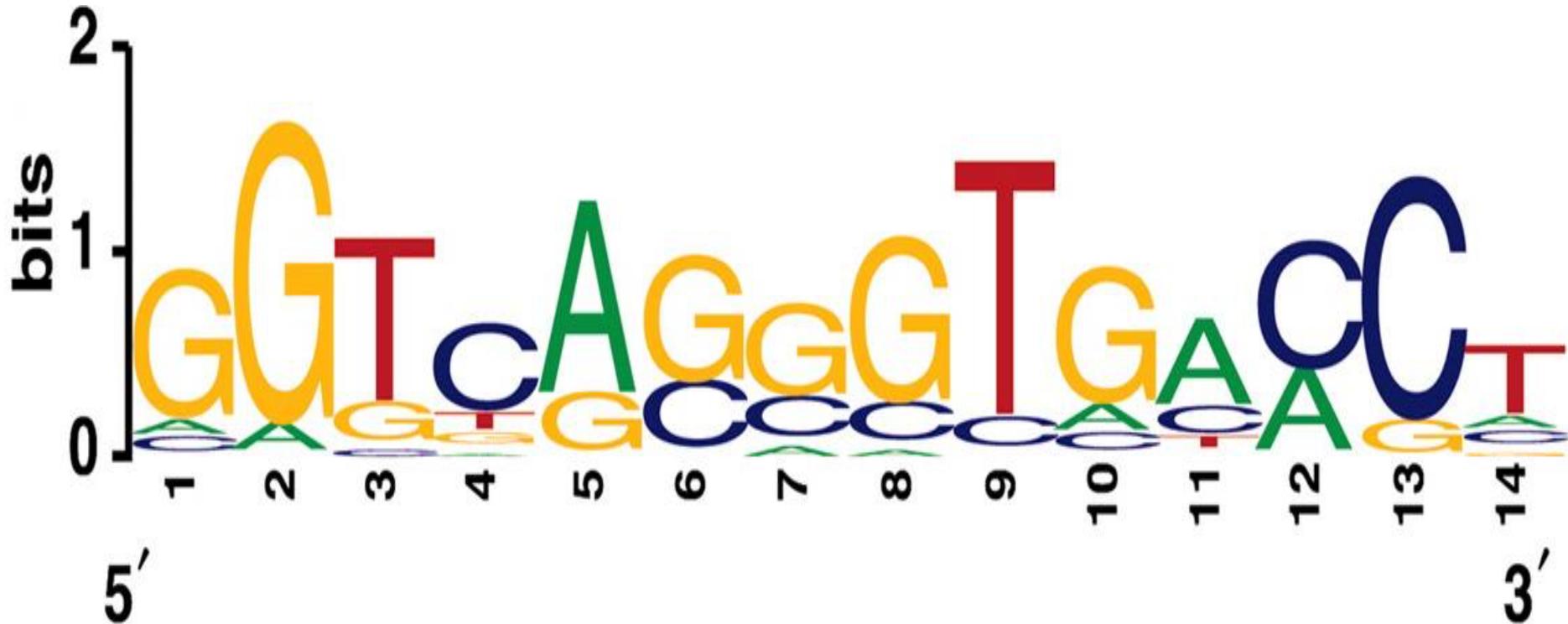
Analysis Techniques

How do they work?

Pfam and SMART databases

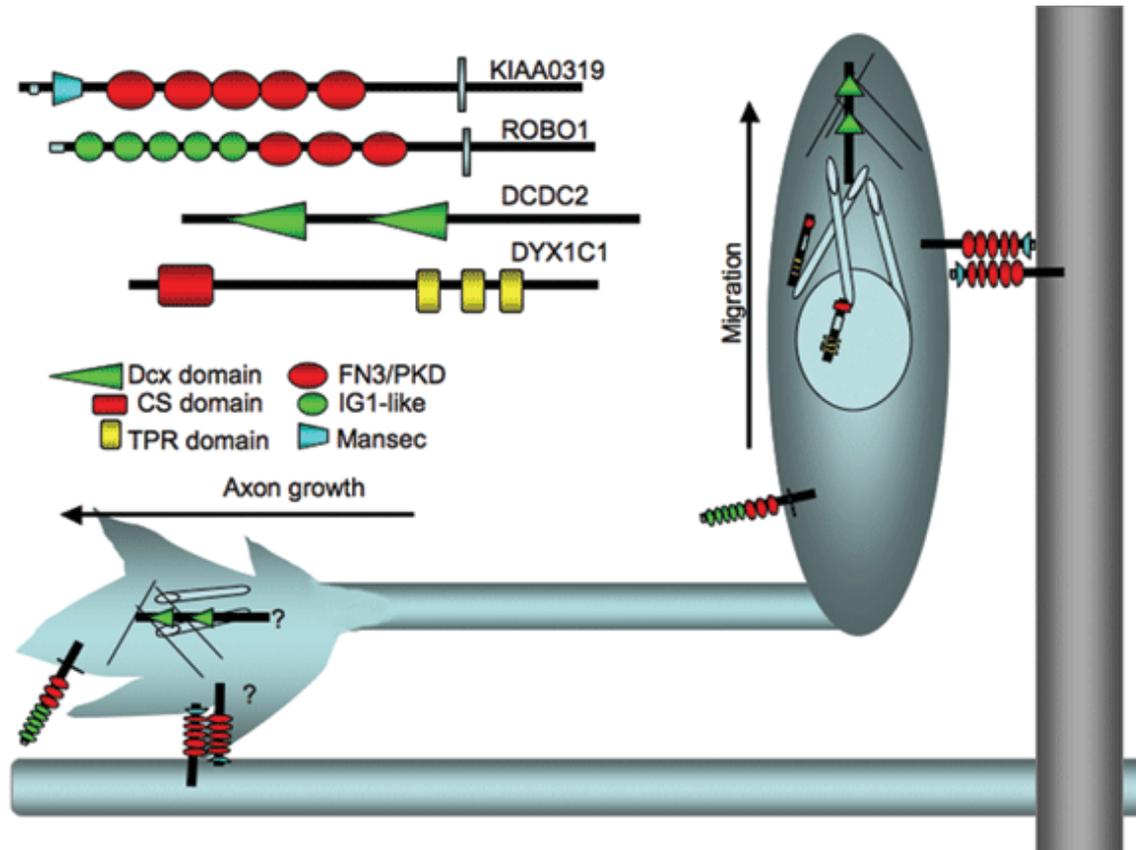


What is a DNA motif?

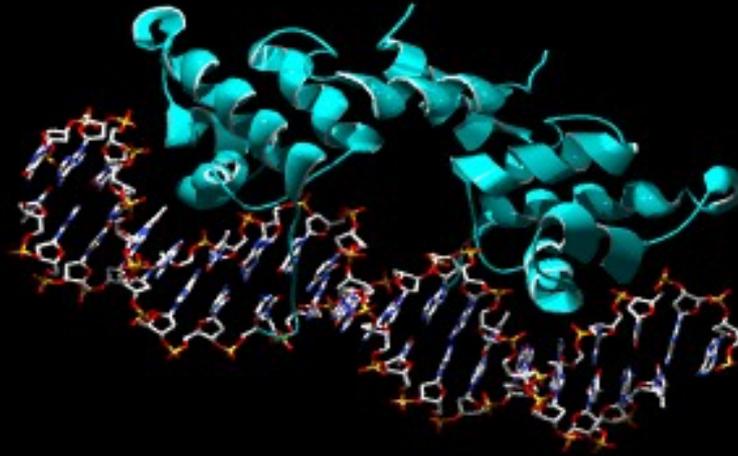


Short, conserved regions of DNA that can be found within domains

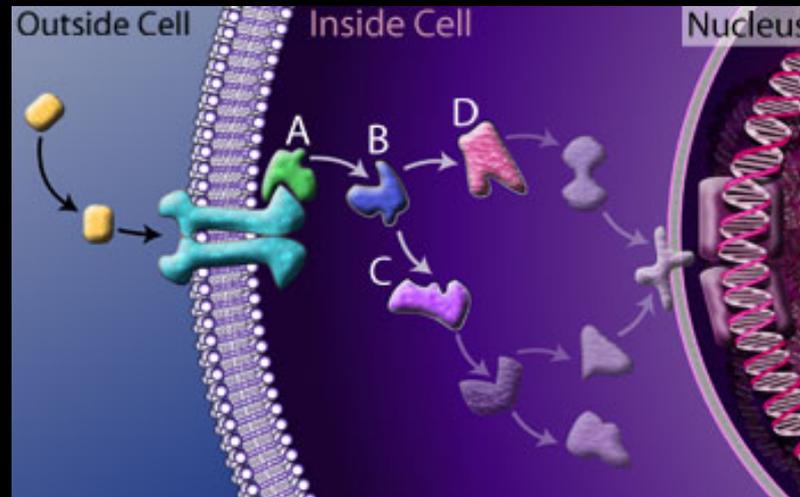
What is a protein domain?



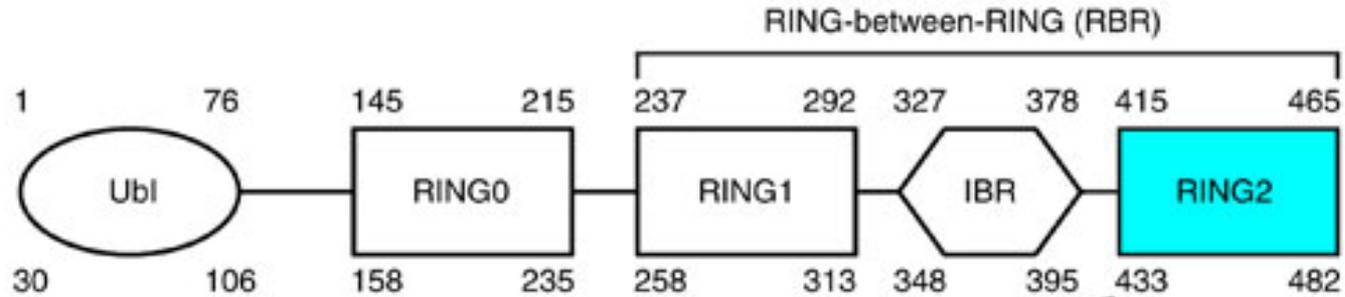
Conserved structural or functional protein segments



What other functions can proteins have that a protein domain can tell you?



How is a domain determined?



b

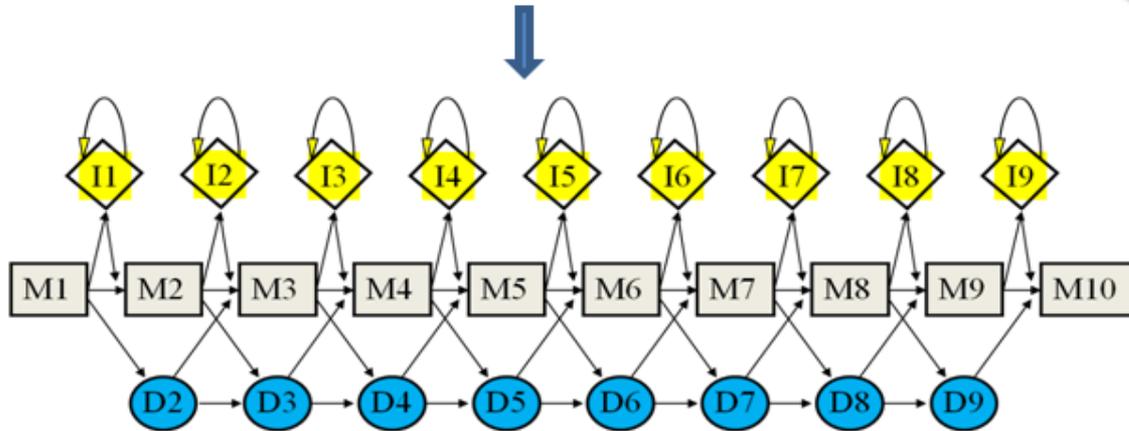
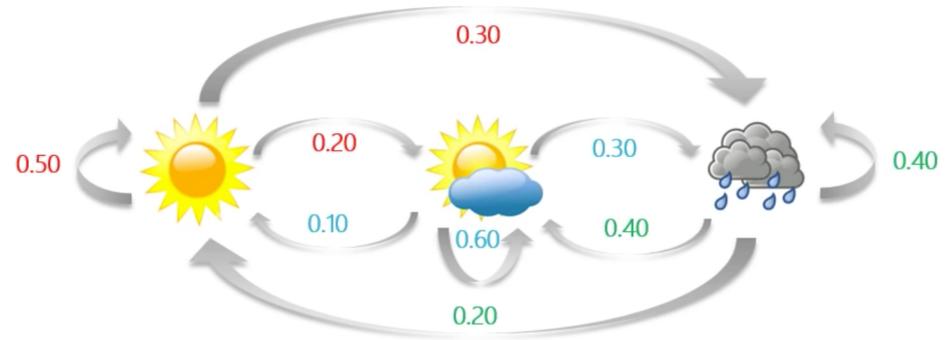


Multiple sequence alignments between homologues

How are multiple sequence alignment quantified?

Multiple sequence alignment

Sequence 1:	F	K	L	L	S	H	C	L	L	V
Sequence 2:	F	K	A	F	G	Q	T	M	F	Q
Sequence 3:	Y	P	I	V	G	Q	E	L	L	G
Sequence 4:	F	P	V	V	K	E	A	I	L	K
Sequence 5:	F	K	V	L	A	A	V	I	A	D
Sequence 6:	L	E	F	I	S	E	C	I	I	Q
Sequence 7:	F	K	L	L	G	N	V	L	V	C



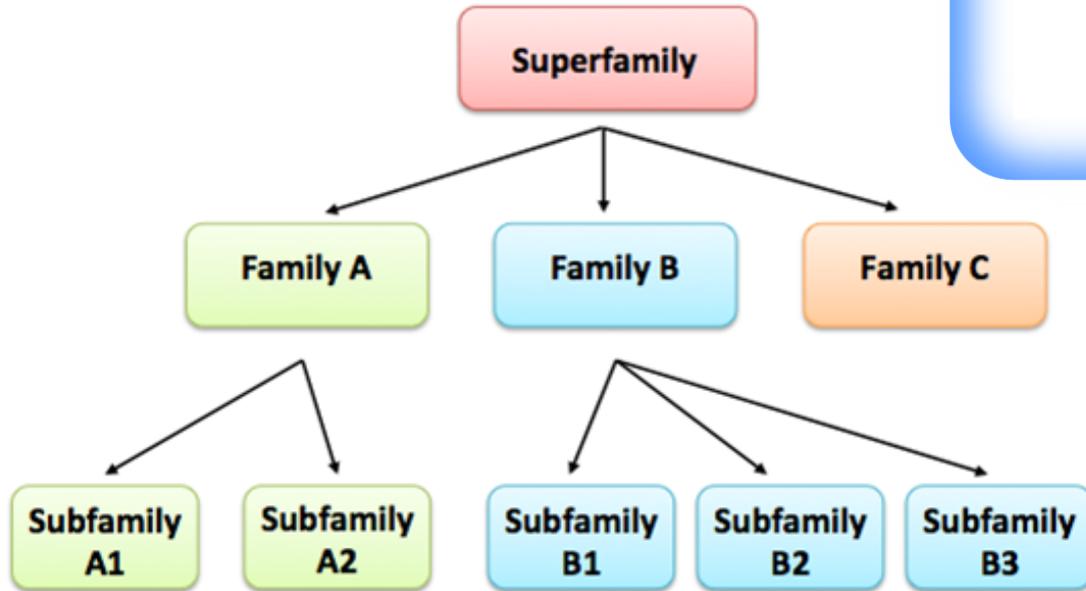
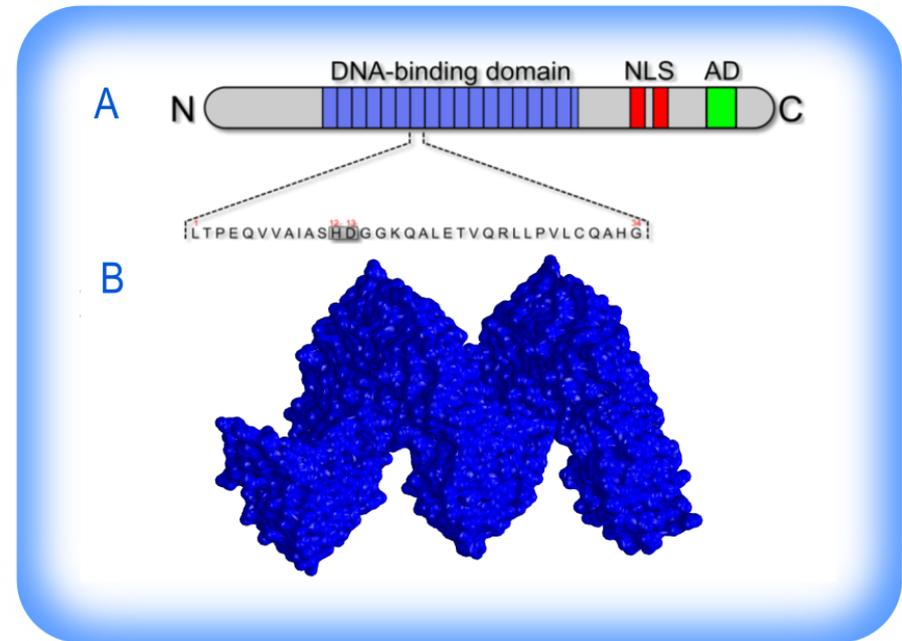
I = insert state

M = match state

D = delete state

Hidden Markov Models (HMMs)

Domain analysis classifies proteins into groups



How do you analyze domains?



Pfam



SMART

Pfam

Advantages	Disadvantages
<p data-bbox="185 358 880 468">Identifies domains with wide spectrum of cellular functions</p> <p data-bbox="127 539 938 589">Can browse clans for homologues</p>	<p data-bbox="1006 419 1846 529">Less sensitive: more false positives and negatives</p>

SMART

Advantages	Disadvantages
<p data-bbox="108 1001 952 1046">More accurate domain identification</p> <p data-bbox="160 1122 904 1172">Domains extensively annotated</p> <p data-bbox="189 1243 875 1293">>100 million protein domains</p>	<p data-bbox="994 1062 1856 1229">Less comprehensive: Mostly identifies domains in signaling, extracellular, and chromatin proteins</p>

How do SMART results compare to Pfam?

Pfam

Homo sapiens



SMART

Homo sapiens

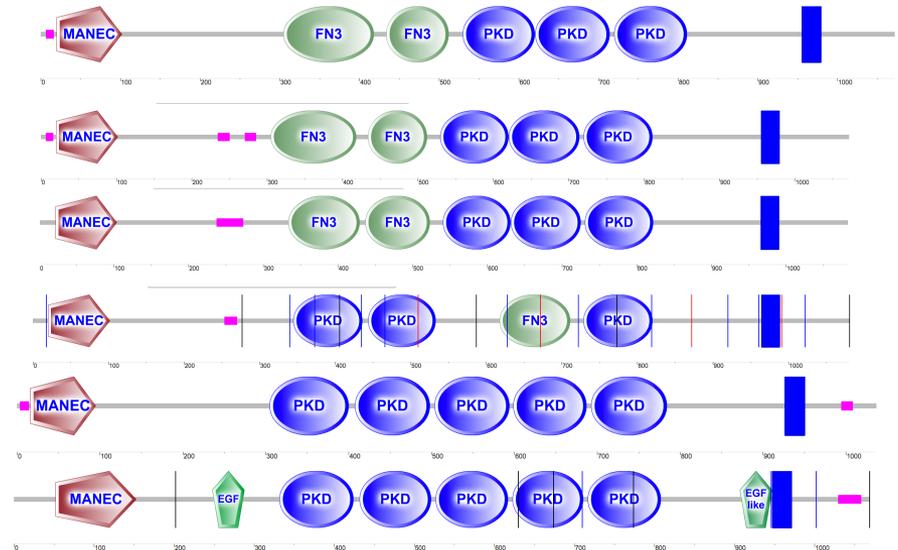
Macaca mulatta

Mus musculus

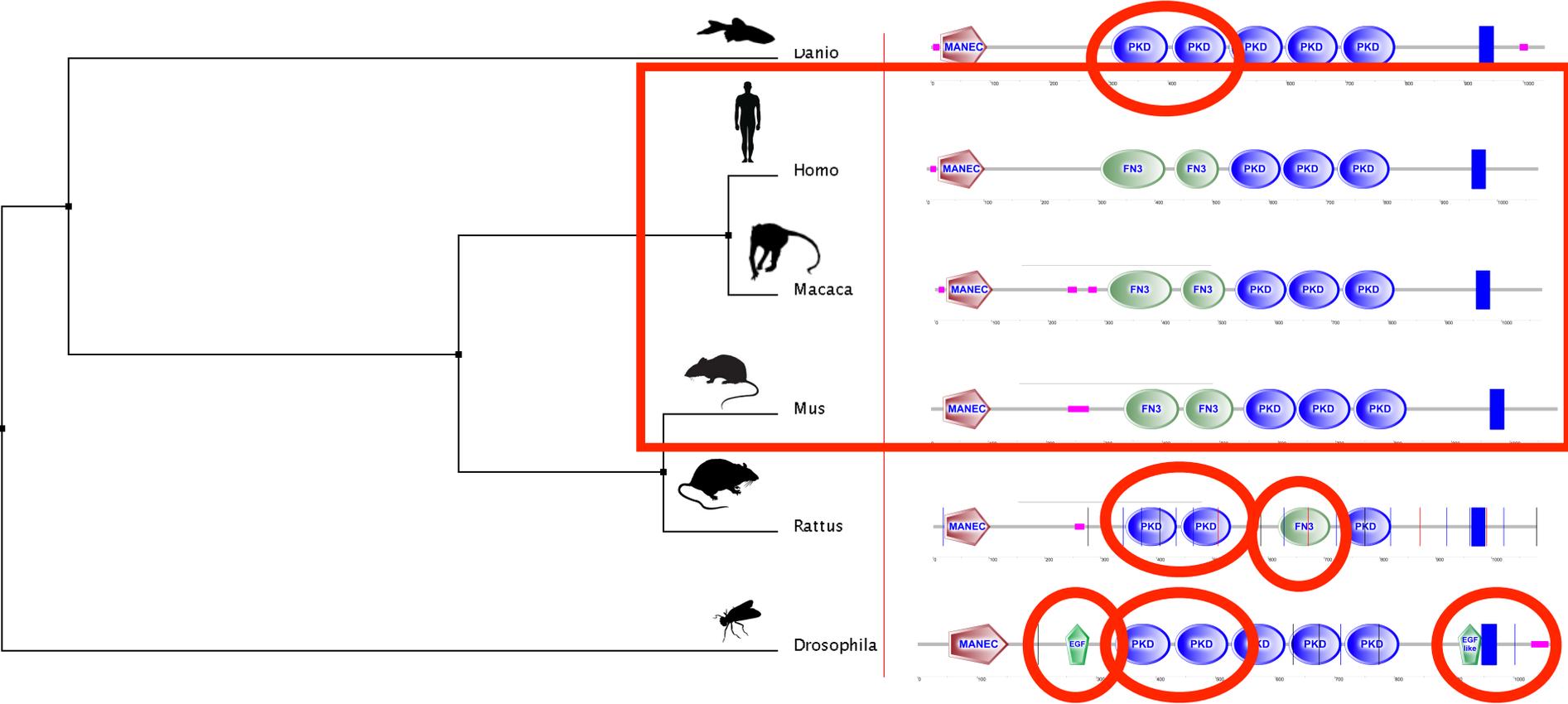
Rattus norvegicus

Danio rerio

Drosophila melanogaster

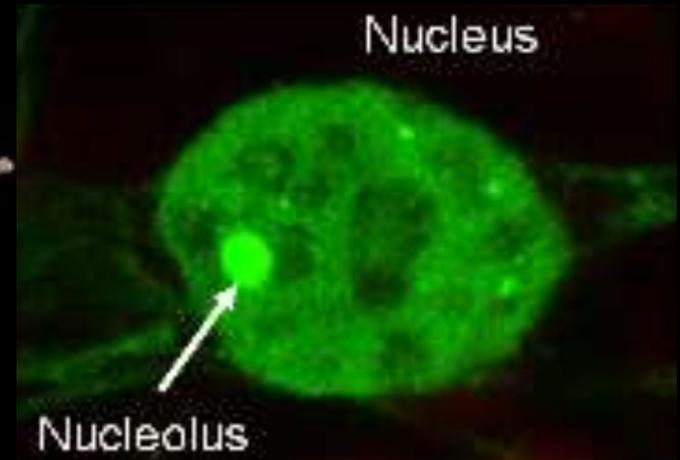


How can you combine phylogeny and domain analysis?



Clustal Omega and SMART

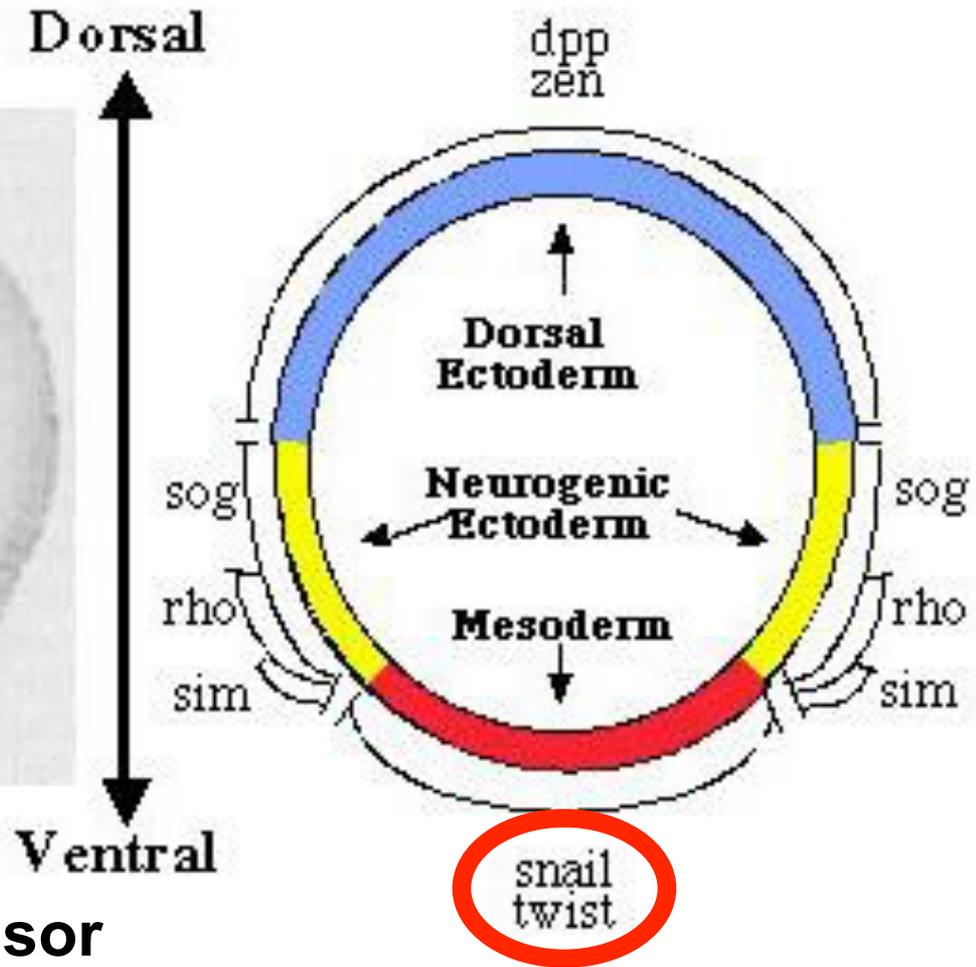
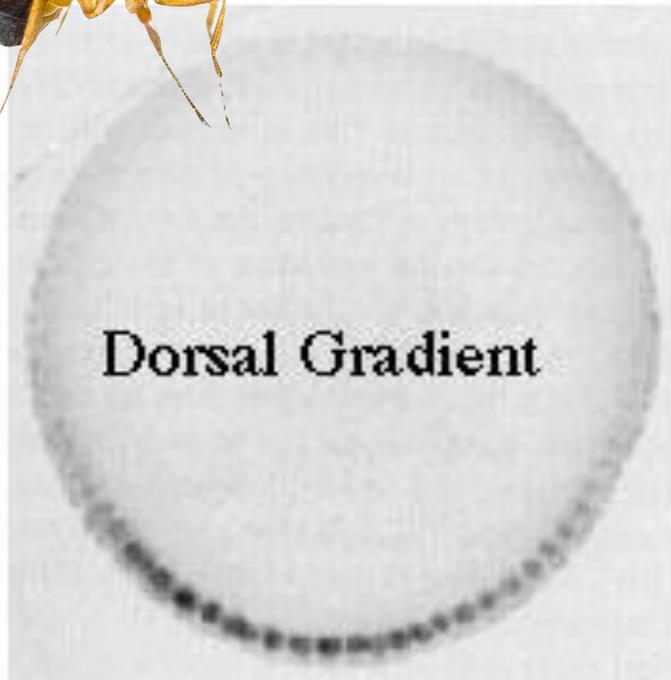
Domain Analysis of the *Nematostella vectensis* SNAIL ortholog reveals unique nucleolar localization that depends on zinc-finger domains



Ada A. Dattoli, Mark A. Hink, Timothy Q. DuBuc, Bram J. Teunisse, Joachim Goedhart, Eric Rottinger, Marten Postma

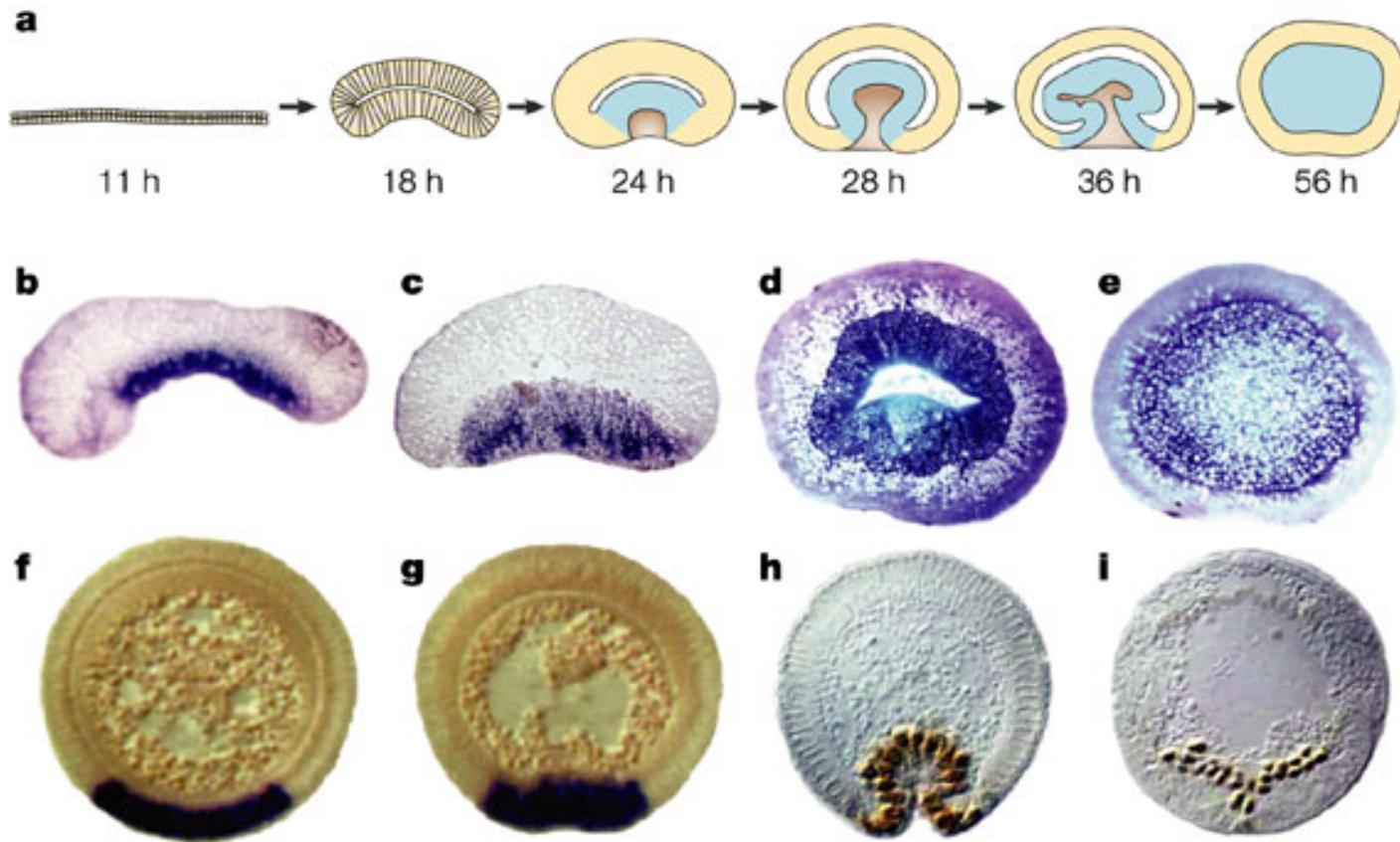
What do Cnidarians look like?

How did the SNAIL protein get it's name?



Transcriptional repressor
First identified in *Drosophila*
Required for mesoderm formation (gastrulation)

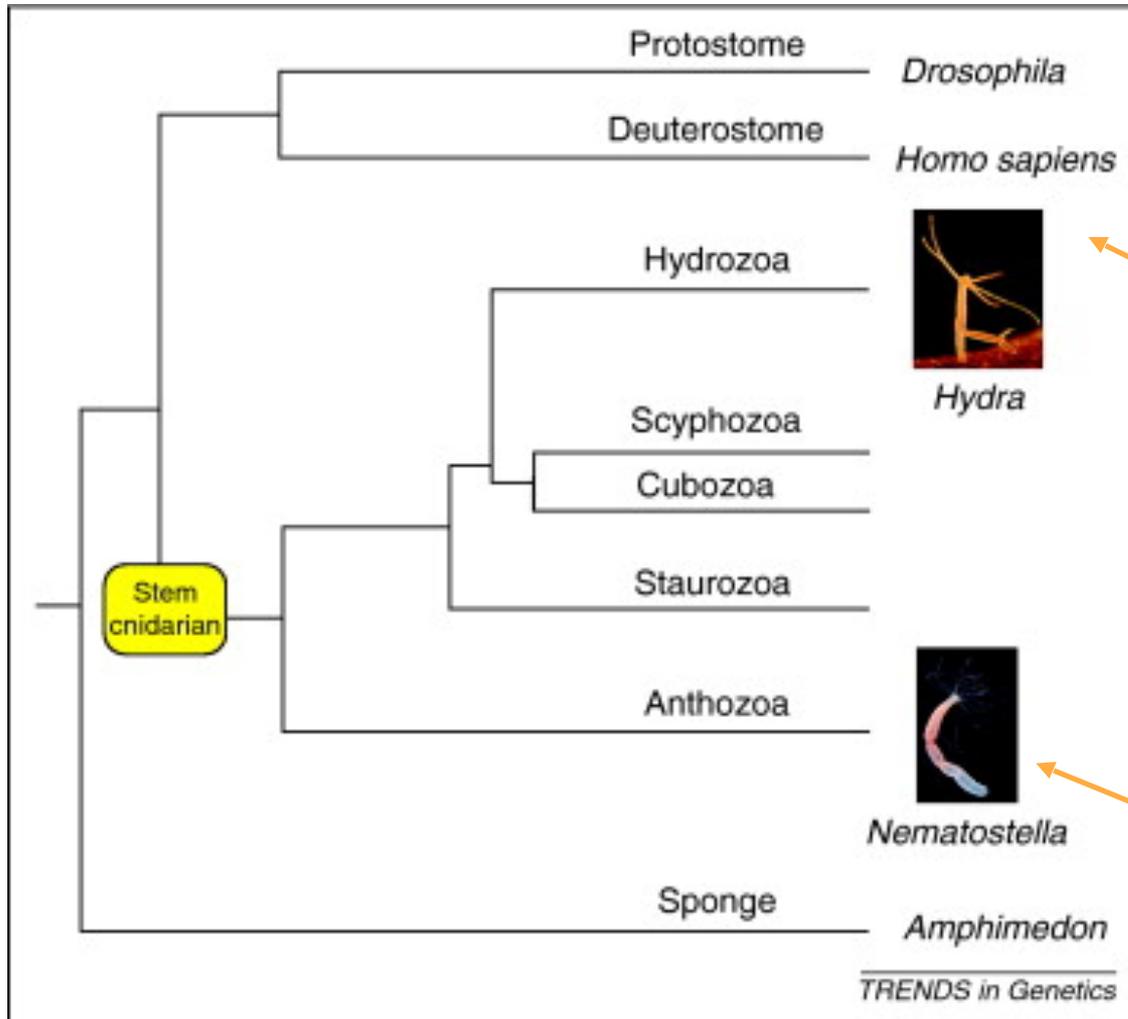
Why are they studying Cnidarians?



Nature Reviews | **Genetics**

Cnidarians don't have a mesoderm layer

Which SNAIL Proteins were studied?



Human SNAIL proteins → *HsSNAIL1*

HsSNAIL2

Cnidarian SNAIL proteins → *NvSNAILA*

NvSNAILB



Fig 1a: What does the SNAIL protein look like?

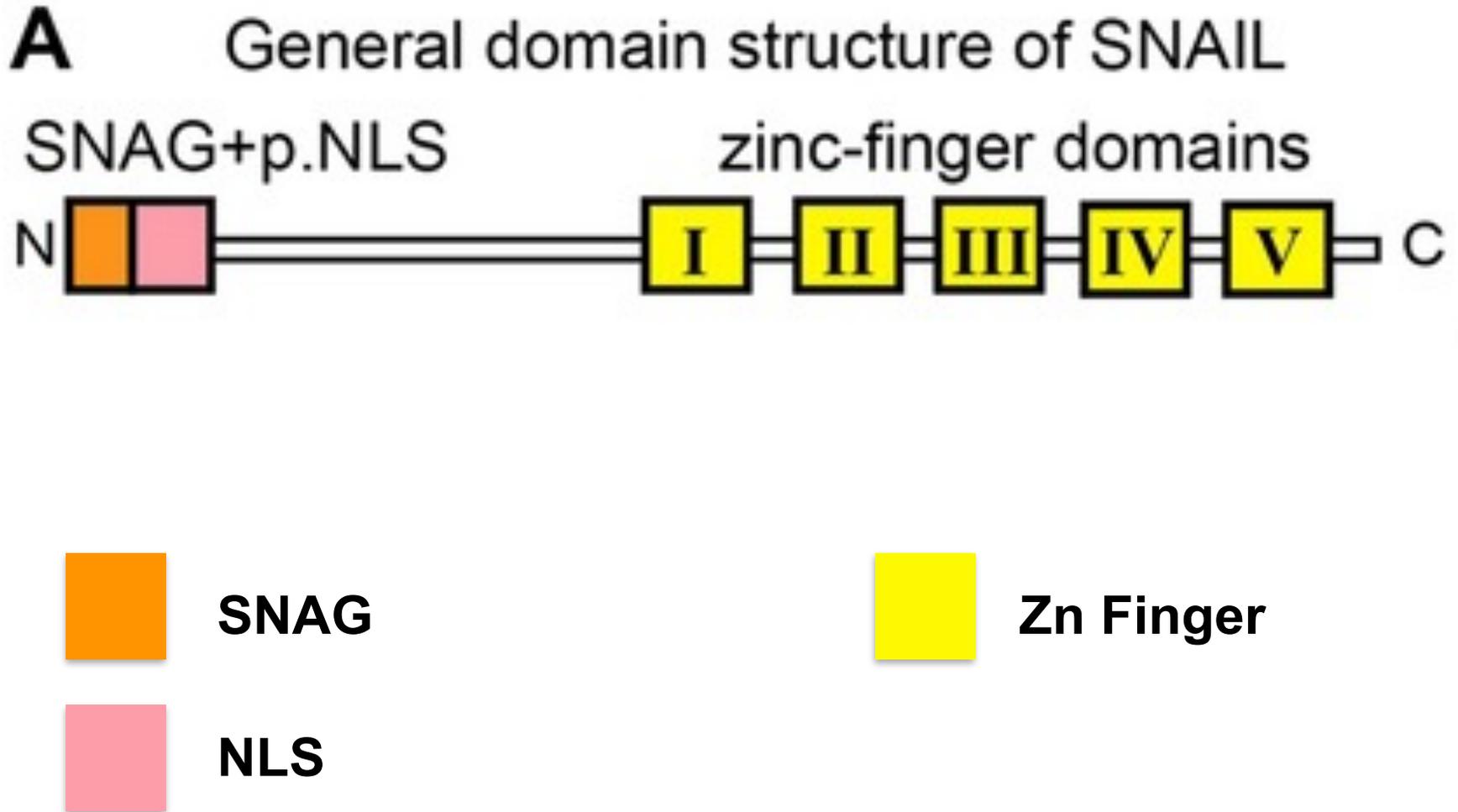
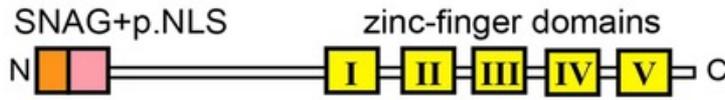


Fig 1: How were the SNAIL homologues determined?

A General domain structure of SNAIL



B

	SNAG										p.NLS									
	1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20
<i>AdSNAIL2</i>	M	P	R	S	F	L	V	K	K	K	P	E	K	W	T	K	H	T	L	D
<i>AdSNAIL1</i>	M	P	K	A	F	L	V	K	K	K	T	E	K	I	K	R	V	P	H	N
<i>PcSNAIL1</i>	M	P	R	S	F	L	V	K	K	K	M	H	L	D	E	C	L	R	Q	Q
<i>AmSNAIL1</i>	M	P	R	S	F	L	V	K	K	K	P	E	K	W	T	K	H	T	L	D
<i>AmSNAIL2</i>	M	P	K	A	F	L	V	K	K	K	T	E	K	I	K	R	V	P	H	N
<i>HmSNAIL1</i>	M	P	R	S	F	L	V	K	Q	F	F	E	K	S	S	N	R	L	P	N
<i>NvSNAILA</i>	M	P	R	S	F	L	V	K	K	T	C	D	K	K	A	L	L	R	N	
<i>NvSNAILB</i>	M	P	R	S	F	L	V	K	T	K	T	E	R	C	S	H	F	D	S	P
<i>HsSNAIL1</i>	M	P	R	S	F	L	V	R	K	P	S	D	P	N	R	K	P	N	Y	S
<i>HsSNAIL2</i>	M	P	R	S	F	L	V	K	K	H	F	N	A	S	K	K	P	N	Y	S
<i>HsSNAIL3</i>	M	P	R	S	F	L	V	K	T	H	S	S	H	R	V	P	N	Y	R	R
<i>XISNAIL1</i>	M	P	R	S	F	L	V	K	K	H	F	S	A	S	K	K	P	N	Y	S
<i>XISNAIL2</i>	M	P	R	S	F	L	V	K	K	H	F	N	S	A	K	K	P	N	Y	G
<i>GgSNAIL1</i>	M	P	R	S	F	L	V	K	K	H	F	S	A	S	K	K	P	N	Y	S
<i>GgSNAIL2</i>	M	P	R	S	F	L	V	K	K	H	F	N	S	S	K	K	P	N	Y	S
<i>MmSNAIL1</i>	M	P	R	S	F	L	V	R	K	P	S	D	P	R	R	K	P	N	Y	S
<i>MmSNAIL2</i>	M	P	R	S	F	L	V	K	K	H	F	N	A	S	K	K	P	N	Y	S
<i>MmSNAIL3</i>	M	P	R	S	F	L	V	K	T	H	S	S	H	R	V	P	N	Y	G	K
<i>DrSNAIL1</i>	M	P	R	S	F	L	V	K	K	Y	F	T	S	K	R	P	N	Y	S	E
<i>DrSNAIL1-like</i>	M	P	R	S	F	L	V	K	K	Y	F	T	N	K	K	P	N	Y	S	E
<i>DrSNAIL3</i>	M	P	R	S	F	L	V	K	K	H	L	T	N	K	K	P	D	Y	G	V
<i>DrSLUG</i>	M	P	R	S	F	L	V	K	K	H	F	N	A	A	K	K	P	N	Y	S
<i>AcSNAIL2-like</i>	M	P	R	S	F	L	V	K	K	H	F	N	A	S	K	K	P	N	Y	S

C

	C2H2 zinc-finger																											
	I					C					C					H												
<i>HsSNAIL2</i>	K	F	Q	C	N	L	C	N	K	T	Y	S	T	F	S	G	L	A	K	H	K	Q	L	H	C	-		
<i>NvSNAILA</i>	K	H	Q	C	H	Q	C	N	K	G	Y	S	T	P	L	G	L	A	K	H	Q	Q	F	H	C	-		
<i>NvSNAILB</i>	K	L	Q	C	P	N	C	M	K	G	F	N	A	L	A	T	L	M	R	H	Q	Y	F	Y	C	P		
	II					C					C					H												
<i>HsSNAIL1</i>	A	F	N	C	K	Y	C	N	K	E	Y	L	S	L	G	A	L	K	M	H	I	R	S	H	T	L		
<i>HsSNAIL2</i>	S	F	S	C	K	Y	C	D	K	E	Y	V	S	L	G	A	L	K	M	H	I	R	T	H	T	L		
<i>NvSNAILA</i>	S	F	T	C	K	H	C	D	K	I	Y	V	S	L	G	A	L	K	M	H	I	R	T	H	T	L		
<i>NvSNAILB</i>	P	F	H	C	K	Y	C	E	K	L	Y	D	S	L	G	A	L	K	M	H	I	R	T	H	T	L		
	III					C					C					H												
<i>HsSNAIL1</i>	P	C	V	C	G	T	C	G	K	A	F	S	R	P	W	L	L	Q	G	H	V	R	T	H	T	G	E	K
<i>HsSNAIL2</i>	P	C	V	C	K	I	C	G	K	A	F	S	R	P	W	L	L	Q	G	H	I	R	T	H	T	G	E	K
<i>NvSNAILA</i>	P	C	K	C	S	I	C	G	K	A	F	N	R	P	W	L	L	Q	G	H	I	R	T	H	T	G	E	K
<i>NvSNAILB</i>	P	C	K	C	K	I	C	G	K	A	F	S	R	P	W	L	L	Q	G	H	V	R	T	H	T	G	E	K
	IV					C					C					H												
<i>HsSNAIL1</i>	P	F	S	C	P	H	C	S	R	A	F	A	D	R	S	N	L	R	A	H	L	Q	T	H	S	D	V	K
<i>HsSNAIL2</i>	P	F	S	C	P	H	C	N	R	A	F	A	D	R	S	N	L	R	A	H	L	Q	T	H	S	D	V	K
<i>NvSNAILA</i>	P	Y	Q	C	T	N	C	K	R	A	F	A	D	R	S	N	L	R	A	H	M	Q	T	H	A	V	V	K
<i>NvSNAILB</i>	P	Y	K	C	T	Q	C	Q	R	A	F	A	D	R	S	N	L	R	A	H	L	Q	T	H	S	D	V	K
	V					C					C					H												
<i>HsSNAIL1</i>	K	Y	Q	C	Q	A	C	A	R	T	F	S	R	M	S	L	L	H	K	H	Q	E	S	G	C	S		
<i>HsSNAIL2</i>	K	Y	Q	C	K	N	C	S	K	T	F	S	R	M	S	L	L	H	K	H	E	E	S	G	C	C		
<i>NvSNAILA</i>	K	Y	S	C	S	R	C	K	K	S	F	S	R	M	S	L	L	V	K	H	E	D	S	G	C	P		
<i>NvSNAILB</i>	K	Y	S	C	K	Q	C	S	K	S	F	S	R	M	S	L	L	L	K	H	-	E	G	S	C	S		

***NvSNAILB* has an incomplete first Zn Finger Domain**

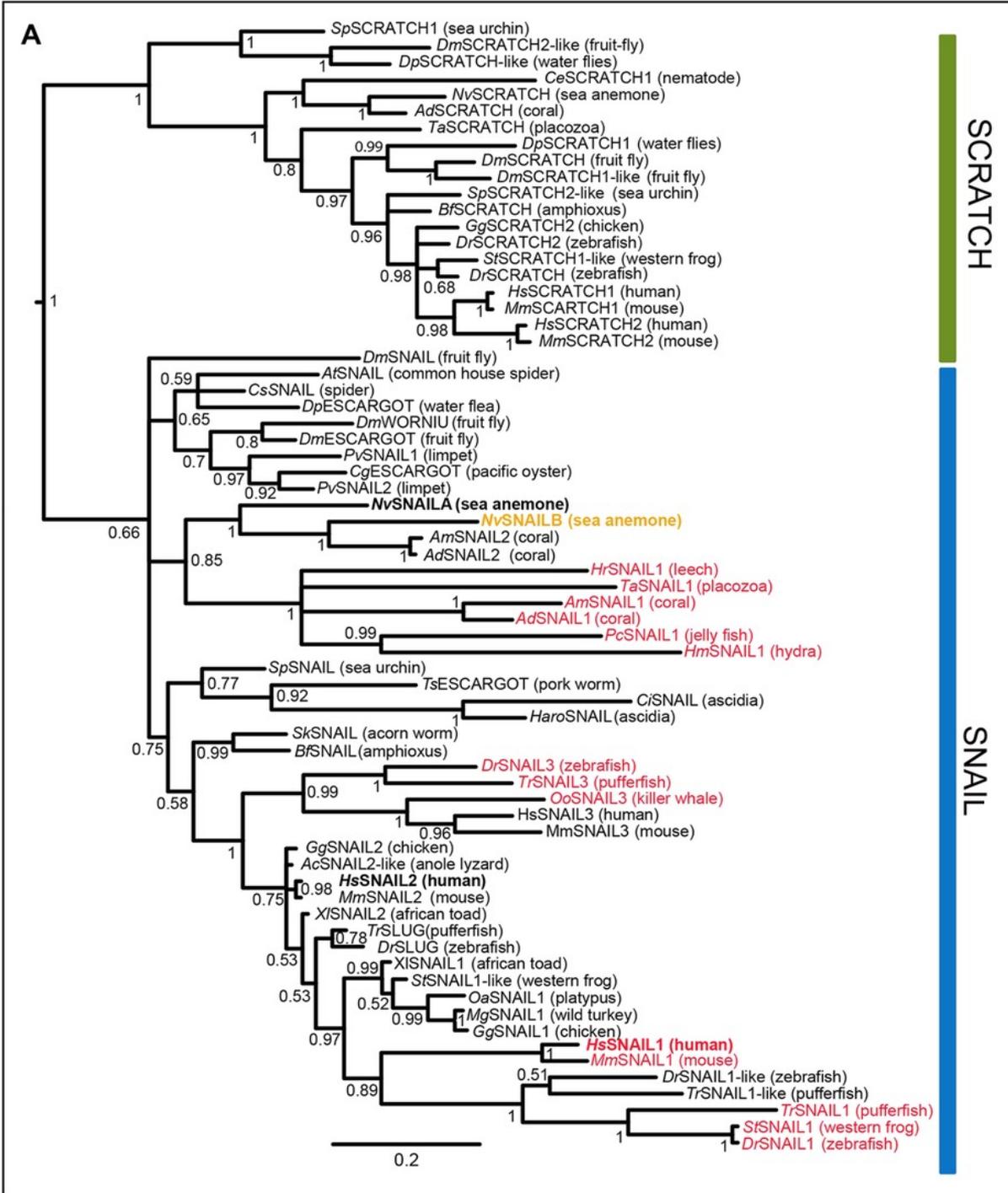
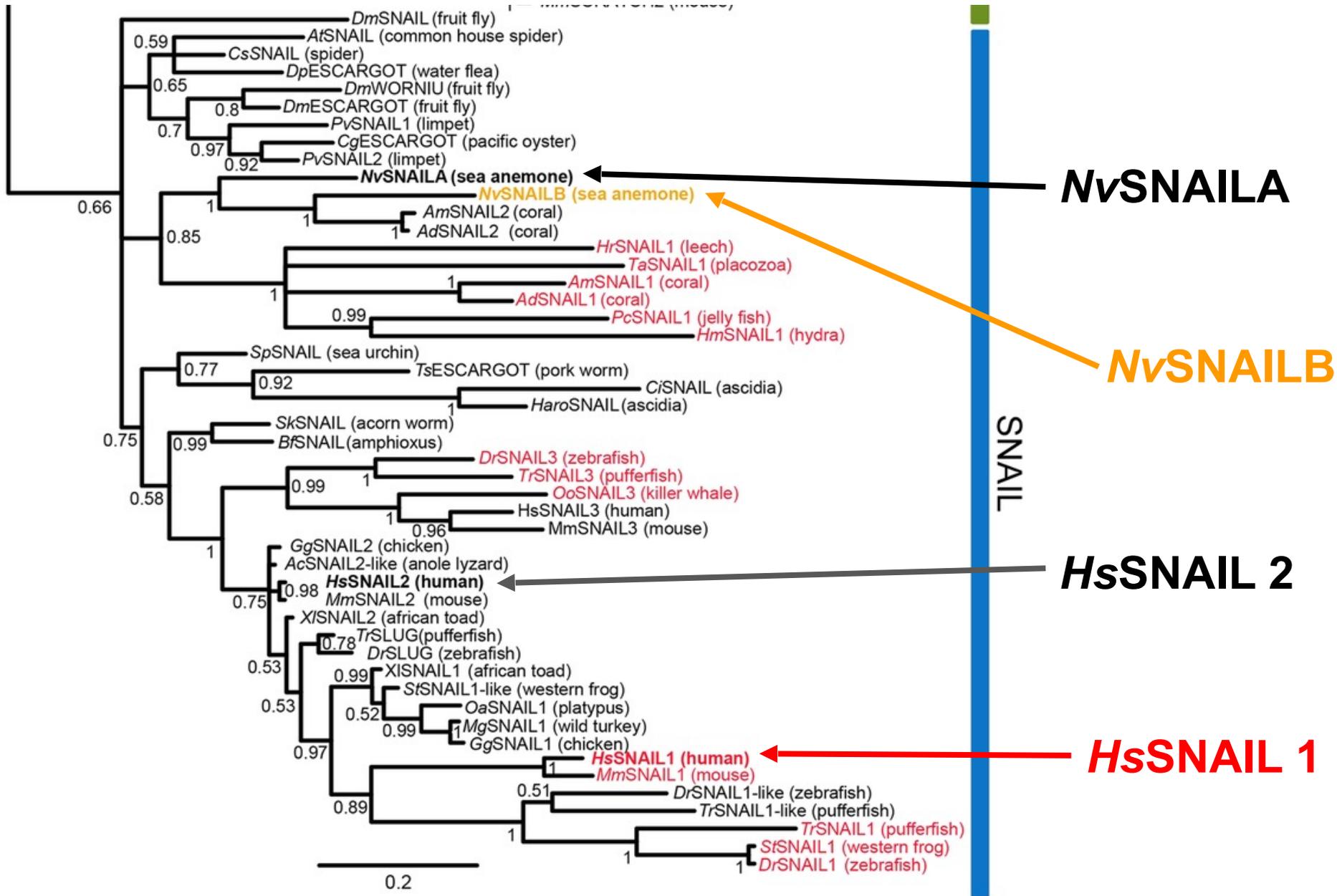


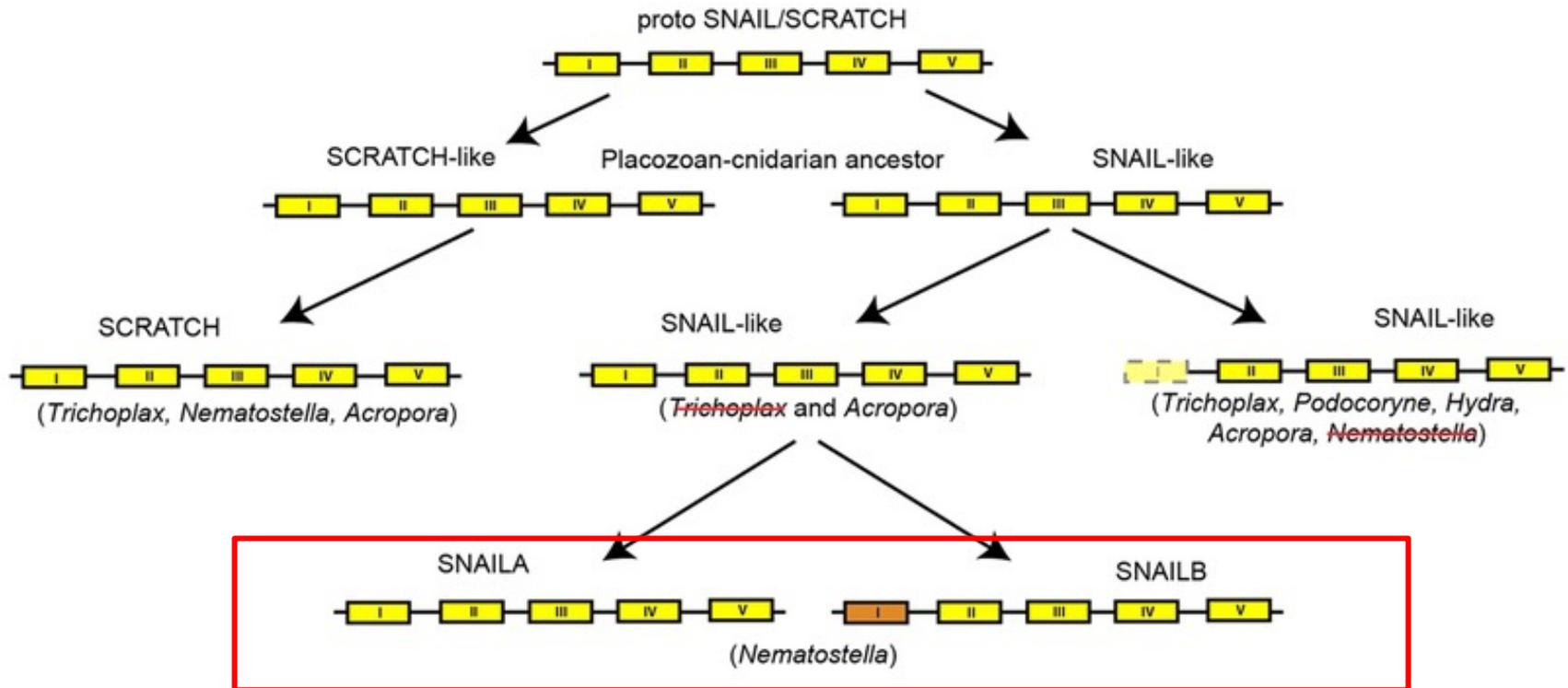
Fig 2a:
How did the
SNAIL
superfamily
evolve?



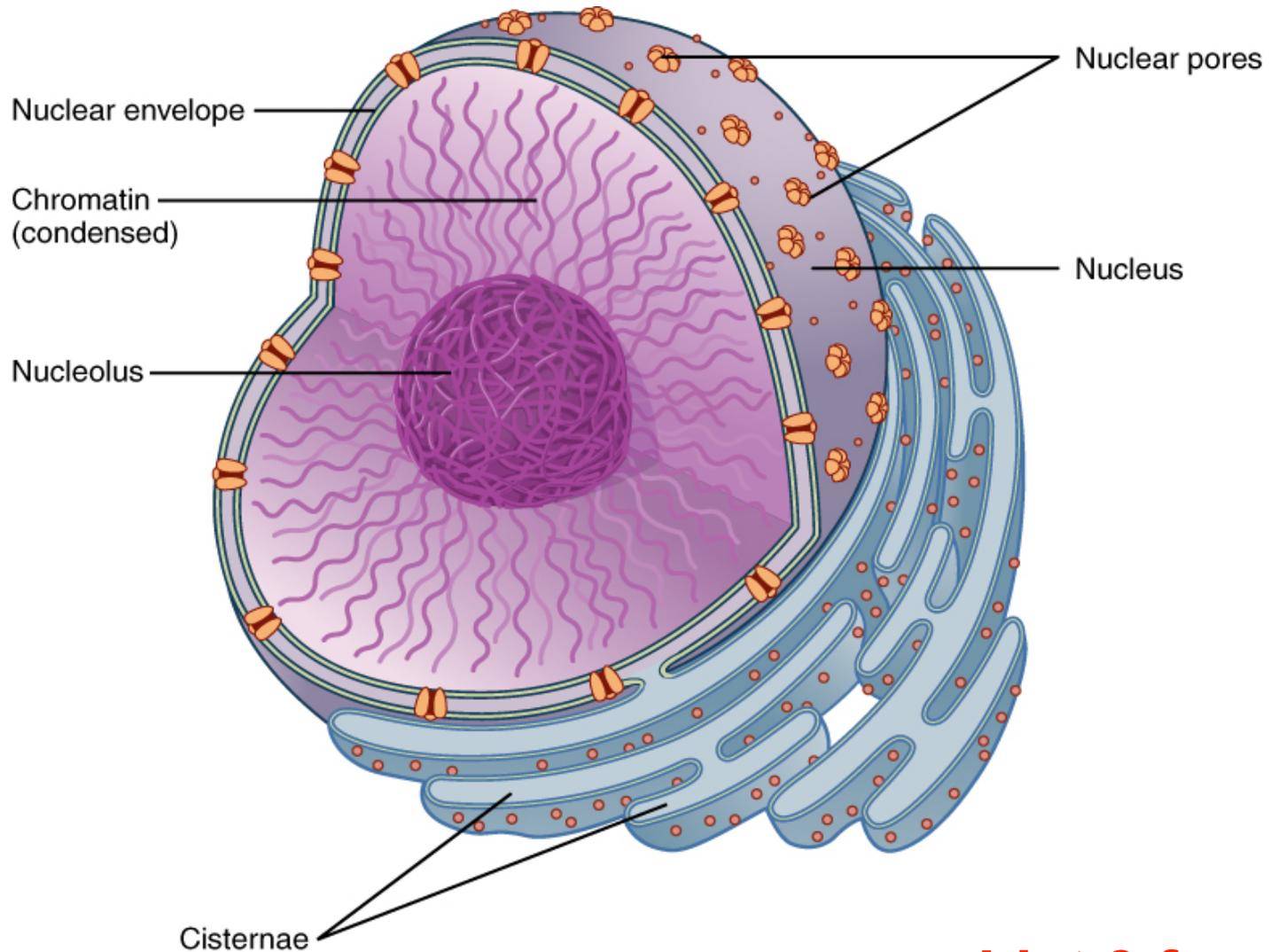
2A: How are the SNAIL proteins related?

Figure 2B: How did the two SNAIL protein types arise?

B



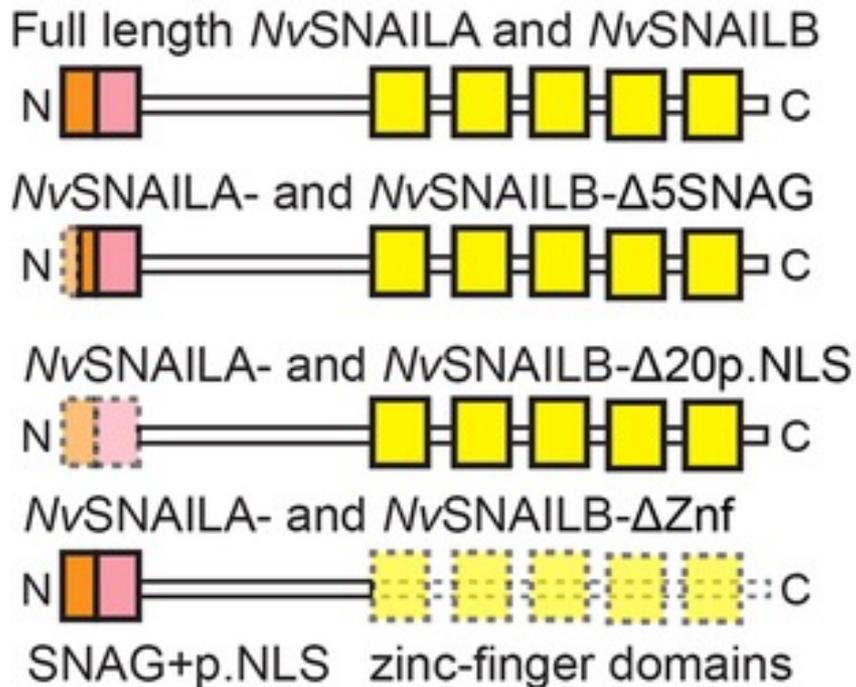
What is a nucleolus?



List 3 functions

Fig 3E: What kinds of mutants were constructed?

E



Deletion Type (Δ)

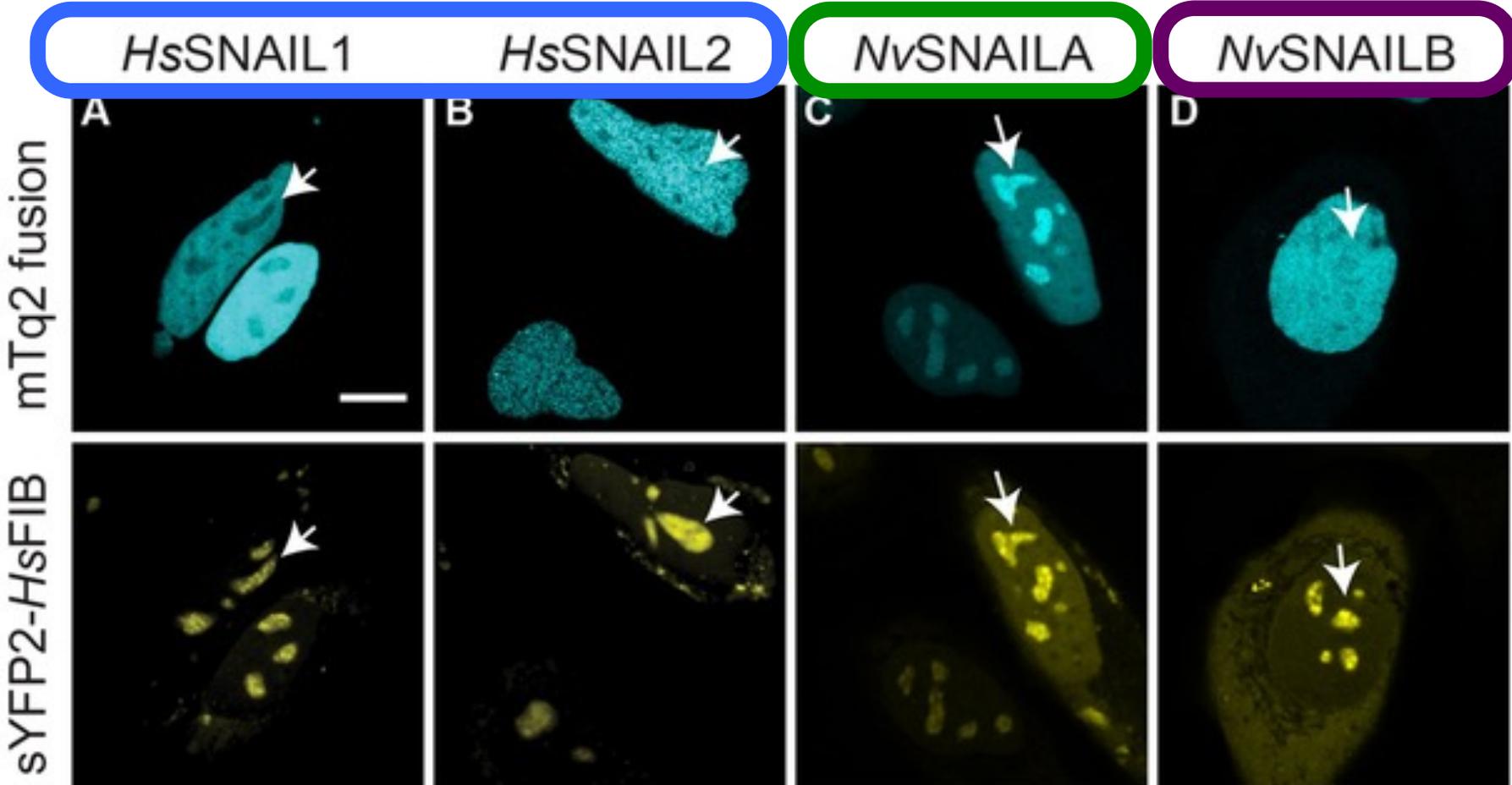
Full Length

First five AAs deleted from SNAG Domain

SNAG domain and putative NLS

All of the Zn-finger domains

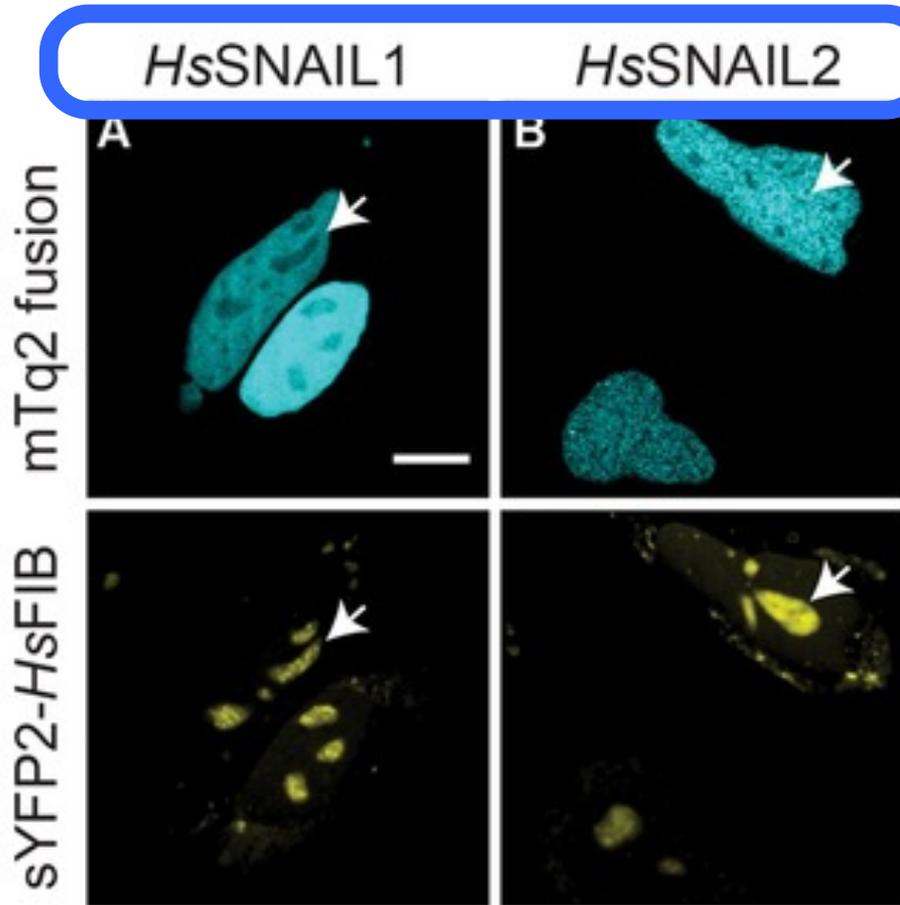
Fig 3A-D: Where does *HsSNAIL1,2* and *NvSNAILA,B* localize?



All exhibit strong nuclear localization

nuclear
nucleolar

Fig 3A-B: How do HsSNAIL1 and 2 normally Localize?



nuclear
nucleolar

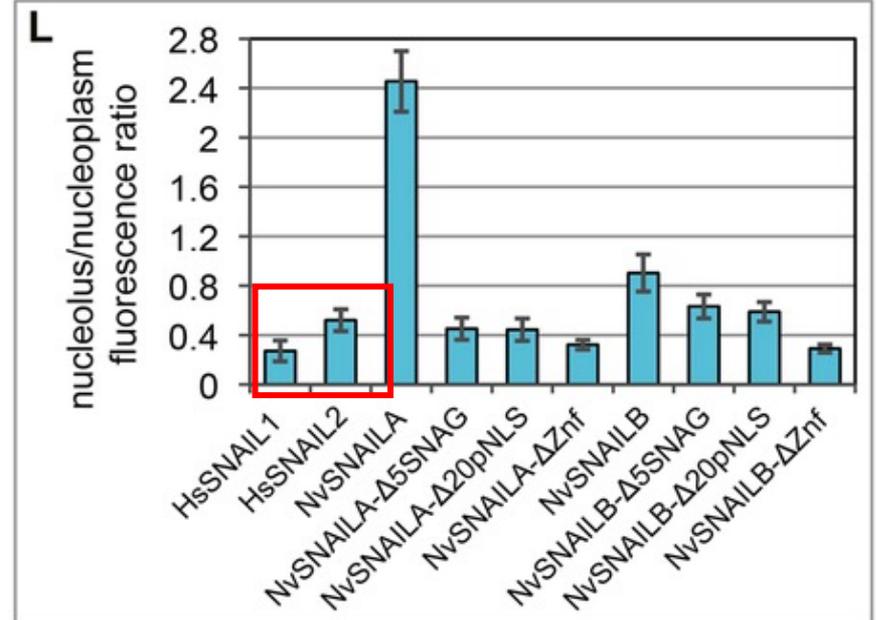
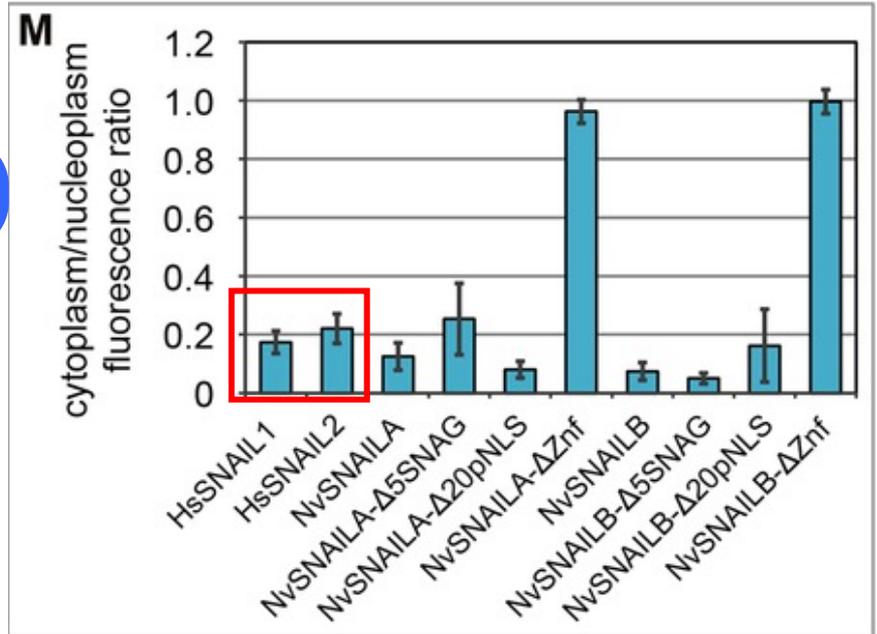
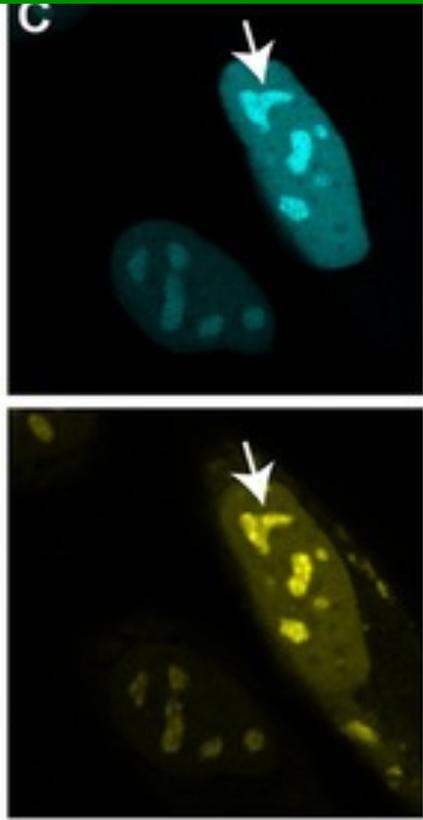


Fig 3C: How does NvSNAILA normally localize?

NvSNAILA

mTq2 fusion
sYFP2-HsFIB



nuclear
nucleolar

Full length NvSNAILA and NvSNAILB



NvSNAILA shows strong Nucleolar Localization

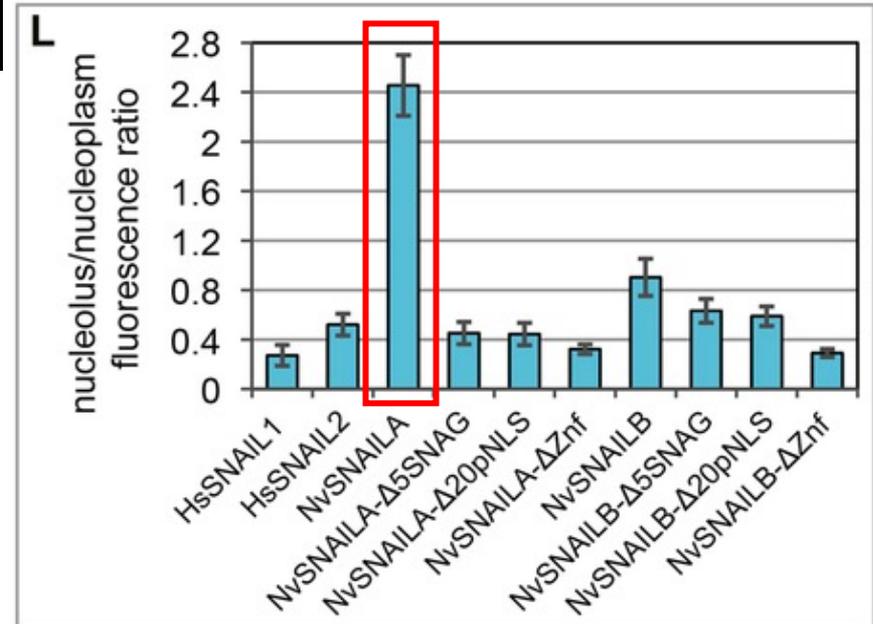
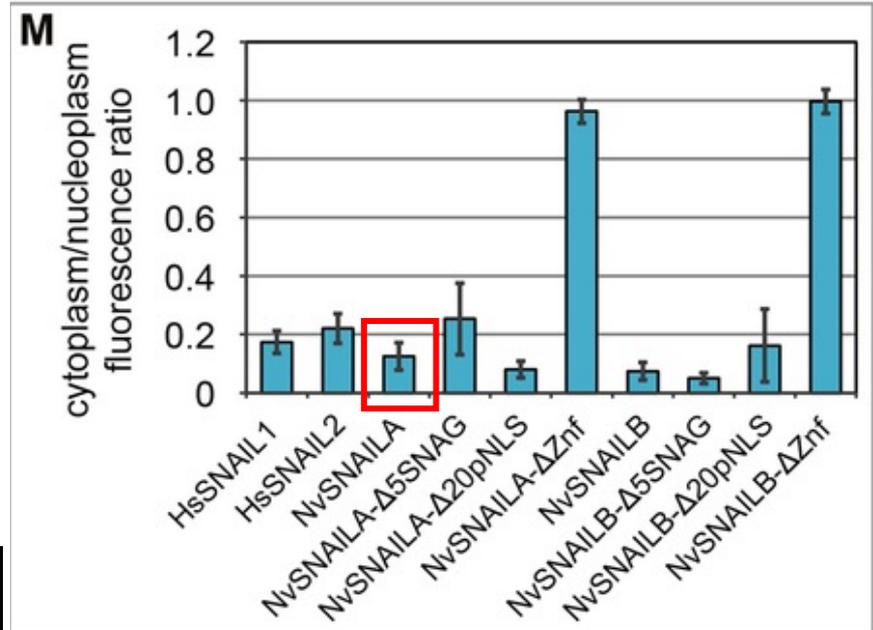
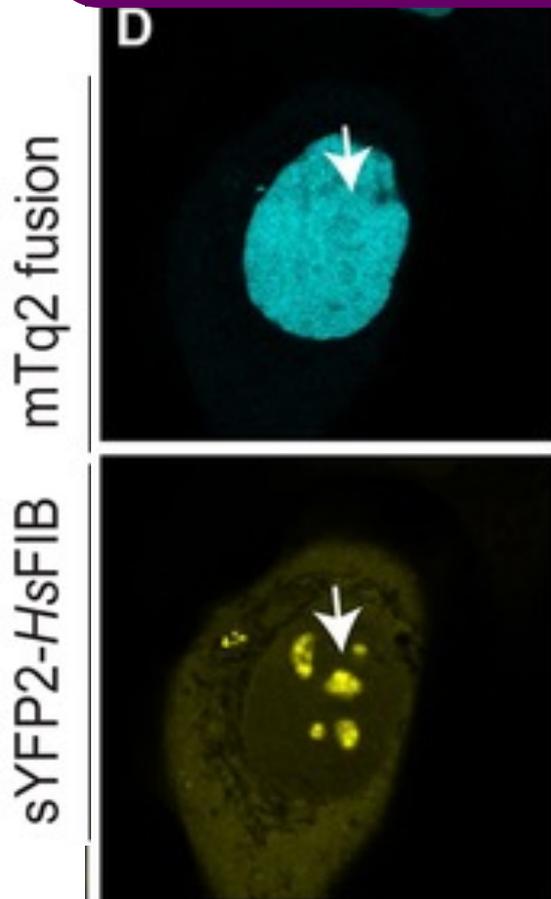


Fig 3D: How does NvSNAILB normally localize?

NvSNAILB



nuclear
nucleolar

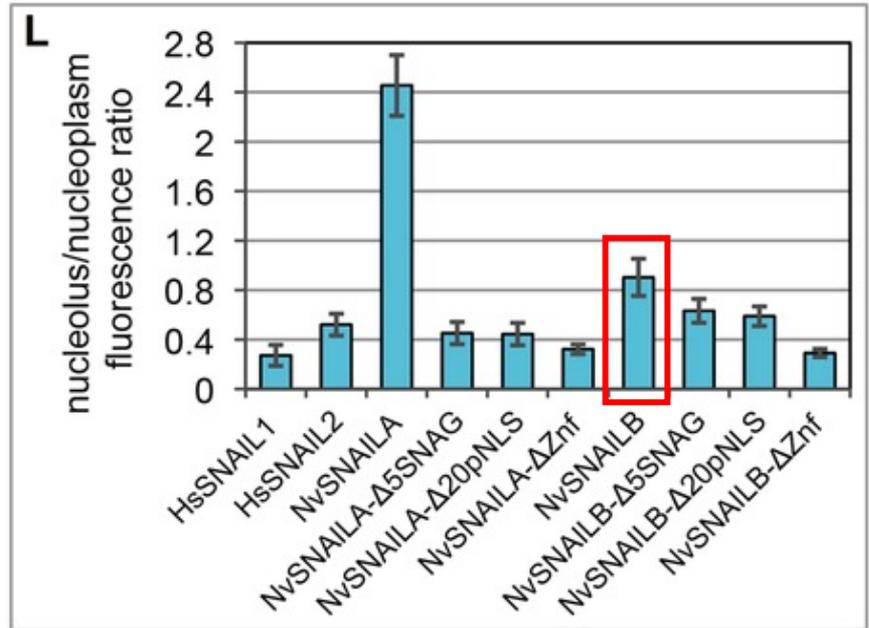
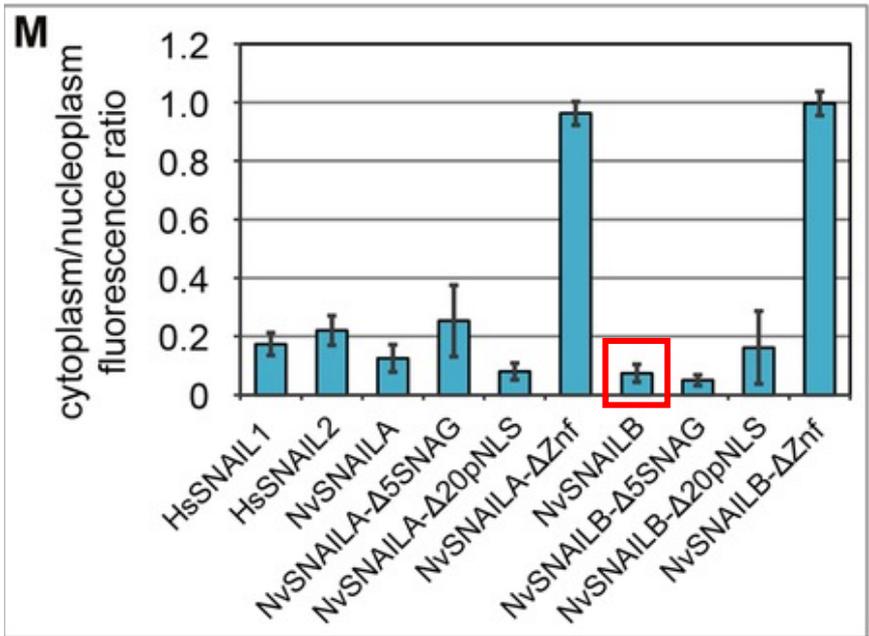
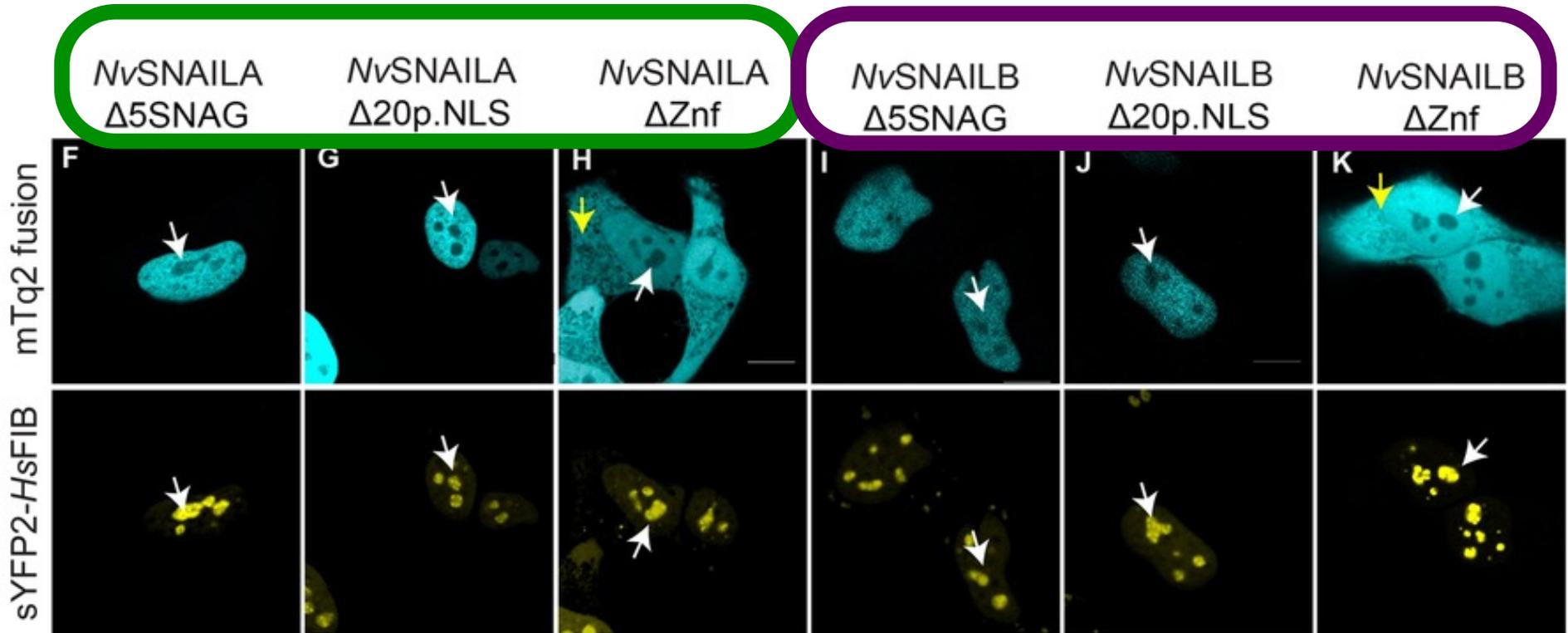


Fig 3F-K: How do the SNAIL mutants localize?

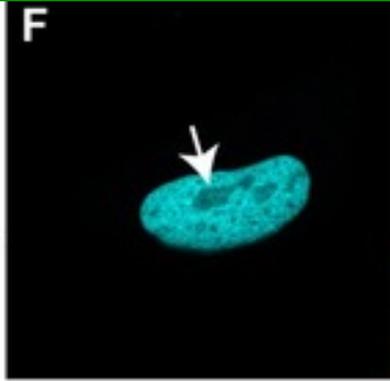


nuclear
nucleolar

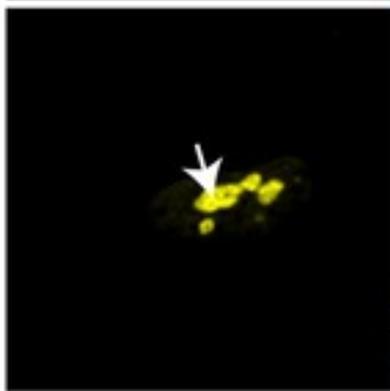
Fig 3F: How does the *NvSNAILA-Δ5SNAG* mutant localize?

NvSNAILA
Δ5SNAG

mTq2 fusion



sYFP2-HsFIB



nuclear
nucleolar

NvSNAILA- and *NvSNAILB*-*Δ5SNAG*

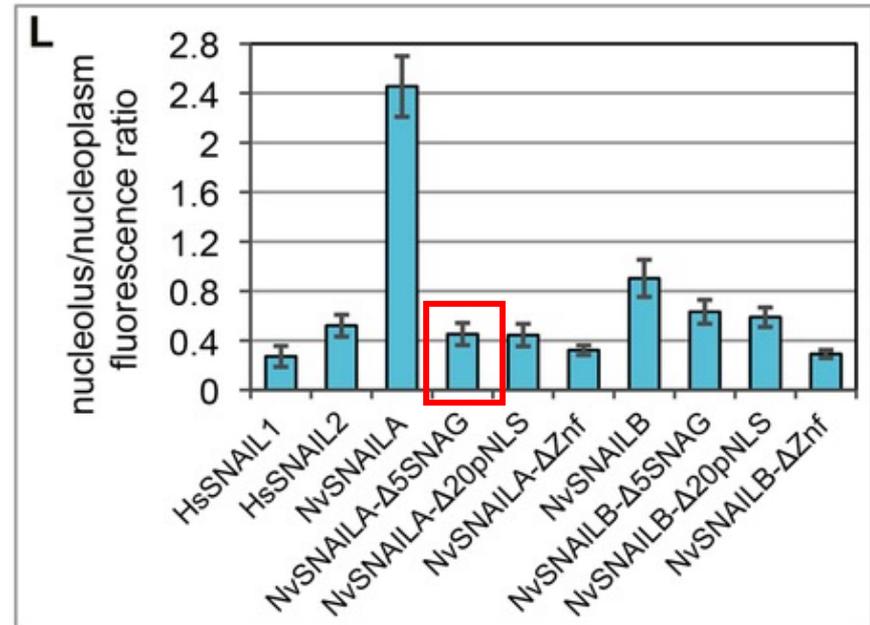
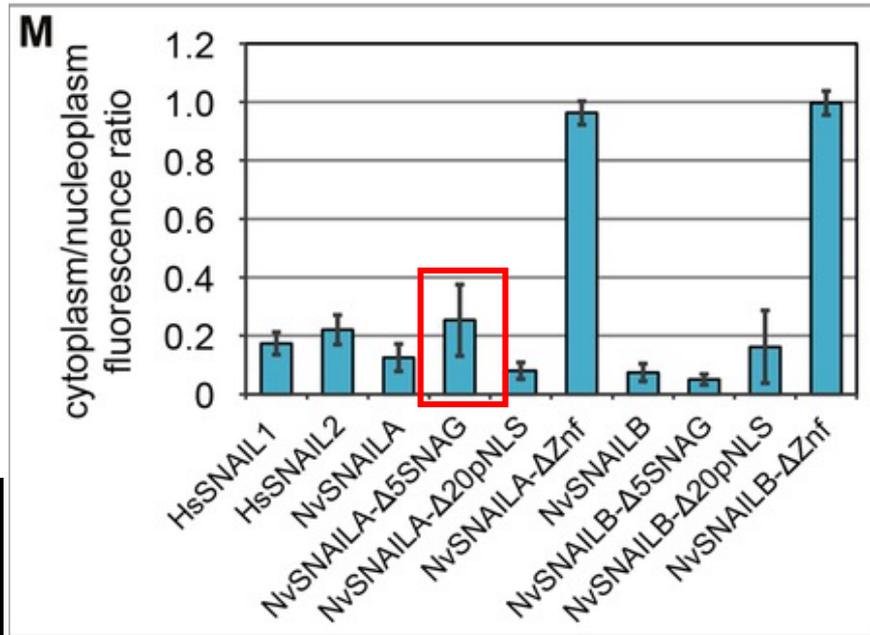
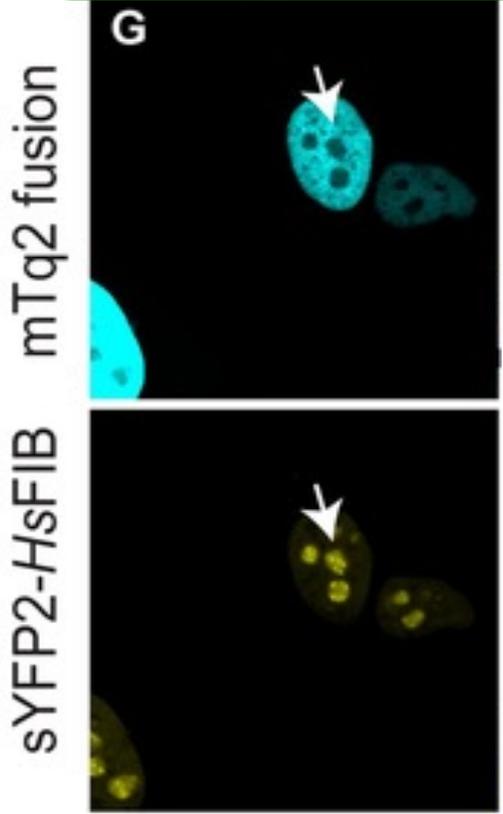


Fig 3G: How does the NvSNAILA-Δ20p.NLS mutants localize?

**NvSNAILA
Δ20p.NLS**



**nuclear
nucleolar**

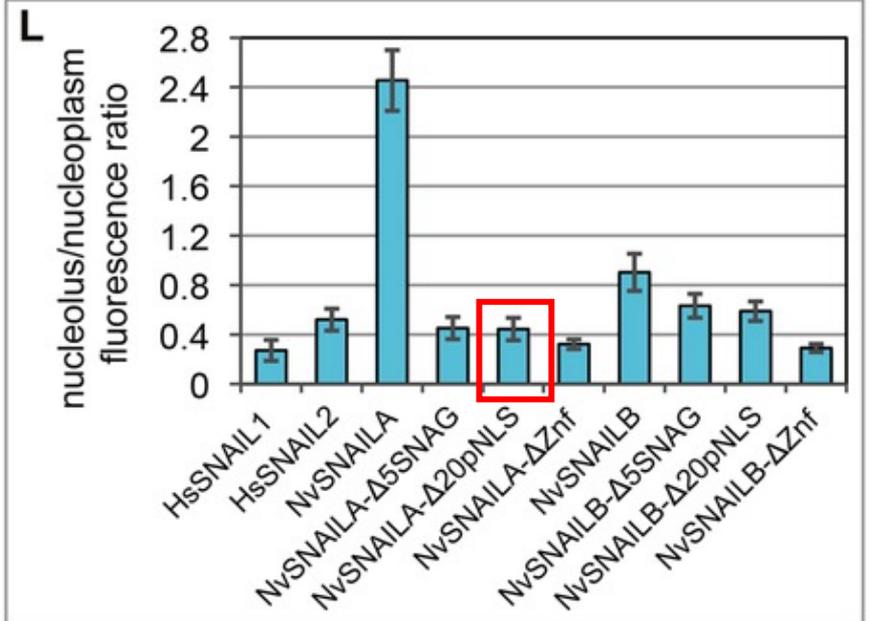
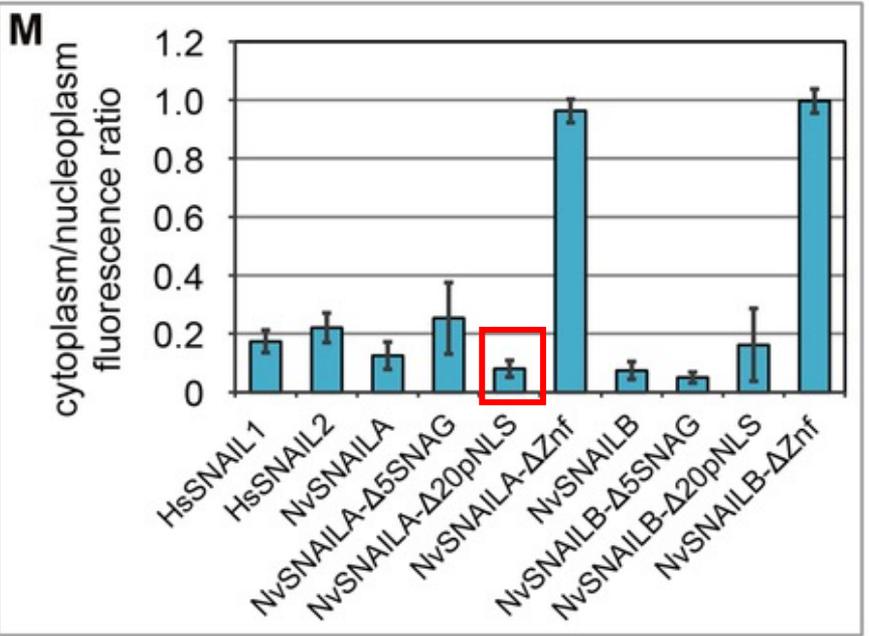
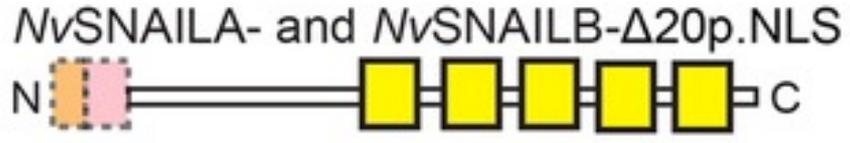
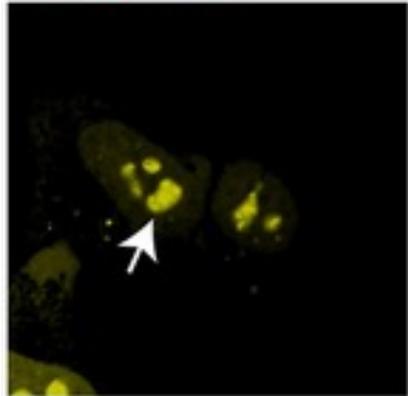
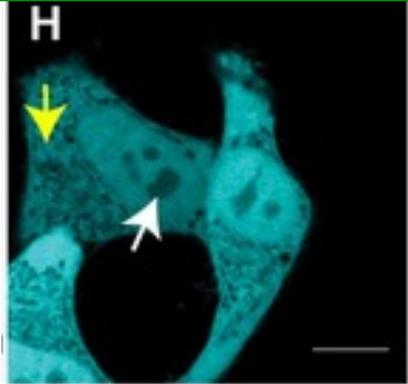


Fig 3H: How does the NvSNAILA-ΔZnf mutant localize?

**NvSNAILA
ΔZnf**

mTq2 fusion



nuclear
nucleolar

NvSNAILA- and NvSNAILB-ΔZnf

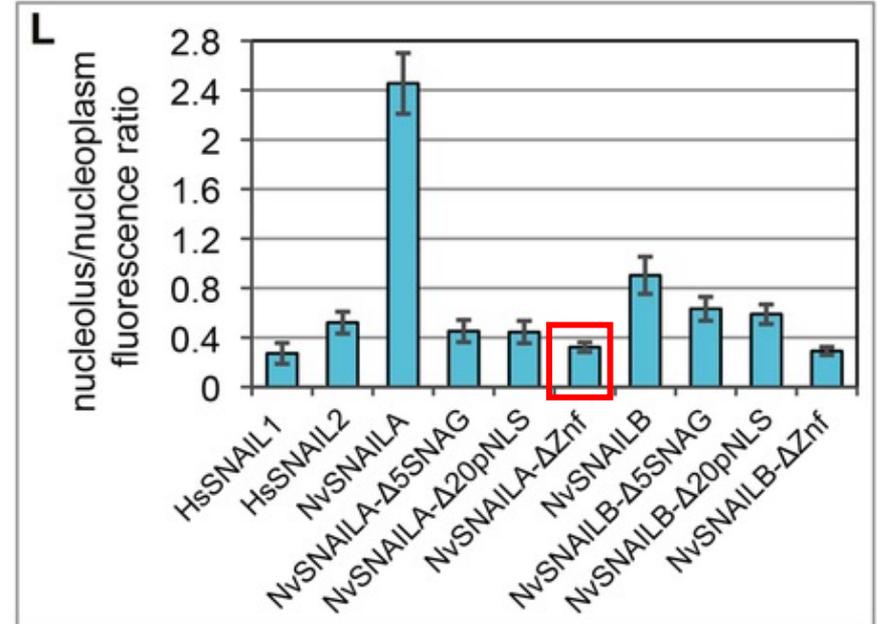
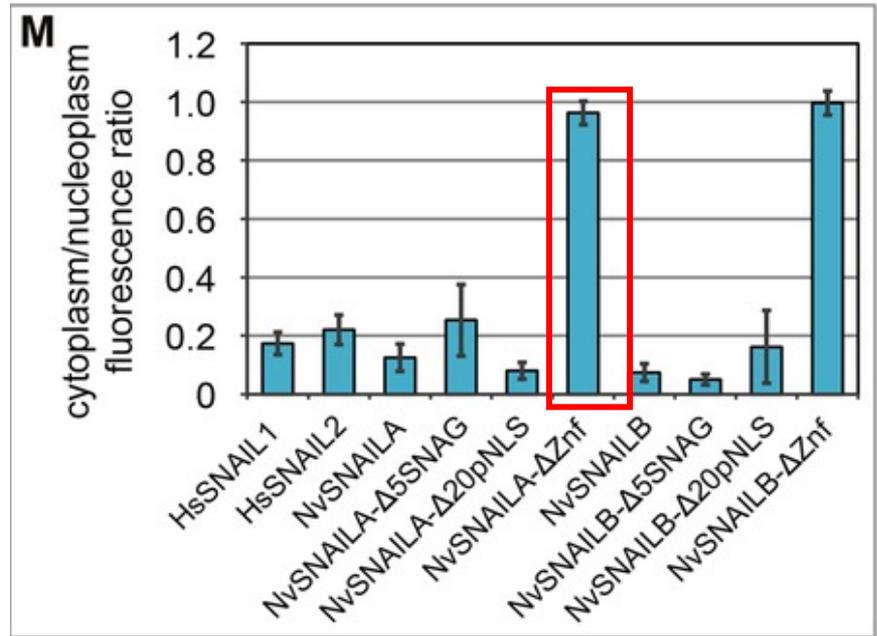
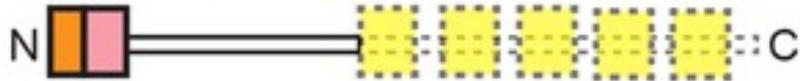
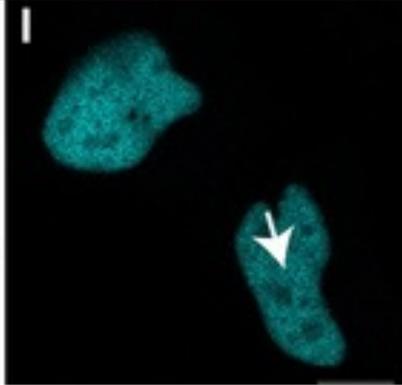


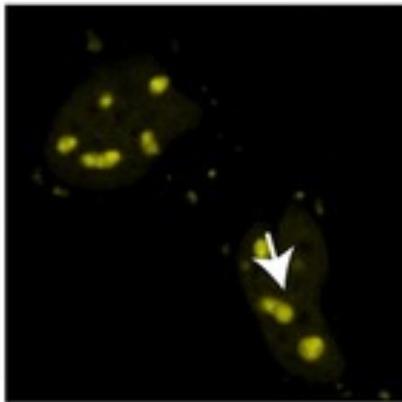
Fig 3I: How does the NvSNAILB-Δ5SNAG mutant localize?

NvSNAILB
Δ5SNAG

mTq2 fusion



sYFP2-HsFIB



nuclear
nucleolar

NvSNAILA- and *NvSNAILB*-Δ5SNAG

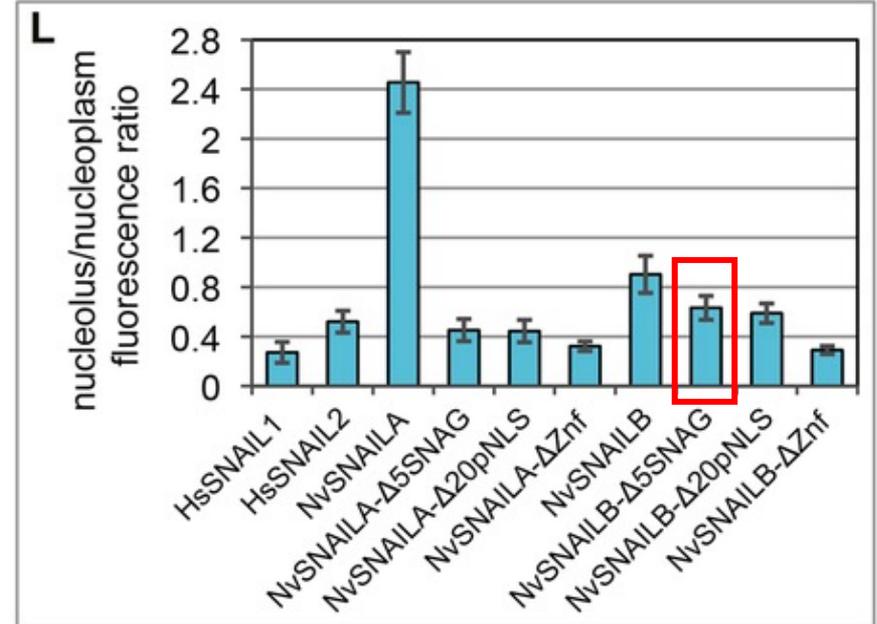
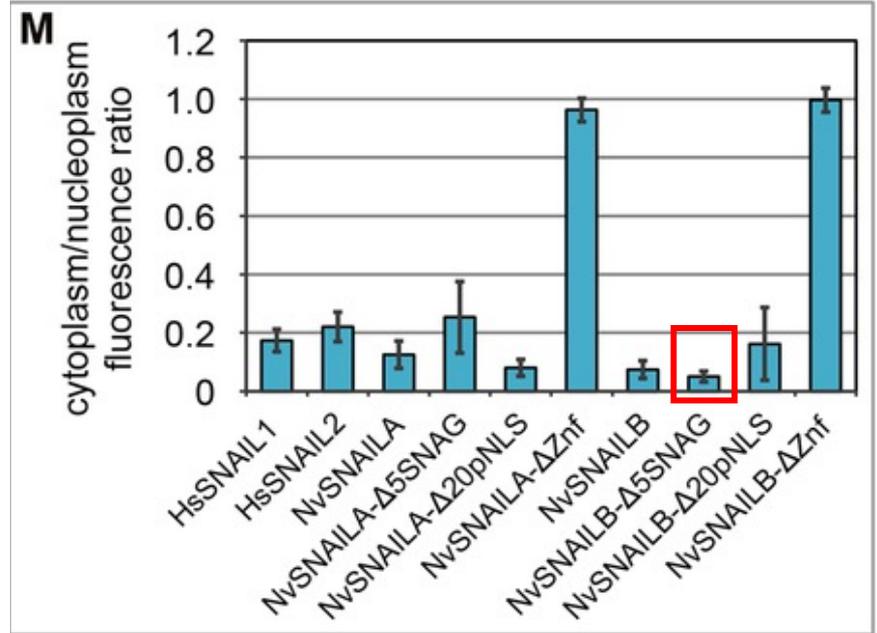


Fig 3J: How does the NvSNAILB-Δ20p.NLS mutant localize?

NvSNAILB
Δ20p.NLS

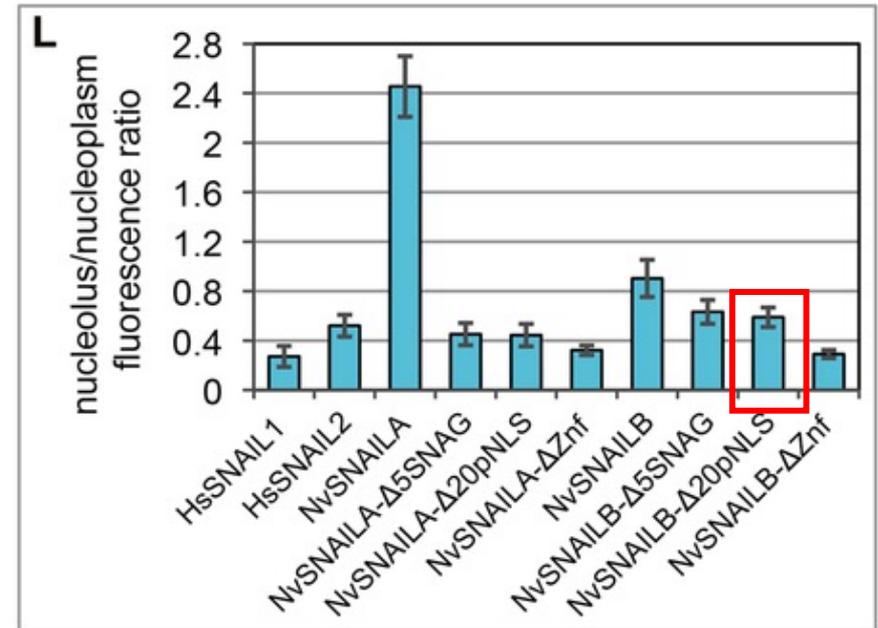
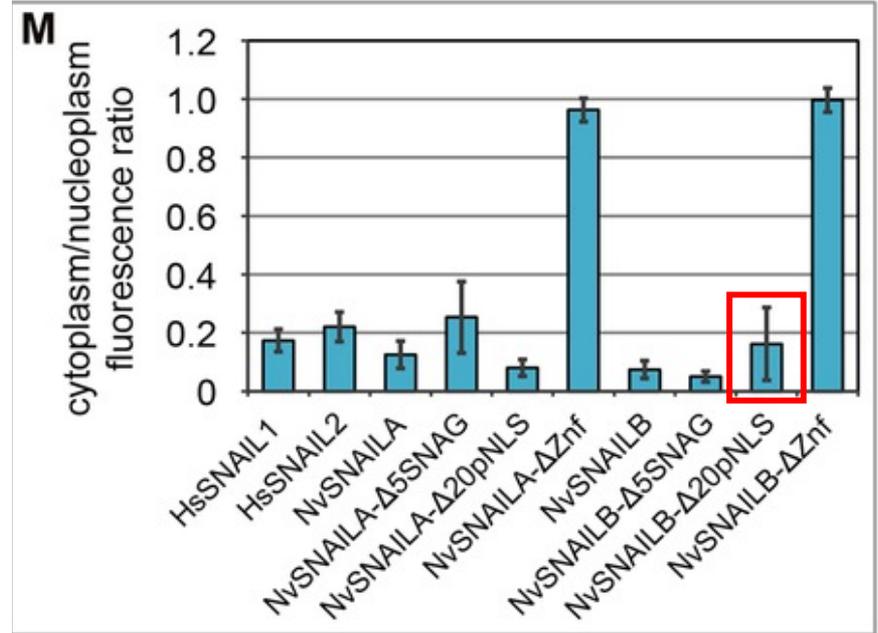
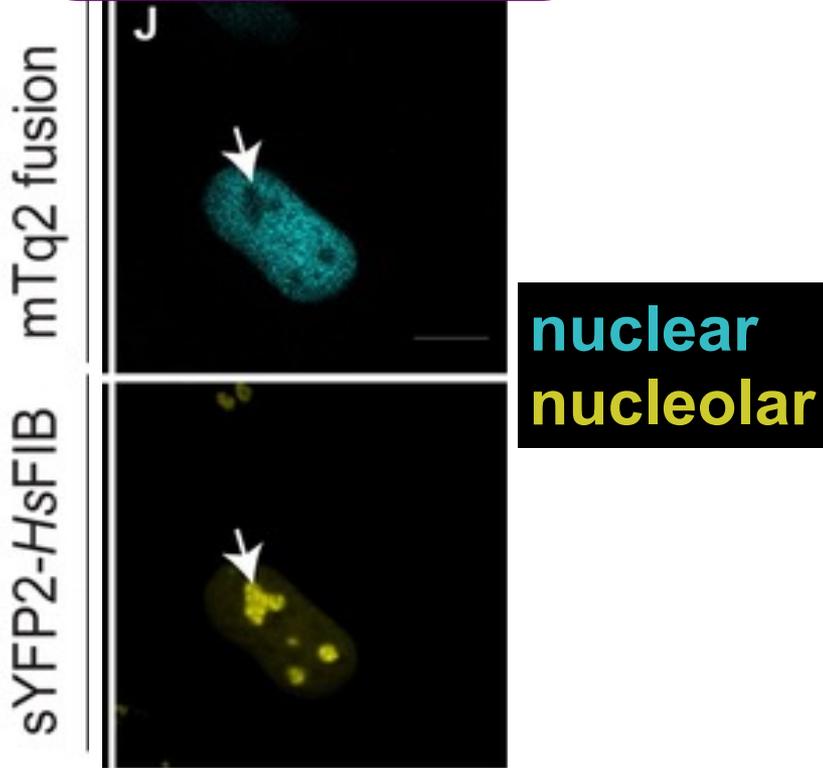
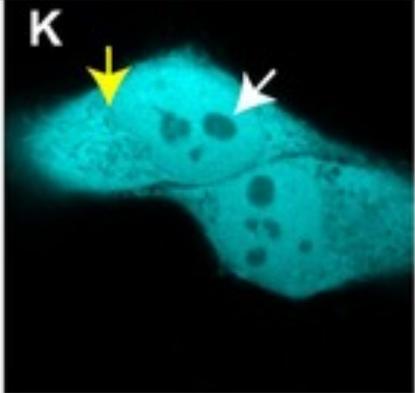


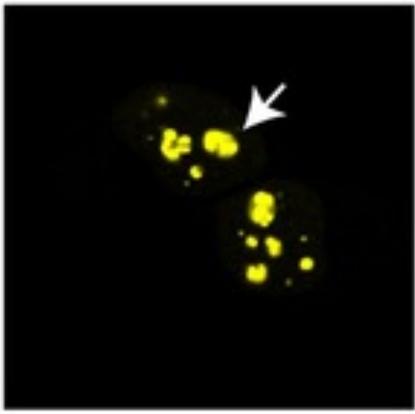
Fig 3K: How does the NvSNAILB-ΔZnf mutant localize?

NvSNAILB
ΔZnf

mTq2 fusion



sYFP2-HsFIB



nuclear
nucleolar

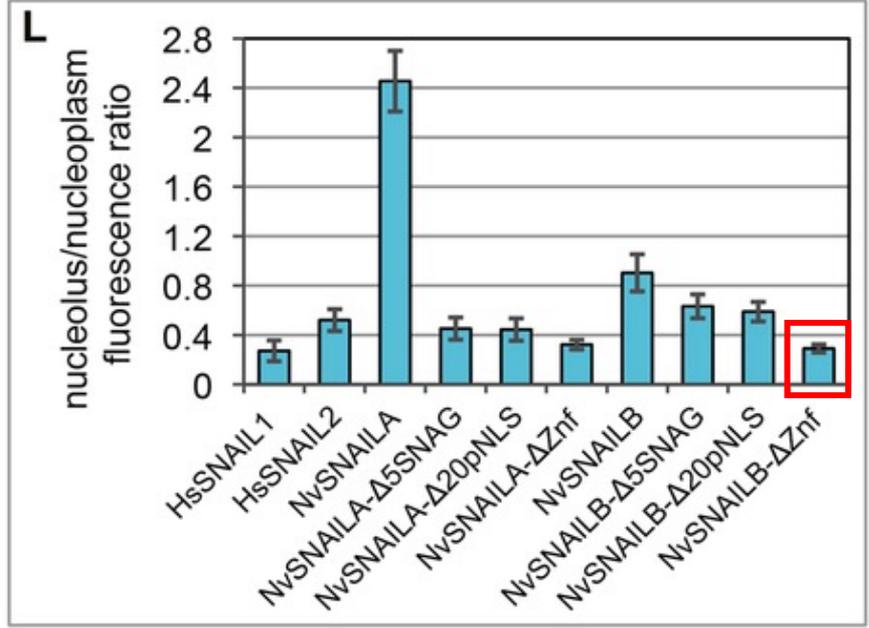
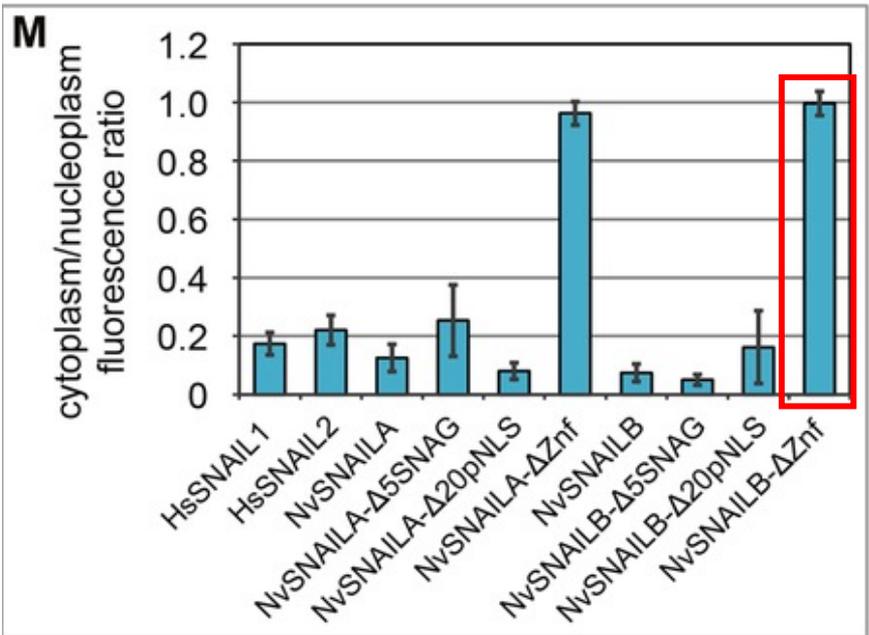
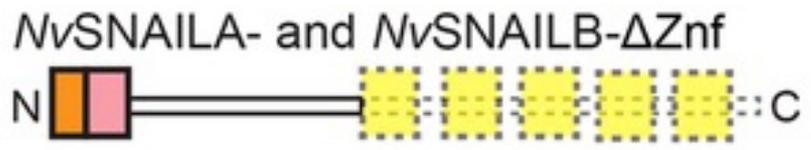
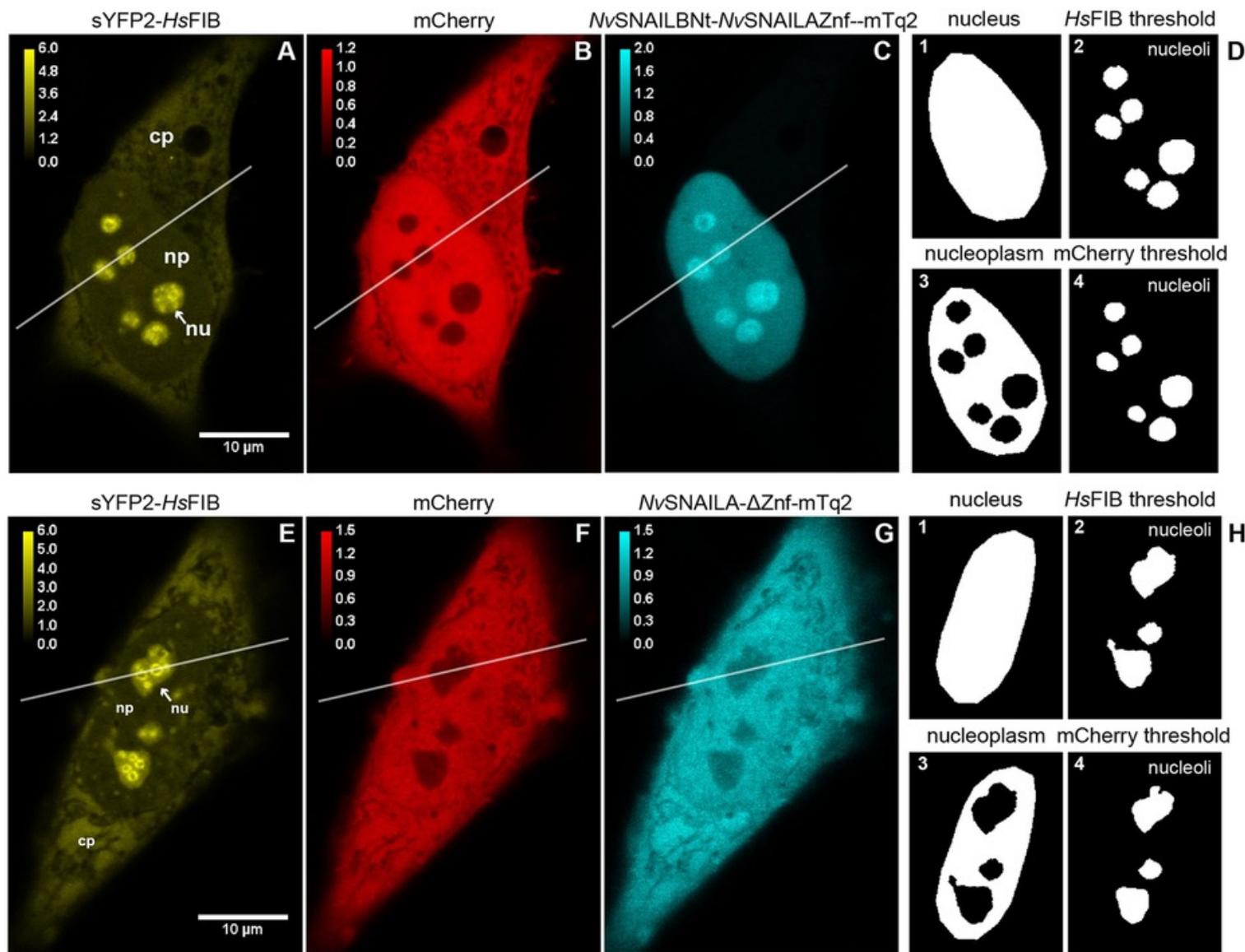
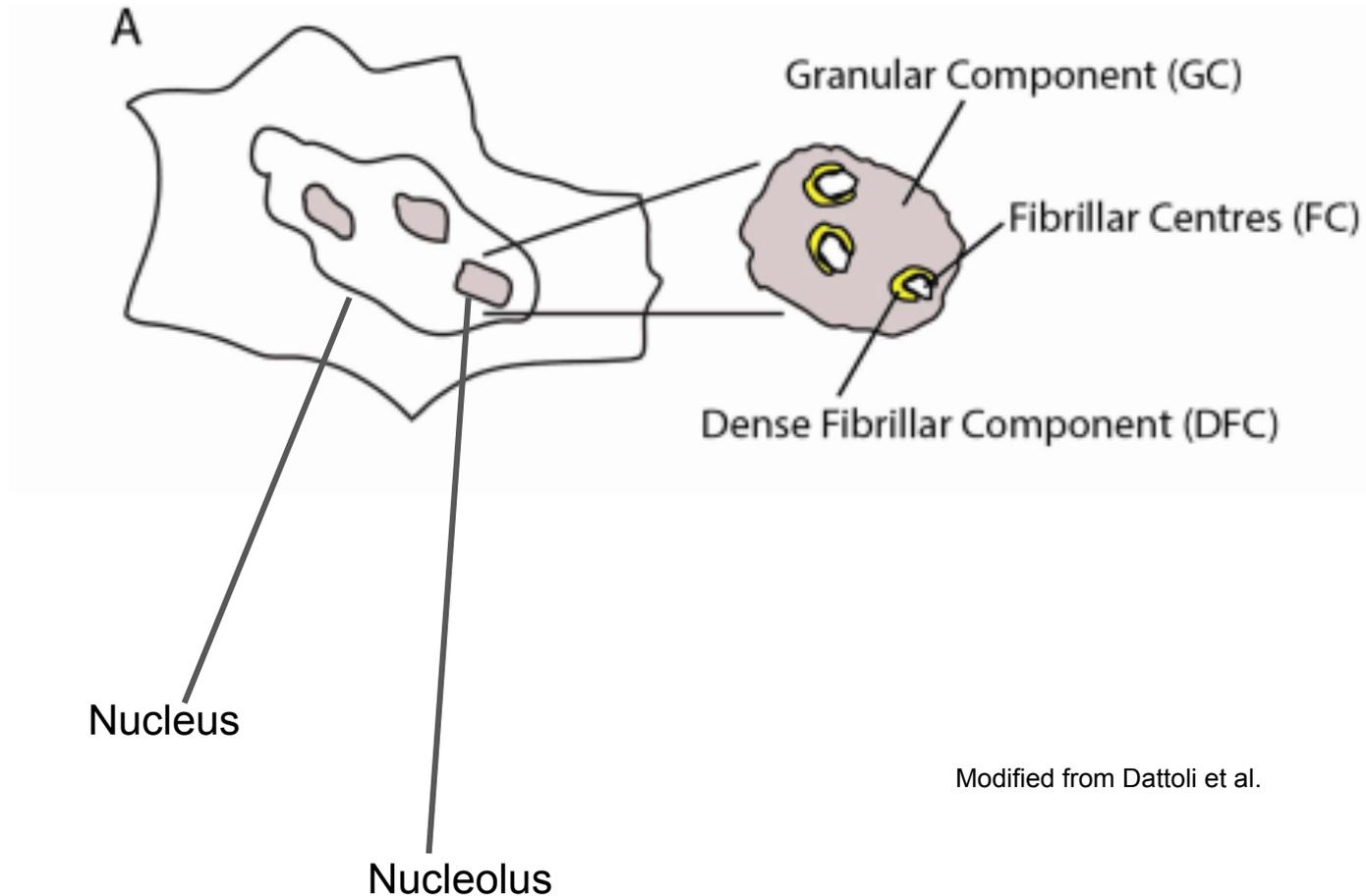


Fig 4A-G: How did they standardize fluorescence levels?



Images used to show normalized fluorescence intensity observed with the 3 different fluorophores

Fig 5A: Where specifically within the nucleolus does *NvSNAILA* localize?



Modified from Dattoli et al.

Fig 5B-K: Where does *NvSNAILA* localize specifically within the nucleolus?

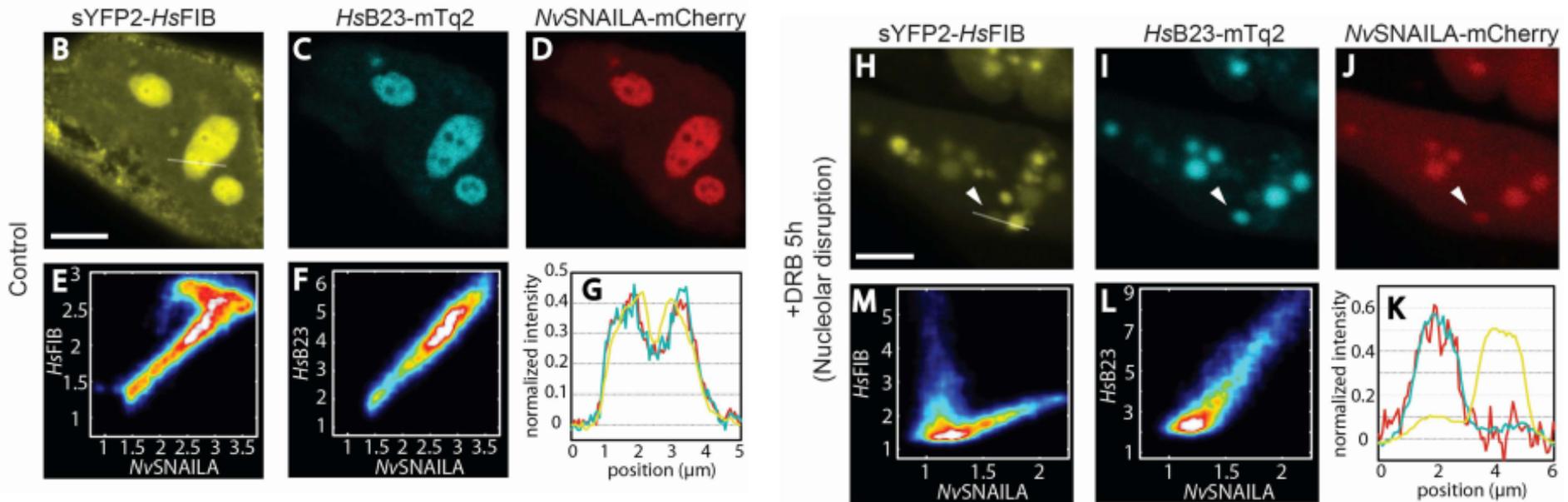


Fig 6A-D: Where does *Nv*SNAILA and B localize in *Nematostella vectensis* embryos?

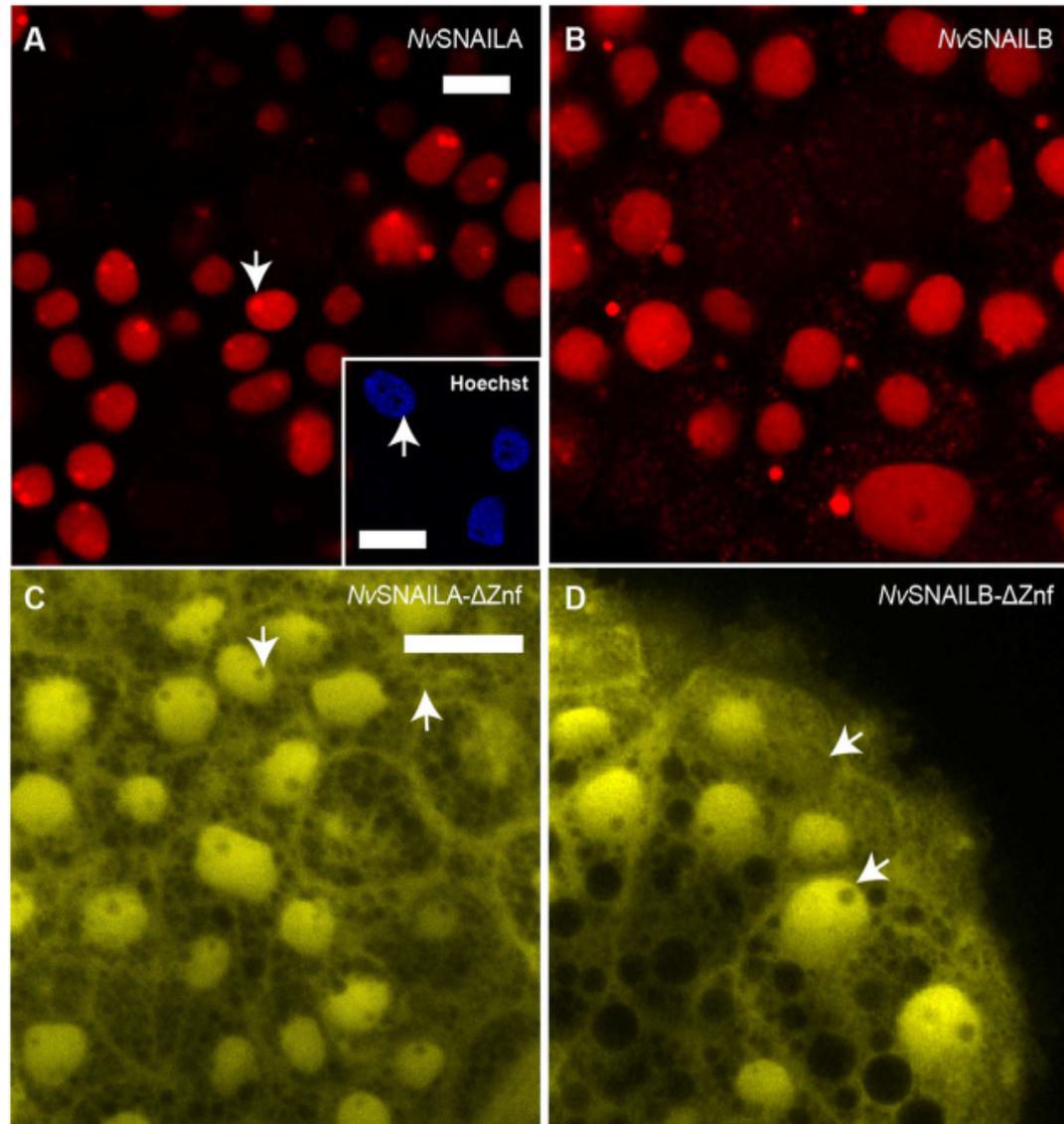
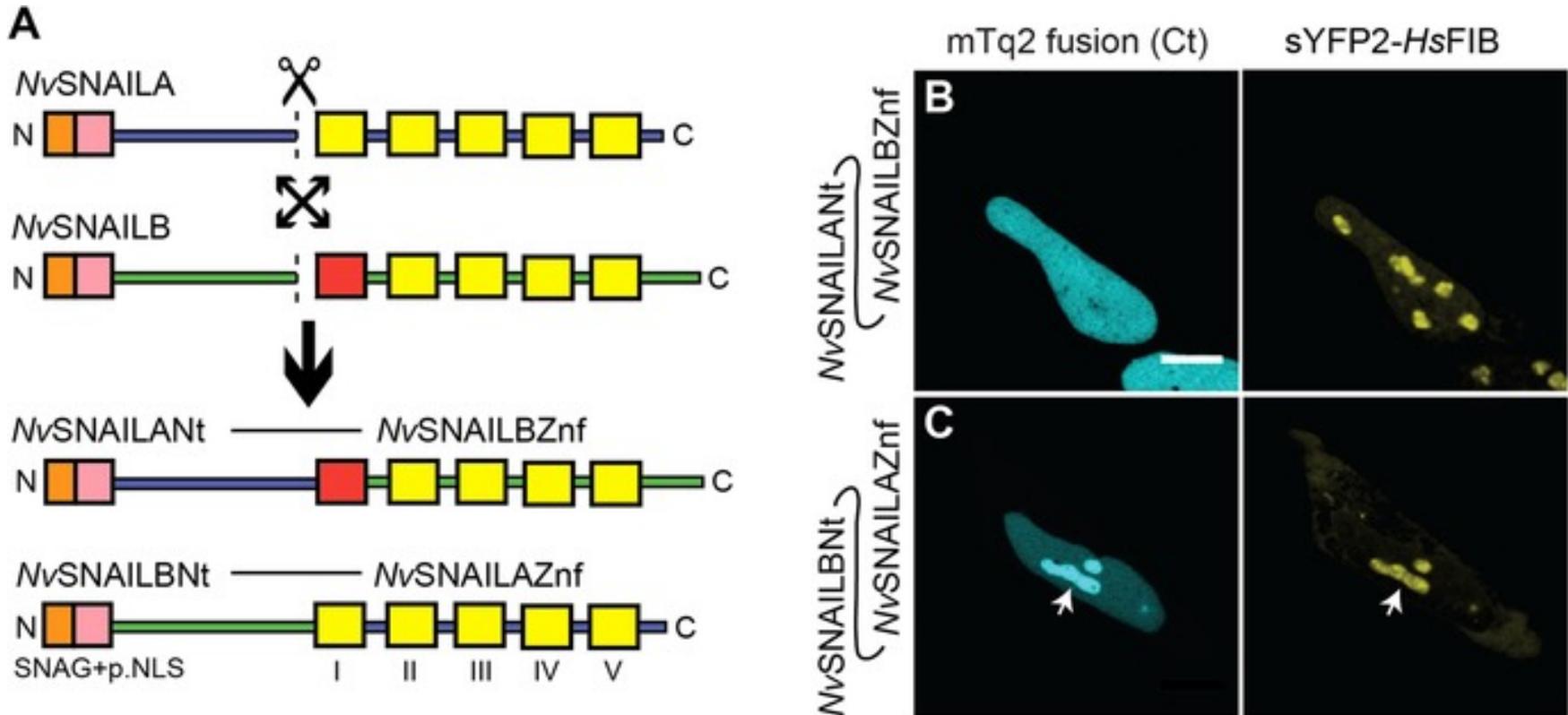


Fig 7A-C: How were they sure that *NvSNAILA*'s localization capacity was due to the Zinc-finger domains?



nuclear
nucleolar

Fig 7D-E: How were they sure that *NvSNAILA*'s localization capacity was due to the Zinc-finger domains?

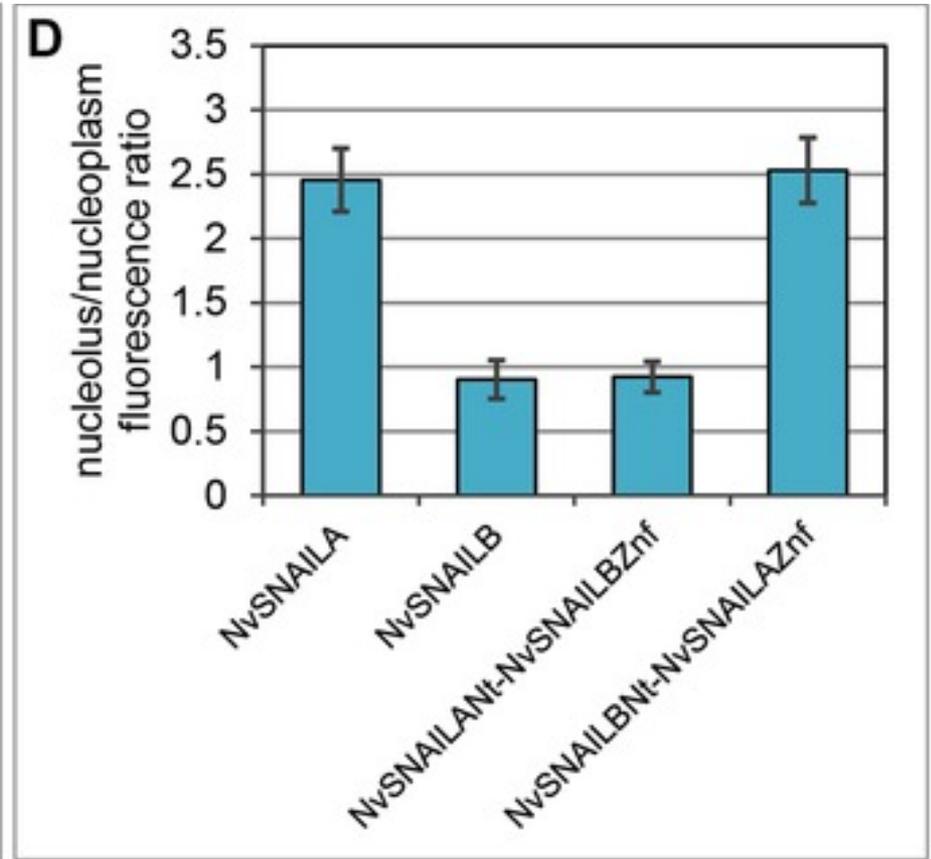
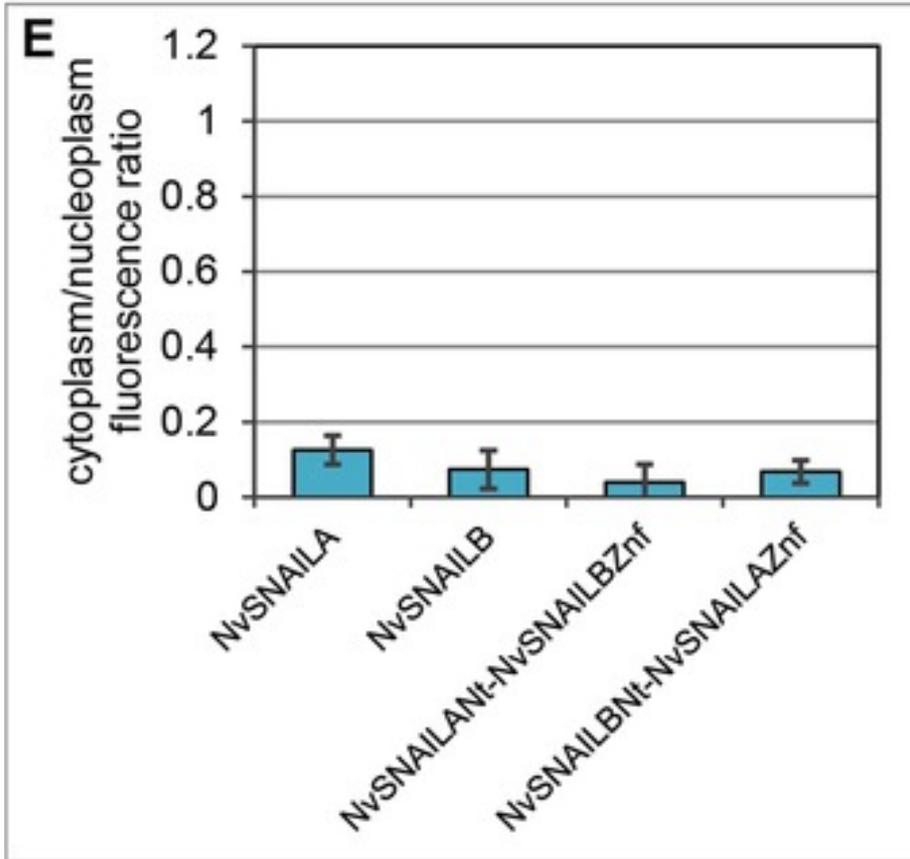
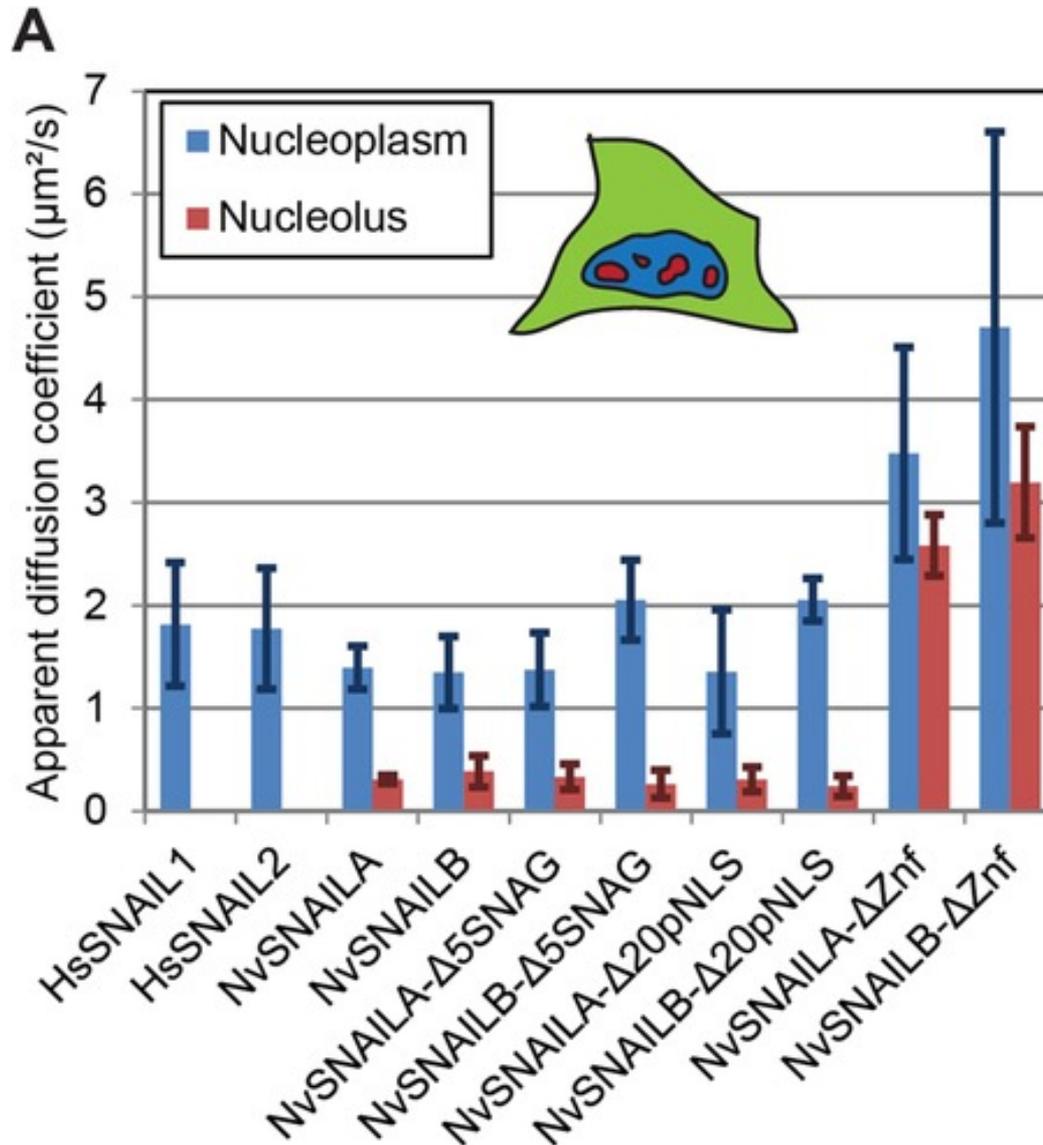
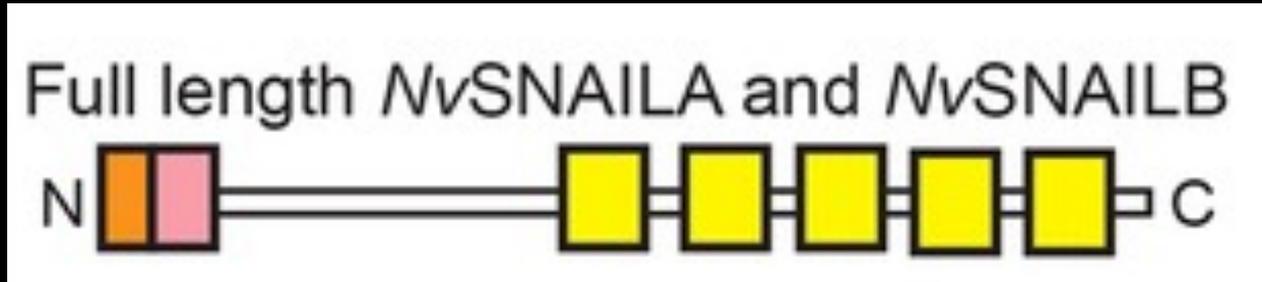


Fig 8A-B: How do the SNAIL protein domains confer function?



In summary

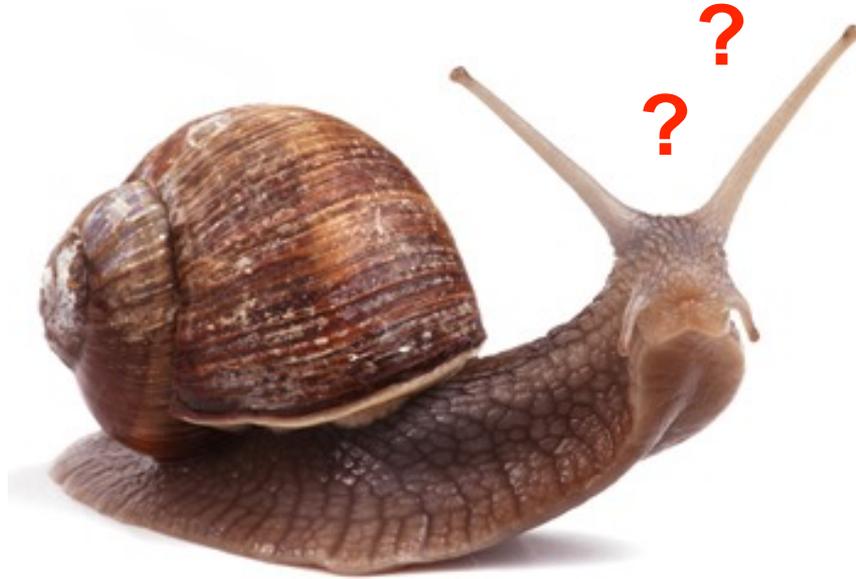


Protein domains are conserved structural and functional regions that can be analyzed with programs such as **Pfam** and **SMART**.

Zn Finger domain loss may have happened multiple times throughout evolutionary history.

The **Zinc Finger domains** play a role in **nuclear** localization, while putative **NLS** does not.

The first **Zinc Finger Domain** of *Nv*SNAILA confers strong **nucleolar** localization capability.



Further Questions?