


**Life on earth is under the influence of basic environmental periodic changes such as day-night cycles or the fluctuation of seasons**

**Organisms have adapted to these rhythms in order to maximally benefit from the limited natural resources**

**The mechanism that keeps track of time, and therefore allows the organism to anticipate upcoming daily changes is termed**

**CIRCADIAN CLOCK**





**Circadian clocks are molecular time-keeping mechanisms that reside in a wide range of cell types in a variety of organisms**

The key feature of a circadian clock is its ability to synchronize (***entrain***) to environmental time cues (so-called ***zeitgebers***, “time-givers”) and to maintain rhythmic function when placed in constant conditions



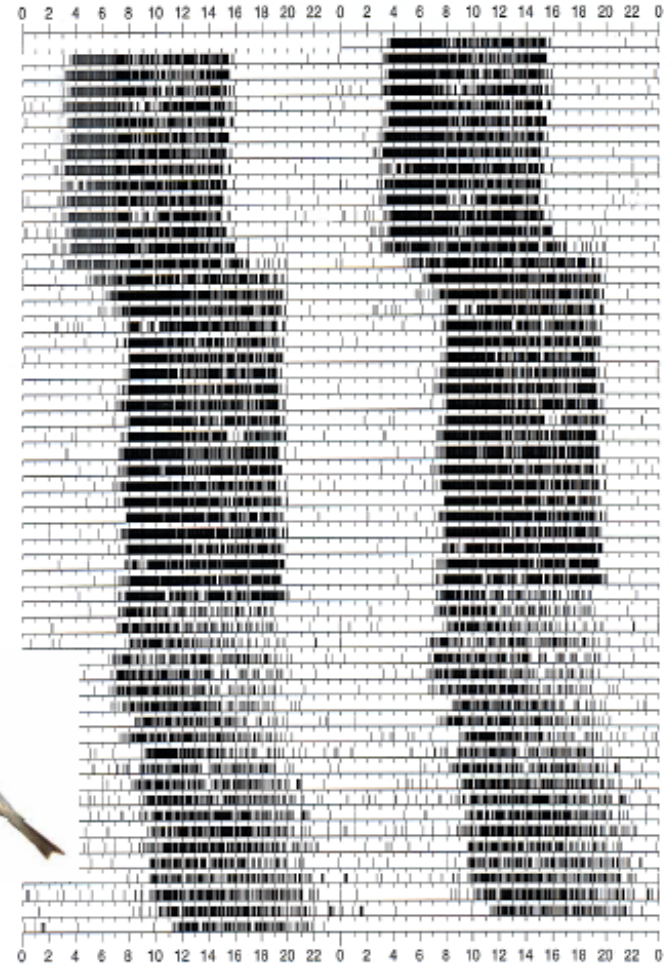
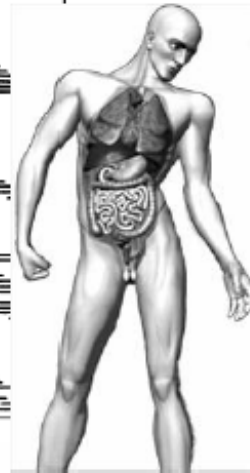
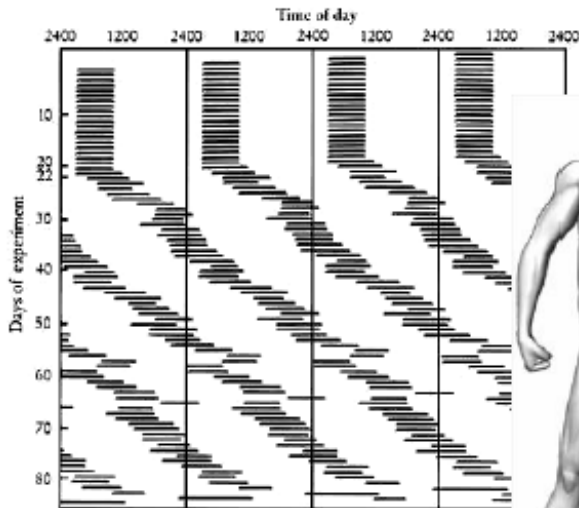
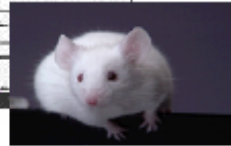
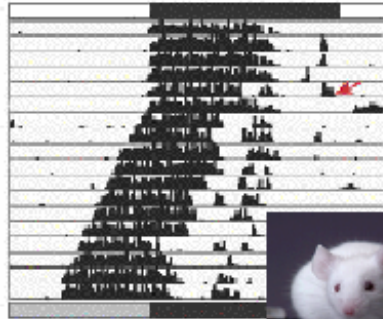
**Endogenous clock systems  
(circadian, circannual, ultradian)  
are present in all organisms known.**

**They have evolved to measure time and to keep  
the organism in entrainment with the environment.**

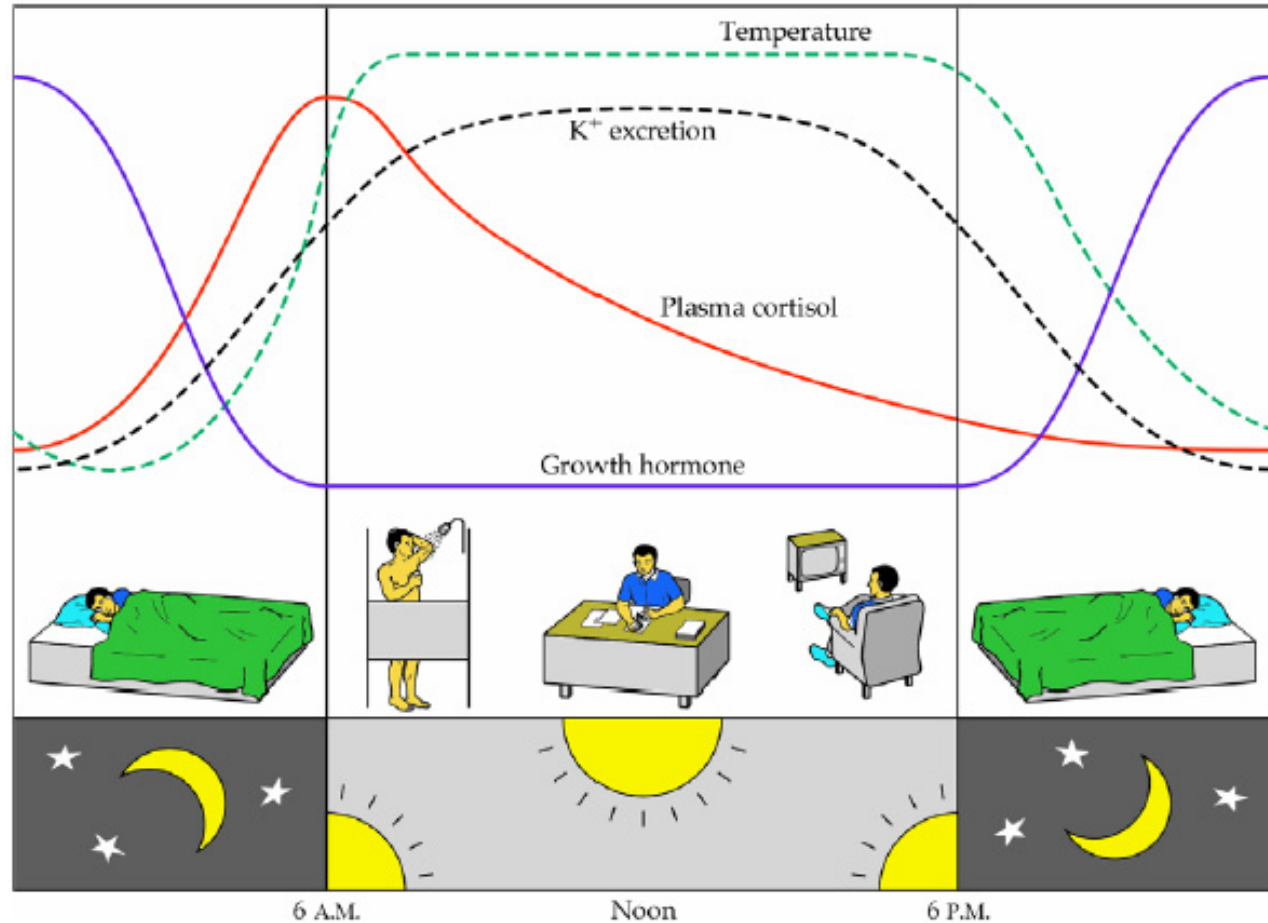
**Light, feeding and temperature are the strongest  
Zeitgebers for circadian rhythms.**



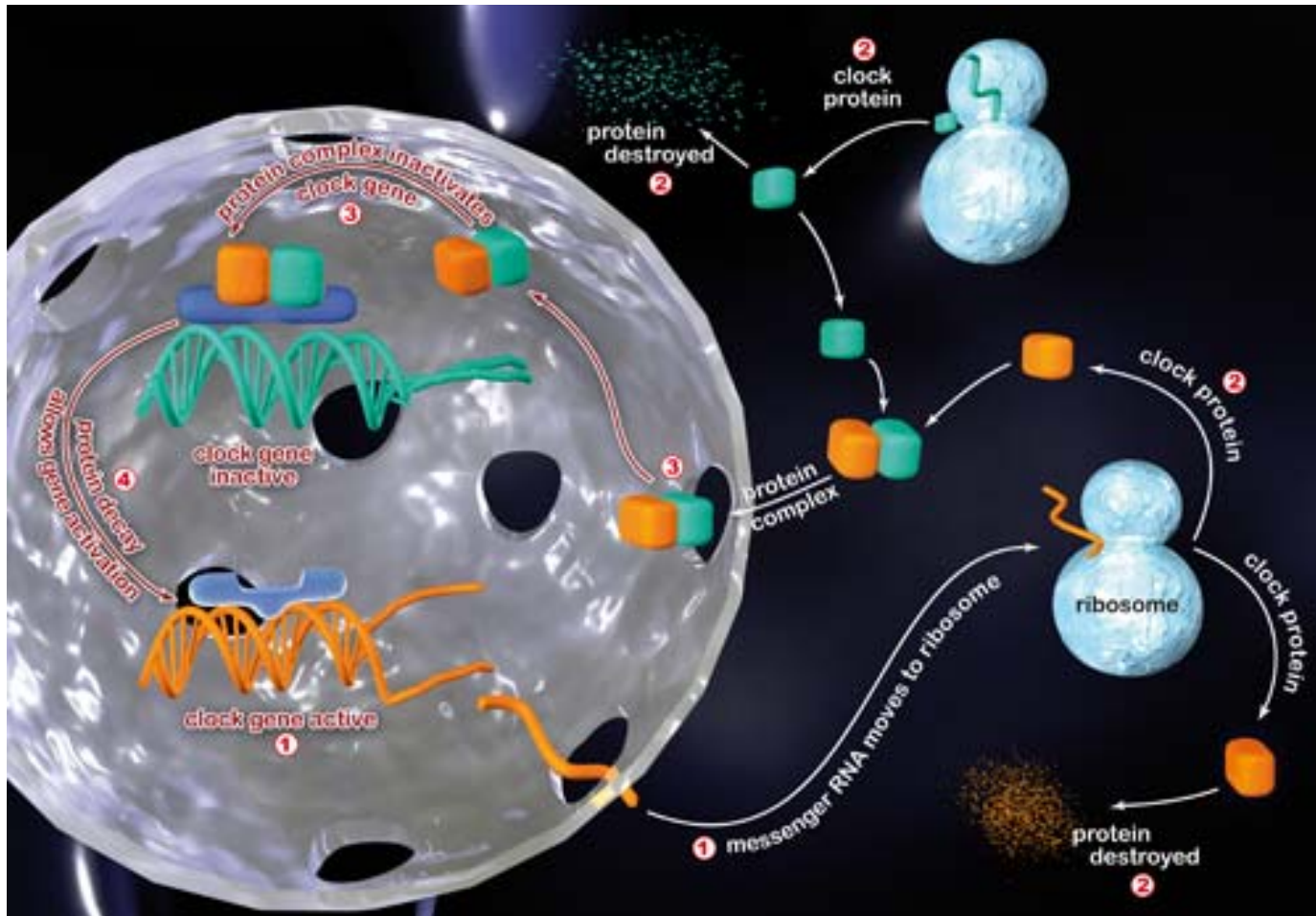
# Biological rhythms are ubiquitous



# Rhythms of body function in humans



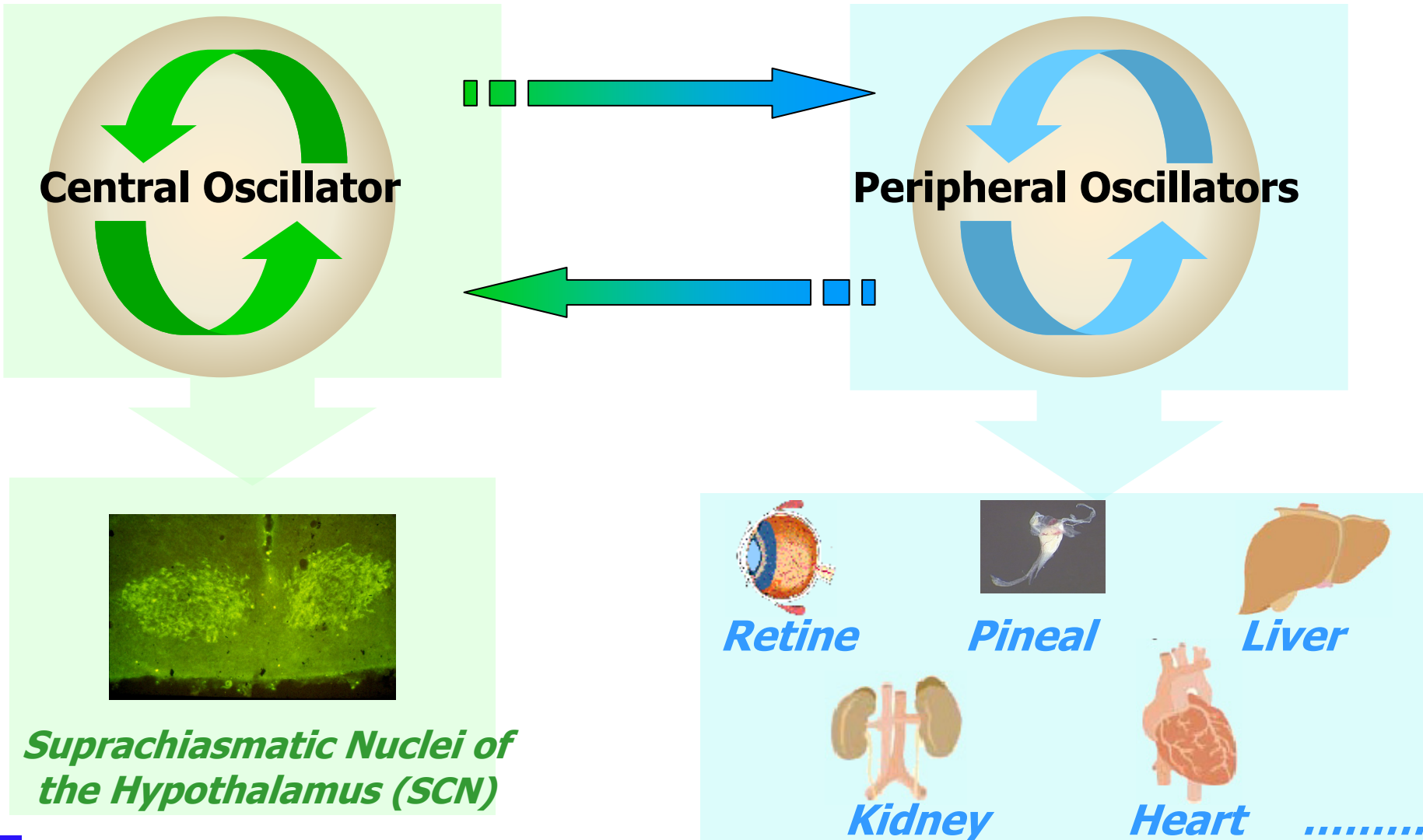
# Circadian rhythms generation consists of interacting positive and negative transcriptional/translational feedback loops





# VERTEBRATE CIRCADIAN SYSTEM

## MULTIOSCILLATORY SYSTEM



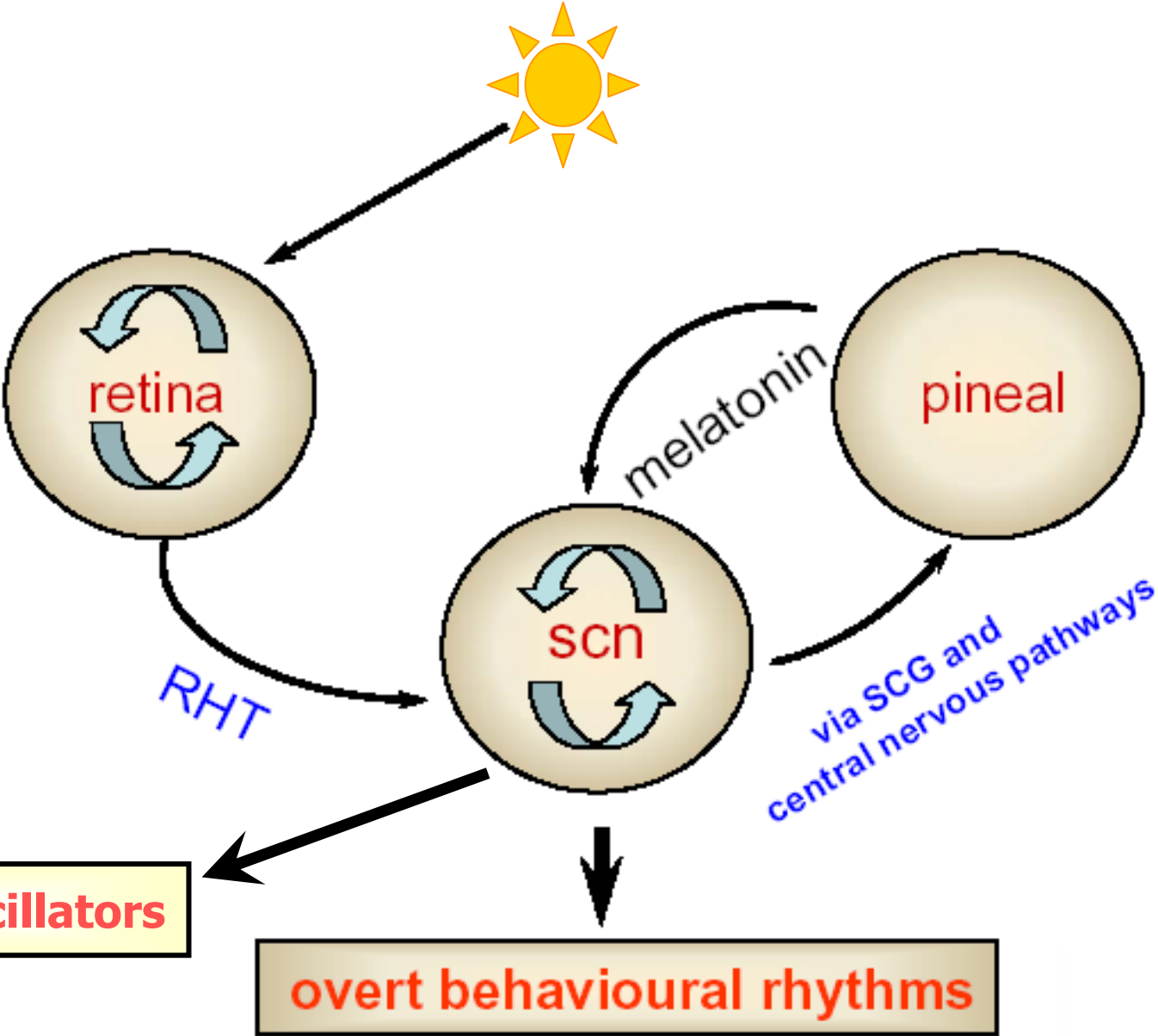
**Key question:**

**How and where is the light signal , the most powerful zeitgeber, detected?**



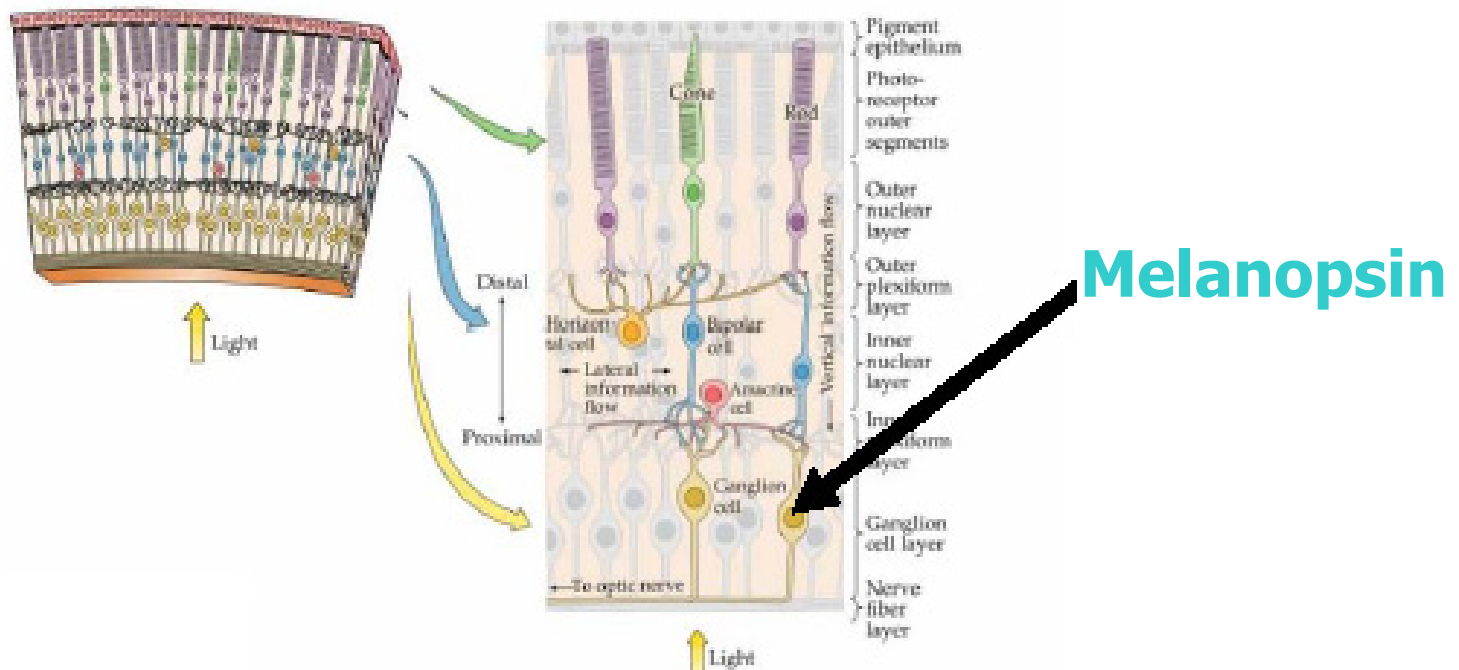


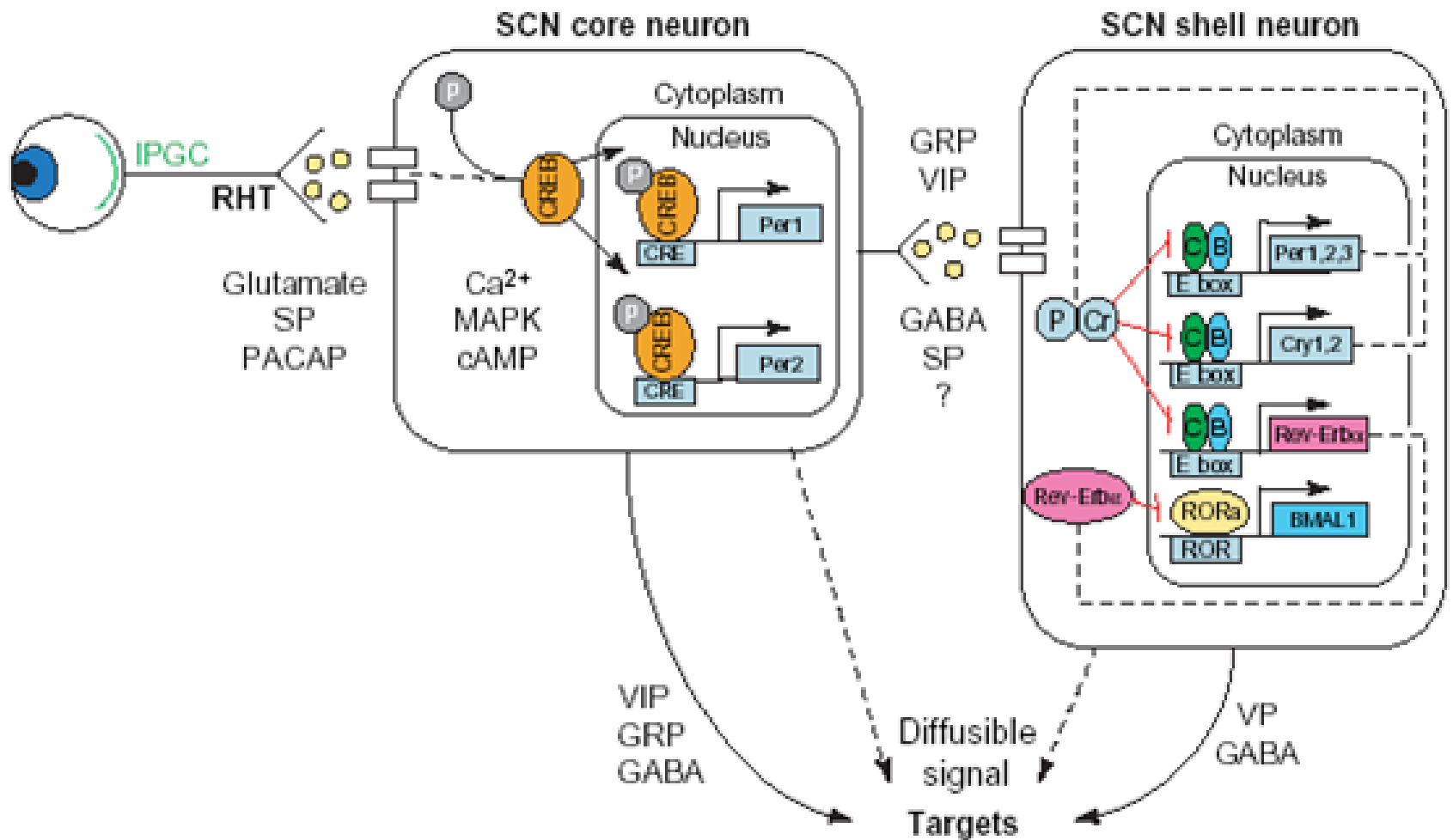
# MAMMALS



# In mammals

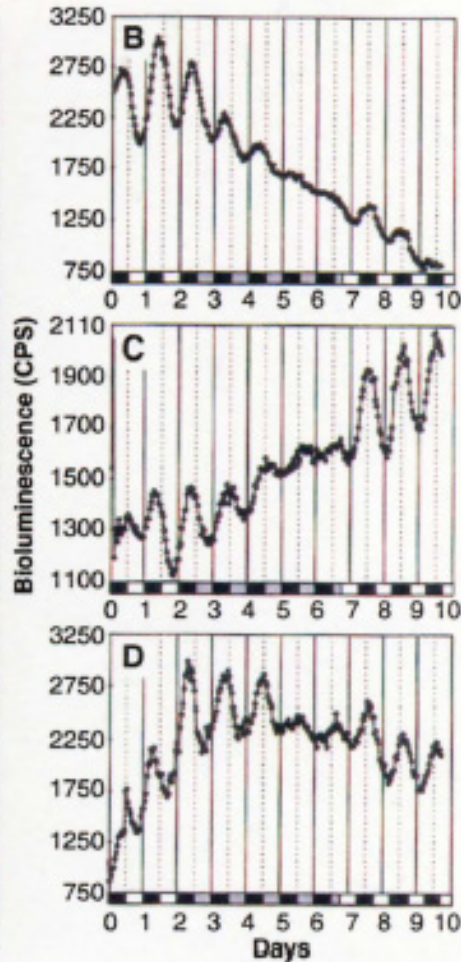
**Light input to the circadian system comes exclusively from the retina, mediated via a subset of ganglion cells that project to the suprachiasmatic nucleus.**





# Light responsive circadian oscillators

## *Drosophila melanogaster*



**Head**

***Period* gene driven bioluminescence**

- present throughout the whole fly

**Thorax**

- rhythms maintained in various body parts in culture

**Abdomen**

## *Drosophila*

**Rhythmic clock gene expression can be found in any tissue**

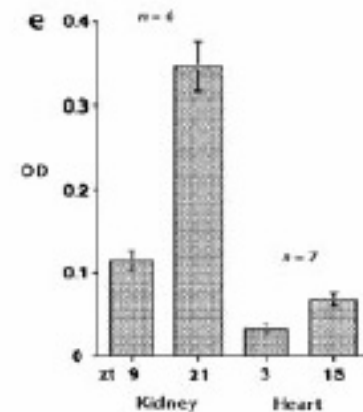
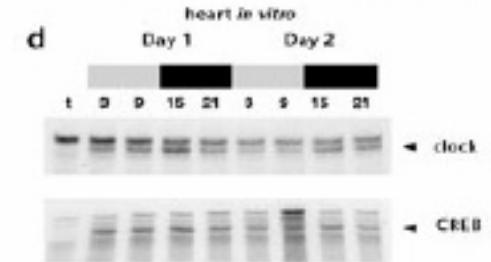
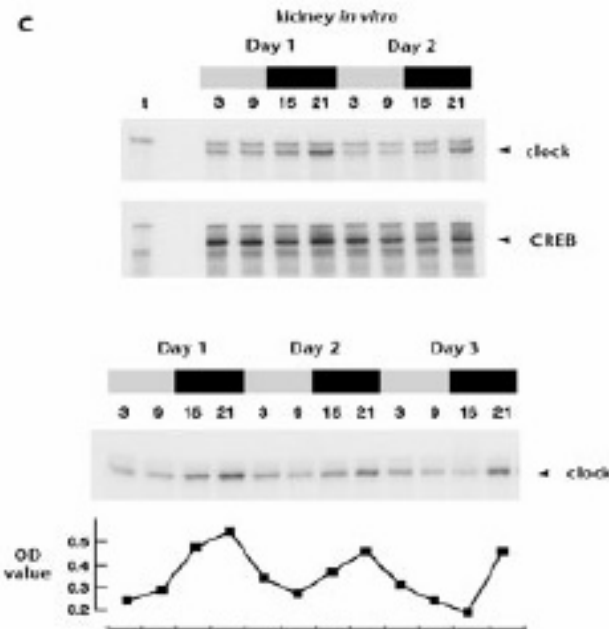
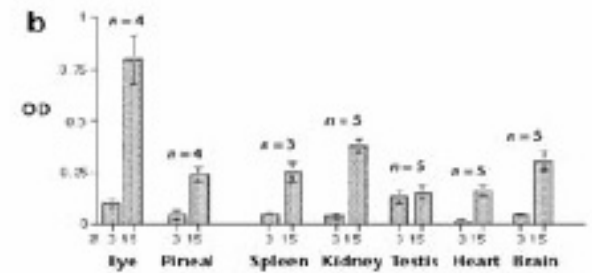
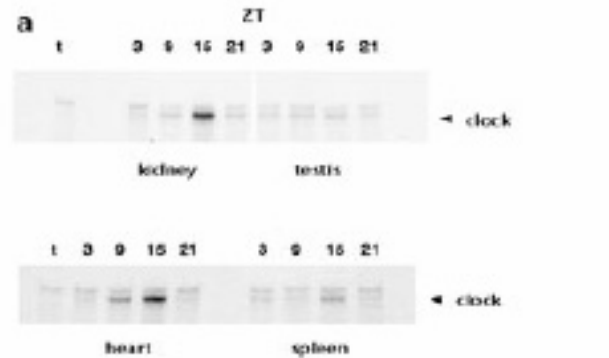
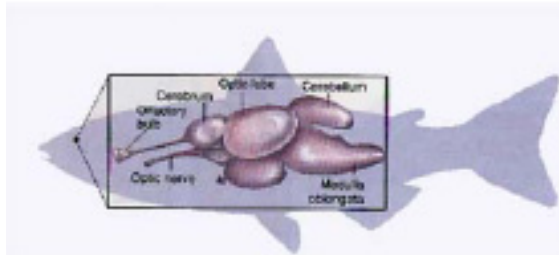
**Rhythmic clock gene expression is maintained in culture**

**All cultured tissues are light responsive**



# Light responsive circadian oscillators

## ZEBRAFISH



# ZEBRAFISH

**Rhythmic clock gene expression can be found in almost any tissue**

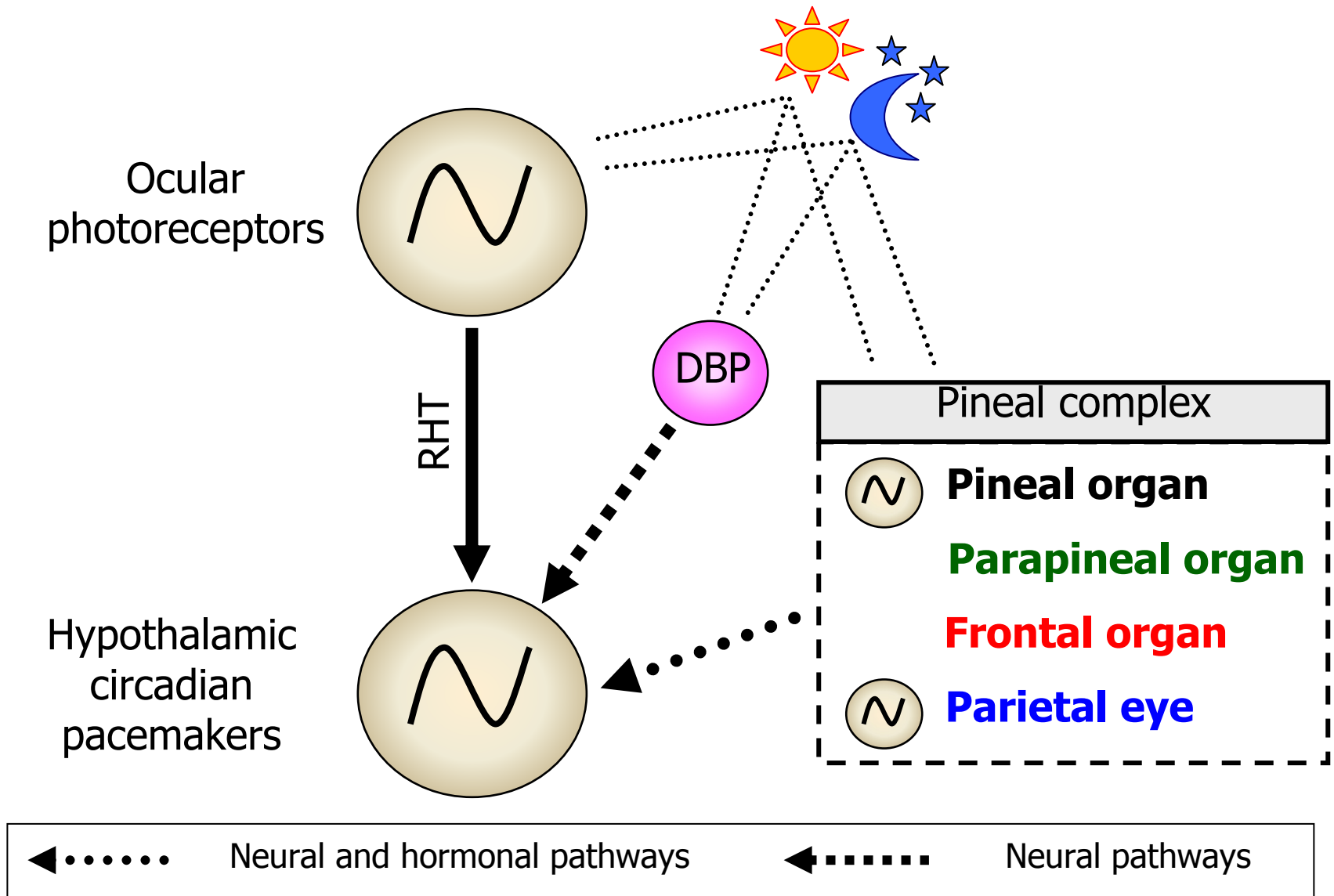
**Rhythmic clock gene expression is maintained in culture**

**All rhythmic tissues are directly light responsive**





# BRAIN CIRCADIAN PHOTORECEPTION



# *Podarcis sicula* (Family Lacertidae)



- Phylogenetic position
- ectotherm hibernator
- interesting model for understanding circadian organization, its evolution, and its variability

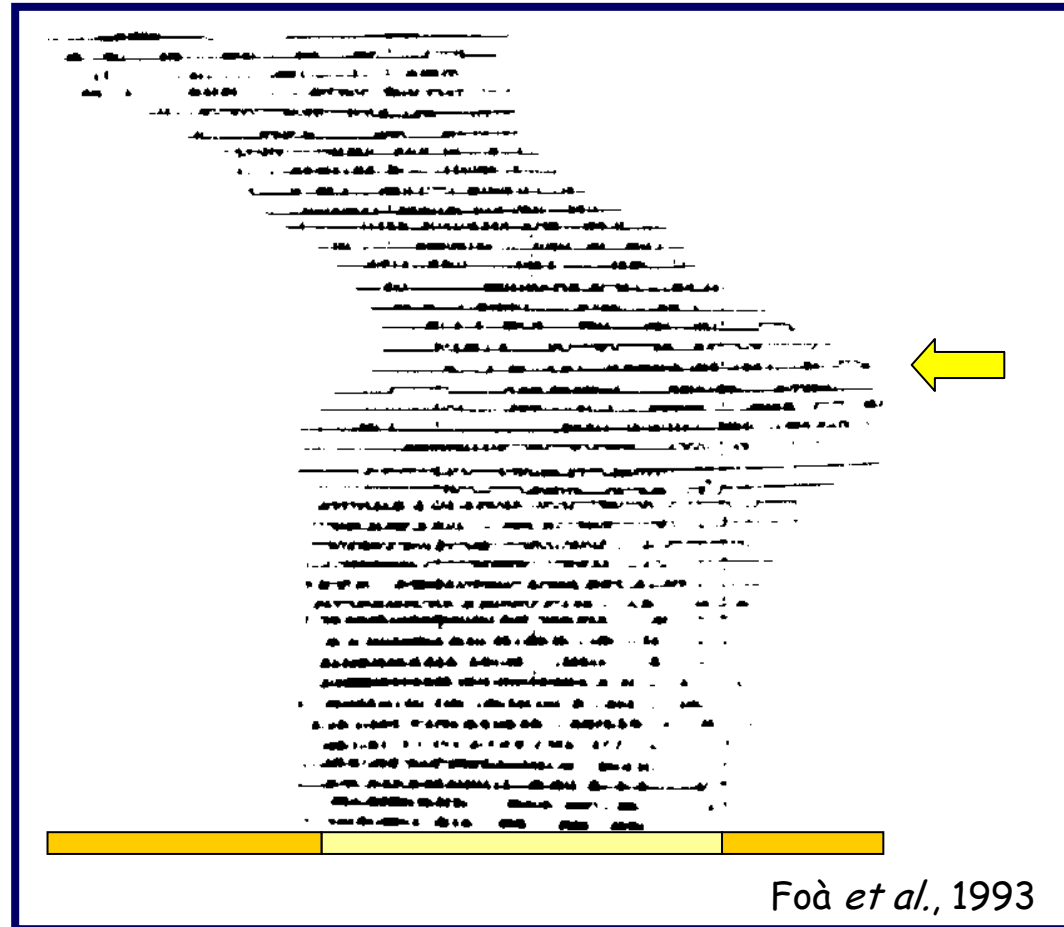


# Entrainment to LD cycles in reptiles

PINX-RETX lizard entrained  
to LD cycles



*Podarcis sicula*



# Opsins isolated and/or detected by in situ hybridization or immunocytochemistry with specific antiserum from extraocular photoreceptive structures

Melanopsin	Amphibians	<i>X. laevis</i>	Provencio et al., 1998
		<i>D. rerio</i>	Bellingham et al., 2002
	Teleosts	<i>G. morhua</i>	Drivenes et al., 2003
Pinopsin	Amphibians	<i>B. japonicus</i>	Yoshikawa et al., 1998
VA opsin (different isoforms)	Teleosts	<i>S. salar</i>	Philp et al., 2000
		<i>D. rerio</i>	Kojima et al., 2000
		<i>C. carpio</i>	Moutsaki et al., 2000
		<i>P. altivelis</i>	Minamoto and Shimizu, 2002
Rhodopsin	Birds	<i>C. livia</i>	Wada et al., 1998
	Teleosts	<i>P. altivelis</i>	Masuda et al., 2003
RH2 opsin	Reptiles	<i>P. sicula</i>	Pasqualetti et al., 2003
tmt-opsin	Teleosts	<i>D. rerio</i>	Moutsaki et al., 2003

Basal region of lateral ventricle

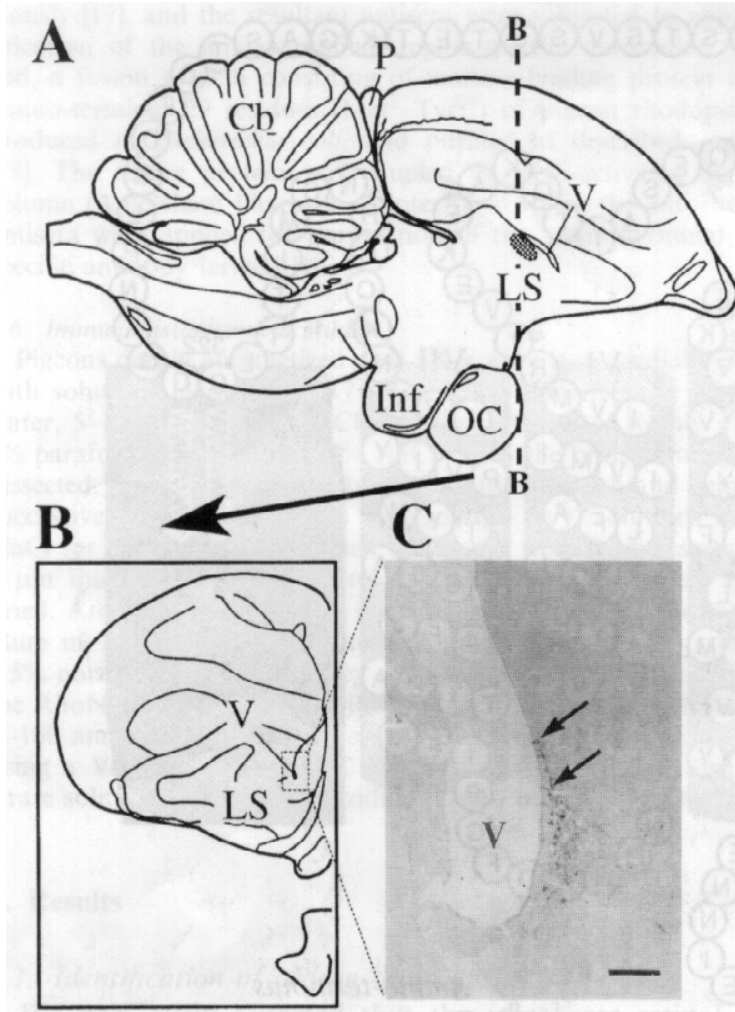
Anterior hypothalamus



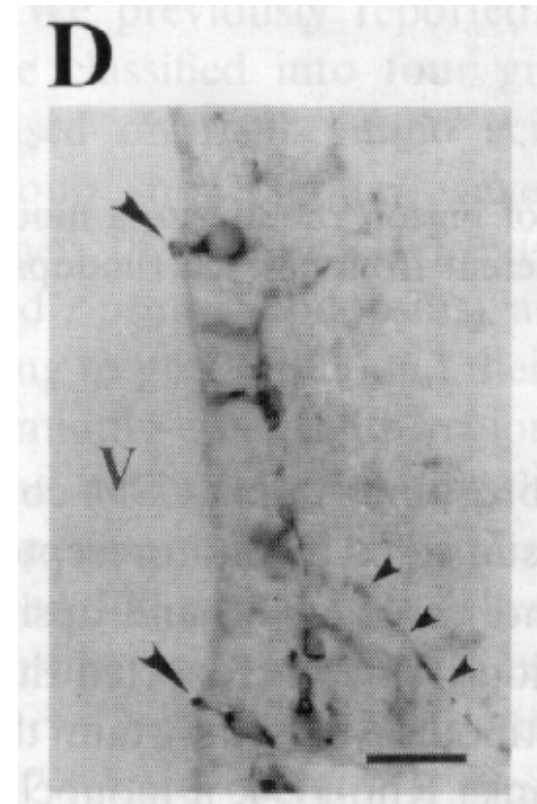


# DEEP BRAIN PHOTORECEPTORS

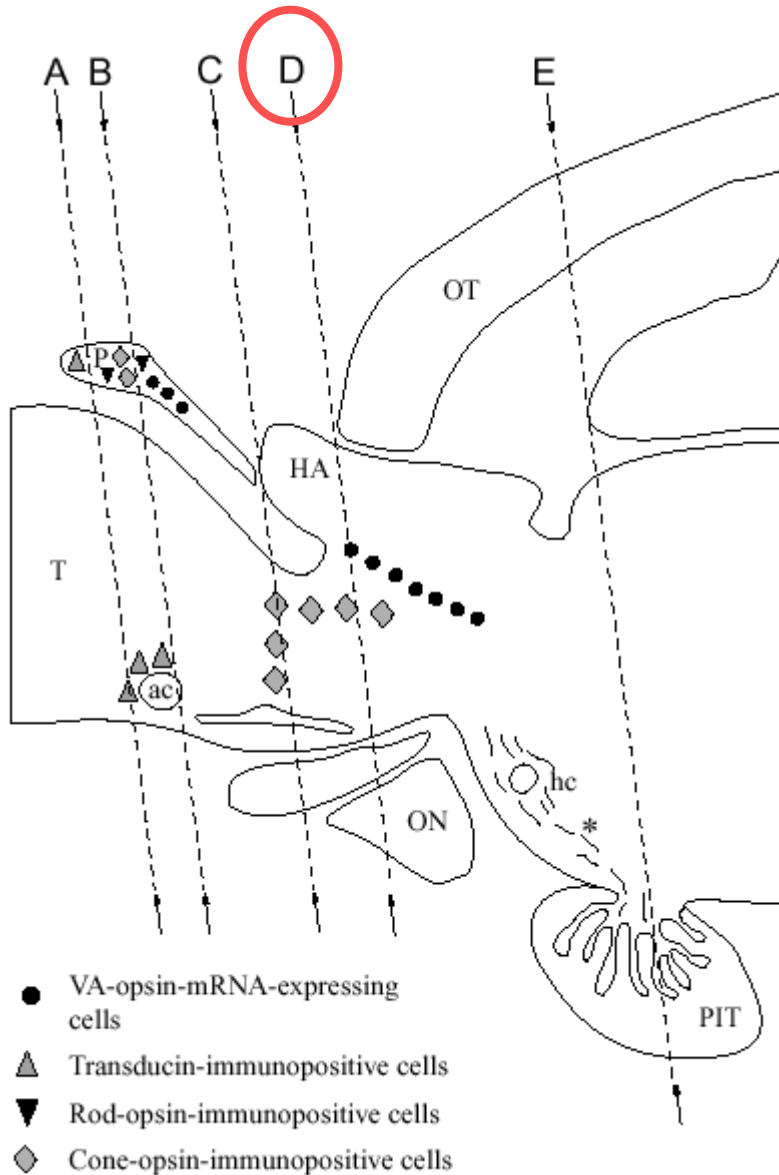
## Basal region of lateral ventricle in *Columba livia*



CSF-contacting neurons labelled with antiserum against rodopsin

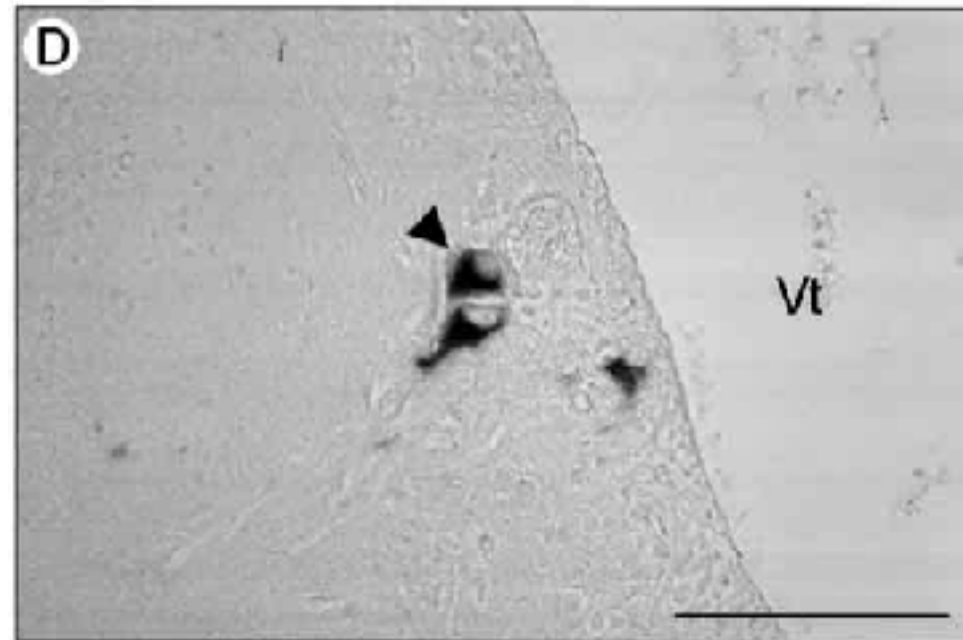


# DEEP BRAIN PHOTORECEPTORS



Anterior hypothalamus  
of *Salmo salar*

Neurosecretory cells labelled with  
antibody against cone-opsin

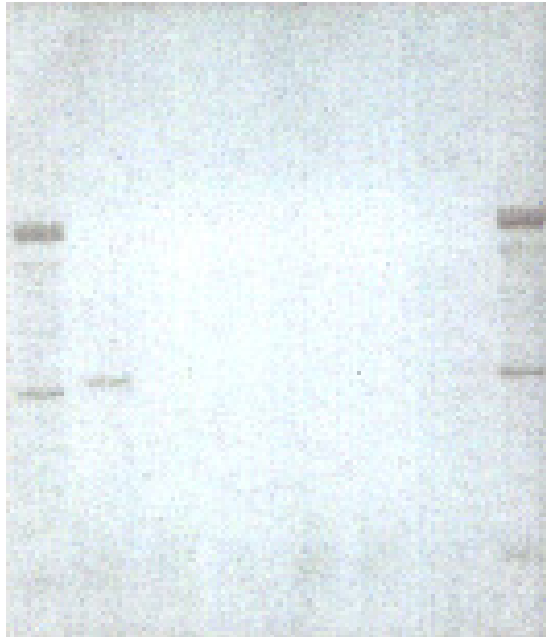


Vt: III ventricle



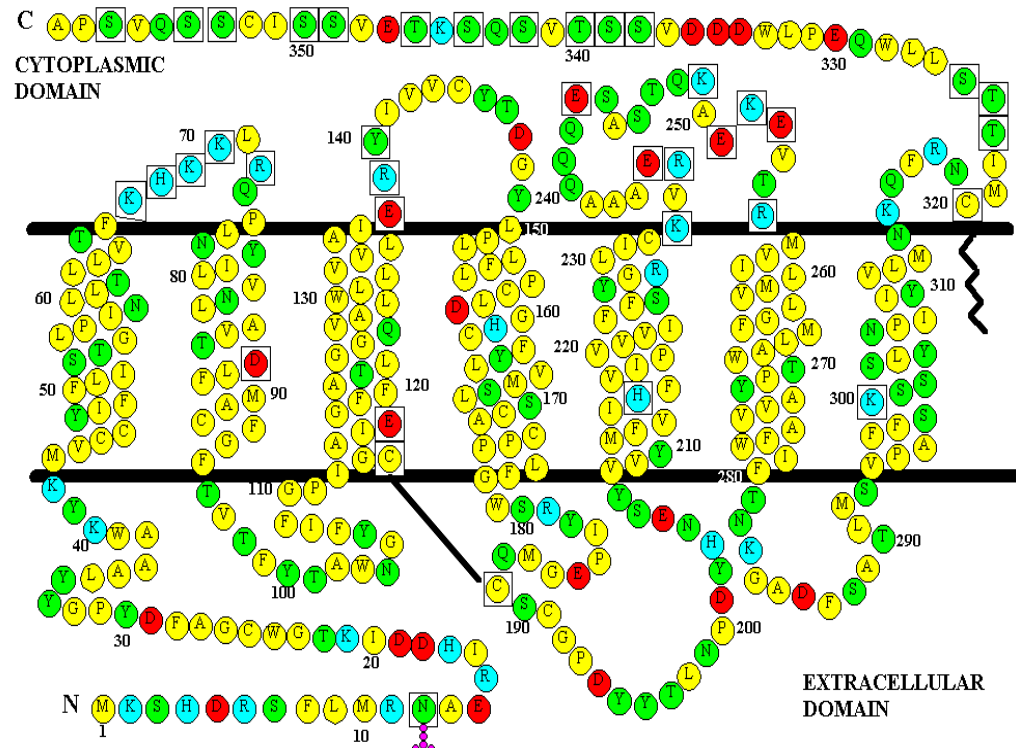
# Identification of brain opsin in the lizard *Podarcis sicula*

L 1 2 3 4 5 6 L



L - ladder  
 1 - brain  
 2 - heart  
 3 - liver  
 4 - skin  
 5 - testicle  
 6 - negative control

RT-PCR in different tissues of lizards



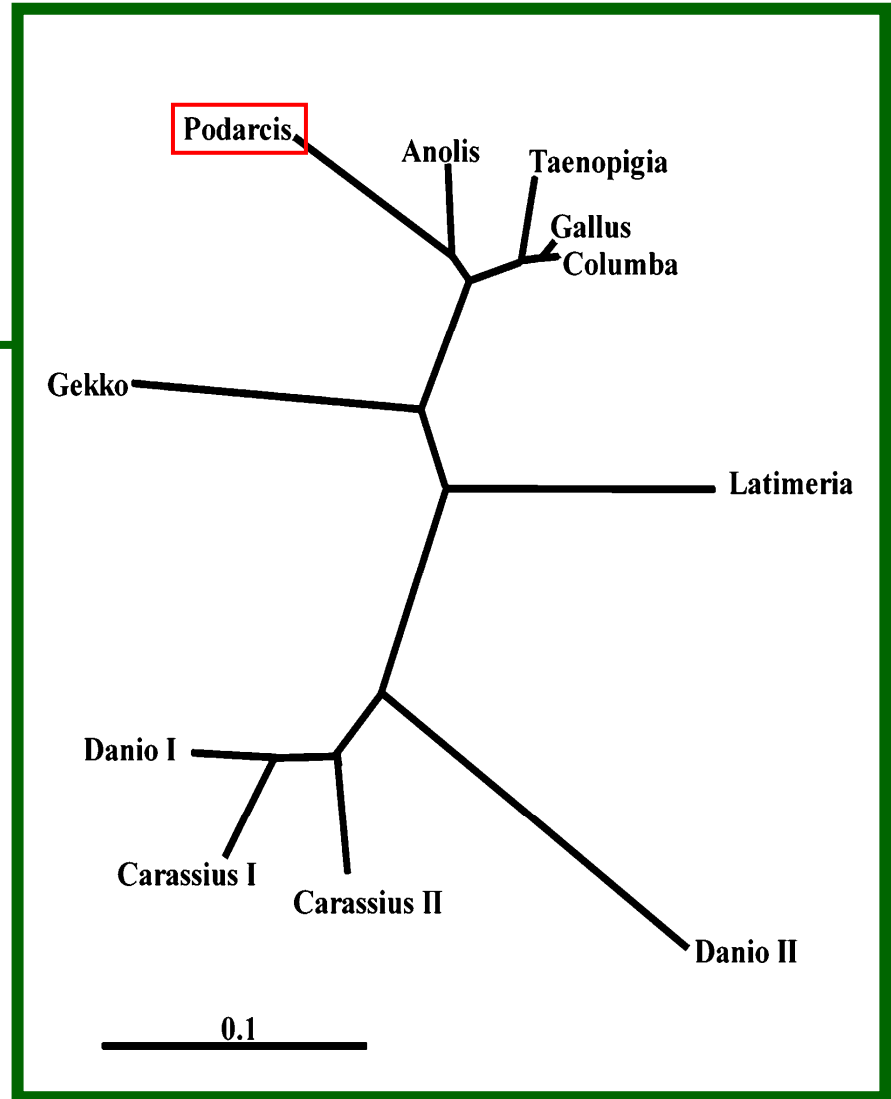
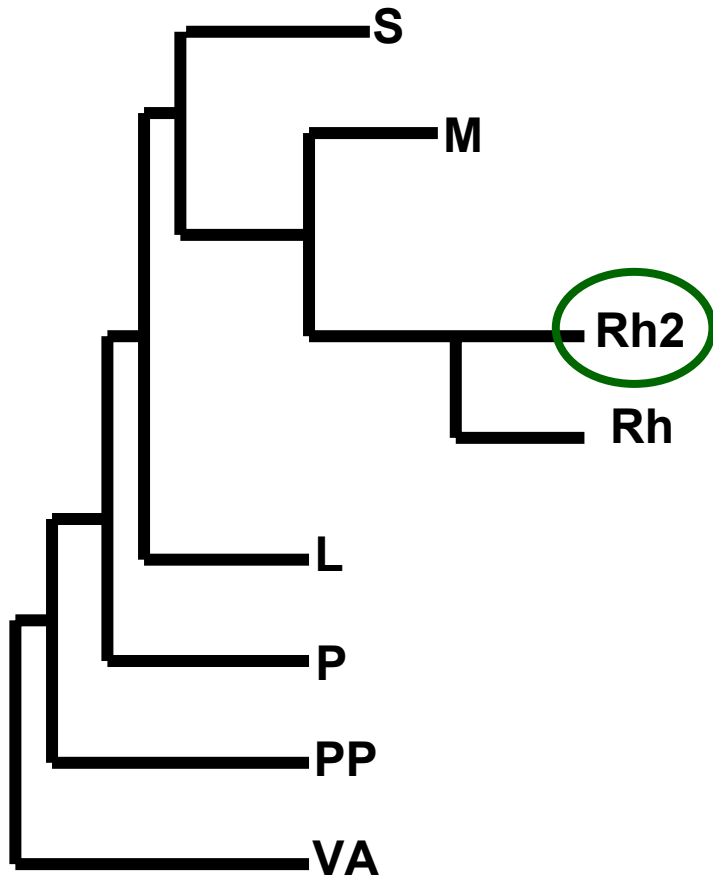
AA Sequence of the brain opsin

Lys300 retinal  
 Glu138-Arg139-Tyr140 transducin

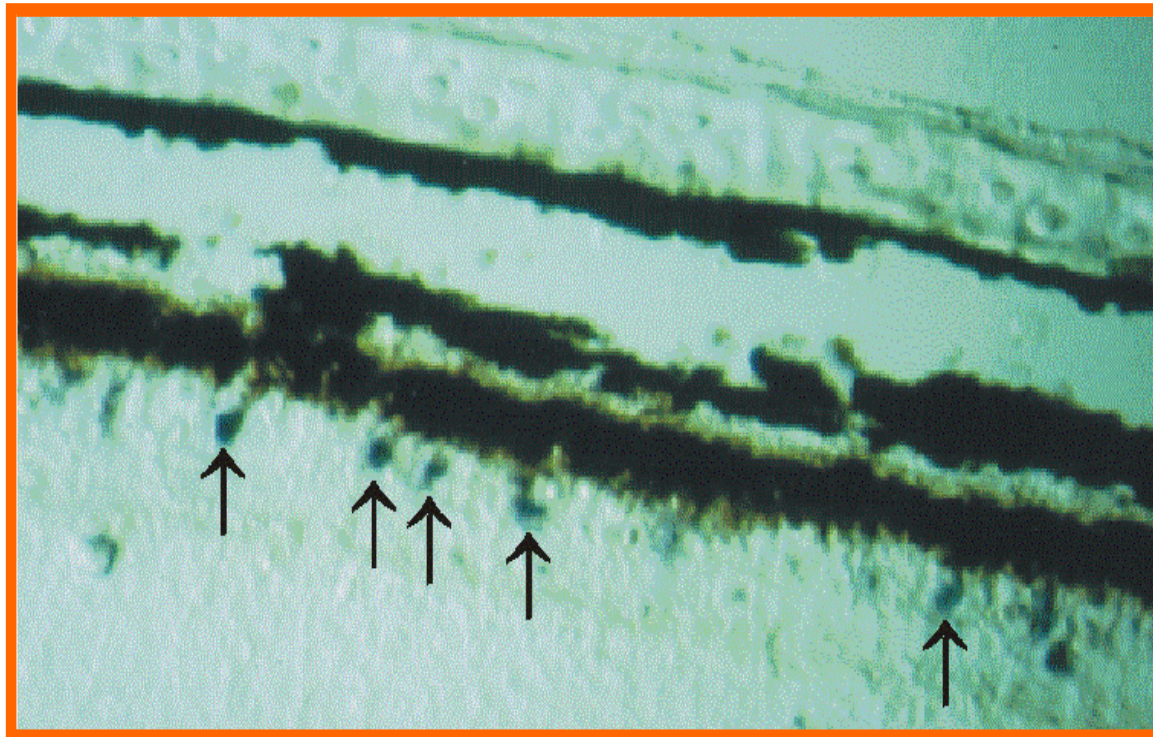
Pasqualetti et al., 2003, Eur, J, Neurosci.



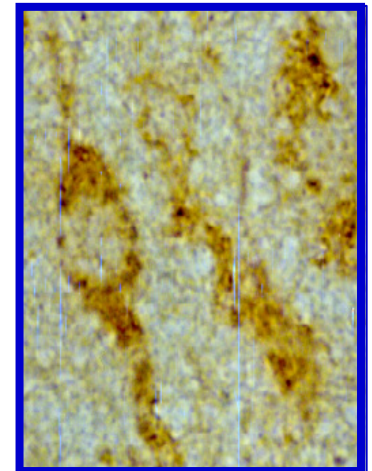
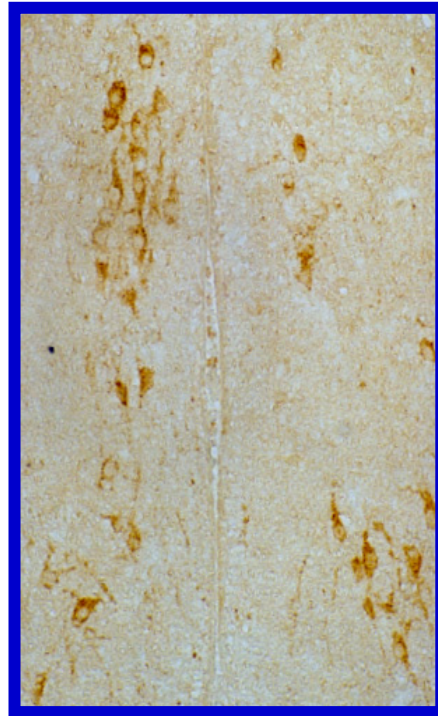
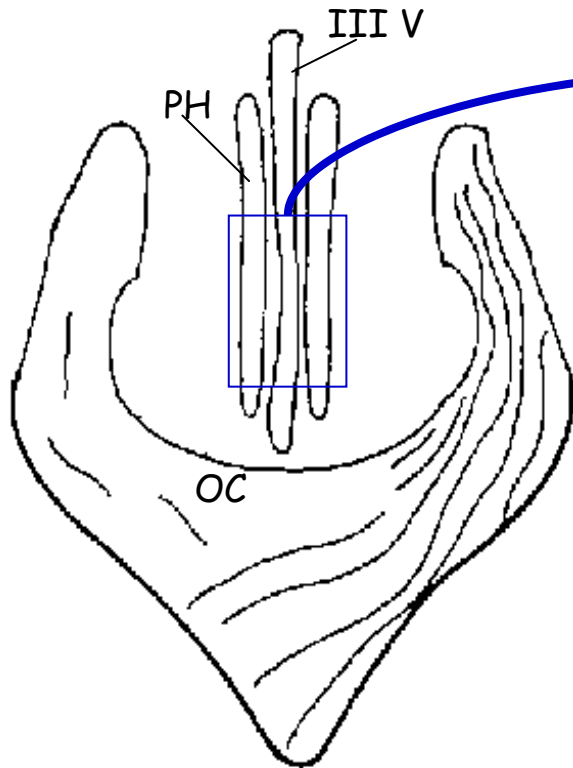
# Rh2 Group rhodopsin-like pigments (470-510 nm)



***Podarcis sicula* retinae  
in situ hybridization of RH2 opsin**



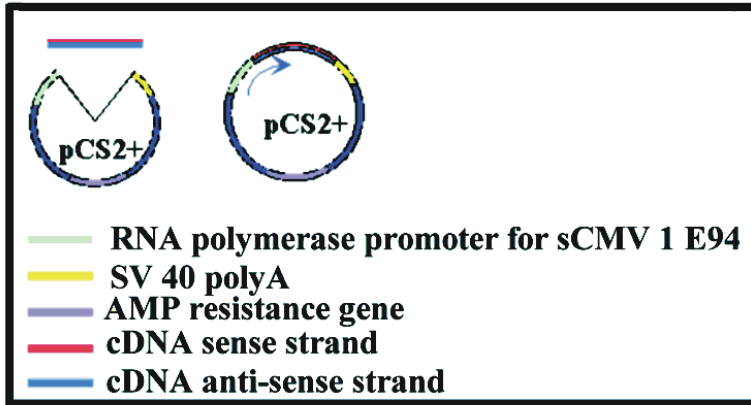
# IMMUNOCYTOCHEMICAL ANALYSIS OF RH2 EXPRESSION IN THE BRAIN



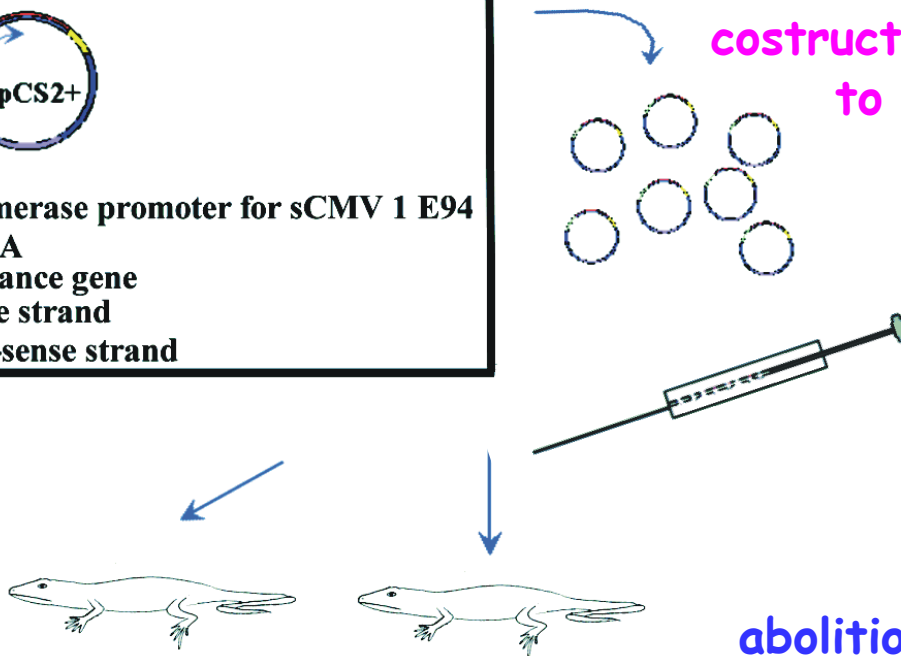
Anti-cone-opsin antiserum CERN 874 (specific for chicken Rh2 opsin) labelled neurons localized into the periventricular area (PH) of the hypothalamus, close to the border of the third cerebral ventricle  
OC: optic chiasm

# ARE BRAIN CONE-OPSINS NECESSARY FOR PHOTIC ENTRAINMENT OF LOCOMOTOR RHYTHMS?

Post-transcriptional inactivation experiments by injecting an eukaryotic expression vector transcribing the antisense cone-opsin RH2 mRNA in the third cerebral ventricle of pinx-retx lizards previously entrained to an LD cycle



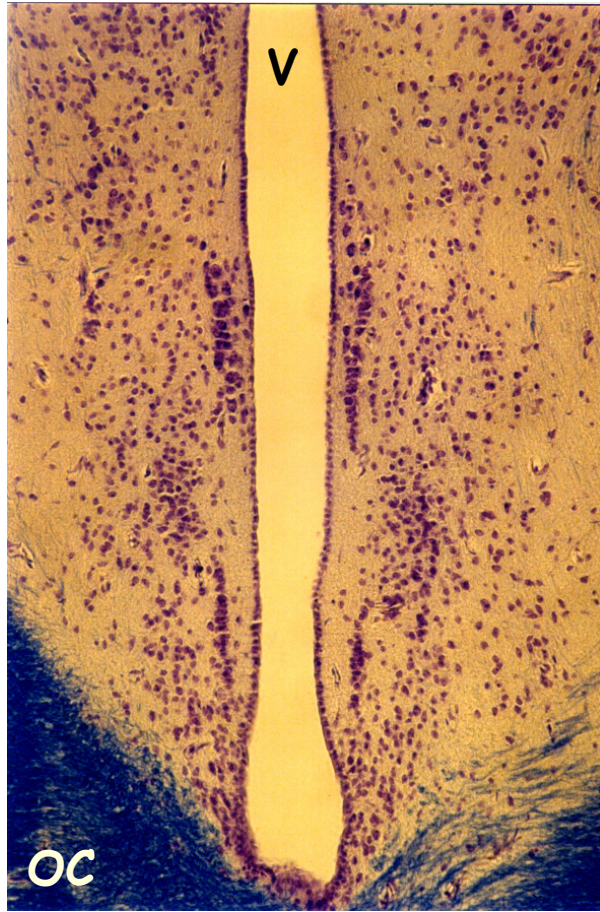
If after the injection with antisense construct, CERN 874 completely failed to label cells within the PH



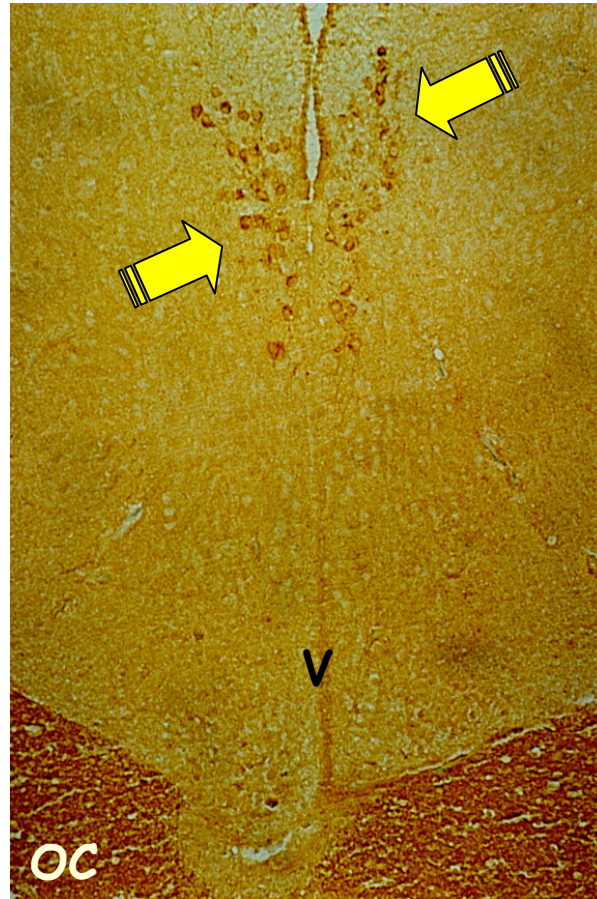
abolition of cone-opsins expression within the PH



# Histology



## Sense



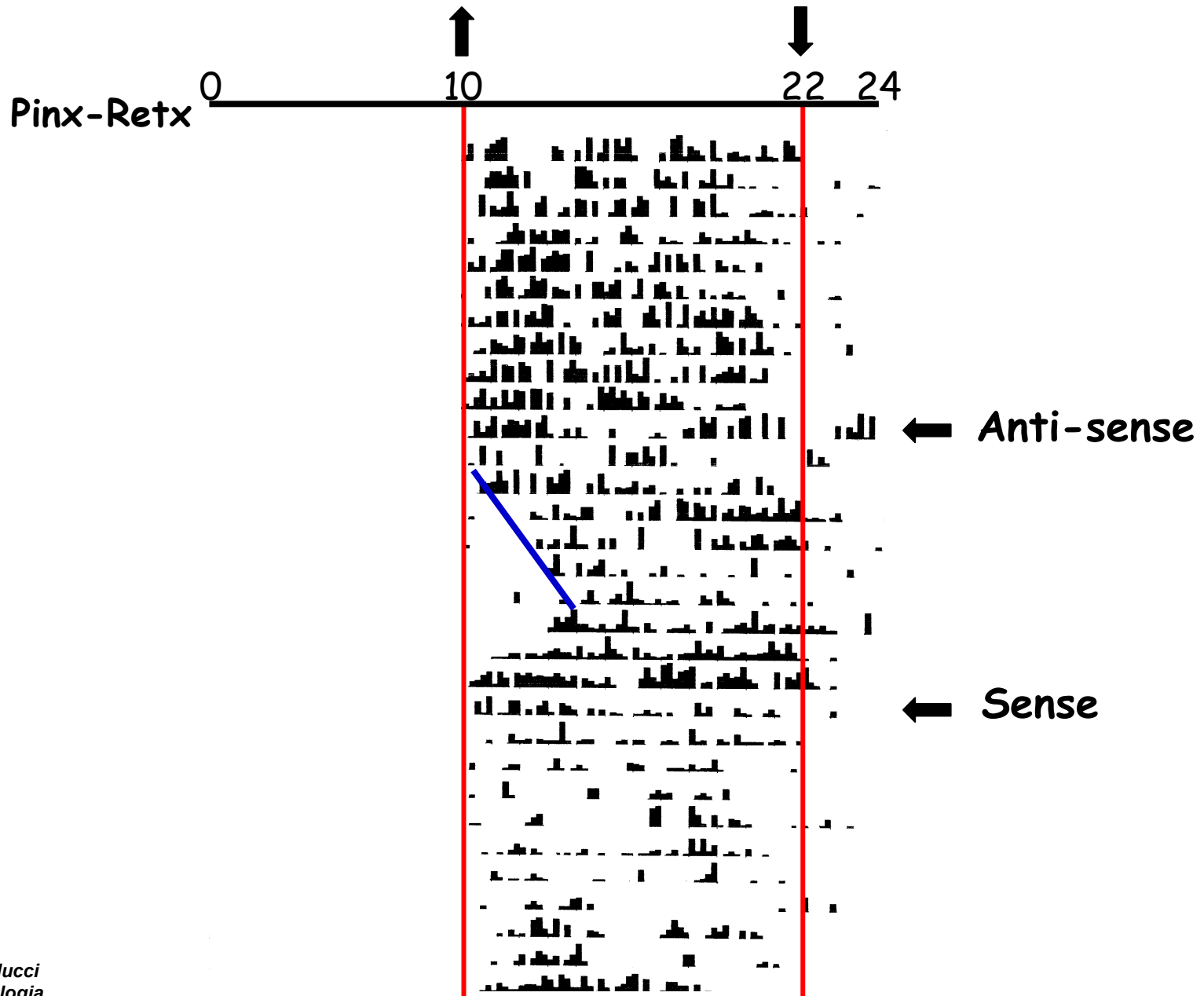
## Antisense



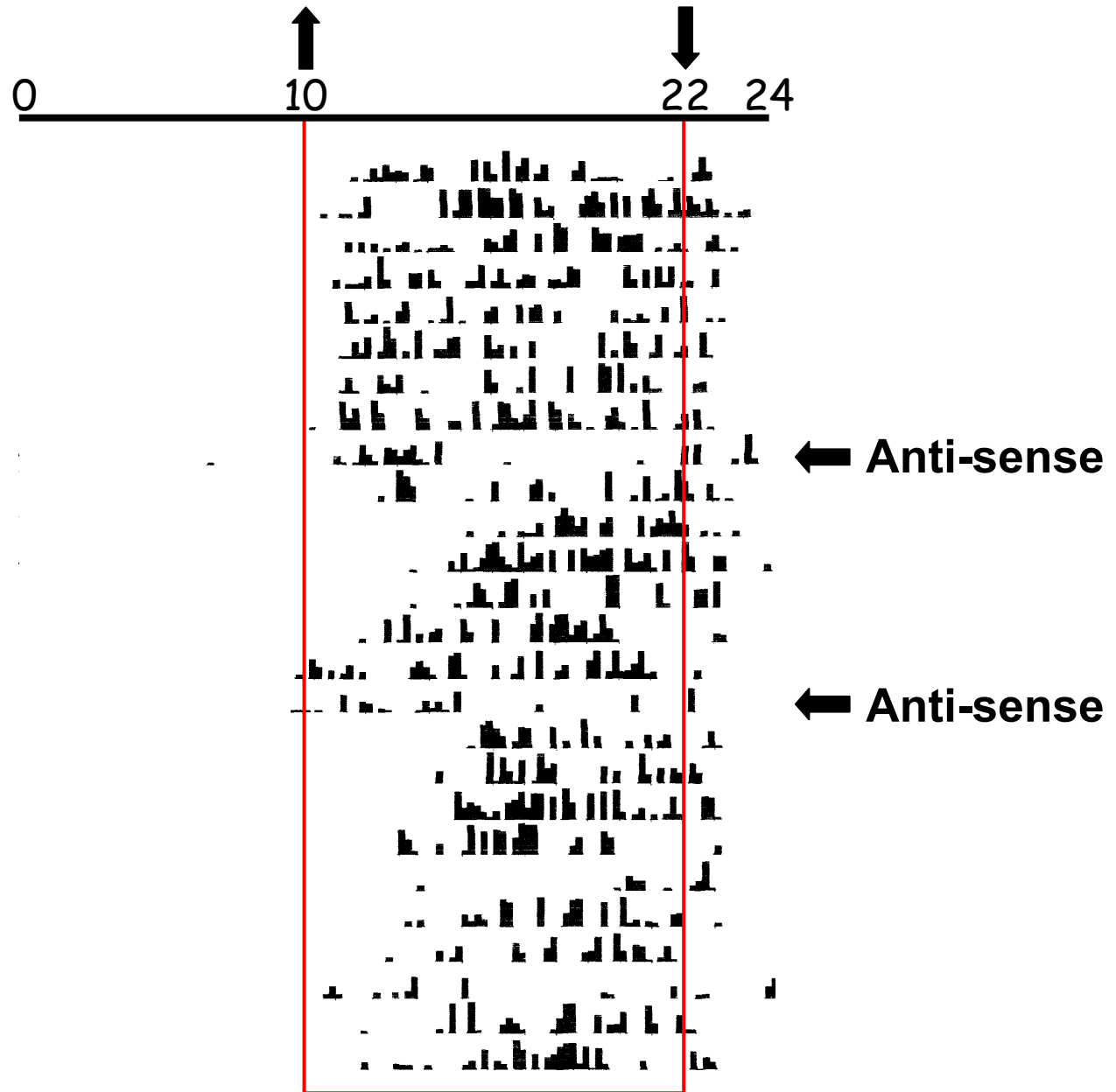
OC: Optic chiasm

V: III ventricle

# BEHAVIOURAL TESTS

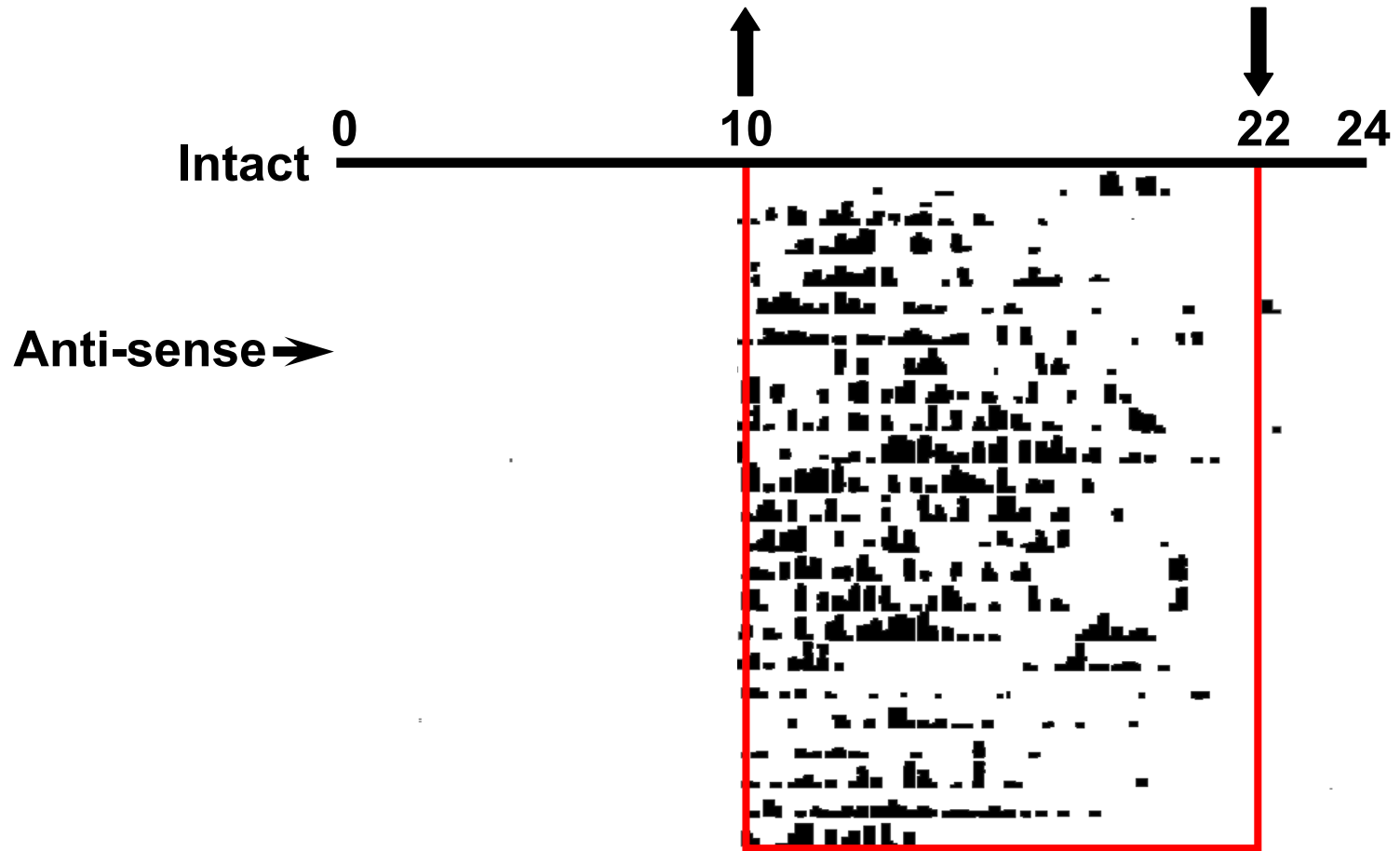


# BEHAVIOURAL TESTS





# BEHAVIOURAL TESTS



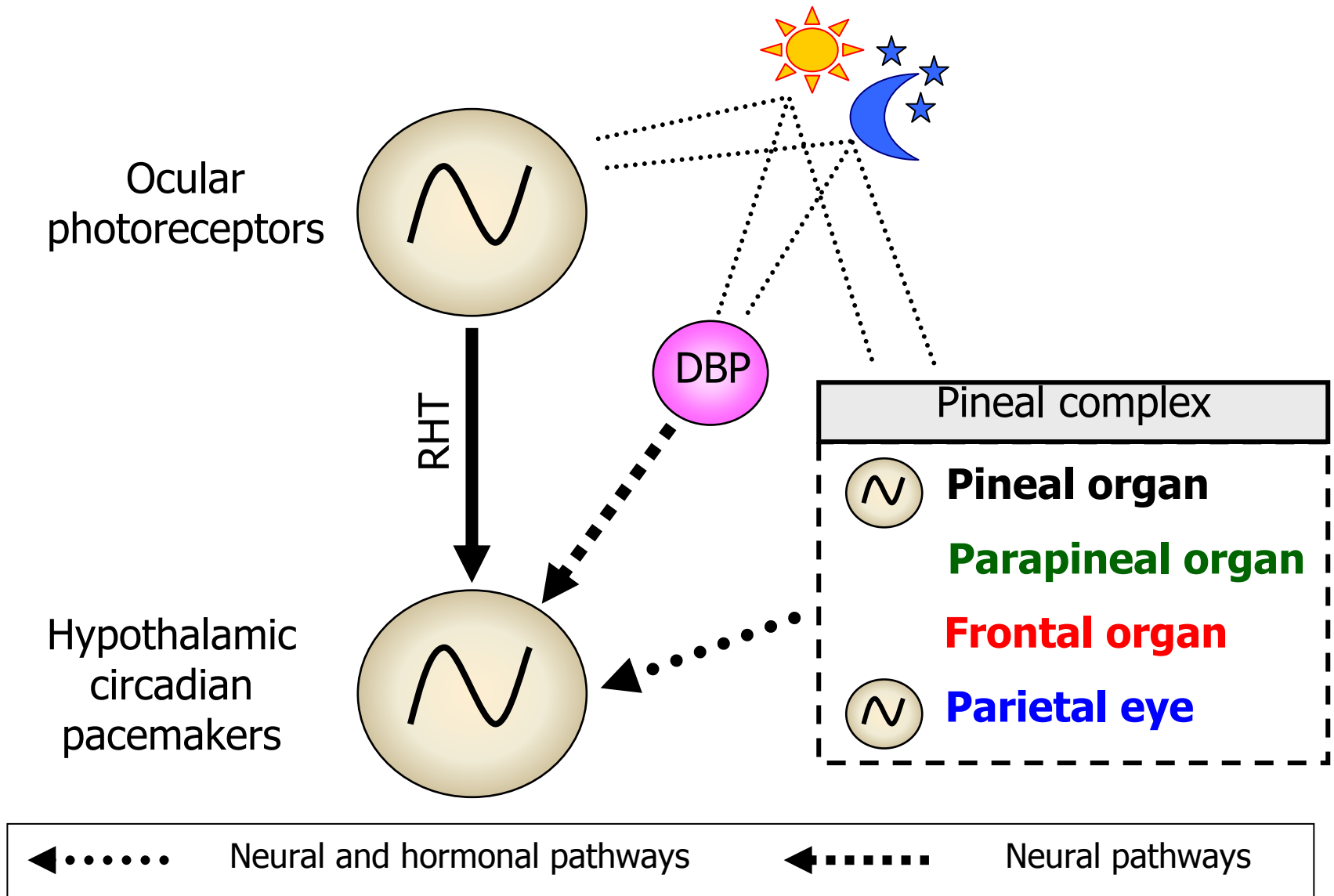
*Posttranscriptional inactivation experiments of endogenous brain cone-opsins mRNA demonstrate, for the first time in a vertebrate, that brain cone-opsins of lizards are part of a true circadian brain photoreceptor participating in photic entrainment of behavioral rhythms*

Experiments don't allow us to exclude the possibility that an antisense construct inactivated cone-opsins different from Ps-RH2, since differences among sequences of cone-opsins are often limited to few amino acids. Furthermore, polyclonal antiserum CERN 874 could recognize other cone-opsins inactivated by antisense treatments

MELANOPSIN      PINOPSIN      RHODOPSIN      ... ..

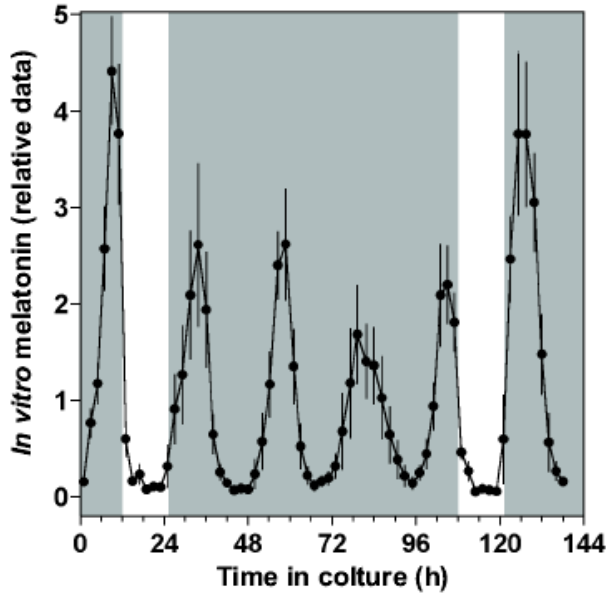


# BRAIN CIRCADIAN PHOTORECEPTION

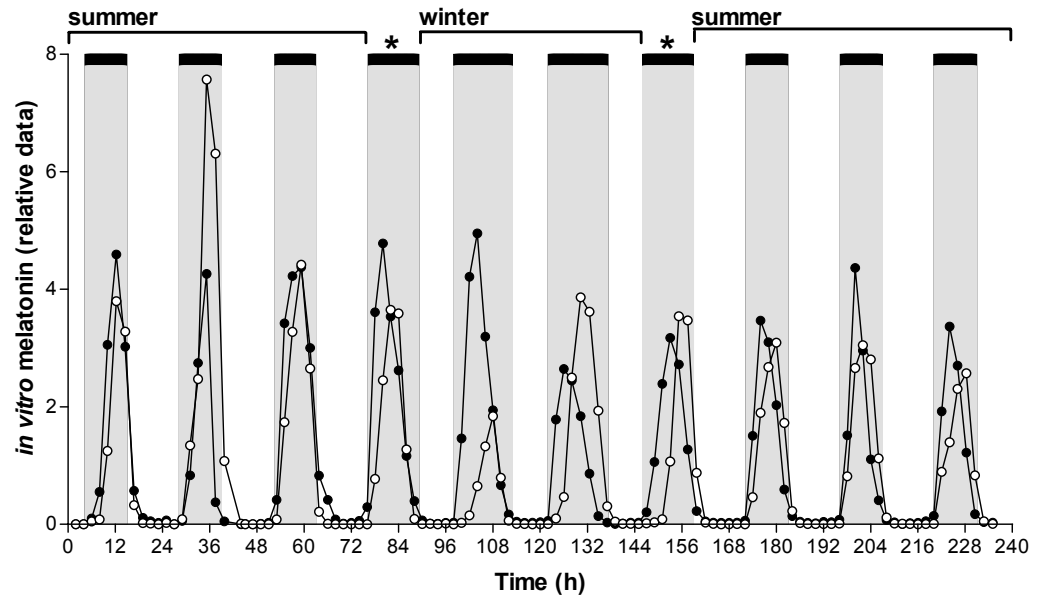
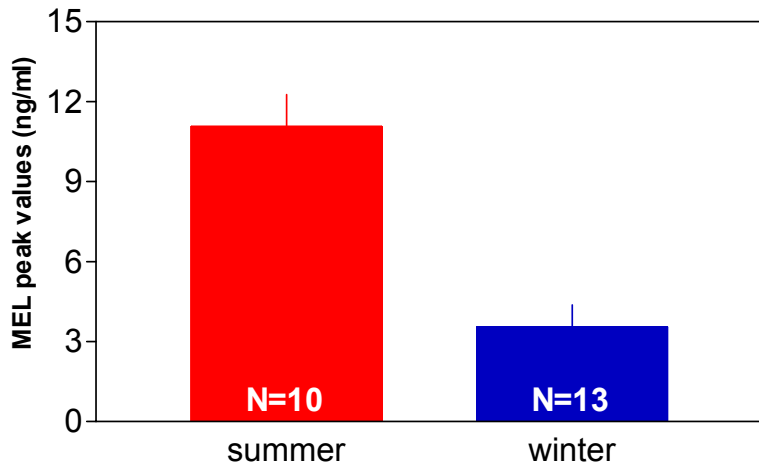
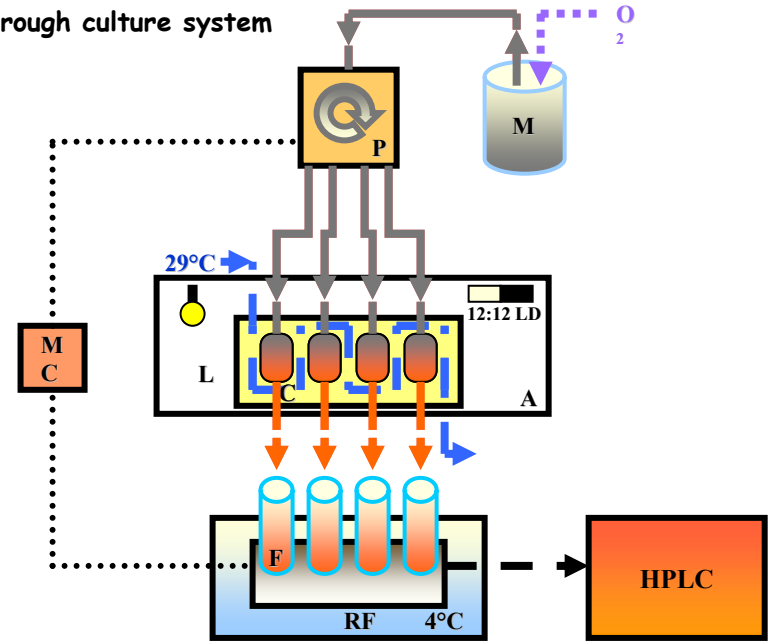


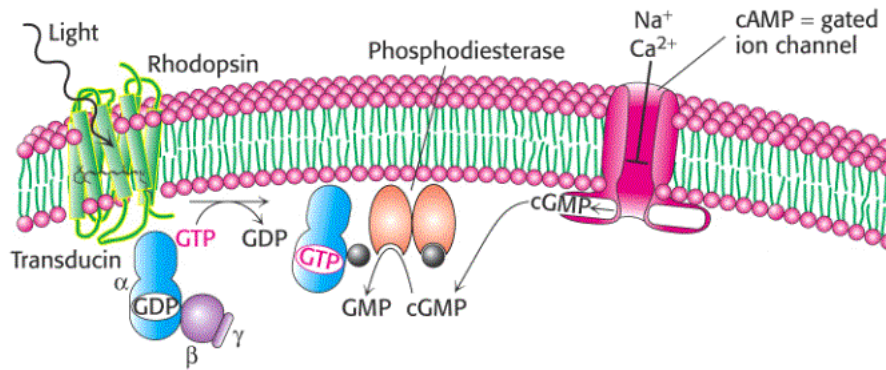


Bertolucci et al., 2003 JBR



Flow through culture system





Identification of  
non visual opsin  
in *Podarcis sicula*

**PINOPSIN**

**MELANOPSIN**

- RT-PCR with primers designed on the conserved regions of orthologs
- 5'-3' RACE for full-length cloning

**Phylogenetic analysis**

Maximun parsimony analysis

**Expression in photoreceptive organs**

# Pinopsin

attgcttgctcagcctcacctgatctctgtcccacatagaagaa  
**atg**caagcttcaaacgcttcatgggttgaggtgagaaacaggaccccaggcccttttgag  
M Q A S **N** A S W V E V R N R T P G P F E  
gggccgcaatggccctacctagccccacagagcacctacatatcagtggtgctcctcatg  
G P Q W P Y L A P Q S T Y I S V A V L M  
gggctggtgggtcatctcggccaccttgggtgaacggcttgggtgattggtgctcctgtccag  
G L V V I S A T L V N G L V I V V S V Q  
ttcaagaagctccggttctcctttgaactacgtcctgggtgaacctggccgtggctgatctc  
F K K L R S P L N Y V L V N L A V A D L  
ttggtcaccttctttgggagcaccatcagcttcgtaaacaatgccagggcttctttatc  
L V T F F G S T I S F V N N A Q G F F I  
tttggccaggcaacctgcaatgtgaaggcttcatgggtctccttaacagggattgtgggt  
F G Q A T **C** E F **E** G F M V S L T G I V G  
ctttggctccttggcgatcctggcctttgagaggtatctcgtgatctgcaaacgggtgggt  
L W S L A I L A F **E R Y** L V I C K P V G  
gatttccgcttcccagcaaggcatgcggtgcttgggtgtgctttcacttgggggtggteg  
D F R F P A R H A V L G C A F T W G W S  
ttcgtctggacgggtcccaccactcctcggatggagcagctacgtccctgaaggcttgaga  
F V W T V P P L L G W S S Y V P E G L R  
acatcctgcgcccccaactggtagctggcggaagcagcaacaacagctatatcatgact  
T S **C** G P N W Y S G G S S N N S Y I M T  
ttgtttgtgacctgctttgcatgacctctcagcacaatcctcttctcctacgccaacttg  
L F V T C F A M P L S T I L F S Y A N L  
ctcatgacgctgcaaacagtagcggctcagcagaaagagcaggagacaactcagagggcc  
L M T L R T V A A Q Q K E Q E T T Q R A  
gagagggaggtgacacgaatggttgcgccatggtggctgcttccctcgtctgctggctg  
E R E V T R M V V A M V A A F L V C W L  
ccctacgcccagcttcgccatggtgggtcgccaccacacaaggacctcgccatacggccagcg  
P Y A S F A M V V A T H **K** D L A I R P A  
ctcgcttccctgccatcatatttctccaagacggcgacagtgtagaaccatcatctat  
L A S L P S Y F S K T A T V Y N P I I Y  
gtcttcatgaacaagcagttccgcagttgcctcctctacaagatgagctgtggccacaga  
V F M N K Q F R S C L L Y K M S **C** G H R  
gcattatcttcccaggacacgacaccagctgggatcagcctgccaggccgcctcaccacc  
A L S S Q D T T P A G I S L P G R L T T  
tcagcttcaaaaggaagcaggaatcaagtgtcgccttcc**tta**acgatgc  
S A S K G S R N Q V S P S

*Gallus gallus* 69%

*Columba livia* 72%

*Anolis carolinensis* 77%

*Phelsuma madagascariensis* 72%

*Bufo japonicus* 68%

**N**: potential site for N-glycosylation (Asn<sup>5</sup>)

**C**: cysteines for an intramolecular disulfide bond (Cys<sup>106</sup>, Cys<sup>183</sup>)

**E**: glutamic acid for retinylidene Schiff-base counterion (Glu<sup>109</sup>)

**K**: lysine for Schiff-base linkage with a chromophore 11-cis-retinal (Lys<sup>273</sup>)

**C**: cysteines for palmitoylation (Cys<sup>316</sup>)

**ERY** motif at position 130-132 necessary to bind the  $\alpha$ -subunit of transducin



acgcggggggcggaagcatggcagaggcgcacgagcagacacagaat  
**atg**ggaactcagcaccgaataaaaagtagatgttcctgatcgtgttctttacactgtaggc  
 M G T Q H R I K V D V P D R V L Y T V G  
 tcctgtgttcttgtcattggttctattggaatcacaggaaatcttcttgtcctctatgca  
 S C V L V I G S I G I T G N L L V L Y A  
 ttttacagcaacaagcgtctgaggacaccggcaaactatttcataatgaattagcggca  
 F Y S N K R L R T P A N Y F I M N L A A  
 agtgattttctgatgtctgcaactcaagctccaatctgctttctcaacagcatgcacaca  
 S D F L M S A T Q A P I C F L N S M H T  
 gaatggatactcggagacatagggttgtaacttttatgtattttgtggggcactccttggga  
 E W I L G D I G **C** N F **Y** V F C G A L F G  
 ataacttcaatgatgactttattagctatttcagttgatcgctactgtgtgattactaag  
 I T S M M T L L A I S V D R Y C V I T K  
 cctctgcagtctataaaaagggtcttcaaagaaacgctcatgcatcattgcctttgtc  
 P L Q S I K R S S K K R S C I I I A F V  
 tggctctactcgctgggatggagtgtatgtcctctctttgggatggagttcctatatacct  
 W L Y S L G W S V C P L F G W S S Y I P  
 gaaggtttgatgatatcttgtacatgggactacgtatcctattctccagcaaacagaagt  
 E G L M I S **C** T W D Y V S Y S P A N R S  
 tataccatgatgttatgttgctttgtgttctttatccctctgataataatcttccactgc  
 Y T M M L C C F V F F I P L I I I F H C  
 tatctattcatgtttctggccattagaagtactggcaggaacgttcagaagttaggatca  
 Y L F M F L A I R S T G R N V Q K L G S  
 acctataaccggaagtcgaatgtttcacagtcagtgaaagtgaaatggaagtttagcaaaa  
 T Y N R K S N V S Q S V K S E W K L A K  
 attgcctttgtggctattgtgggtgtttgttctgtcctggctcctcctatgctta  
 I A F V A I V V F V L S W S P Y A

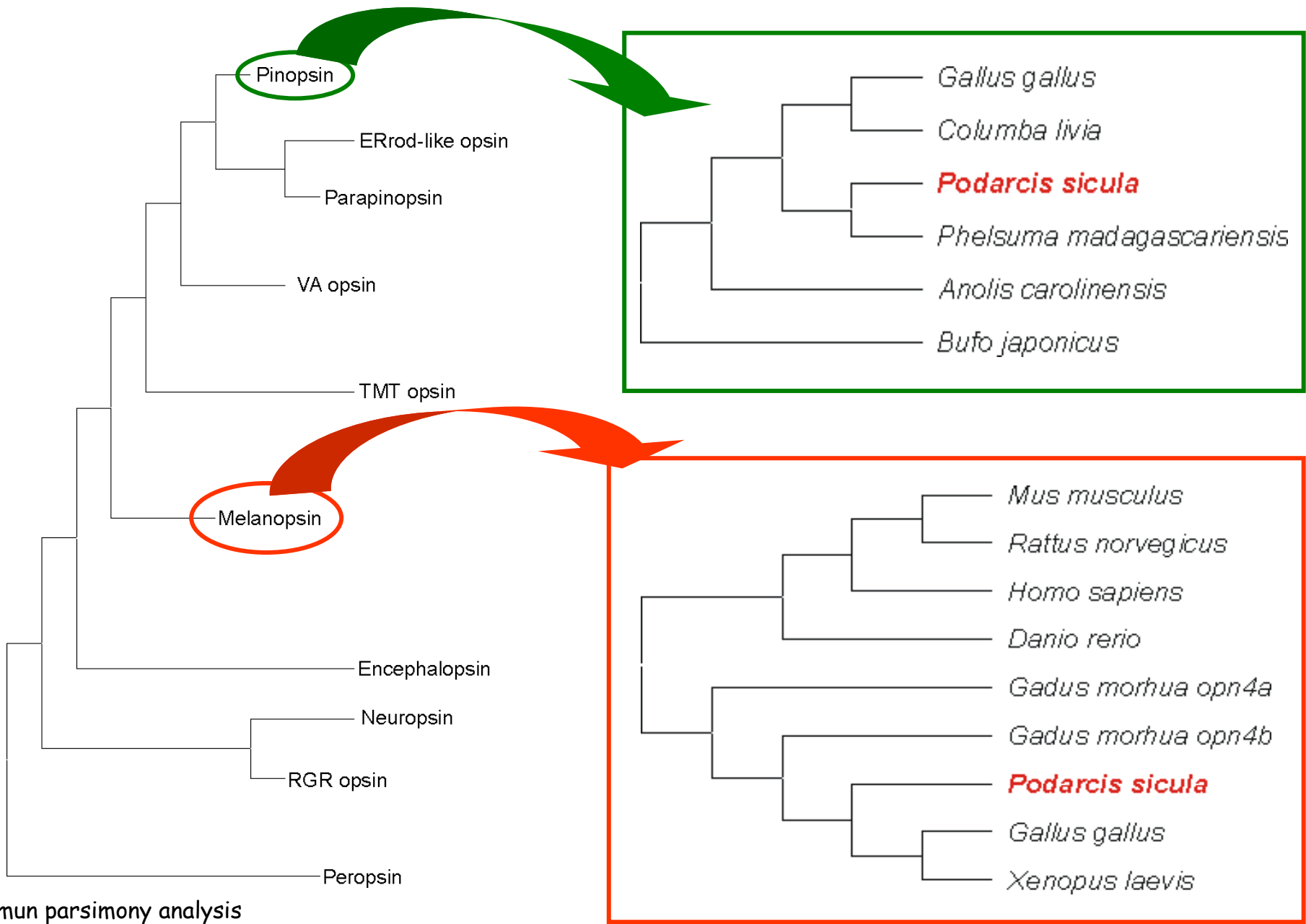
# Melanopsin

<i>Homo sapiens</i>	53%
<i>Mus musculus</i>	54%
<i>Rattus norvegicus</i>	53%
<i>Gallus gallus</i>	80%
<i>Xenopus laevis</i>	75%
<i>Gadus morhua</i>	64%
<i>Danio rerio</i>	48%
<i>Branchiostoma floridae</i>	40%

**C:** Conserved sites disulfide bridge formation (Cys89, Cys167)

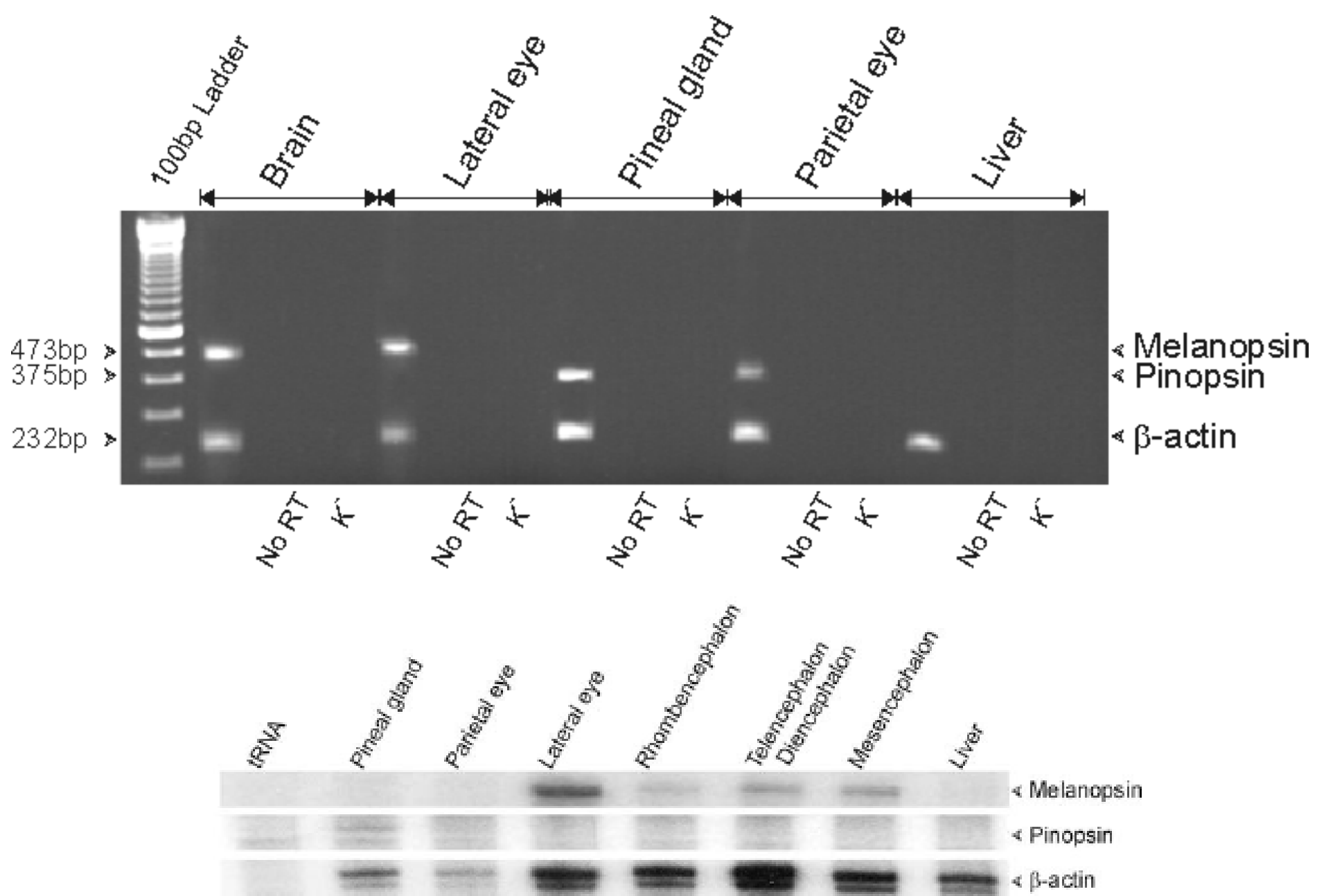
**Y:** aromatic residue conserved in all invertebrates (Tyr92)

NON VISUAL OPSIN



Maximun parsimony analysis





# EVOLUTION IN THE DARK

