

Transporters

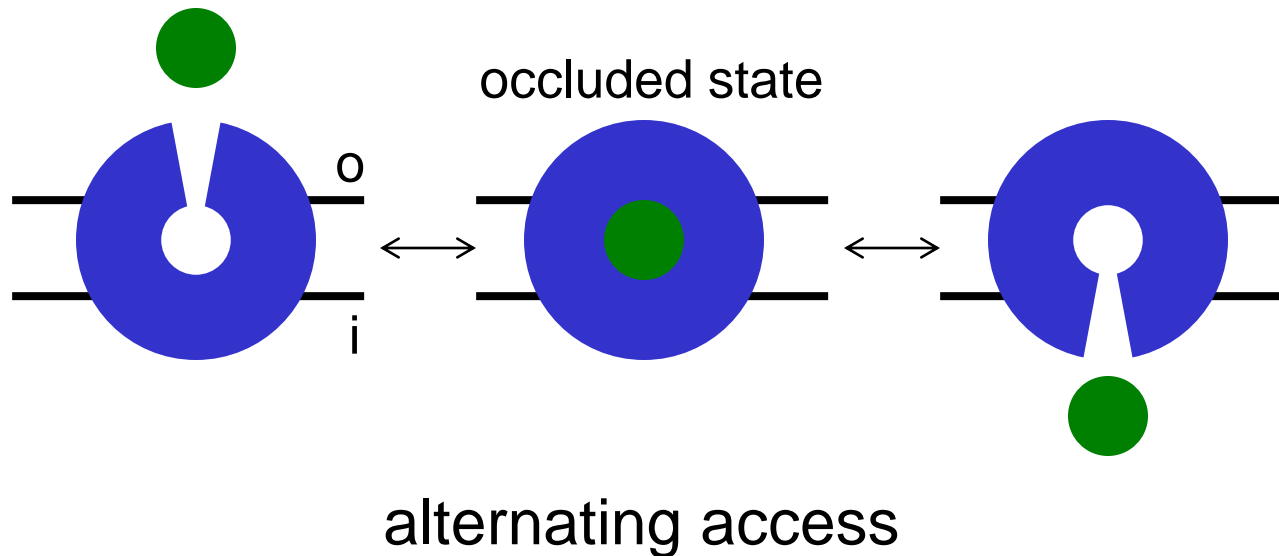
different from channels:

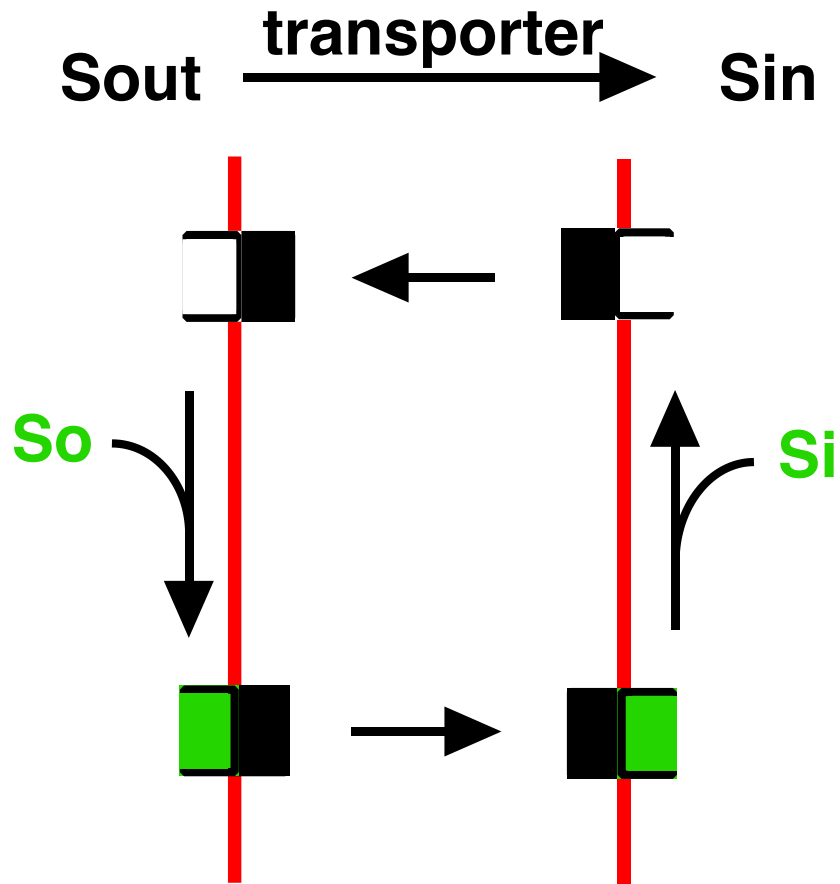
speed

saturation

concentration

--not exactly

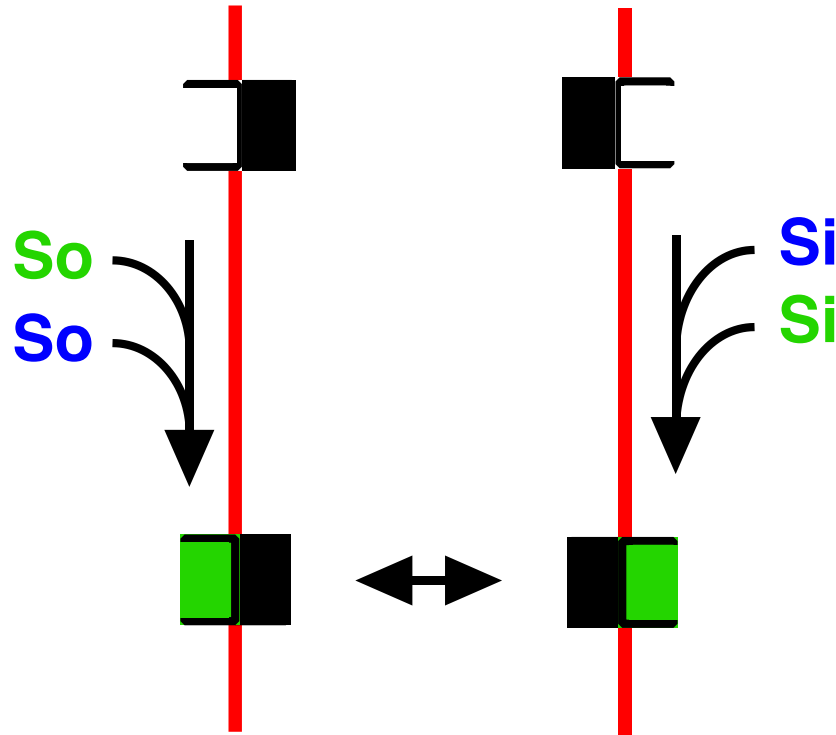




movement of unloaded carrier crucial for net transport

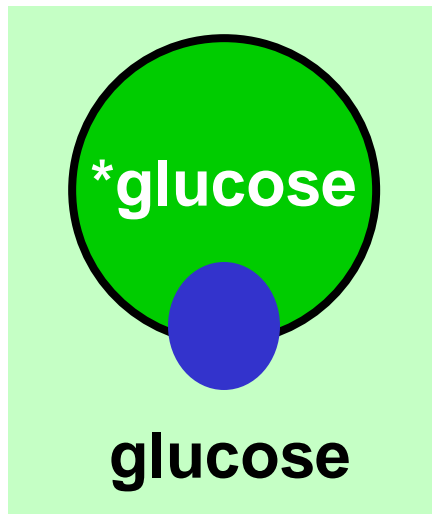
exchange

$S_{out} + S_{in}$ $\xrightarrow{\text{transporter}}$ $S_{in} + S_{out}$



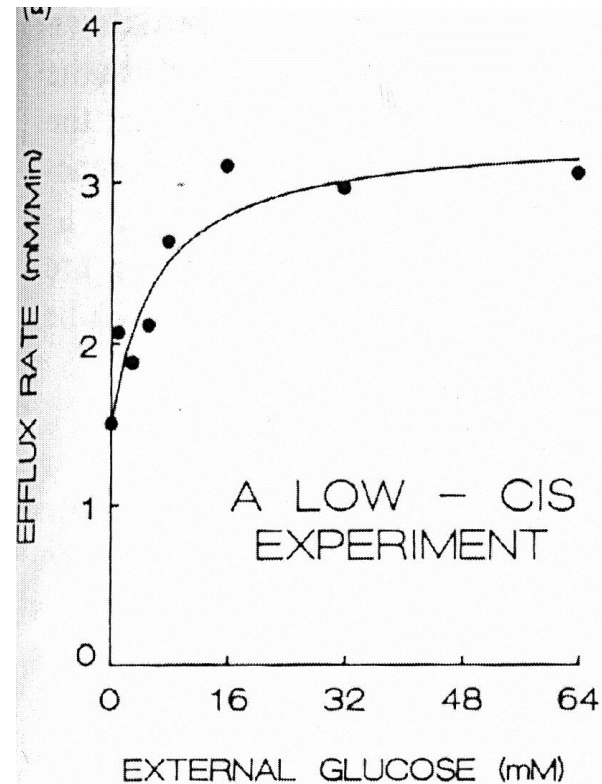
--no net flux

membranes preloaded with ^{14}C -glucose
diluted into medium with unlabelled glucose

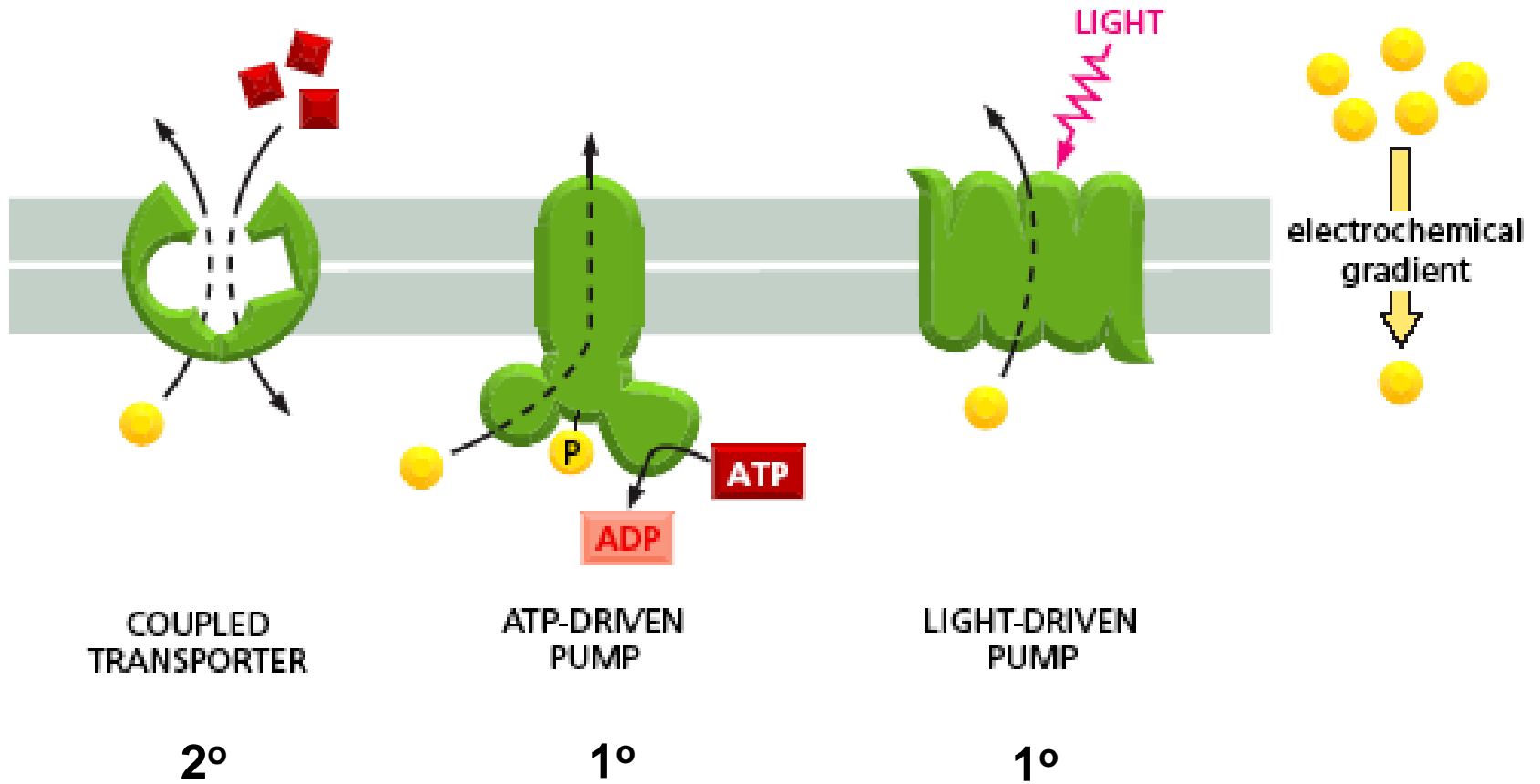


only loaded carrier moves
—not unloaded
translocation steps slow
--channel cannot do this

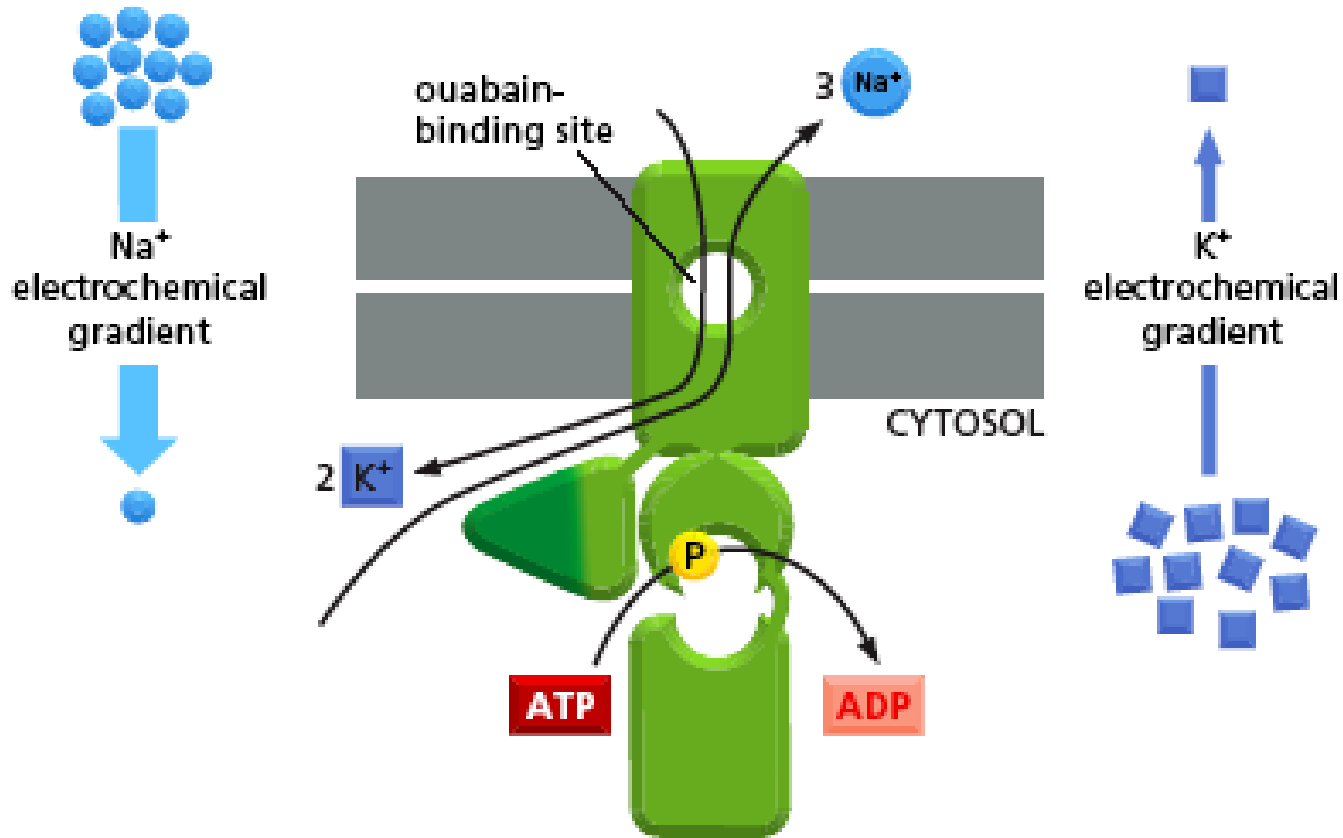
heteroexchange
obligate exchangers cannot do net flux
amphetamines may release monoamines this way (exchange-diffusion)



different classes

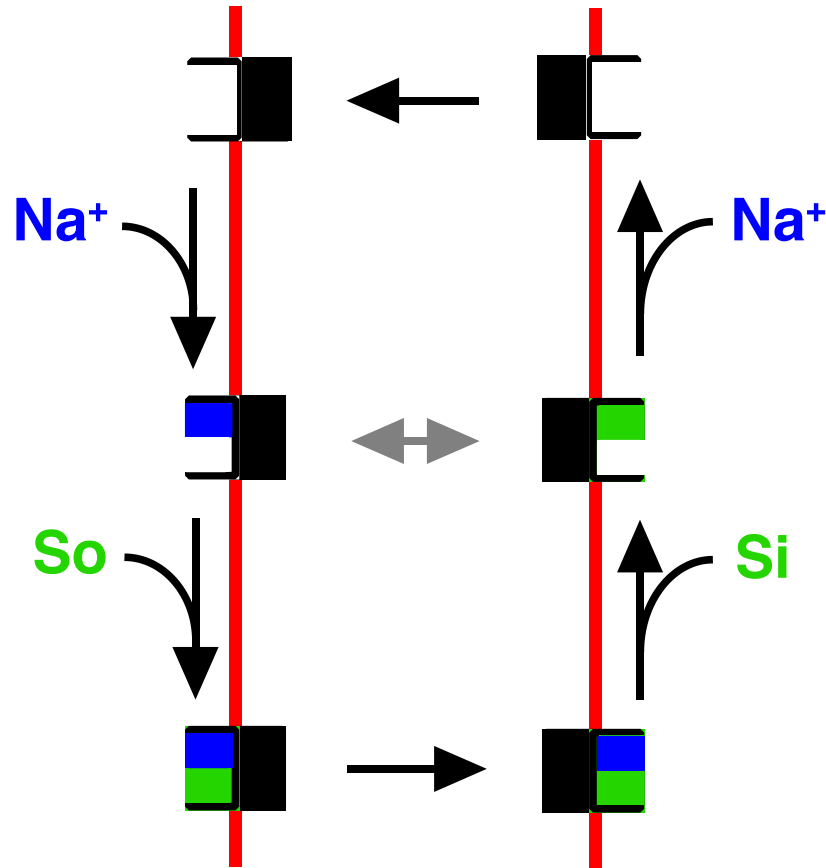


P-type ATPase: Na⁺/K⁺-ATPase



P-type involve phosphorylated intermediate
how can phosphorylation outside membrane
trigger movement of ions across membrane?

ionic coupling



coupling rules: Na^+ can only move with S --why?
AND gate, ~coincidence detector

equilibrium



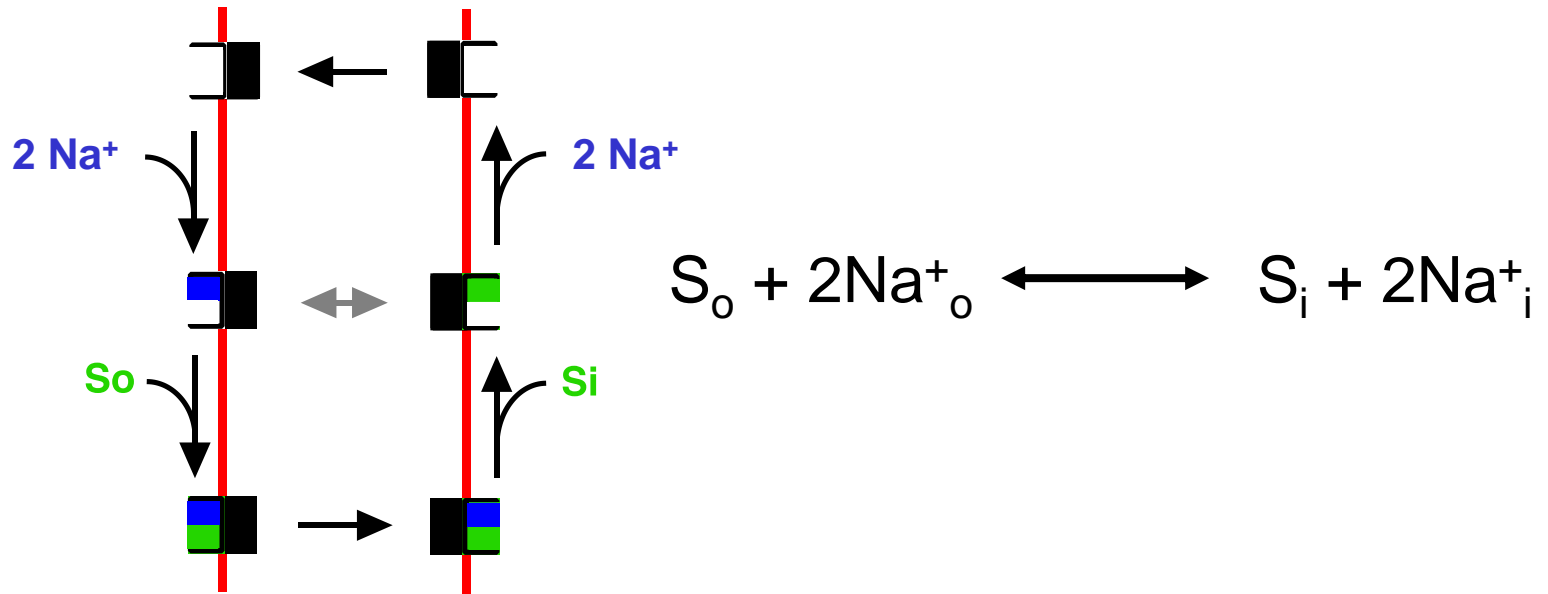
stoichiometry of 1 Na⁺ : 1 S and ~12-fold Na⁺ gradient
will generate ? gradient of S

at equilibrium, equal rates in and out of cell

$$[Na^+]_o \times [S]_o = [Na^+]_i \times [S]_i$$

$$[Na^+]_o / [Na^+]_i = [S]_i / [S]_o$$

what if it is an exchanger?



if coupling involves $2 Na^+ : 1 S$, then

$$[Na^+]_o^2 \times [S]_o = [Na^+]_i^2 \times [S]_i$$

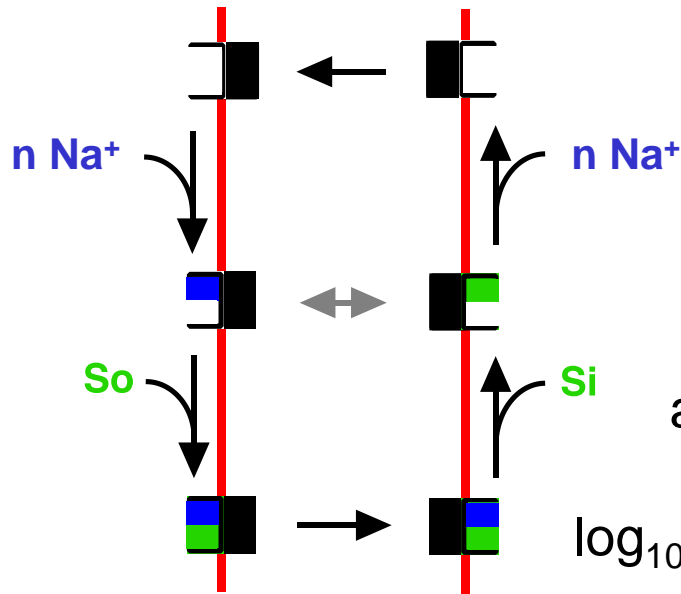
$$([Na^+]_o / [Na^+]_i)^2 = [S]_i / [S]_o \text{ or}$$

$$\log_{10} (S_{in}/S_{out}) = 2 \log_{10} (Na^+_{out}/Na^+_{in})$$

why not just make the stoichiometry very high?

what if net flux involves charge movement?

electrogenic transport (transport that moves net charge)



negative resting membrane potential
augments chemical gradient for Na^+ by

$$- z_T \Delta \Psi / 60 \text{ mV}$$

where $z_T =$ net charge moved and $\Delta \Psi$ is Vm

added to the concentration gradient,

$$\log_{10} (S_{in}/S_{out}) = n \log_{10} (\text{Na}^+_{out}/\text{Na}^+_{in}) - z_T \Delta \Psi / 60 \text{ mV}$$

where $n = \# \text{Na}^+$ ions cotransported

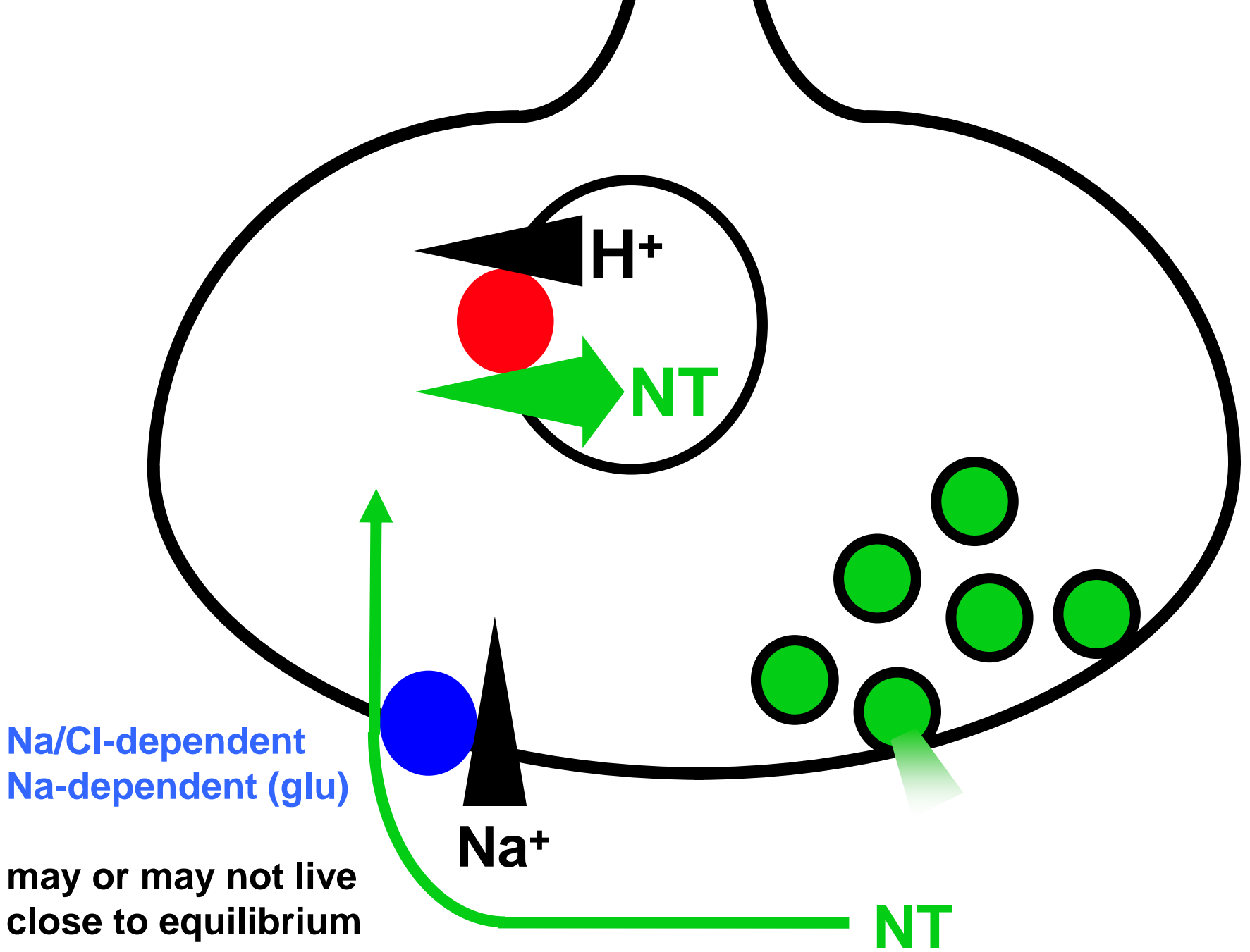
--power of membrane potential

equation changes for different ionic coupling

ionic coupling determines direction of flux

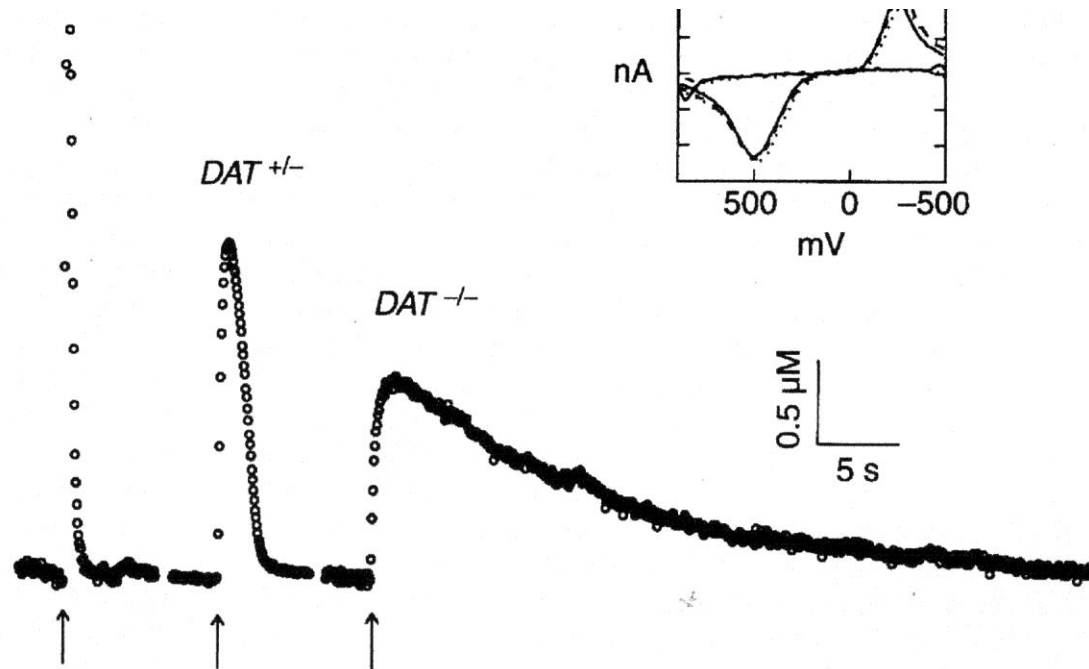
magnitude of gradient (can exceed $10^6:1$)

regulation by membrane potential



reuptake: Na/Cl-dependent transport

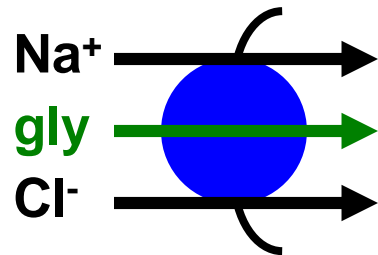
striatal slice
voltammetry



DAT KO:

impaired **rate** of dopamine *clearance* in striatum
also, KO has 95% decrease in dopamine stores!
--crucial role in *recycling*

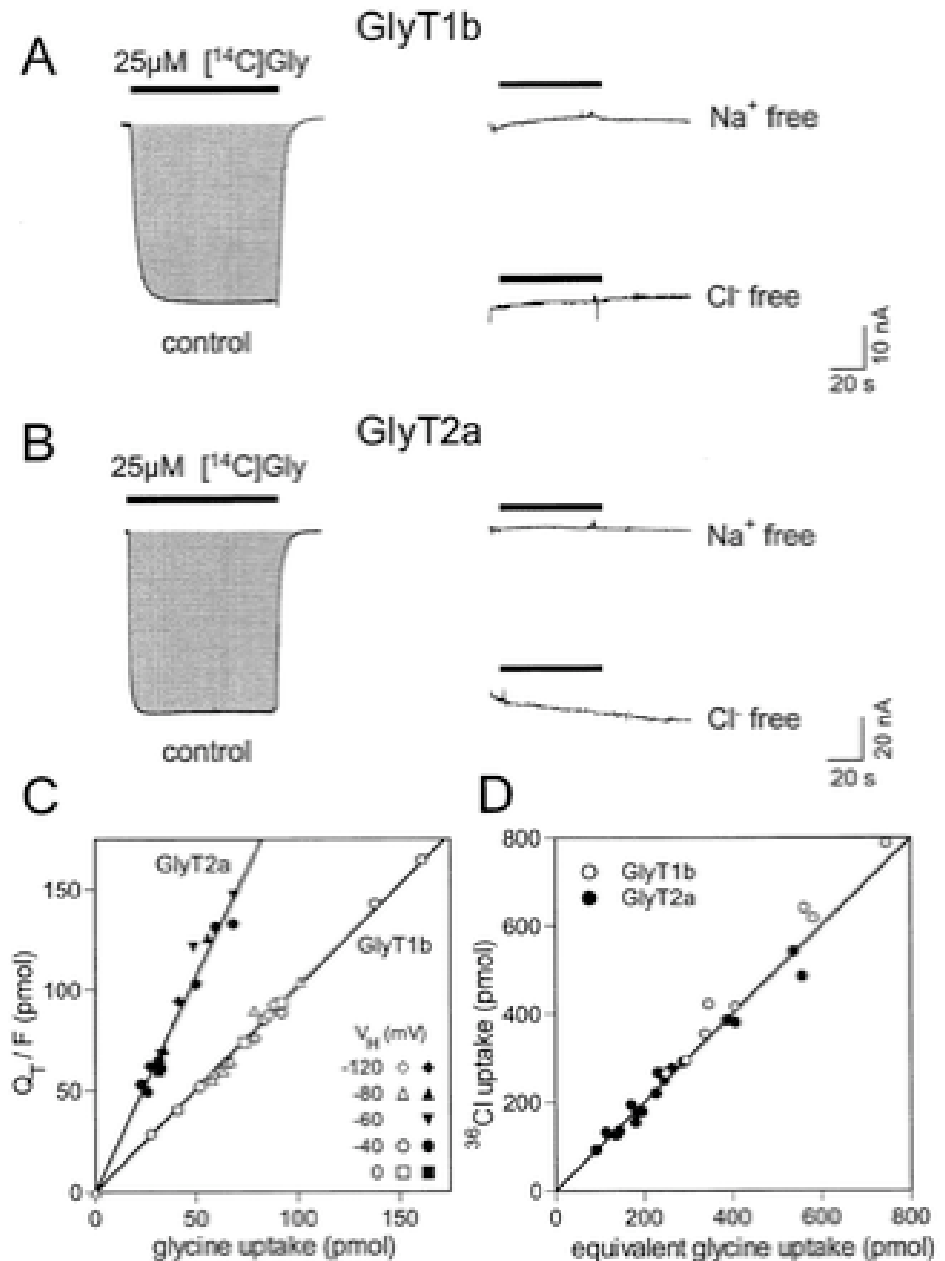
glycine transport



electrogenic transport
 produces currents: **rate**
 --depend on Na^+ , Cl^-

defined by gly addition
 --strictly rectifying

can measure charge:flux
 using labeled glycine, Cl
 suggests fixed stoichiometry
 (Roux and Supplisson, 2000)



for electrogenic glycine transport,

$$\log_{10} (\text{gly}_{\text{in}}/\text{gly}_{\text{out}}) = m \log_{10} (\text{Na}^+_{\text{out}}/\text{Na}^+_{\text{in}}) + n \log_{10} (\text{Cl}^-_{\text{out}}/\text{Cl}^-_{\text{in}}) - z_T \Delta\Psi / 60 \text{ mV}$$

$$z_T \Delta\Psi / 60 \text{ mV} = \log_{10} \frac{\text{Na}^+_o{}^m \times \text{Cl}^-_o{}^n \times \text{gly}_o}{\text{Na}^+_i{}^m \times \text{Cl}^-_i{}^n \times \text{gly}_i}$$

$$\Delta\Psi = \frac{60 \text{ mV}}{z_T} \log_{10} \frac{\text{Na}^+_o{}^m \times \text{Cl}^-_o{}^n \times \text{gly}_o}{\text{Na}^+_i{}^m \times \text{Cl}^-_i{}^n \times \text{gly}_i} = E_{\text{rev}}$$

--like Nernst equation:

$$E_{\text{Na}} = 60 \text{ mV} \log_{10} \text{Na}^+_o/\text{Na}^+_i$$

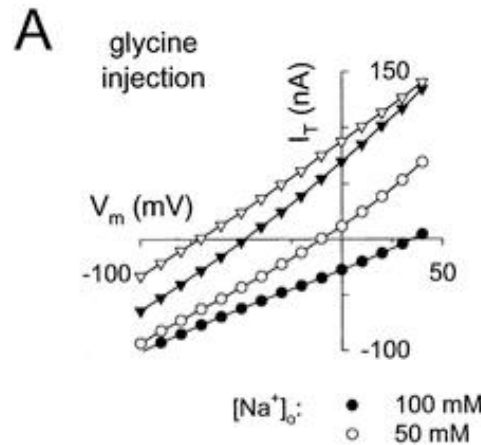
what are the differences?

$$E_{rev} = \frac{60 \text{ mV}}{(m_{Na} - n_{Cl})} \log \frac{Na^+_o{}^m \times Cl^-_o{}^n \times gly_o}{Na^+_i{}^m \times Cl^-_i{}^n \times gly_i}$$

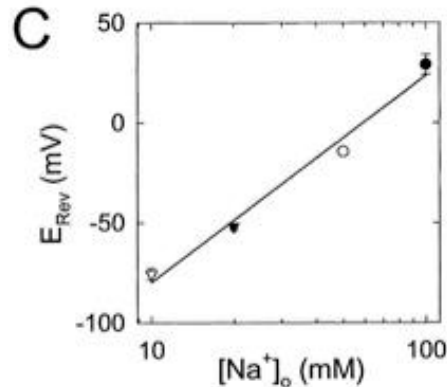
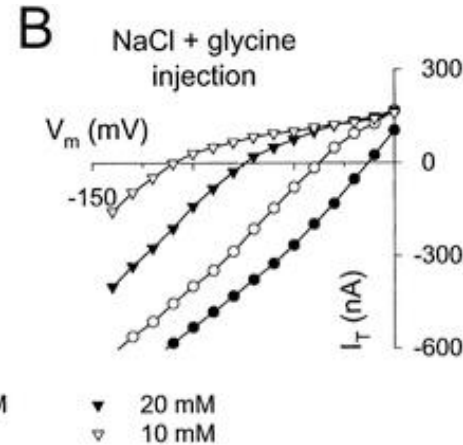
assume n = 1

can use E_{rev} at different ionic gradients to determine n, p
 BUT S-induced currents rectify--need them to reverse

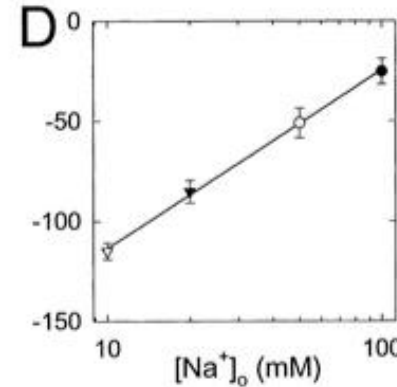
GlyT1



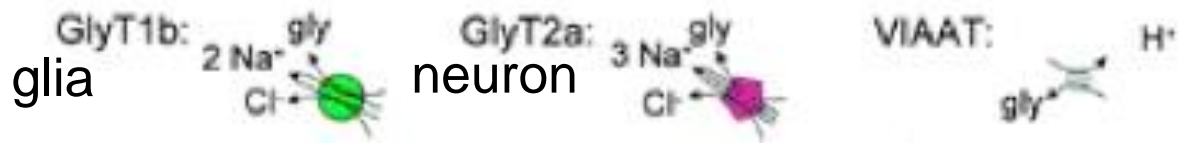
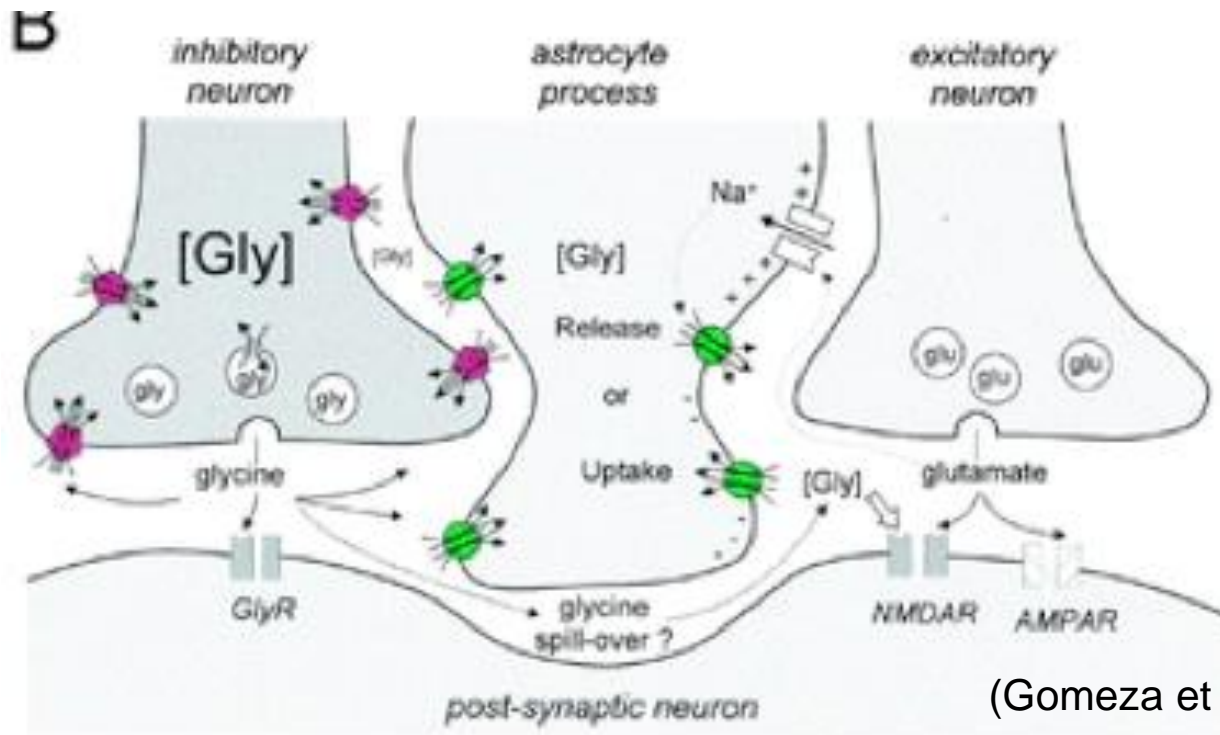
GlyT2



m = ?



m = ?



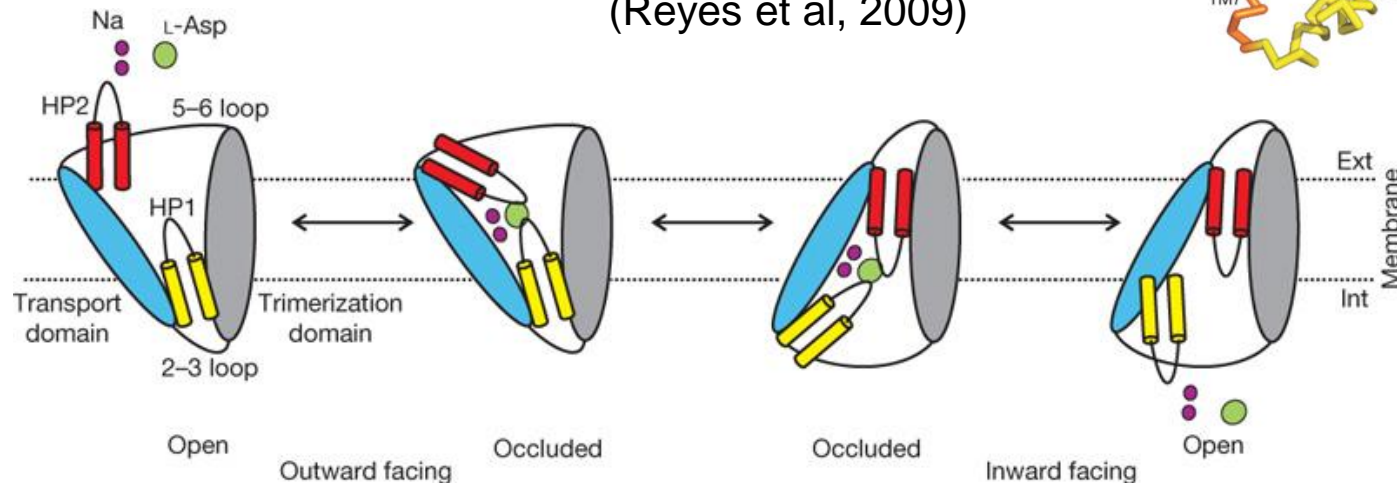
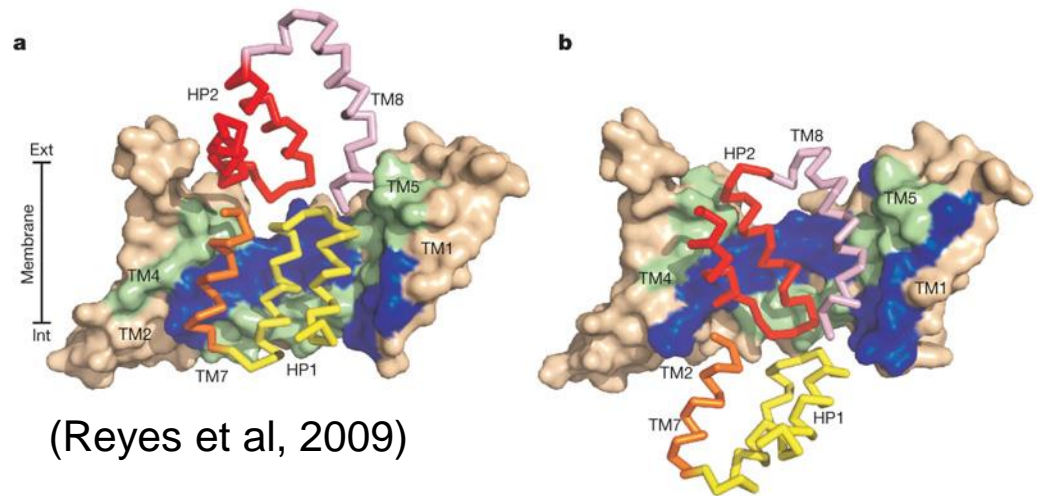
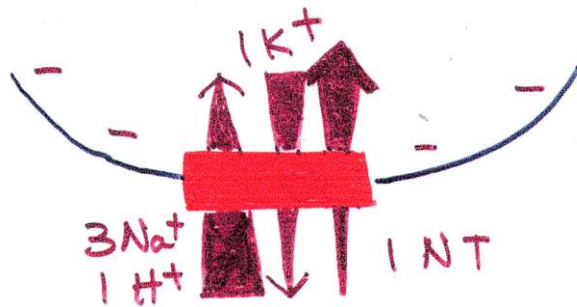
GlyT1 (2 Na^+) allows higher gly_o than GlyT2 (3 Na^+)
 --to activate NMDA-type glutamate receptors?

GlyT1 KO: excess glycine (excess inhibition)--main role clearance
 GlyT2 KO: resembles GlyR KO (startle)--main role packaging
 differences in ionic coupling can also confer transfer between cells

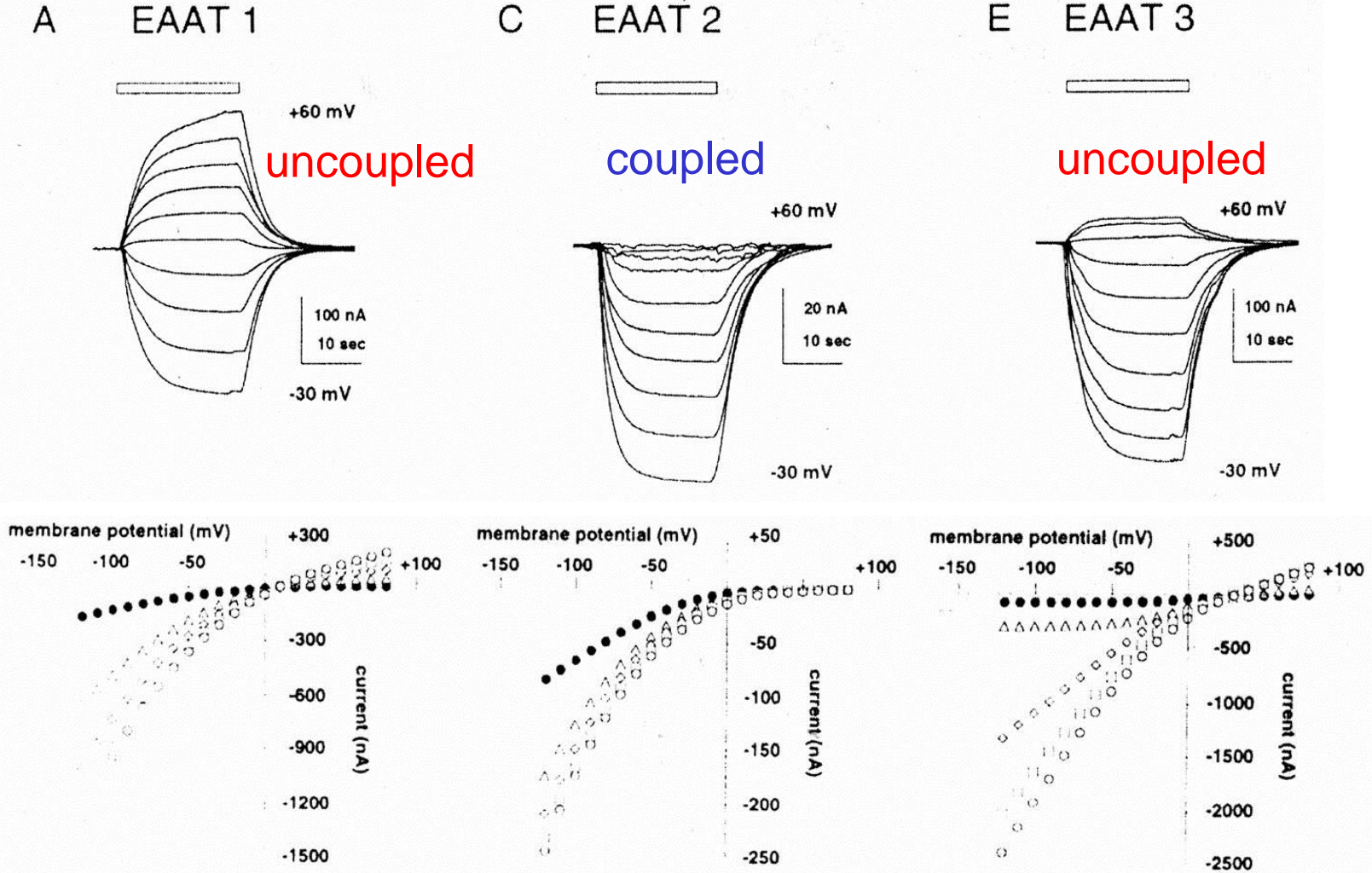
can these transporters release neurotransmitter? how?

excitatory amino acid transporters (EAATs)

rigid body motion



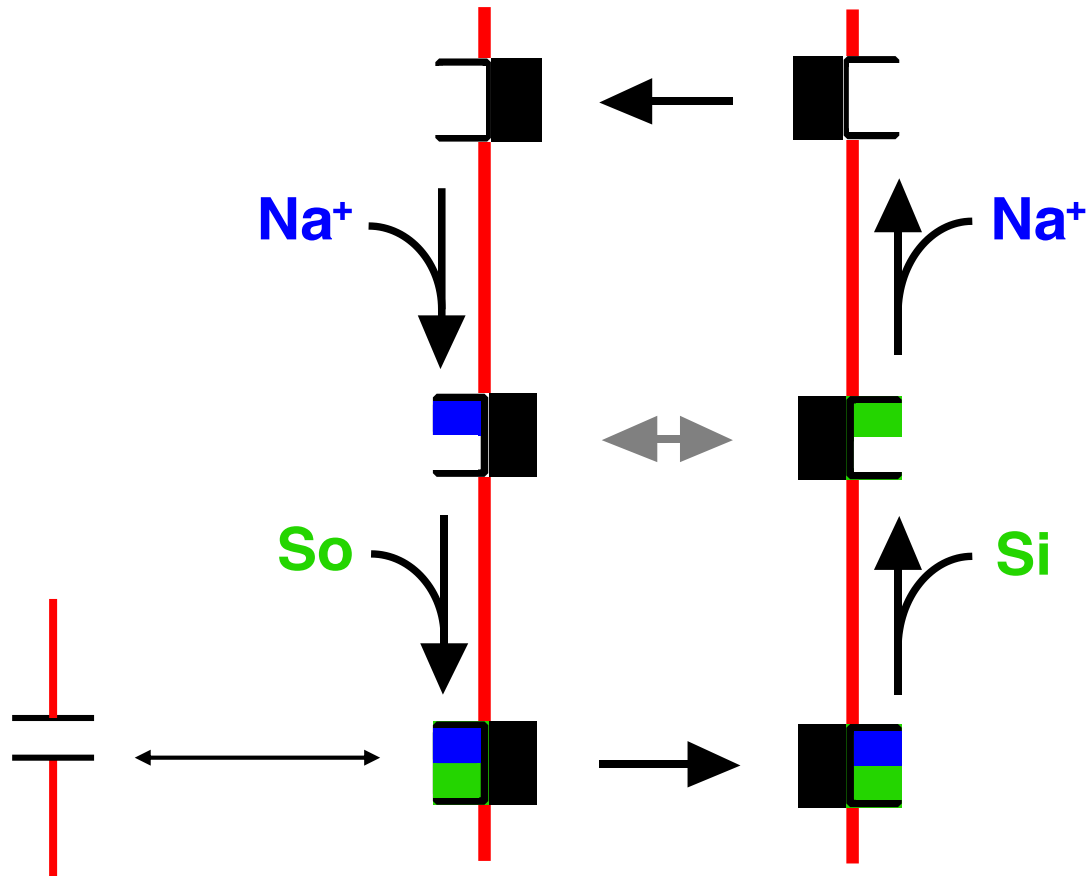
little effect on kinetics of EPSC (buffer—translocation too slow)
 controls activation of perisynaptic receptors, spillover
longer-term effects (seizures, degeneration): equilibrium



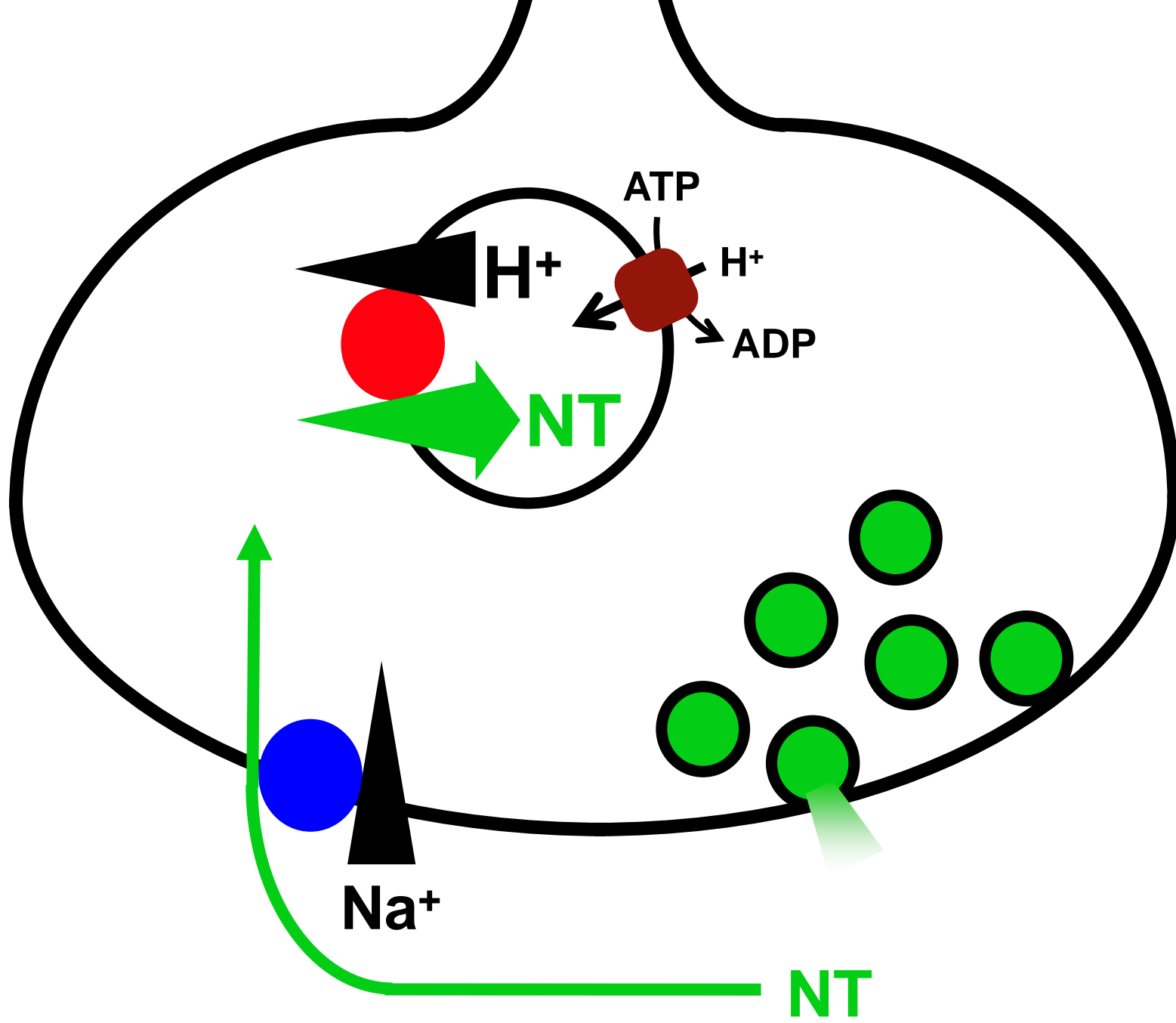
glutamate-induced currents can reverse
 --glutamate-gated chloride channel (receptor)
WHY?

(Wadiche et al, 1995)

some transporters also behave like channels
EAATs can behave as glu-gated chloride channels



transport cycle can gate an ion channel
?evolutionary intermediate

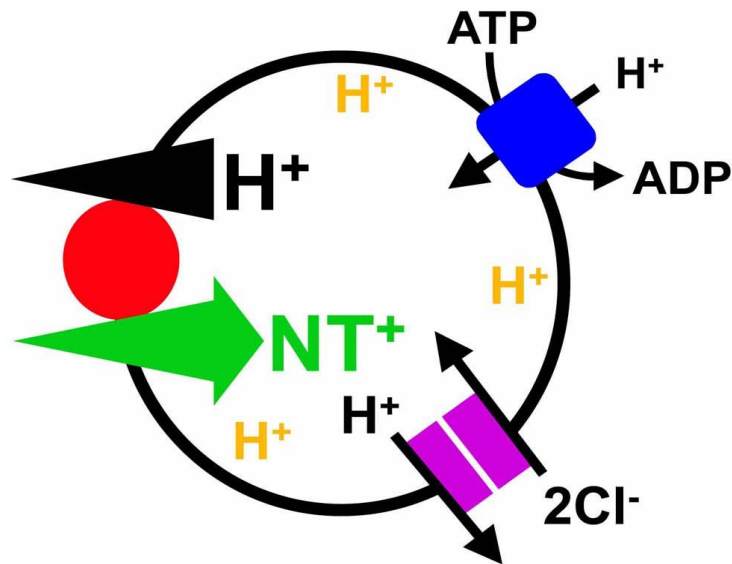


vesicular neurotransmitter transporters

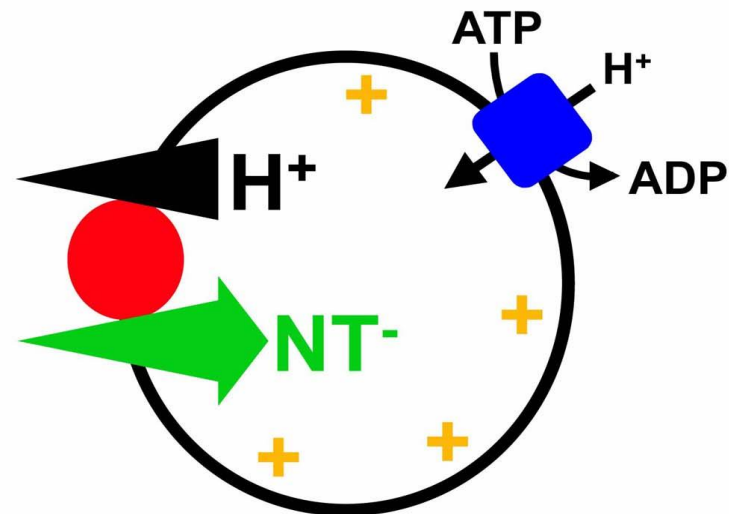
monoamines
acetylcholine

GABA

glutamate



$\Delta\text{pH} > \Delta\psi$



$\Delta\psi > \Delta\text{pH}$

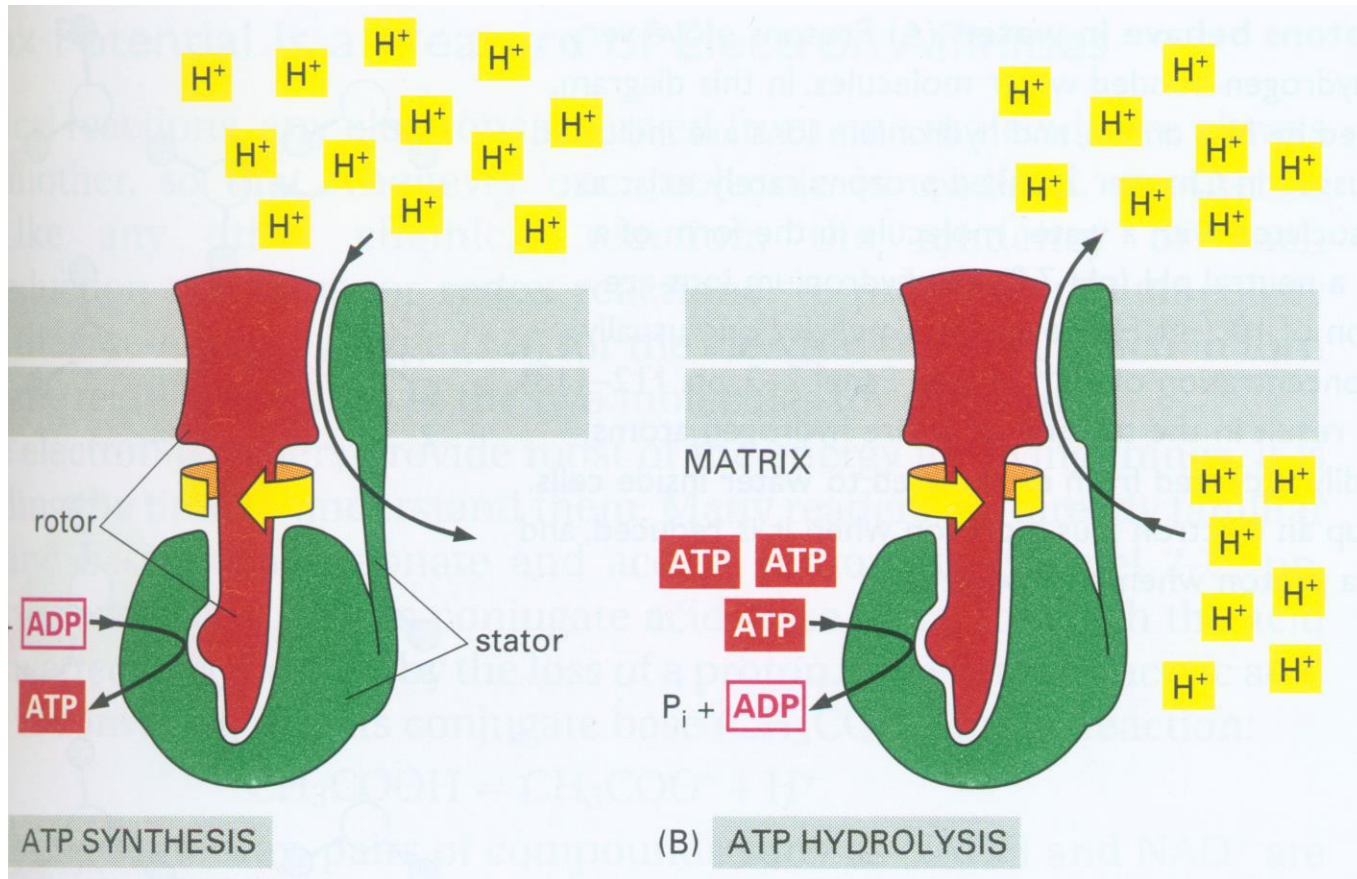
neurotransmitter per vesicle at **equilibrium**

determines location, affinity of receptors activated
depends on H^+ electrochemical gradient: **H^+ ideal**

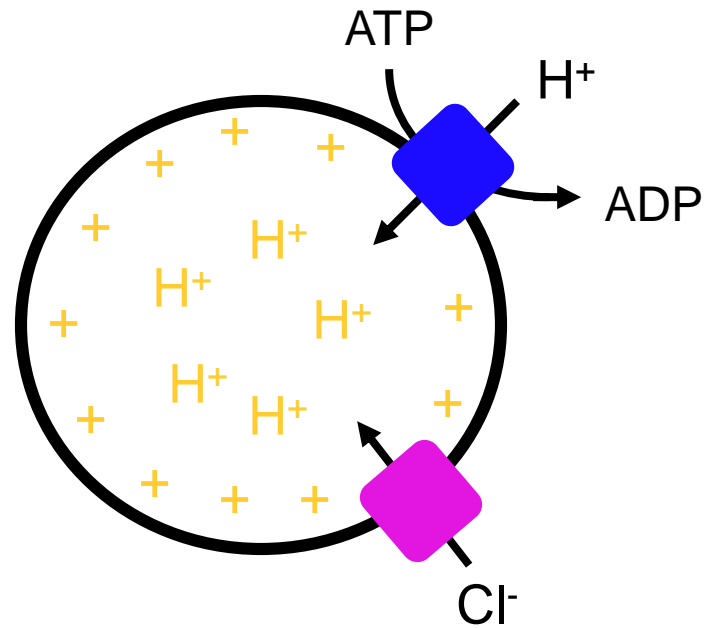
different NT depend on different components (ΔpH and $\Delta\psi$)

F0/F1 ATP synthase

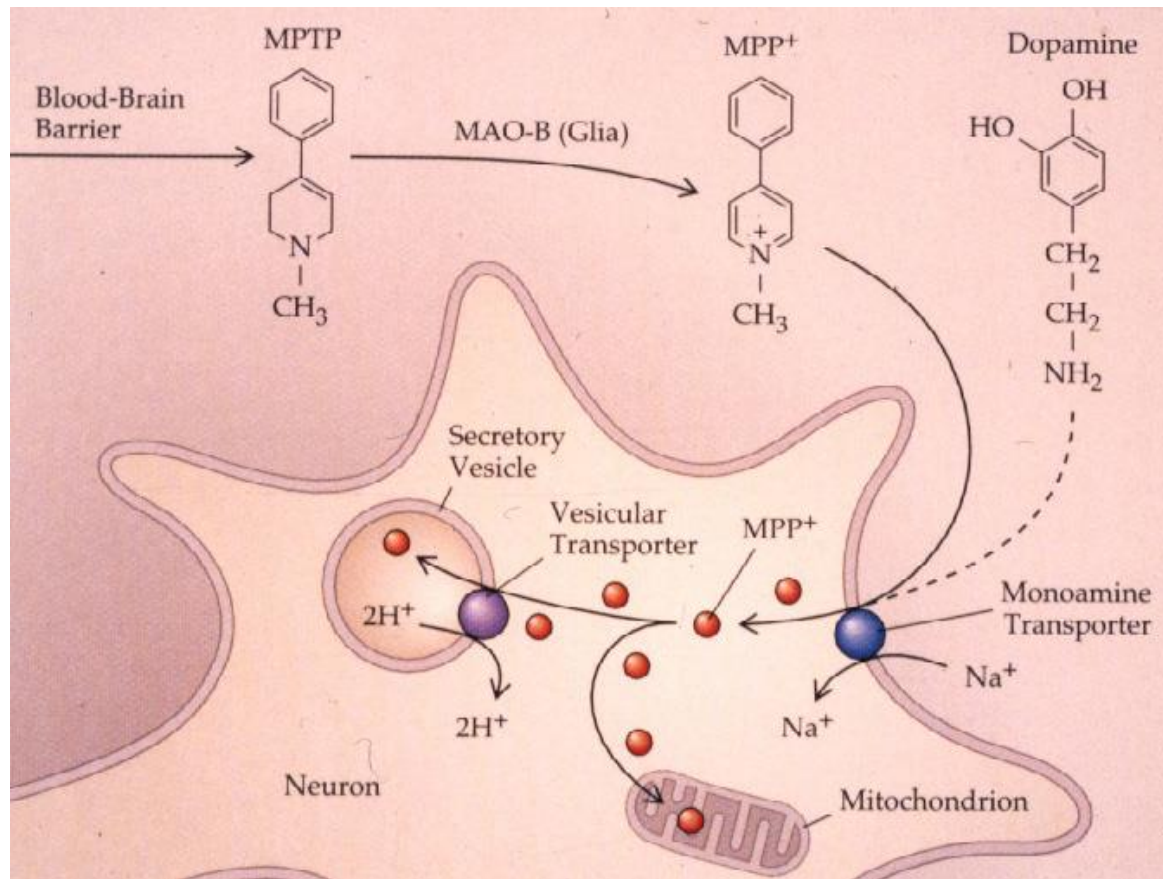
vacuolar H⁺-ATPase



$$\Delta\mu_{H^+} = \Delta pH + \Delta\psi$$



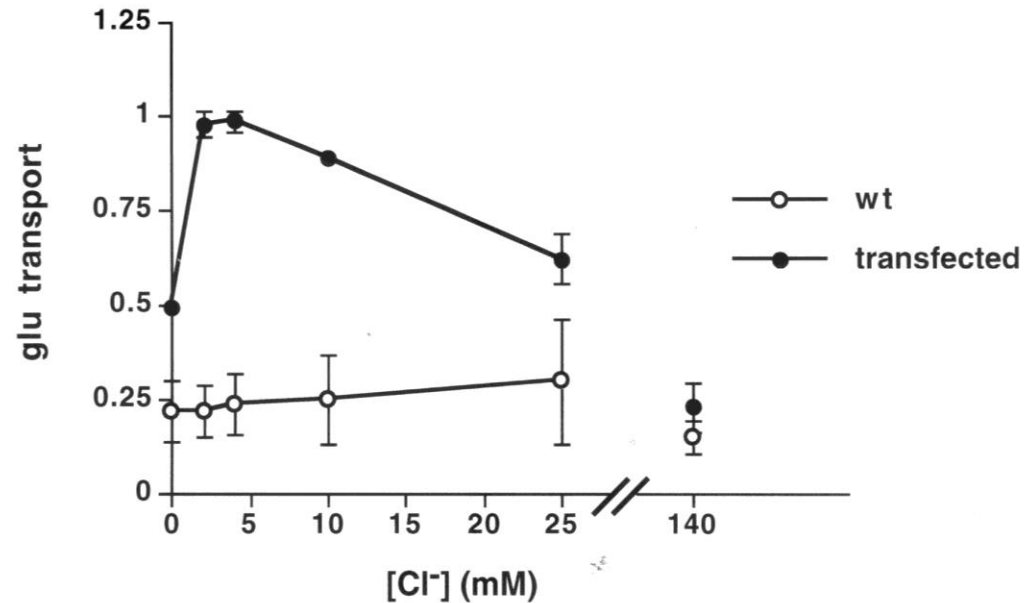
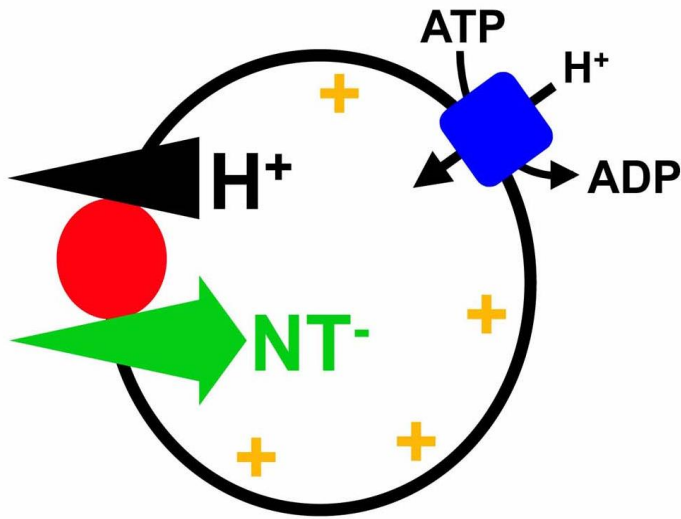
Cl⁻ entry dissipates $\Delta\psi$
cation efflux another way
how to create $\Delta\psi$?



(Liu et al., 1993)

VMAT protects against MPP⁺ toxicity
 ?role in Parkinson's?
 Km ~1 μM (high apparent affinity)

vesicular glutamate transport



(Bellocchio et al., 2000)

originally identified as Na/Pi cotransporter
depends primarily on $\Delta\psi$
low apparent affinity (K_m 1-3 mM)
allosteric activation by chloride (2-10 mM)
defines glutamate neurons

glutamate corelease with dopamine

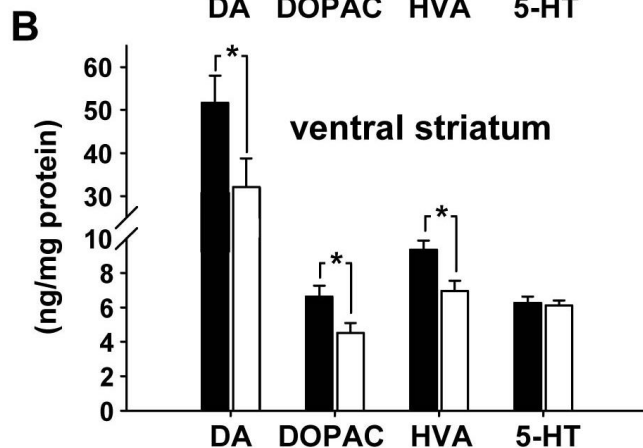
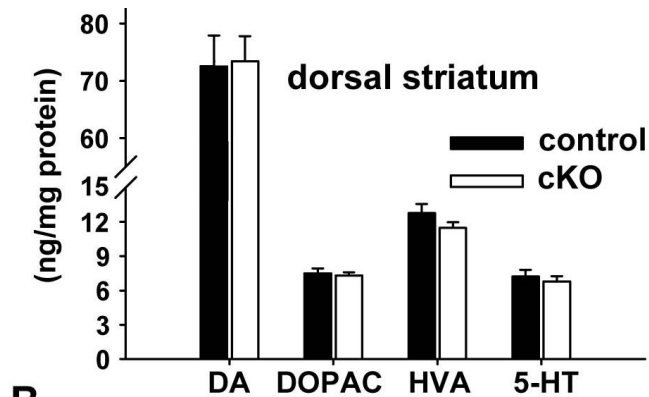
dopamine neurons form glutamatergic autapses *in vitro*

express high levels of VGLUT2 *in vitro*

VTA dopamine neurons express VGLUT2 *in vivo*

especially early in life

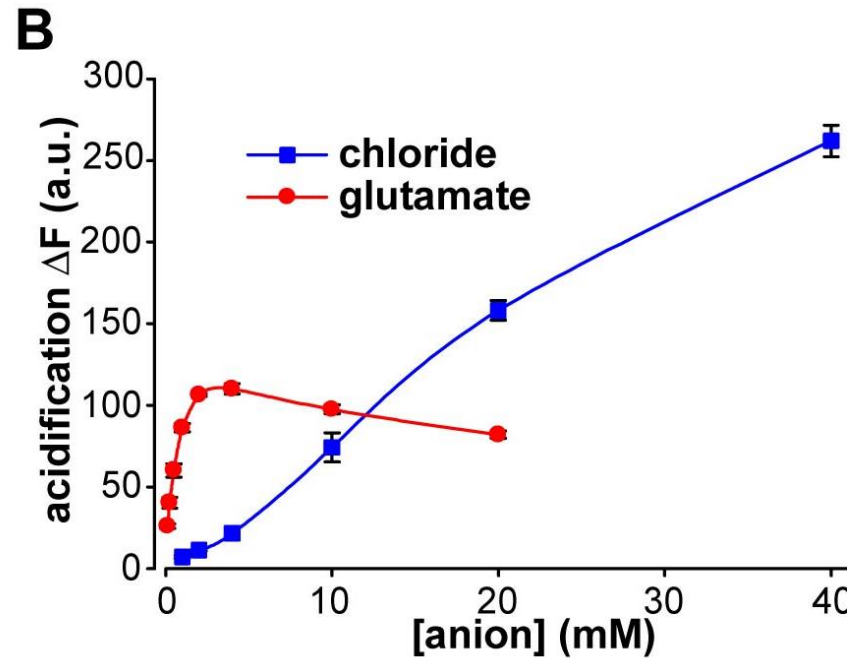
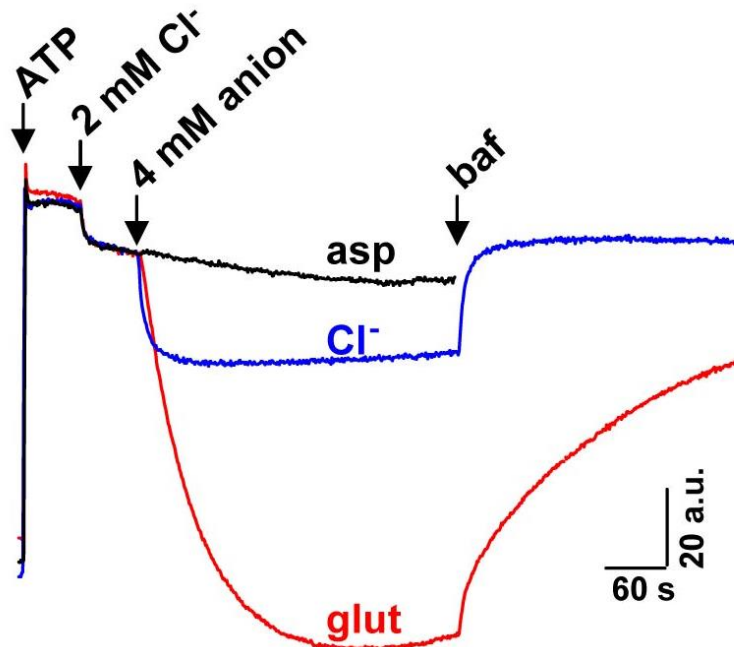
DAT-cre:VGLUT2--



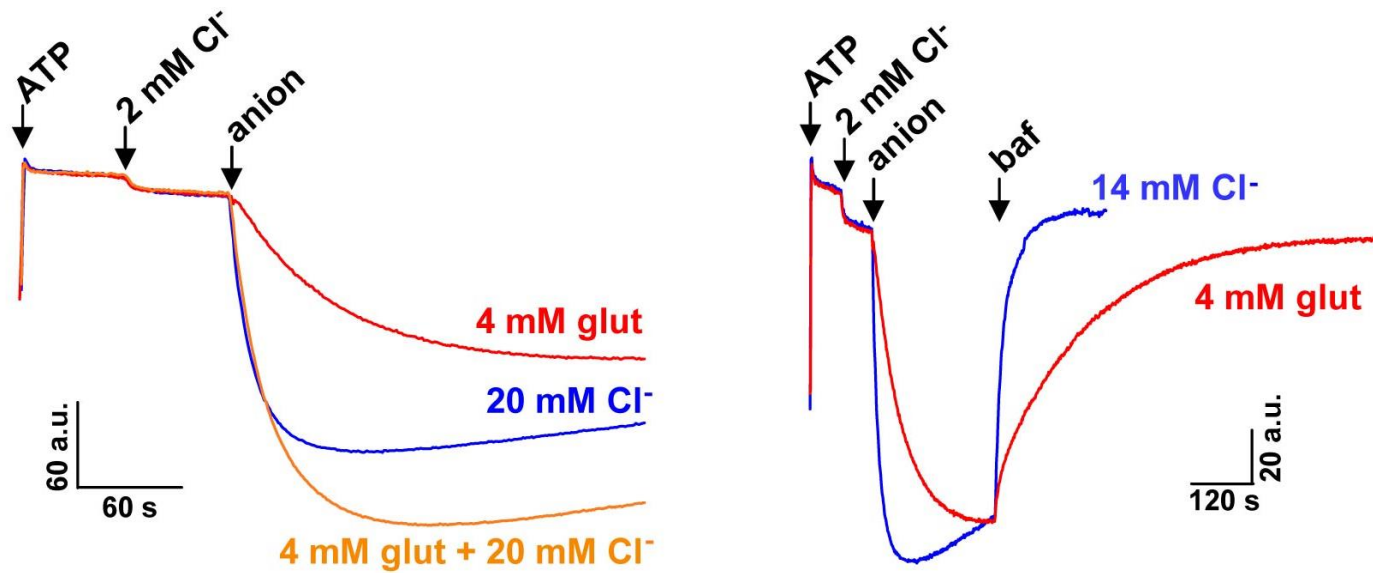
dopamine stores reduced ~35%
selective for ventral striatum
--consistent with localization
of VGLUT2 to VTA

(Hnasko et al., 2010)

acidification: acridine orange

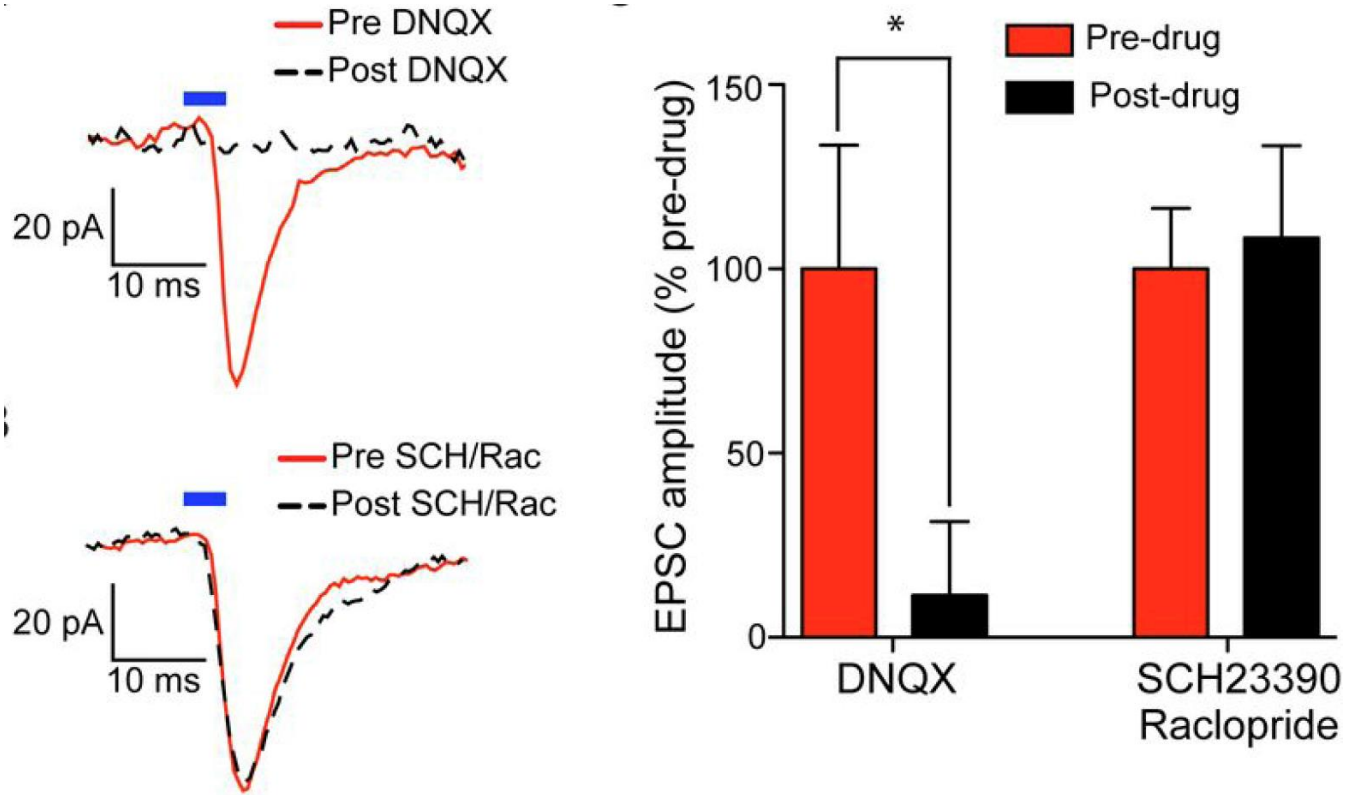


--glutamate also acidifies synaptic vesicles
corelease with other neurotransmitters widespread

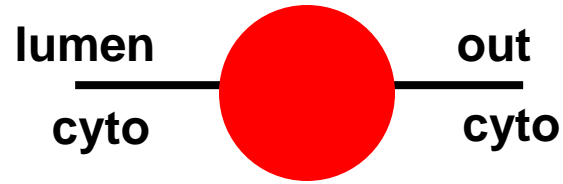


glutamate and Cl⁻ have additive effects on ΔpH
 vesicles acidified with glu retain ΔpH longer
 accounts for dopamine storage promoted by glu

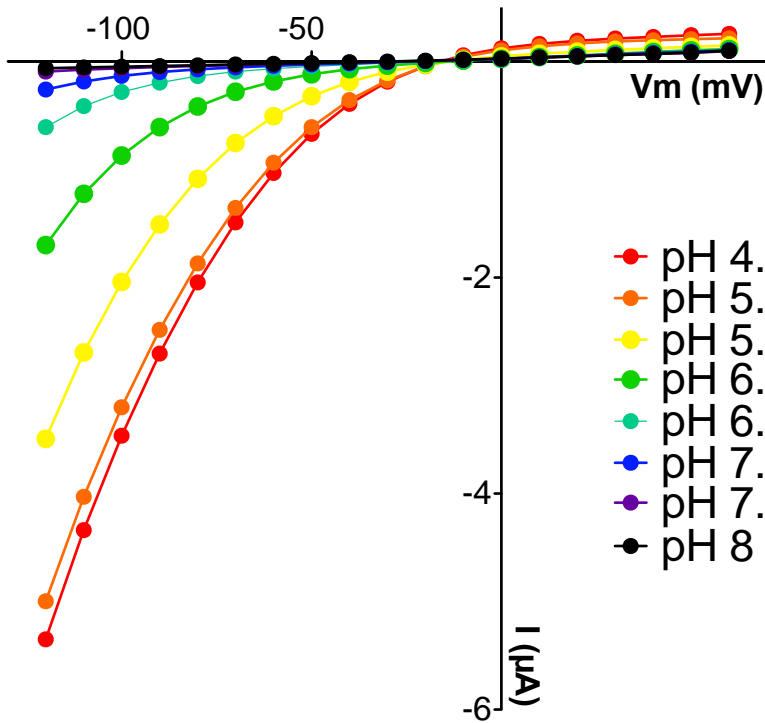
optogenetics: cChR2 in DAT-cre mice



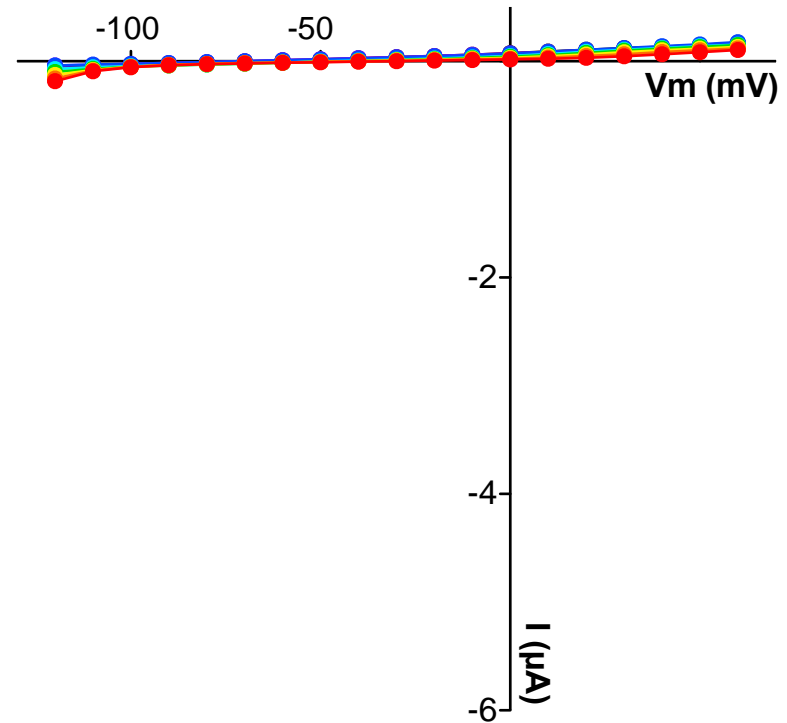
glutamate released by dopamine neurons
also acts as an independent signal
?same or different synaptic vesicles?



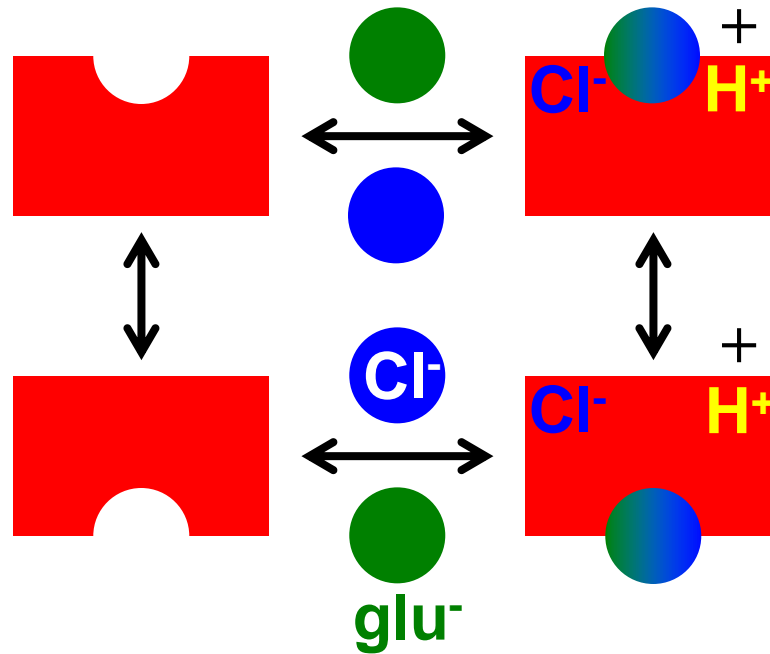
VGLUT2



H₂O



low pH_o activates an inwardly rectifying Cl^- current



glutamate and Cl⁻ permeate through similar pathway
 both driven by $\Delta\psi$: seems counterproductive
 both allosterically activated by luminal Cl⁻ and H⁺:
 role of allosteric activation?

VGLUT function

H⁺ pump-dependent activation

Cl⁻, glu dissipate $\Delta\psi$, increase ΔpH

--make it impossible to disentangle

roles of driving force and allosteric activation

predict huge effects on equilibrium and rate of SV filling

voltage clamp would solve this problem

--but how to record from a vesicle transporter?

1) misexpress transporter at plasma membrane

2) record directly from endosomes

--chloride and glutamate conductances

--allosteric activation by H⁺ as well as Cl⁻ (both sides)

Reading: The Synapse, pp. 147-170

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