

# **Transcriptomics-Based Screening for Pharmacological Treatments to Defer Aging**

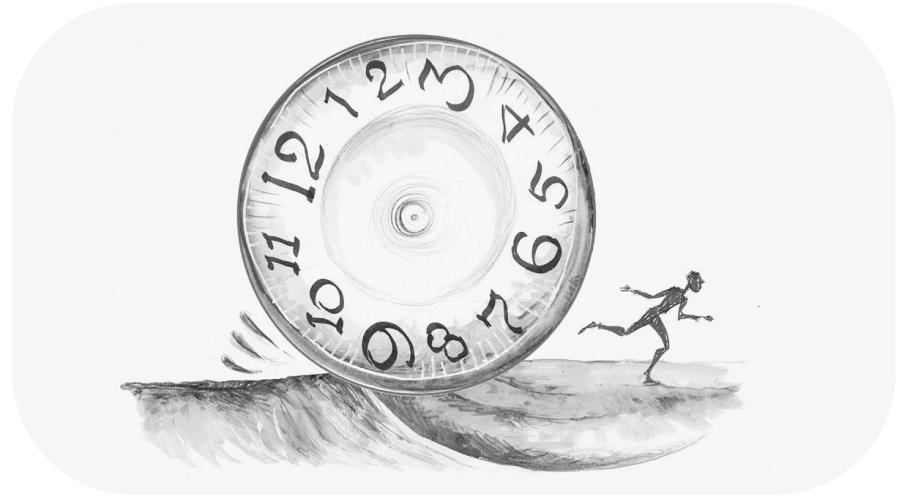
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Kashuk E., Kubenko K., Vaulin. N.



# Background

- **Aging is** associated human **morbidity** and mortality and there are no way to revert it



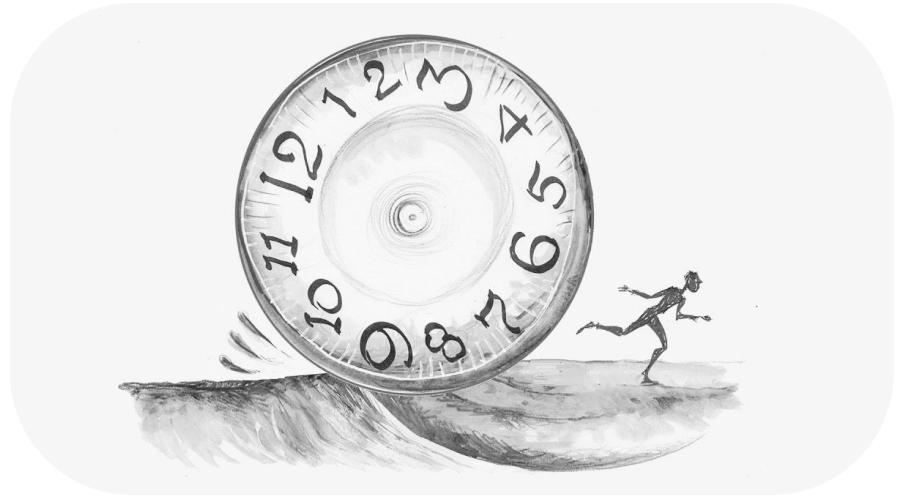
# Background

- **Aging is** associated human **morbidity** and mortality and there are no way to revert it
- However, scientists search for treatments that can **slow down aging**

**Magic pill**



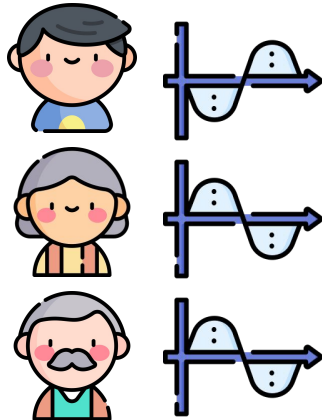
**Aging**



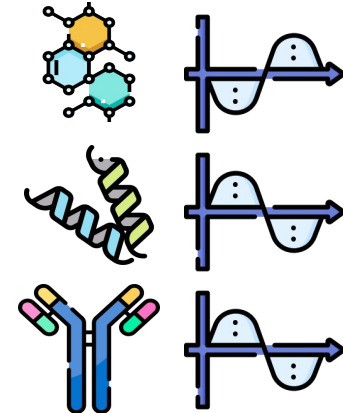
# Background

## 2 transcriptomics datasets

Their own RNA-seq  
~ **Age**



Connectivity Map (CMap)  
~ **Treatment**

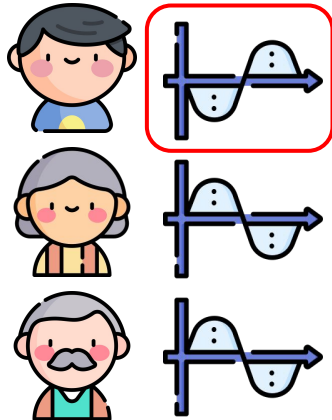


**Transcriptomics-Based Screening Identifies Pharmacological Inhibition of Hsp90 as a Means to Defer Aging**

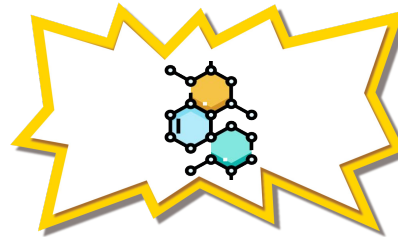
# Background

## 2 transcriptomics datasets

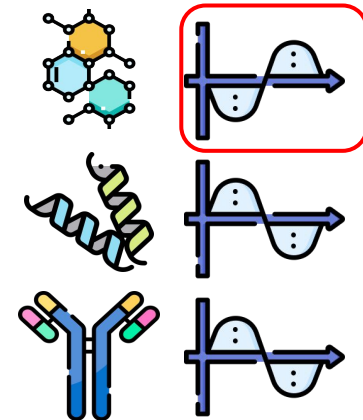
Their own RNA-seq  
~ **Age**



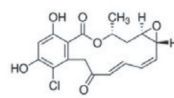
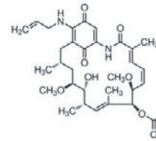
Anti-aging drug



Connectivity Map (CMap)  
~ **Treatment**



## Transcriptomics-Based Screening Identifies Pharmacological Inhibition of Hsp90 as a Means to Defer Aging



HSP90

**E**

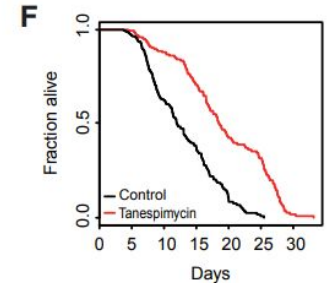
Fraction alive

Days

— Control

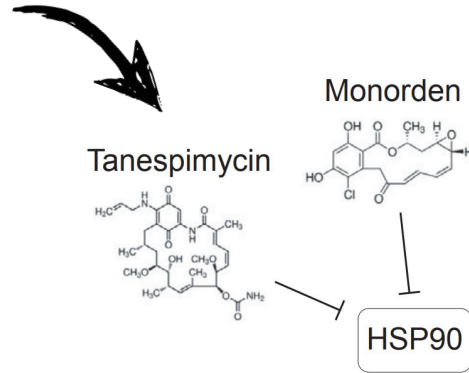
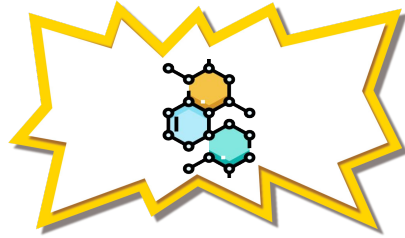
— Monorden

Days	Control (Fraction alive)	Monorden (Fraction alive)
0	1.0	1.0
10	0.9	0.95
20	0.7	0.9
25	0.4	0.8
30	0.0	0.5
35	0.0	0.2
40	0.0	0.0



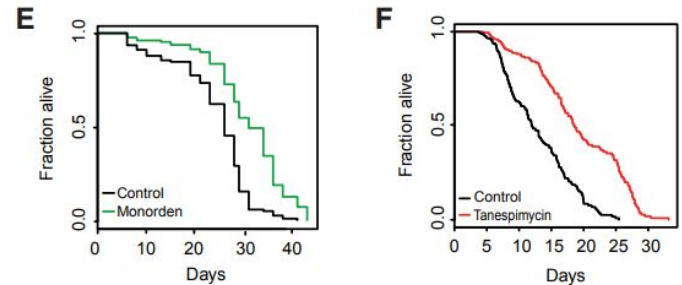
## Transcriptomics-Based Screening Identifies Pharmacological Inhibition of Hsp90 as a Means to Defer Aging

# Spoiler

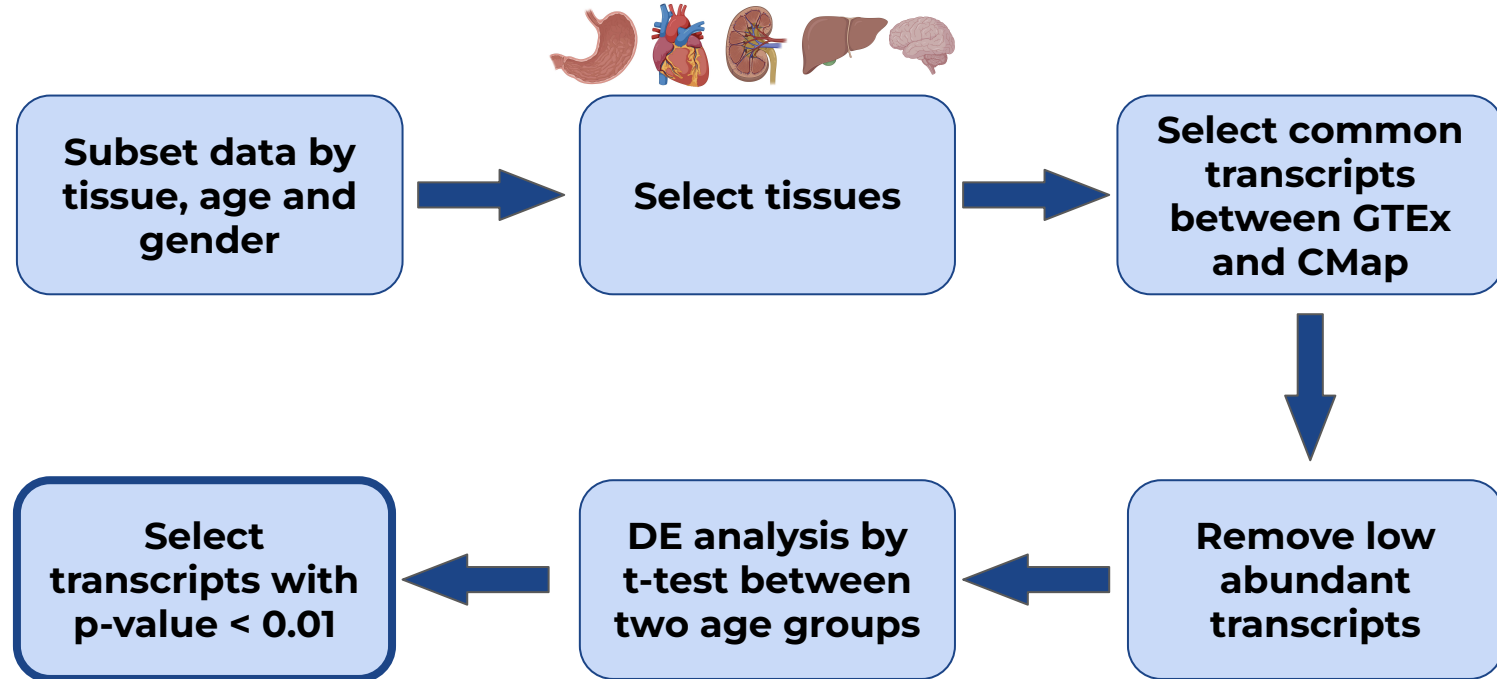


But we gonna do **our own** research!

Authors identified several molecules as potential geroprotectors. Most promising of them (Hsp90 inhibitors) were tested in *C.elegans*

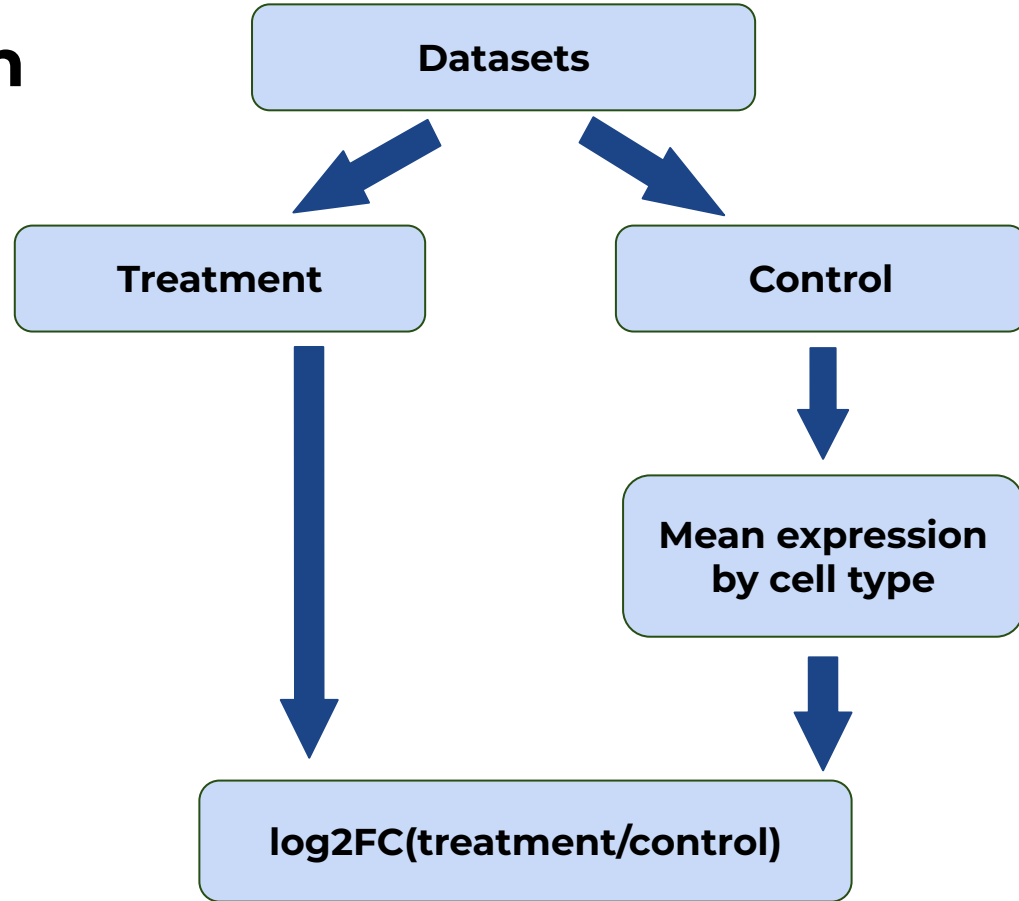
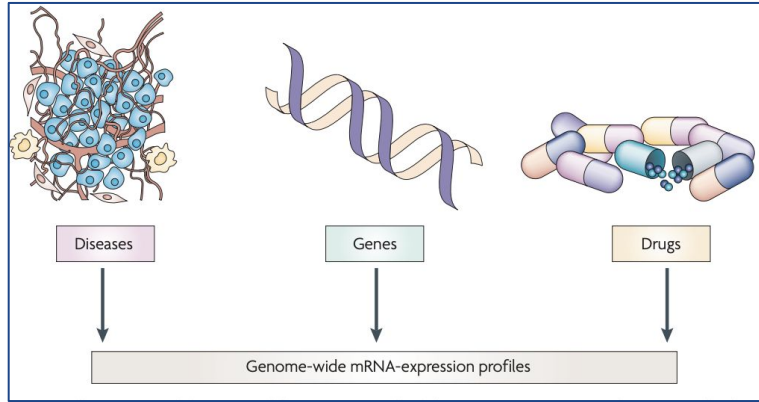


# Data processing





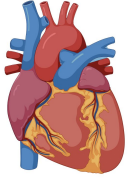
# CMap data preparation



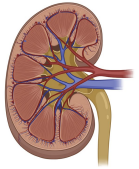
# Random Forest model generation and selection



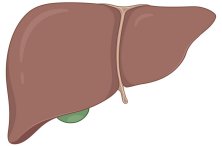
Brain



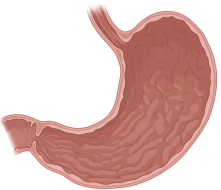
Heart



Kidney



Liver



Stomach

young



old

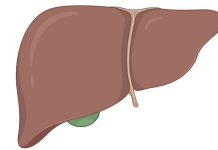
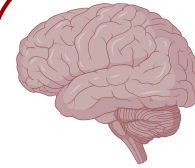


Random  
Forest  
fitting



Selecting  
models with  
 $AUC > 0.75$

**Best age-classifiers**



1

Generating 'drug-induced' transcriptomes for age classification



Prototypical  
'middle-aged'  
transcriptomes

applying CMap  
fold changes



'Drug-induced'  
transcriptomes

Average transcriptome  
between compared  
old and young  
datasets

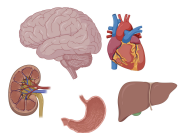
2

Application of age  
classifiers to  
'drug-induced'  
transcriptomes



**Geroprotectors**

# The list of geroprotectors



**100 treatments**  
(out of 3546)



**L-ergothioneine**



**3546 treatments**  
(out of 3546)



**11 compounds**

olanzapine

XL-888

ritonavir

thioguanine

L-ergothioneine

nafamostat

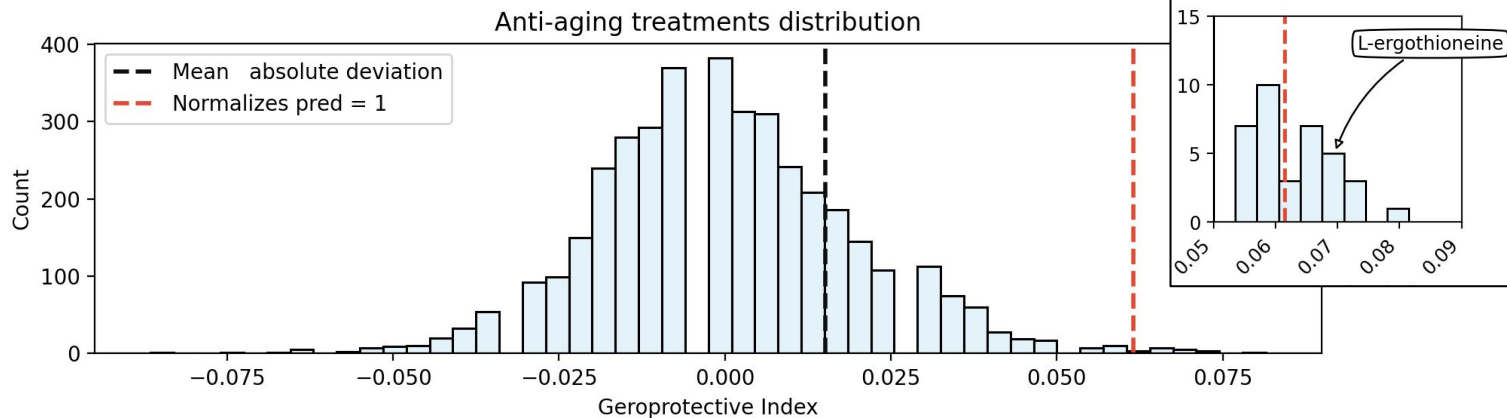
ochratoxin-a

etofylline-clofibrate

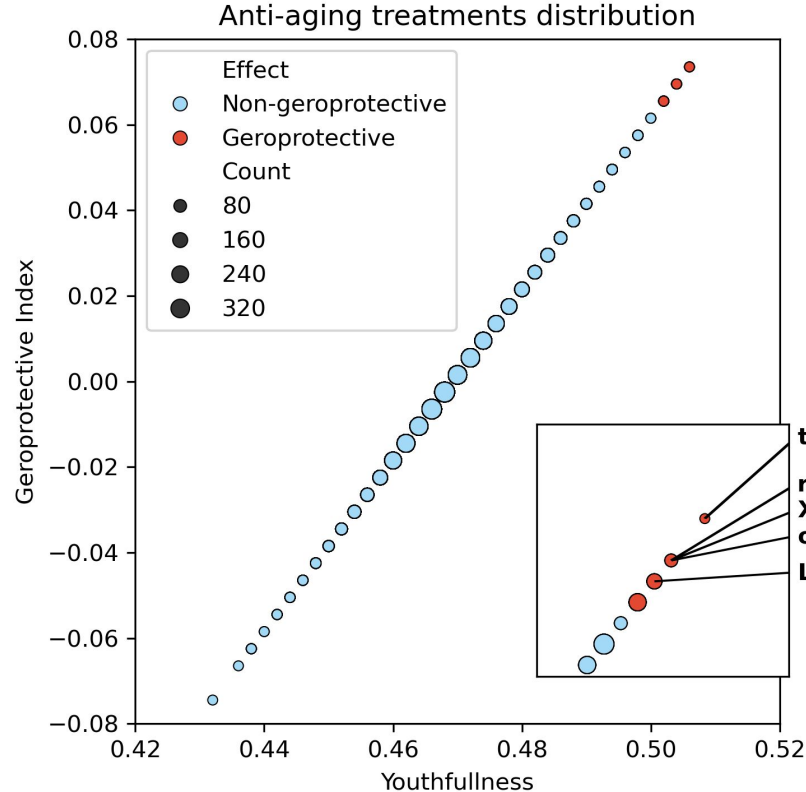
tebipenem

E-2012

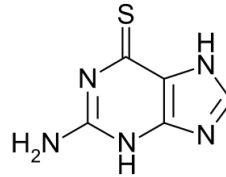
EMF-sumo1-7



# The list of geroprotectors



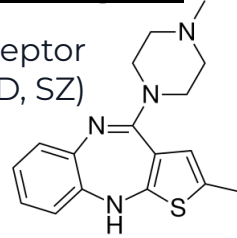
## Thioguanine



Incorporates into DNA and **inhibits synthesis**. Used in the treatment of **leukaemia**.

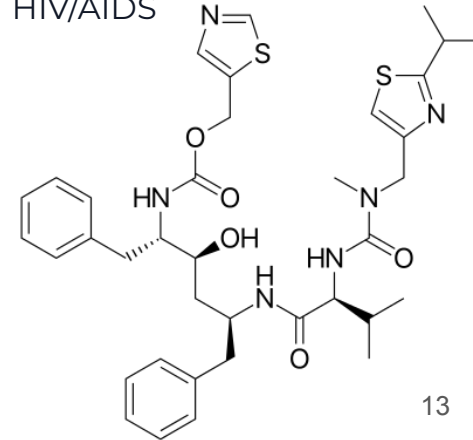
## Olanzapine

Serotonin-dopamine-receptor antagonist, **antipsychotic** (BD, SZ)



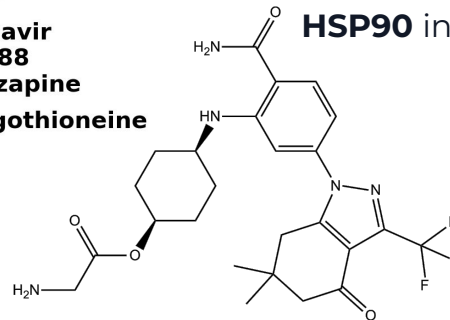
## Ritonavir

Proteases inhibitor, used for HIV/AIDS



## XL-888

**HSP90** inhibitor



thioguanine  
ritonavir  
XL-888  
olanzapine  
L-ergothioneine

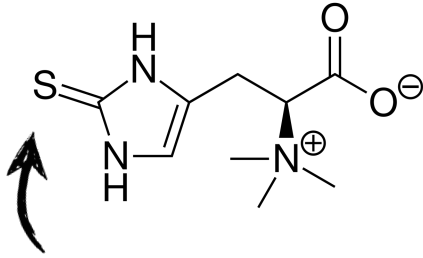
# Result validation: L-ergothioneine

Is



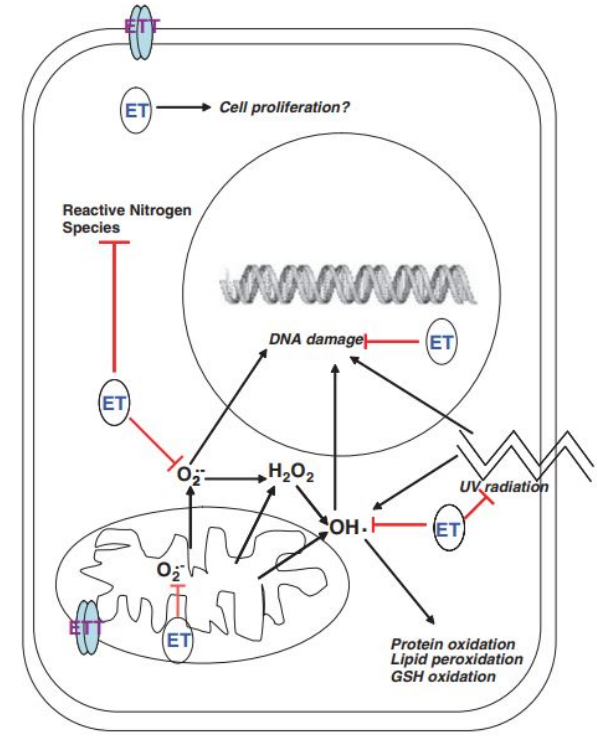
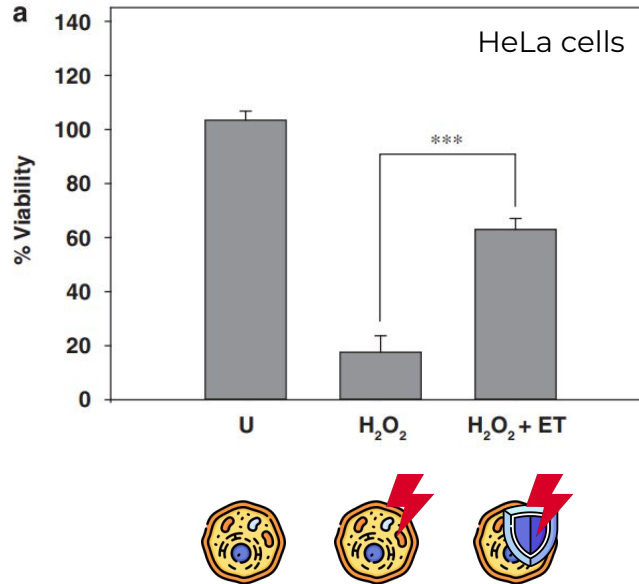
a real geroprotection candidate?

# L-ergothioneine function



**Sulphur** group (can be oxidised and reduced)

May protect against reactive oxygen species



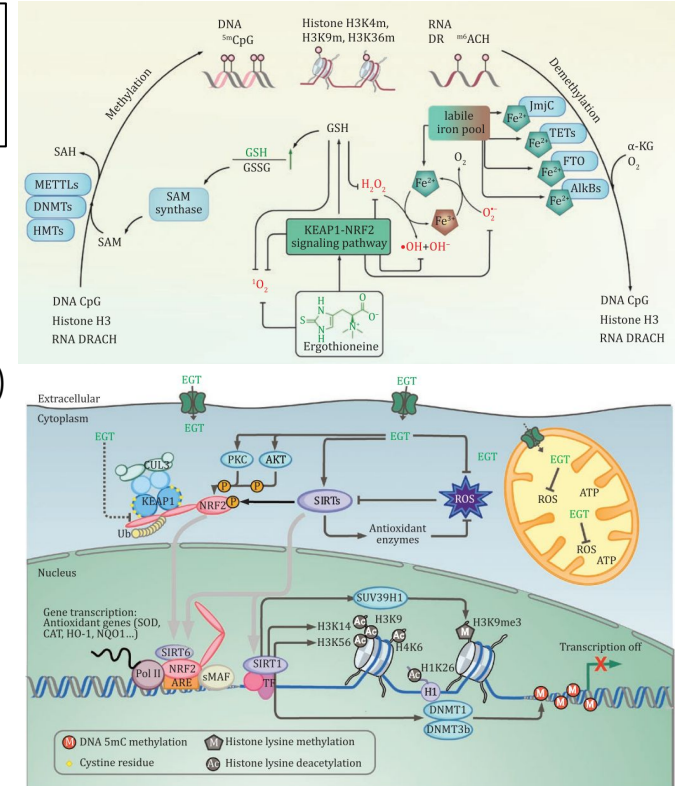
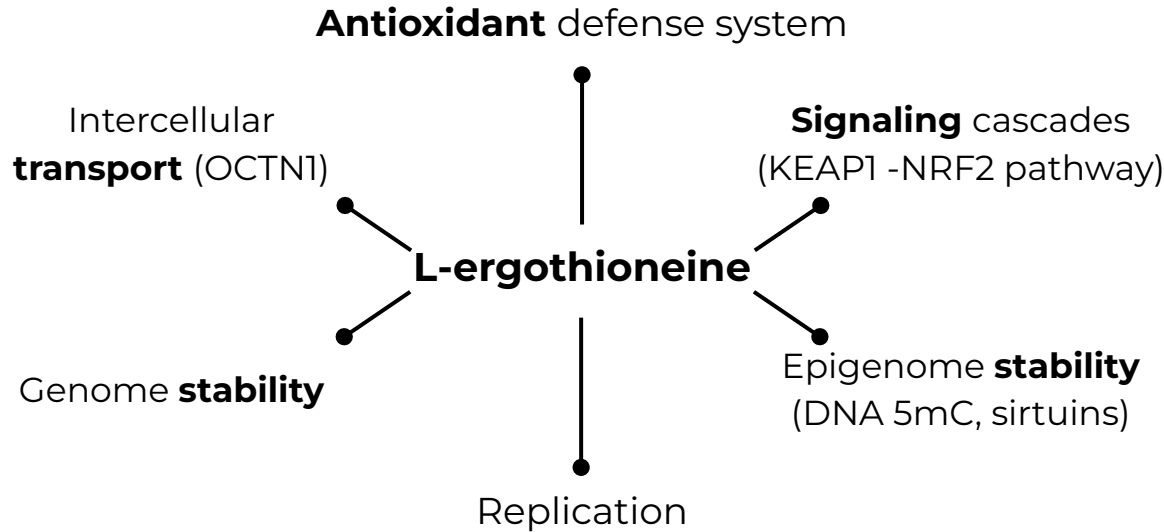
## L-ergothioneine in the network of aging hallmarks

# Ergothioneine and its congeners: anti-ageing mechanisms and pharmacophore biosynthesis

Protein Cell, 2023, XX, 1-16

<https://doi.org/10.1093/procel/pwad048>  
Advance access publication 10 August 2023

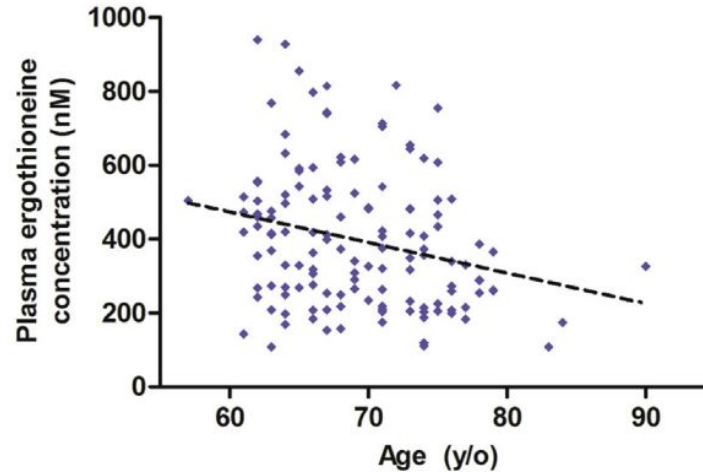
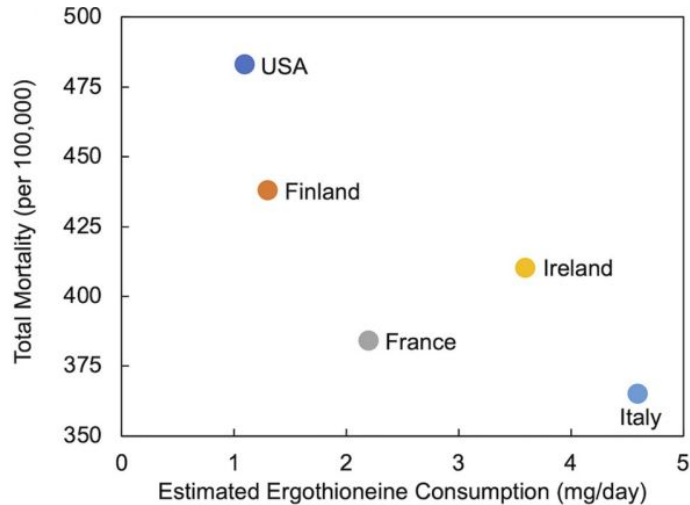
Li Chen<sup>1,2,†</sup>, Liping Zhang<sup>2,†</sup>, Xujun Ye<sup>1,\*</sup>, Zixin Deng<sup>1,2,\*</sup>, Changming Zhao<sup>1,2,\*</sup>





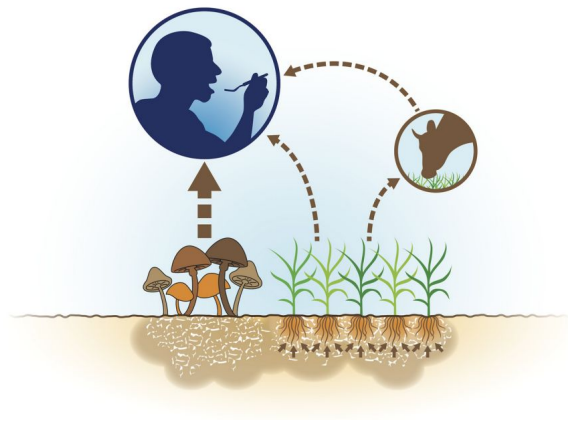
# L-ergothioneine in real life

- L-ergothioneine is correlated with mortality and life expectancy
- L-ergothioneine concentration in plasma decreases with aging



# L-ergothioneine sources

L-ergothioneine is mainly present in **mushrooms**



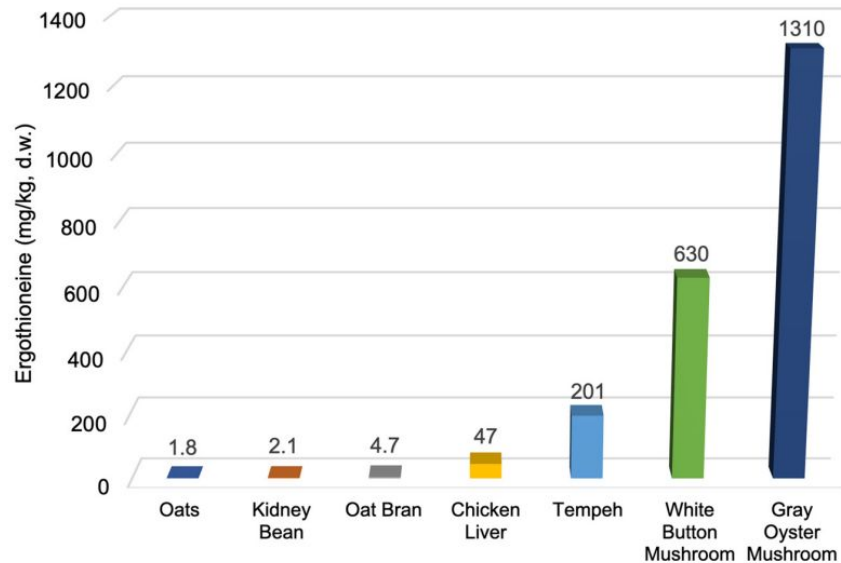
Life Extension, Essential Youth L-эрготионеин, 5 мг, 30 вегетарианских капсул

Артикул товара: H18969

В наличии

3 760 Р

Добавить в корзину



# Conclusions

1. We built transcriptomic **age-classification** model
2. We applied this model to **identify treatments** that result in a "young" transcriptome
3. We additionally investigated the **role of L-ergothioneine in the aging**

## Drawbacks

Why do our results differ from the paper?

1. We used **reduced data** to save time. We selected some organs and ran a sample of 100 treatments (out of 3546 total) on them. We also ran our model for all treatments, but only for the brain.
2. Our compounds-**ranking** procedure differs from the paper due to lack of ran models
3. We suspect we used a slightly different **version of CMap** dataset. In our data we had 3546 treatments against 1309 in the paper.



# Team



**Ksenia Kubenko**

Transcriptome data processing  
and CMap data preparation  
(mean expression by cell type  
calculation)



**Ekaterina Kashuk**

Random Forest model  
generation and selection,  
selection of geroprotectors



**Nikita Vaulin**

CMap data preparation (log2FC  
calculation), literature review  
on geroprotectors

All authors contributed to the presentation and aging

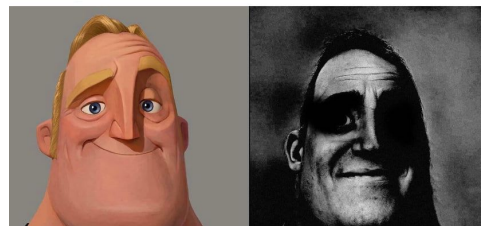
# Geroprotectors table

old	yng	treatment	normalized_pred	geroprotective_index
0.498	0.502	EMF-sumo1-7	1.004	0.0656
0.496	0.504	ochratoxin-a	1.008	0.0670
0.498	0.502	nafamostat	1.004	0.0656
0.496	0.504	<i>L-ergothioneine</i>	1.008	0.0670
<b>0.494</b>	<b>0.506</b>	<b>olanzapine</b>	<b>1.012</b>	<b>0.0740</b>
<b>0.494</b>	<b>0.506</b>	<b>XL-888</b>	<b>1.012</b>	<b>0.0740</b>
0.498	0.502	E-2012	1.004	0.0656
<b>0.494</b>	<b>0.506</b>	<b>ritonavir</b>	<b>1.012</b>	<b>0.0740</b>
<b>0.49</b>	<b>0.51</b>	<b>thioguanine</b>	<b>01.02</b>	<b>0.0815</b>
0.498	0.502	tebipenem	1.004	0.0656
0.496	0.504	etofylline-clofibrate	1.008	0.0670

# CMap

Слайд про CMap

Работа с CMap



	chds	version	dimensions	rhds					cids
1	#1.3								
2	10	6	2	5					
3	id	pr_gene_symbol	pr_is_lmark	LPROT001_A375 6H_X1_B20:B03	LPROT001_A375 6H_X1_B20:B05	LPROT001_A375 6H_X1_B20:B07	LPROT001_A375 6H_X1_B20:B09	LPROT001_A375 6H_X1_B20:B11	LPROT001_A375 6H_X1_B20:B13
4	pert_id	-666	-666	DMSO	DMSO	BRD-K52313696	BRD-K52313696	BRD-K52313696	BRD-K77908580
5	pert_iname	-666	-666	DMSO	DMSO	tacedinaline	tacedinaline	tacedinaline	entinostat
6	cell_id	-666	-666	A375	A375	A375	A375	A375	A375
7	pert_time	-666	-666	6	6	6	6	6	6
8	pert_dose	-666	-666	-666	-666	2	2	2	2
9	5720	PSME1	1	8.7980	8.9395	8.9561	9.4491	9.1994	9.2937
10	55847	CISD1	1	9.8349	9.5334	9.7543	9.9203	9.8904	10.0666
11	7416	VDAC1	1	12.5431	12.3479	12.4662	12.5892	12.7088	12.6397
12	10174	SORBS3	1	8.1017	8.5660	8.3563	8.6225	8.5800	8.3666
13	25803	SPDEF	1	11.0651	11.0922	11.3566	11.1527	11.0557	10.7427
14	466	ATF1	1	6.6887	6.8780	6.6684	7.1662	6.7492	6.7492
15	6676	SPAG4	1	3.5195	3.8840	3.6936	3.4822	3.2684	3.7300
16	1870	E2F2	1	4.5798	4.6260	4.4156	4.2644	4.4611	4.4611
17	6009	RHEB	1	11.7969	12.3234	12.0216	11.9772	12.0580	12.0216
18	3480	IGF1R	1	8.9370	10.0307	9.5279	10.0616	8.9231	8.6343

column  
metadata

rids

row metadata

data matrix

# CMap

Ksusha's means  
(2466225, 3)

	Transcript	sample	value
0	100017	AML001	6.847573
1	100019	AML001	6.603014
2	100037258	AML001	6.761893

CMap  
(214958468, 4)

	Transcript	variable	value	sample
0	12558	AML001_CD34_24H_X1_F1B10:E03	4.6286	AML001
1	11308	AML001_CD34_24H_X1_F1B10:E03	6.5425	AML001
2	12560	AML001_CD34_24H_X1_F1B10:E03	4.4462	AML001

```
%%time
```

```
cmap_means_merged = cmap_melted.merge(means_melted, on=['Transcript', 'sample']).compute()
```

```
CPU times: user 58.8 s, sys: 41.3 s, total: 1min 40s
```

```
Wall time: 59.3 s
```

