

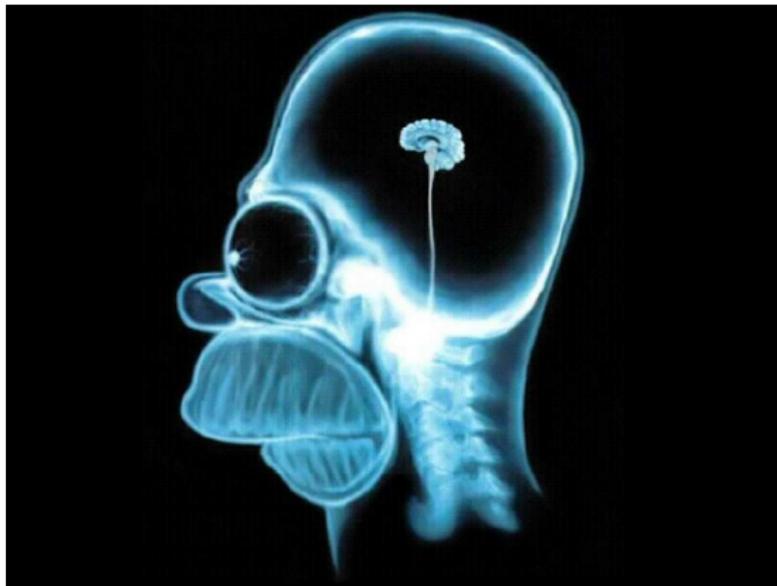
Modeling the role of AMPA receptor trafficking in the expression of long-term potentiation/depression

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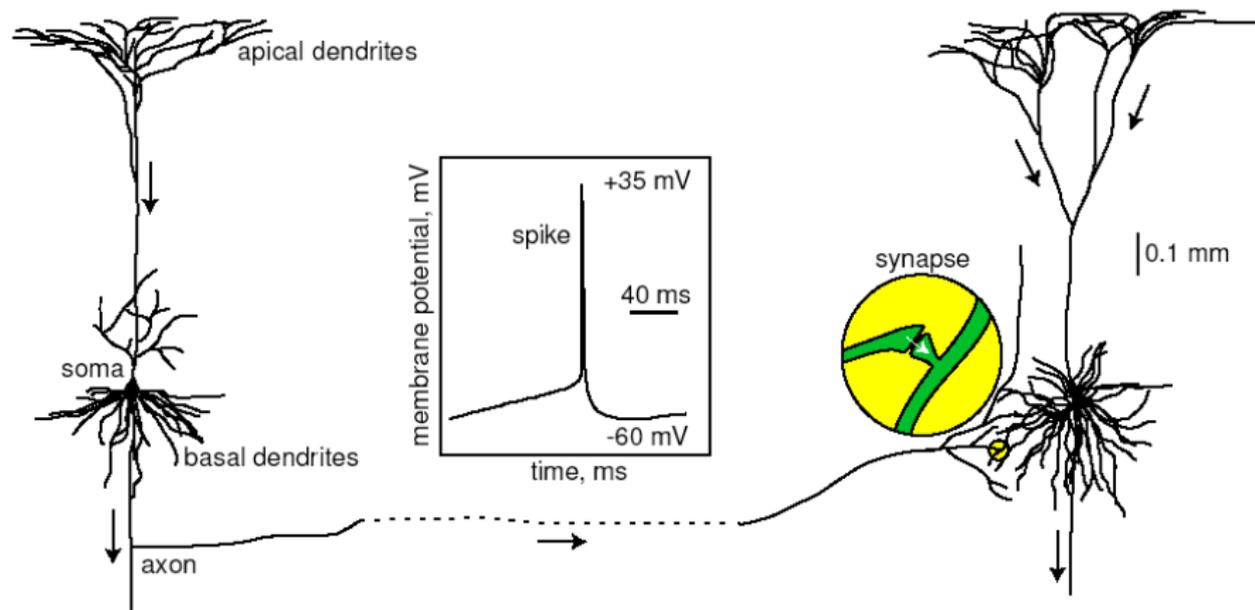
October 8, 2009

The amazing brain

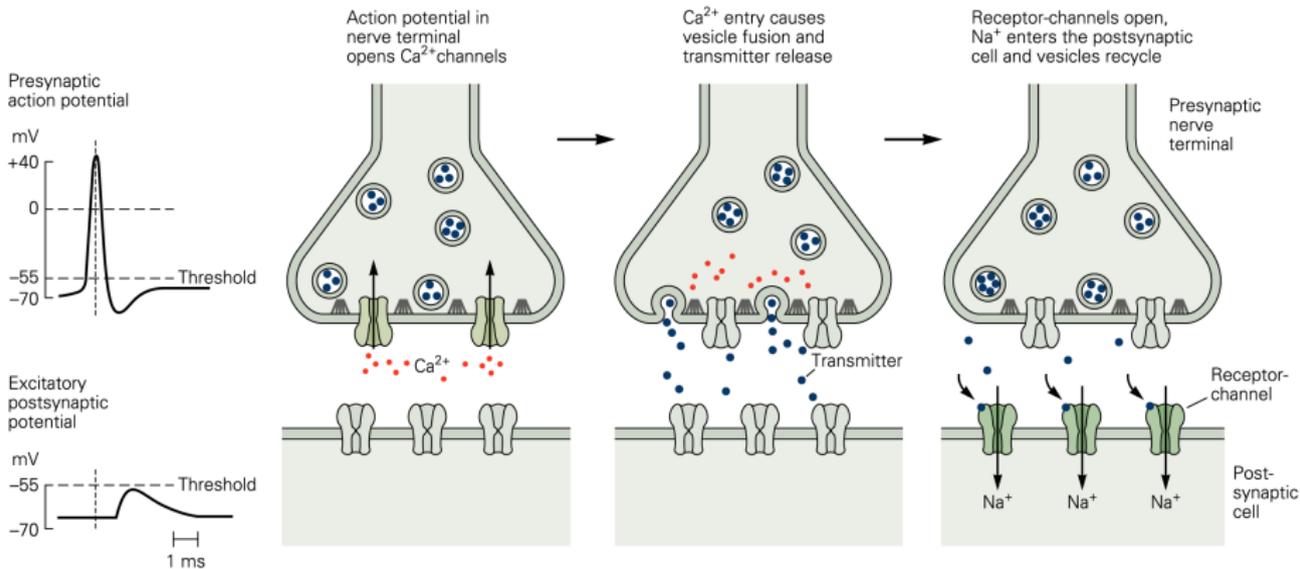


- 10^{11} neurons
- 10 – 10,000 synapses/neuron
- regulates body, behavior
- can learn, remember
- conscious experience

Neurons communicate at synapses

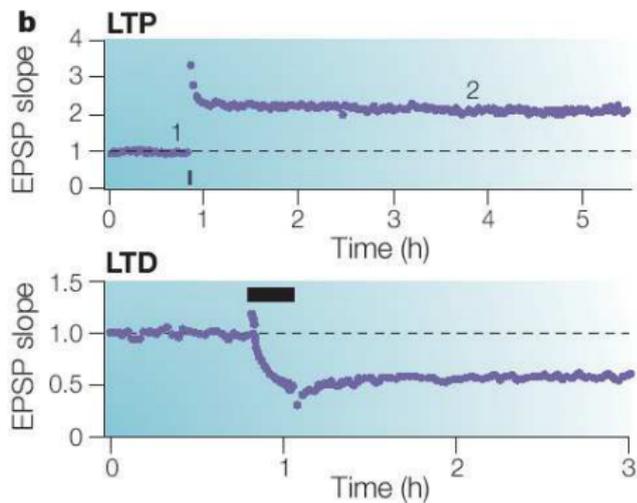
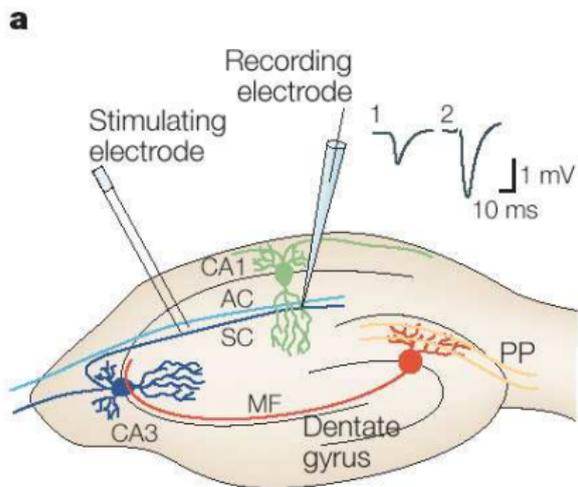


Communication at a synapse



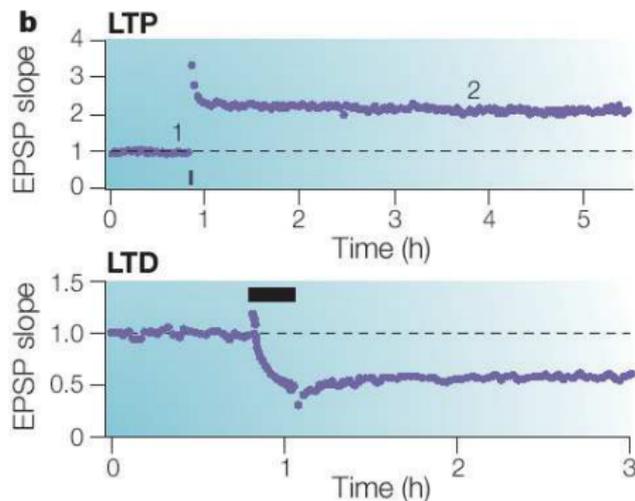
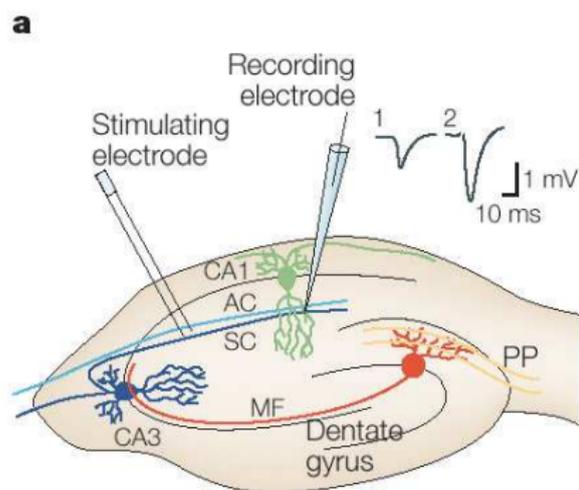
Kandel, Schwartz & Jessel (2000)

Synapses can “learn” – synaptic plasticity



Collingridge et al., *Nat Rev Neurosci* (2004)

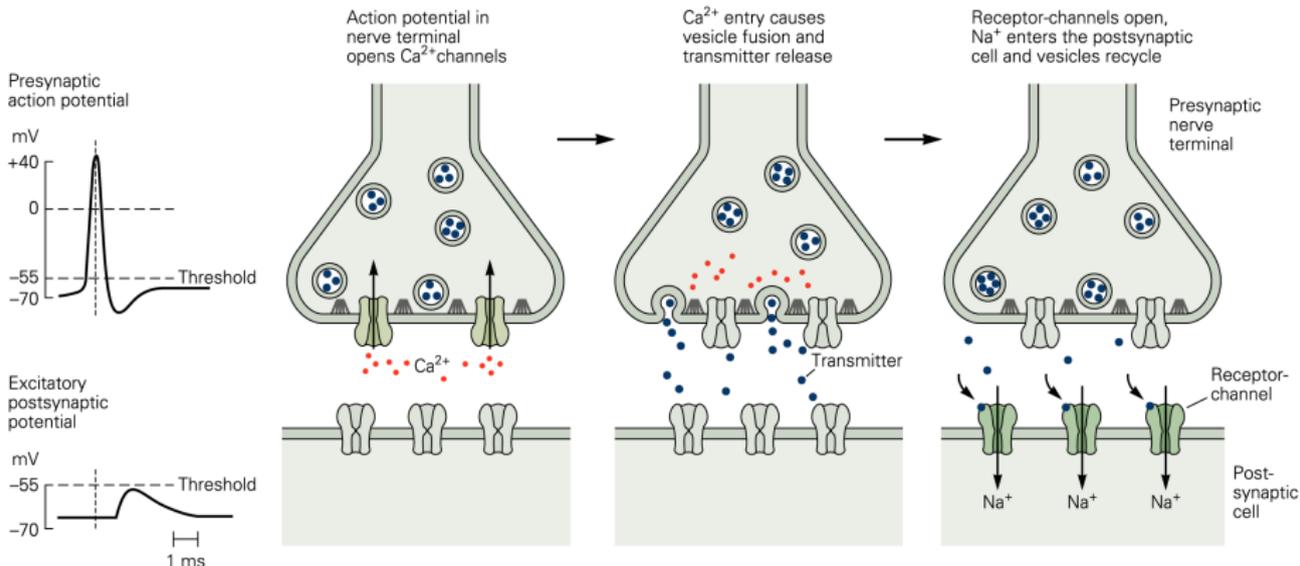
Synapses can “learn” – synaptic plasticity



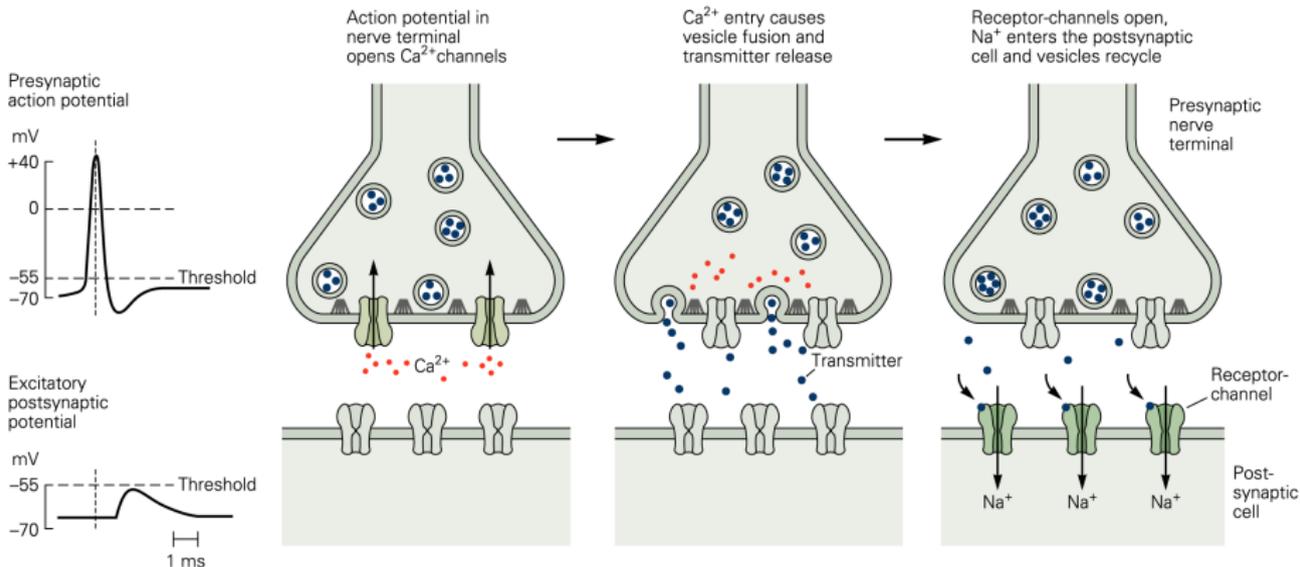
Collingridge et al., *Nat Rev Neurosci* (2004)

- LTP = long-term potentiation (strengthens synapse)
- LTD = long-term depression (weakens synapse)

How/where does synaptic plasticity occur?

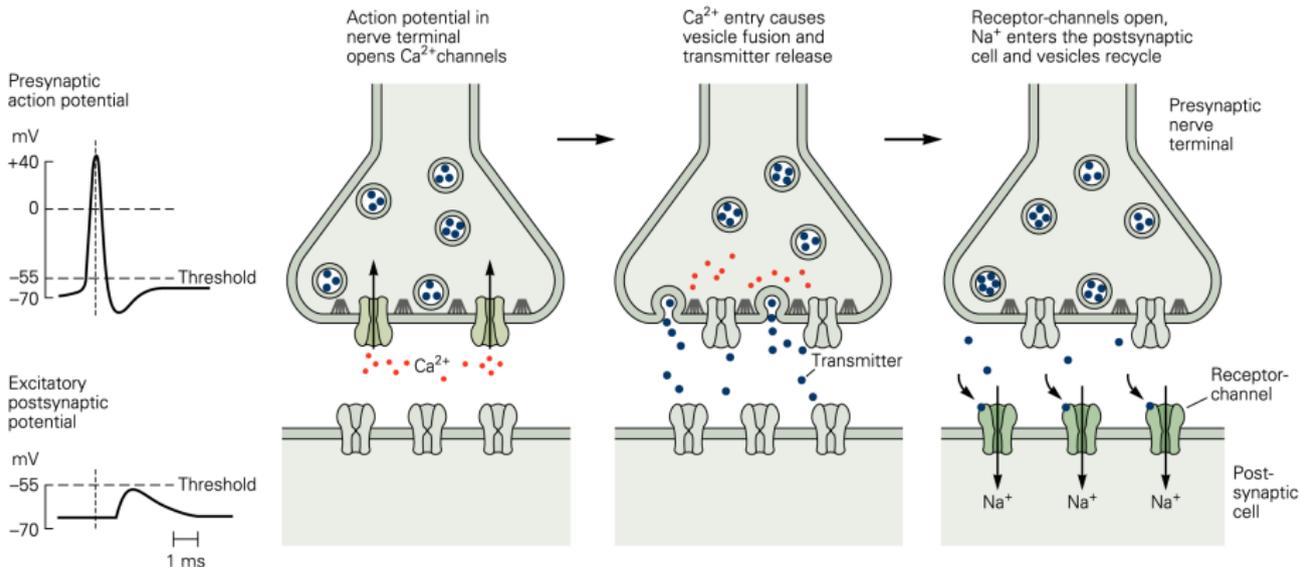


How/where does synaptic plasticity occur?



- Two major hypotheses:

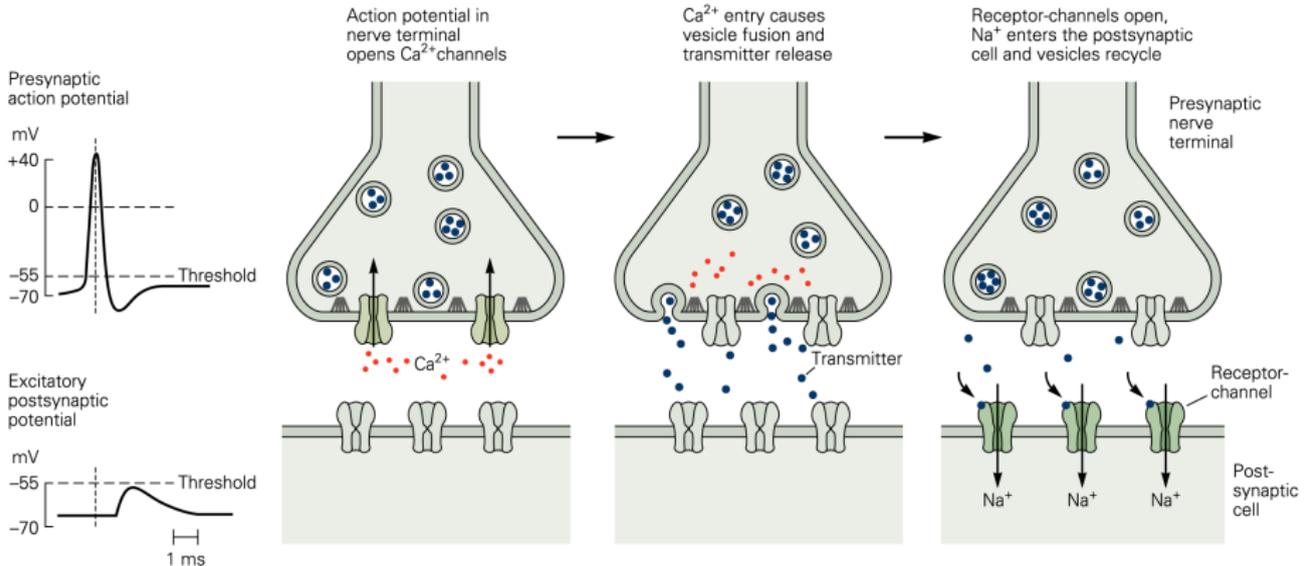
How/where does synaptic plasticity occur?



- Two major hypotheses:

- ① Presynaptic: change in the number of vesicles/probability of release

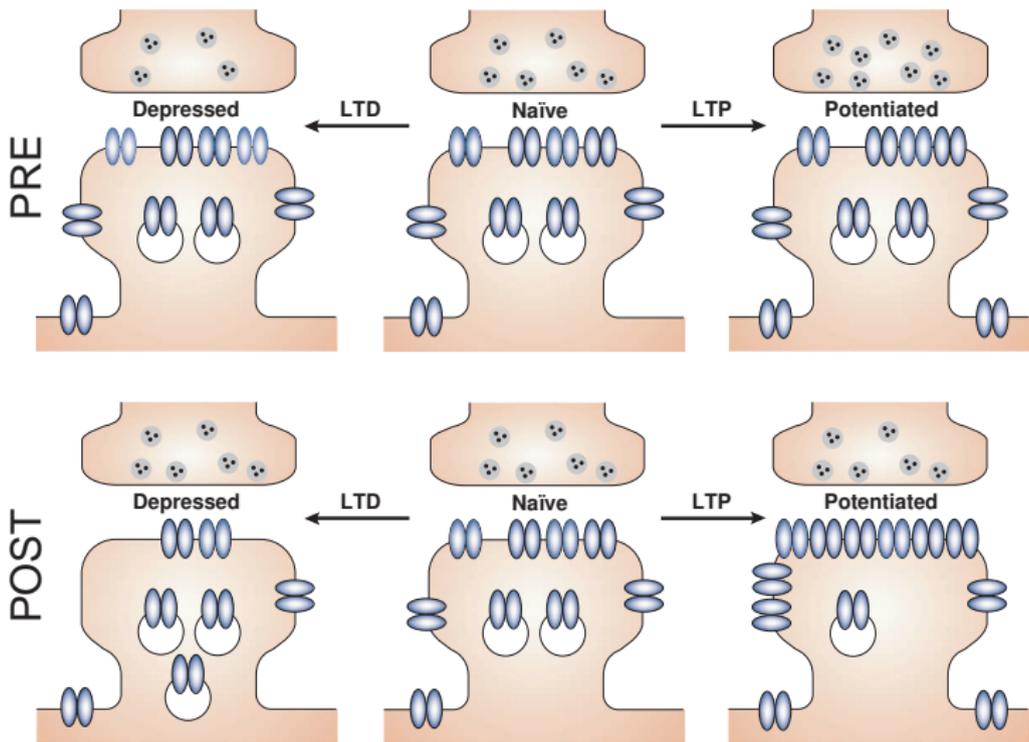
How/where does synaptic plasticity occur?



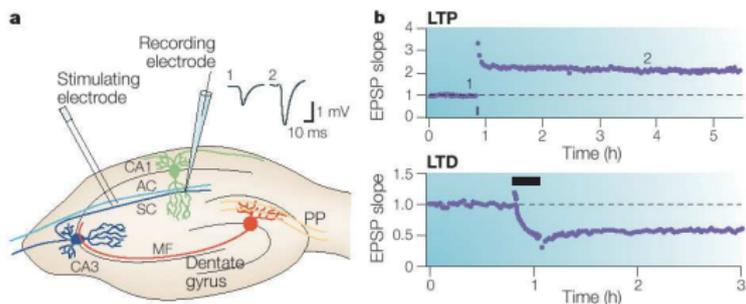
- Two major hypotheses:

- ① Presynaptic: change in the number of vesicles/probability of release
- ② Postsynaptic: change in the number/conductance of receptors

Presynaptic vs. postsynaptic mechanisms



Presynaptic vs. postsynaptic debate



...long-term plasticity...is expressed overwhelmingly via presynaptic changes in reliability of transmitter release.

–Enoki et al., *Neuron* (2009)

Therefore, LTP is the recruitment of new [receptors] to synapses...

–Kerchner & Nicoll, *Nat. Rev. Neurosci.* (2008)

Can modeling help?

Can modeling help?

That depends on who you talk to!

Can modeling help?

That depends on who you talk to!

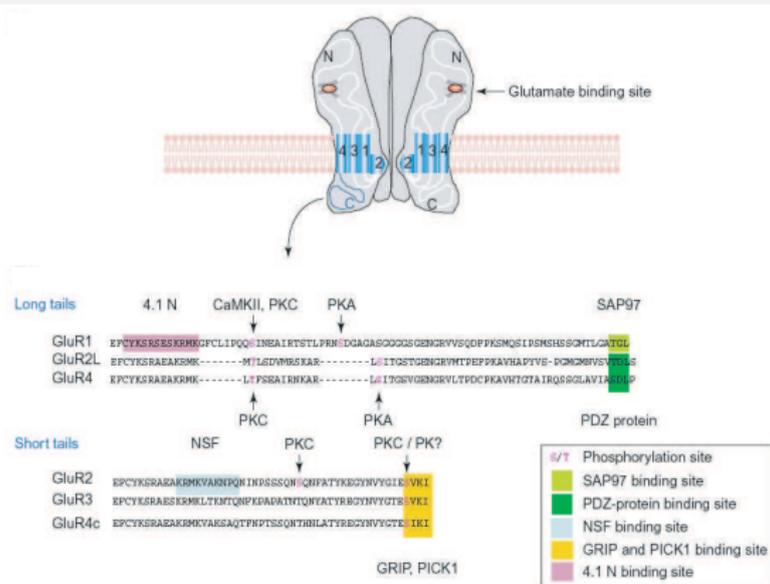
Question to answer...

Can LTP/LTD data be explained by postsynaptic receptor trafficking alone?

Outline for rest of talk

- Introduce AMPA receptors
- Describe AMPA receptor trafficking
- Propose model of AMPA receptor trafficking
- Present results from model
- Conclusions & future directions

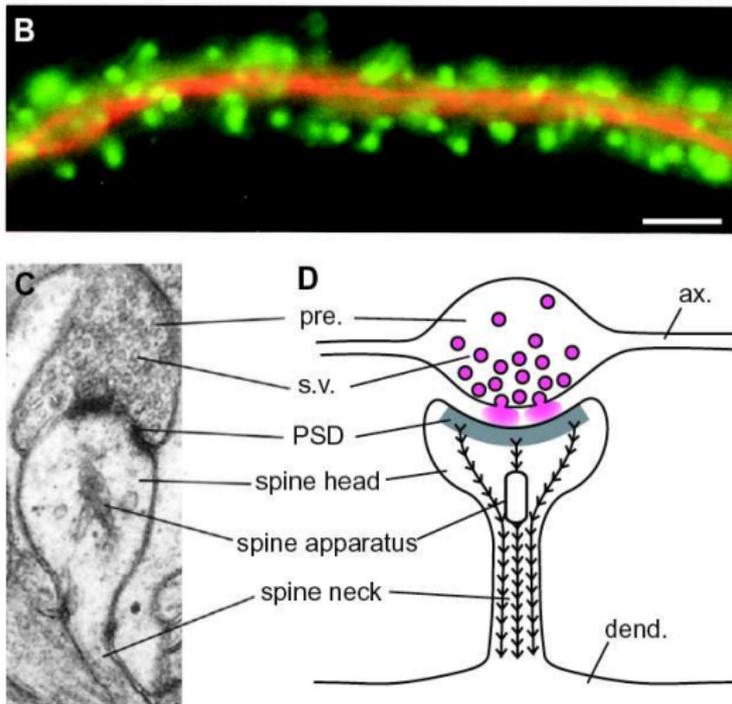
AMPA receptors



Huganir & Song, *Nat. Rev. Neurosci.* (2002)

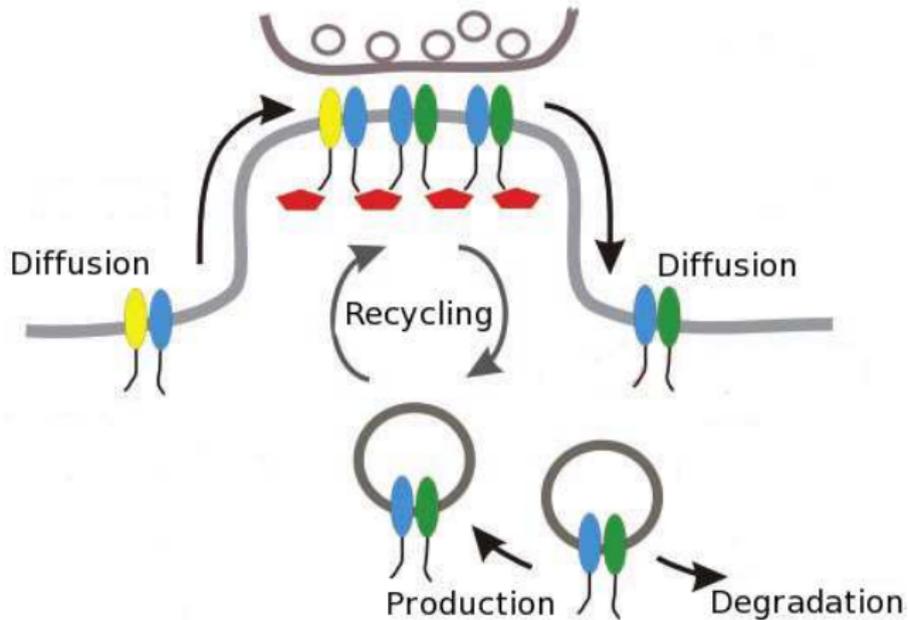
- Responsible for excitable synaptic transmission in CNS
- Formed from four subunits: GluR1 to GluR4
- Dominant heteromers: GluR1/2 and GluR2/3

Excitable synapses located in dendritic spines

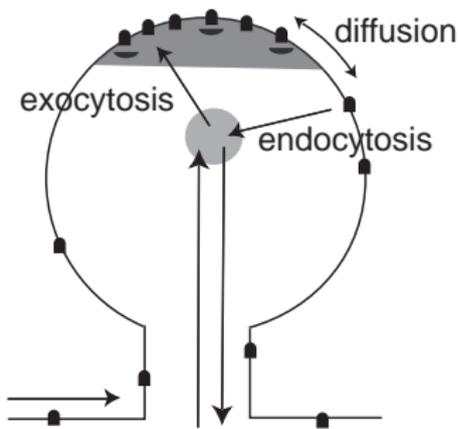


Matus, *Science* (2000)

AMPA receptor trafficking at a spine



Model of AMPA receptor trafficking at a single spine



- AMPA receptor
- scaffolding protein

Cottrell et al., *J. Neurophysiol.* (2000)
 Sorra & Harris, *Hippocampus* (2000)
 Ehlers, *Neuron* (2000)
 Passafaro et al., *Nat. Neurosci.* (2001)
 Groc et al., *Nat. Neurosci.* (2004)

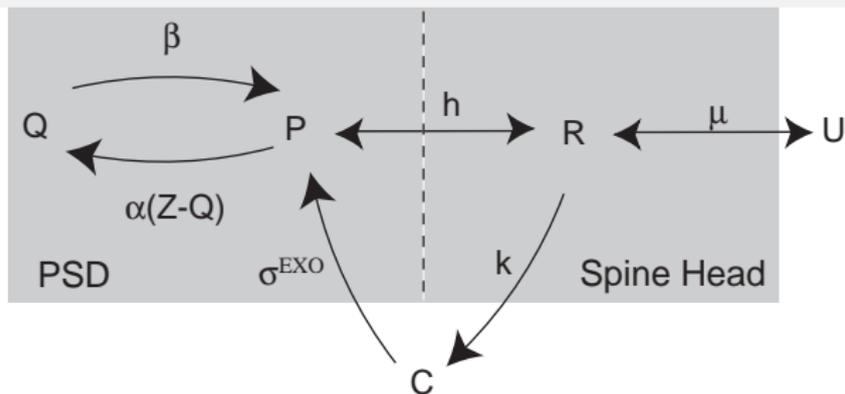
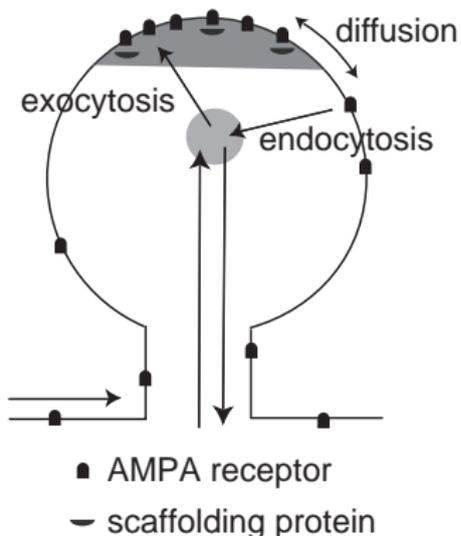
Time constants

- Exo/endocytosis: 10-30 min
- Diffusion: 10 s
 - Surface area of PSD: $0.1 \mu\text{m}^2$
 - Surface area of spine head: $1 \mu\text{m}^2$
 - Diffusion coefficient: $0.01\text{-}0.1 \mu\text{m}^2/\text{s}$
- Binding/unbinding to scaffolding: unknown
- Production/degradation: unknown

Other constants

- Intracellular AMPAR number: 100-500
- AMPAR concentration in dendrite: $10\text{-}100/\mu\text{m}^2$
- Scaffolding concentration: unknown

State diagram



Variables

P = free AMPAR concentration in PSD

Q = bound AMPAR concentration in PSD

R = AMPAR concentration in spine head

Constants

C = intracellular AMPAR number

U = AMPAR concentration in dendrite

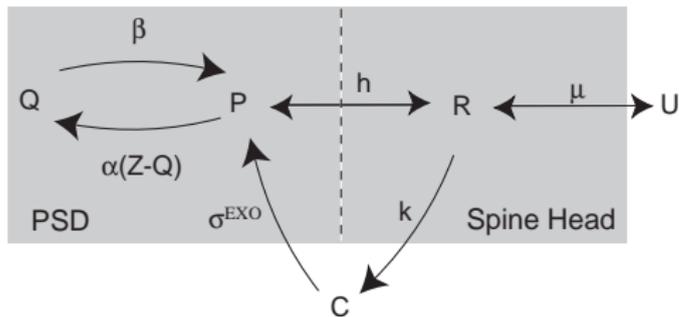
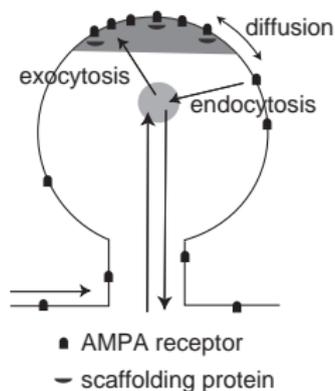
Z = scaffolding protein concentration

Model equations

PSD free:
$$\frac{dP}{dt} = h(R - P) - \alpha(Z - Q)P + \beta Q + \sigma^{\text{EXO}} \frac{C}{a}$$

PSD bound:
$$\frac{dQ}{dt} = \alpha(Z - Q)P - \beta Q$$

Spine head:
$$\frac{dR}{dt} = \mu(U - R) - h(R - P) - kR$$



BAE & Bressloff, *J. Neurosci.* (2006)

Parameter values

Known parameter values:

- Rate of exocytosis, endocytosis: $\sigma^{\text{EXO}}, k = 10^{-3}/\text{s}$
- Surface area of PSD, spine head: $a = 0.1 \mu\text{m}^2, A = 1 \mu\text{m}^2$
- Hopping rate between PSD and spine head: $h = 0.1/\text{s}$
- Hopping rate between spine head and dendrite: $\mu = 0.005/\text{s}$
 - Ashby et al., *J. Neurosci.* (2006)
- Intracellular AMPA receptor number: $C = 100$
- AMPA receptor concentration in dendrite: $U = 20/\mu\text{m}^2$

Unknown parameter values:

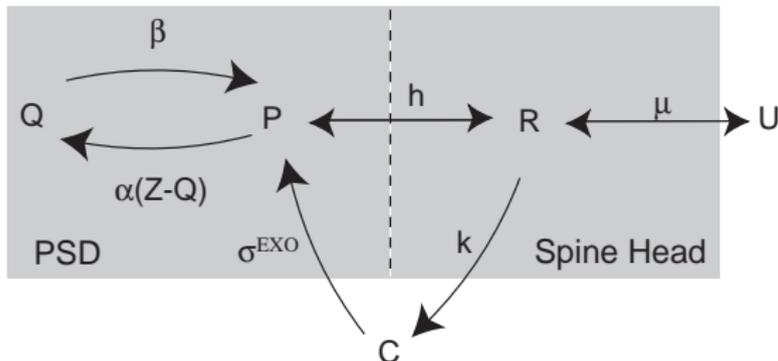
- Binding/unbinding rates α and β
 - Constitutive recycling $\sim 10 - 30 \text{ min} \Rightarrow \alpha = \beta = 10^{-3}/\text{s}$
- Scaffolding protein concentration Z
 - Approx. half AMPARs in PSD are bound $\Rightarrow Z = 200/\mu\text{m}^2$

Steady-state

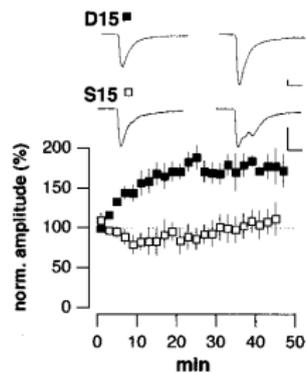
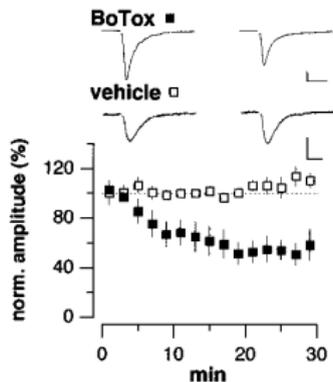
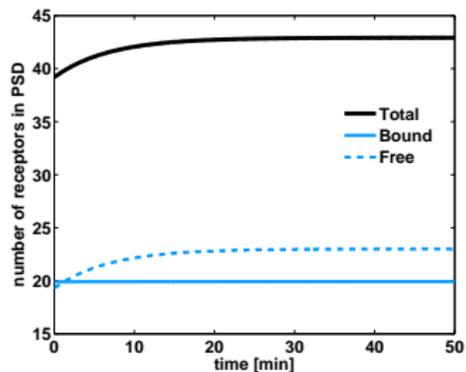
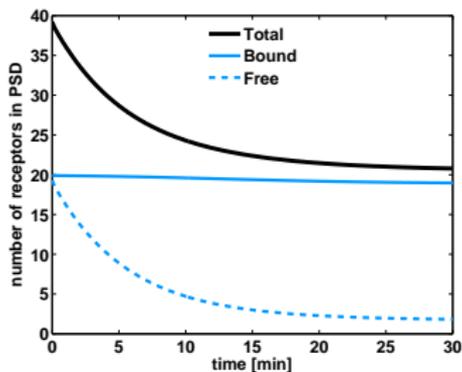
PSD free:
$$P = R + \frac{\sigma^{\text{EXO}} C}{h} \frac{C}{a} \approx 193/\mu\text{m}^2$$

PSD bound:
$$Q = \frac{\alpha P}{\alpha P + \beta} Z \approx 199/\mu\text{m}^2$$

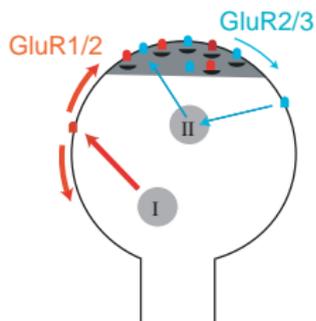
Spine head:
$$R = \frac{\mu U + \sigma^{\text{EXO}} \frac{C}{a}}{\mu + k} \approx 183/\mu\text{m}^2$$



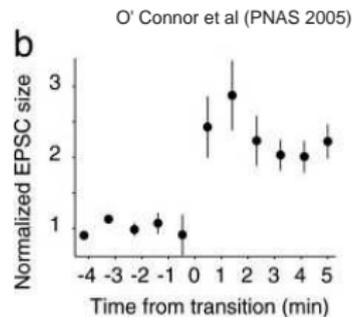
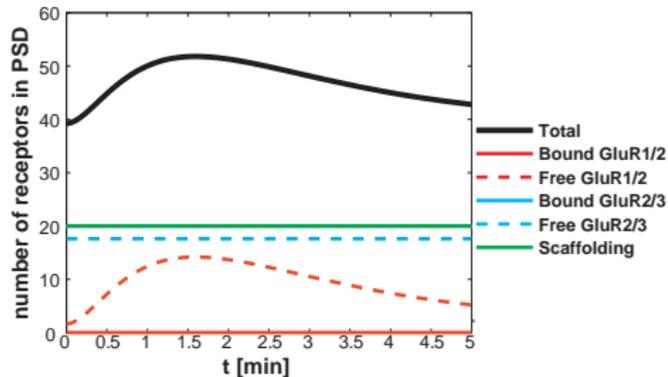
Block exo/endocytosis

Luscher et al., *Neuron* (1999)

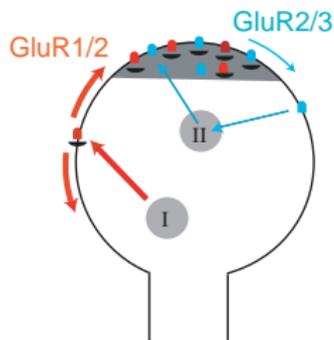
LTP simulation



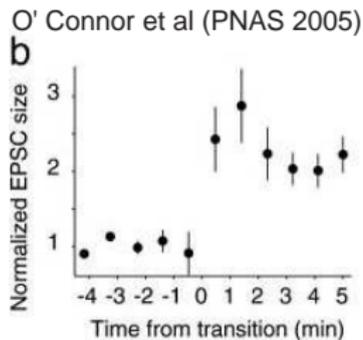
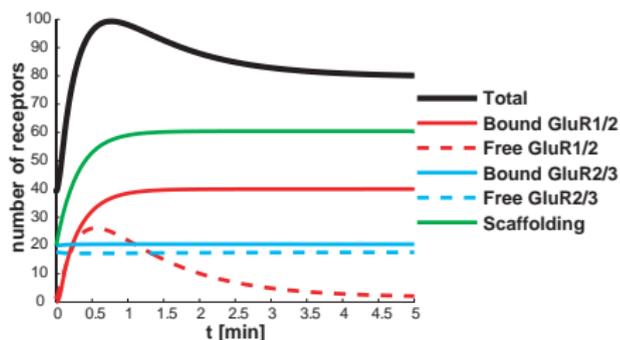
- Activation of GluR1/2 intracellular pool
- Rapid insertion of receptors into ESM



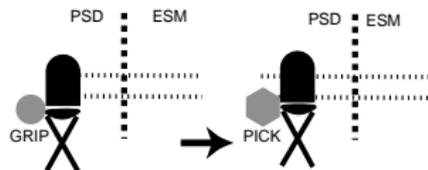
LTP simulation with increase in scaffolding



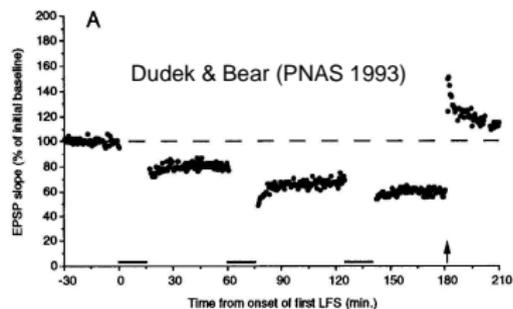
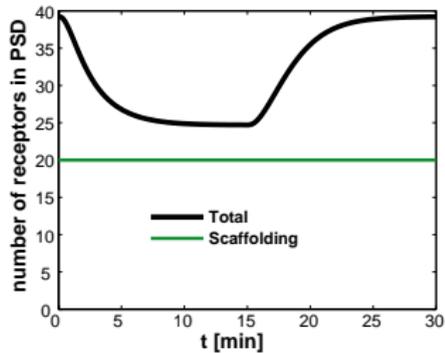
- Activation of GluR1/2 intracellular pool
- Rapid insertion of receptors into ESM
- AMPARs transport slot proteins into PSD



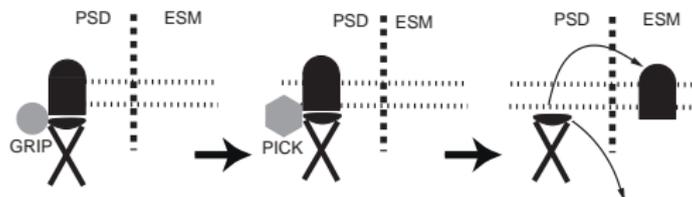
LTD simulation



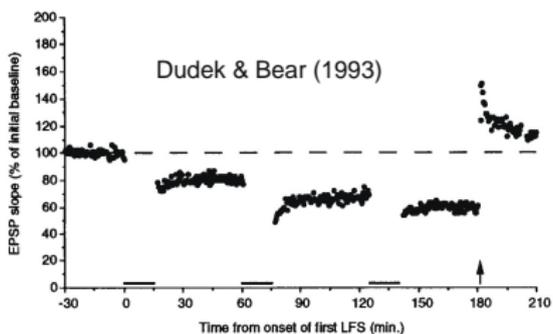
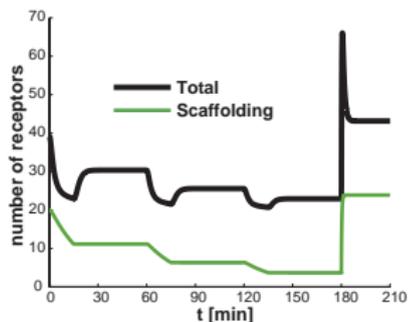
- Switch from AMPA-GRIP to AMPA-PICK receptor-protein complexes
- Rapid unbinding from PSD and trafficking to ESM followed by endocytosis.



LTD simulation with decrease in scaffolding



- Switch from AMPA-GRIP to AMPA-PICK receptor-protein complexes
- Rapid unbinding from PSD and trafficking to ESM followed by endocytosis.
- Unbound scaffolding proteins are degraded.



Conclusions

Can LTP/LTD data be explained by postsynaptic receptor trafficking alone?

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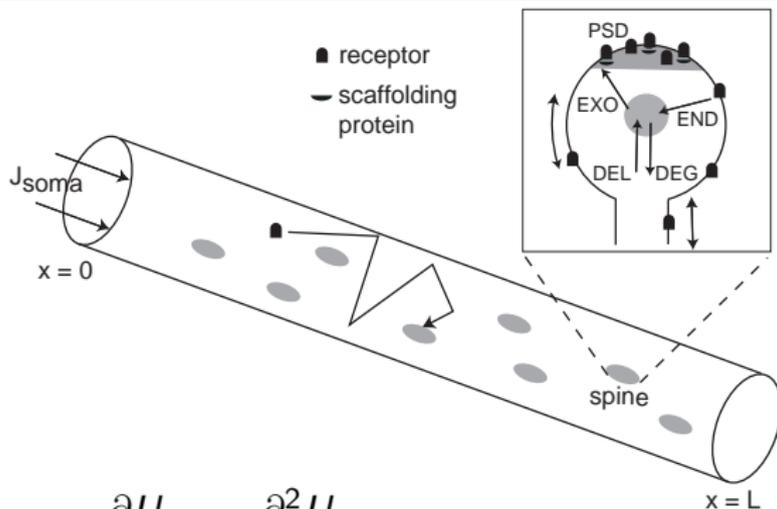
- LTP/LTD data can be reproduced within our model of AMPA receptor trafficking

Conclusions

Can LTP/LTD data be explained by postsynaptic receptor trafficking alone?

- LTP/LTD data can be reproduced within our model of AMPA receptor trafficking
 - LTP requires increase in scaffolding (Shi et al., *Cell* 2001)
 - LTD requires decrease in scaffolding (Colledge et al., *Neuron* 2003)

Future directions – AMPAR trafficking along dendrite



$$\frac{\partial U}{\partial t} = D \frac{\partial^2 U}{\partial x^2} - \eta(x)\mu(x)(U - R)$$

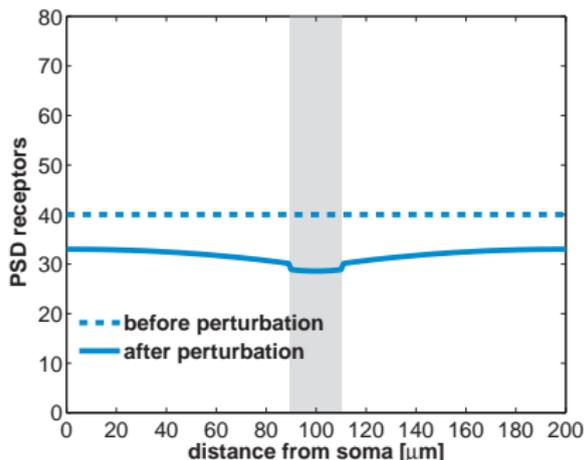
$$-D \left. \frac{\partial U}{\partial x} \right|_{x=0} = J_{\text{soma}}, \quad \left. \frac{\partial U}{\partial x} \right|_{x=L} = 0.$$

Bressloff, BAE, Ward, *SIAM J Appl Math* (2008)
BAE & Bressloff, *J Comput Neurosci* (2008)

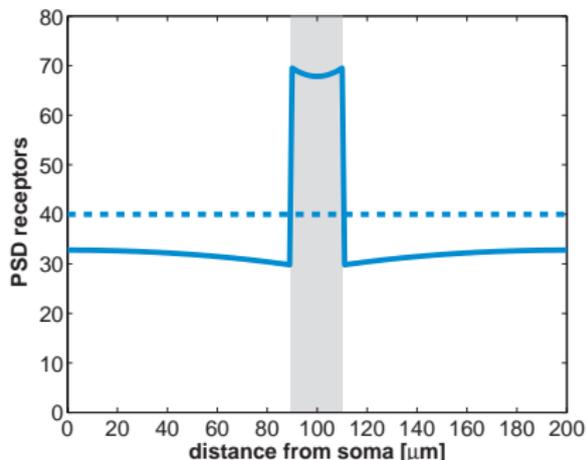
Future directions – AMPAR trafficking along dendrite

Consequences of diffusive coupling

10-fold reduction in
rate of exocytosis
in gray region

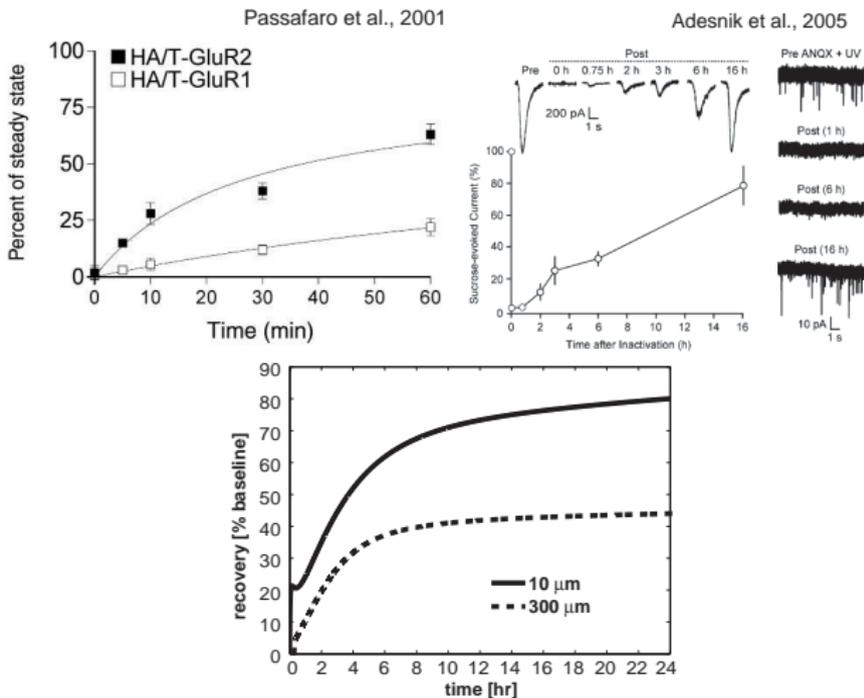


10-fold increase in
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Future directions – AMPAR trafficking along dendrite

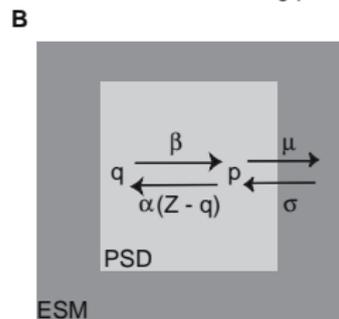
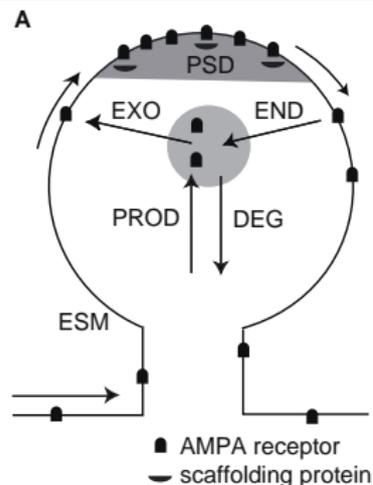
Fast or slow recycling?



Future directions – stochastic models

$$\frac{dp}{dt} = -\alpha(Z - q)p + \beta q - \mu p + \sigma$$

$$\frac{dq}{dt} = \alpha(Z - q)p - \beta q$$

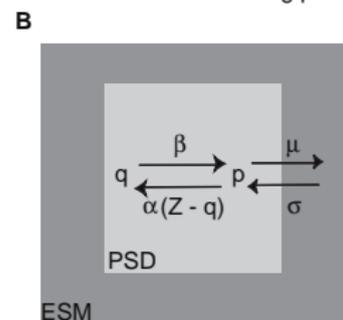
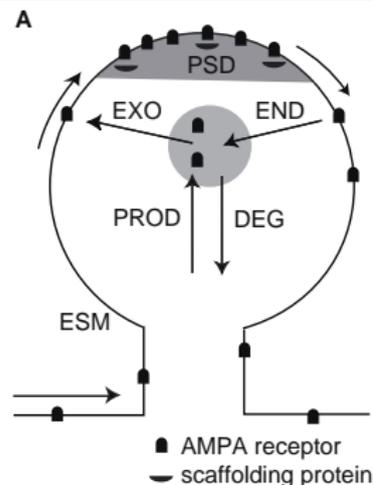


Future directions – stochastic models

$$\frac{dp}{dt} = -\alpha(Z - q)p + \beta q - \mu p + \sigma$$

$$\frac{dq}{dt} = \alpha(Z - q)p - \beta q$$

$$P_{n,m}(t) = \text{Prob}\{n \text{ unbound}, m \text{ bound at time } t\}$$



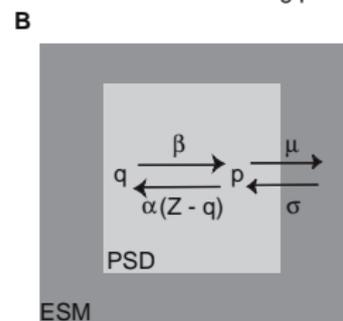
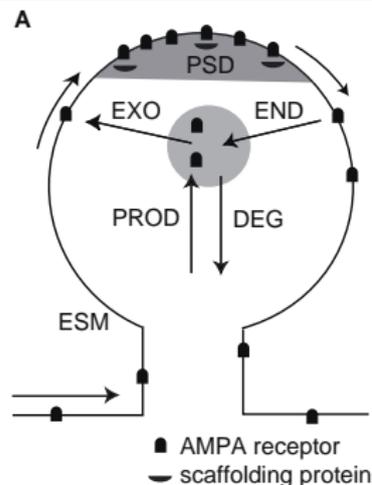
Future directions – stochastic models

$$\frac{dp}{dt} = -\alpha(Z - q)p + \beta q - \mu p + \sigma$$

$$\frac{dq}{dt} = \alpha(Z - q)p - \beta q$$

$$P_{n,m}(t) = \text{Prob}\{n \text{ unbound}, m \text{ bound at time } t\}$$

$$\begin{aligned} \frac{dP_{n,m}}{dt} = & \sigma P_{n-1,m} + \mu(n+1)P_{n+1,m} \\ & + \alpha(n+1)[Z - (m-1)]P_{n+1,m-1} \\ & + \beta(m+1)P_{n-1,m+1} \\ & - [\sigma + \mu n + \alpha n(Z - m) + \beta m]P_{n,m} \end{aligned}$$



Future directions – stochastic models

$$\frac{dp}{dt} = -\alpha(Z - q)p + \beta q - \mu p + \sigma$$

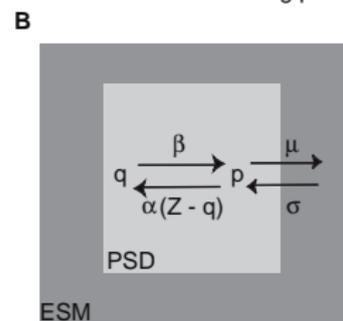
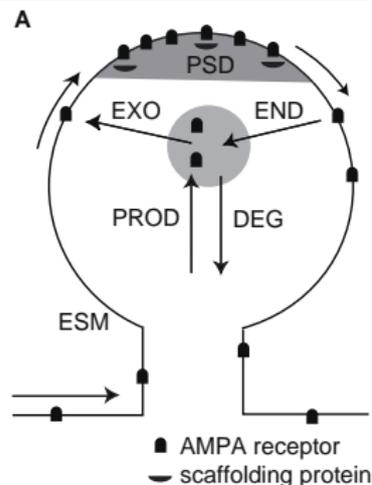
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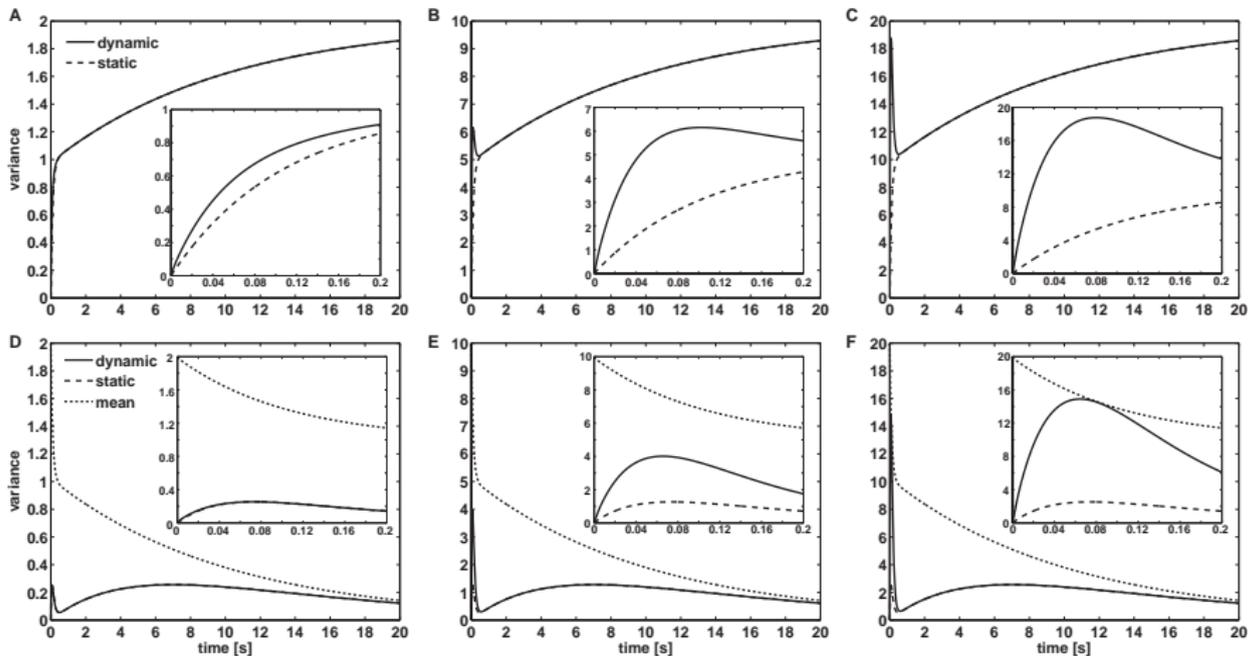
$$\begin{aligned} \frac{dP_{n,m}}{dt} = & \sigma P_{n-1,m} + \mu(n+1)P_{n+1,m} \\ & + \alpha(n+1)[Z - (m-1)]P_{n+1,m-1} \\ & + \beta(m+1)P_{n-1,m+1} \\ & - [\sigma + \mu n + \alpha n(Z - m) + \beta m]P_{n,m} \end{aligned}$$

$$\text{stochastic gate : } 0 < \mu_{open} \xrightleftharpoons[\gamma+]{\gamma-} \mu_{closed} = 0$$

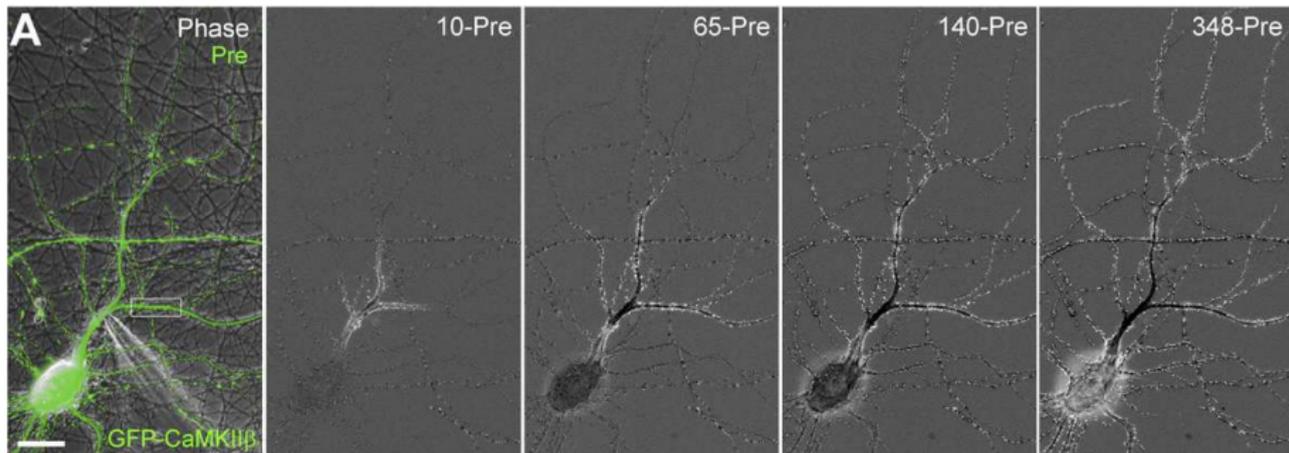
$$\sigma(t) = C\mu(t) \quad (C \text{ bath conc.})$$



Future directions – stochastic models



Future directions – trafficking of other proteins



Rose et al., *Neuron* (2009)

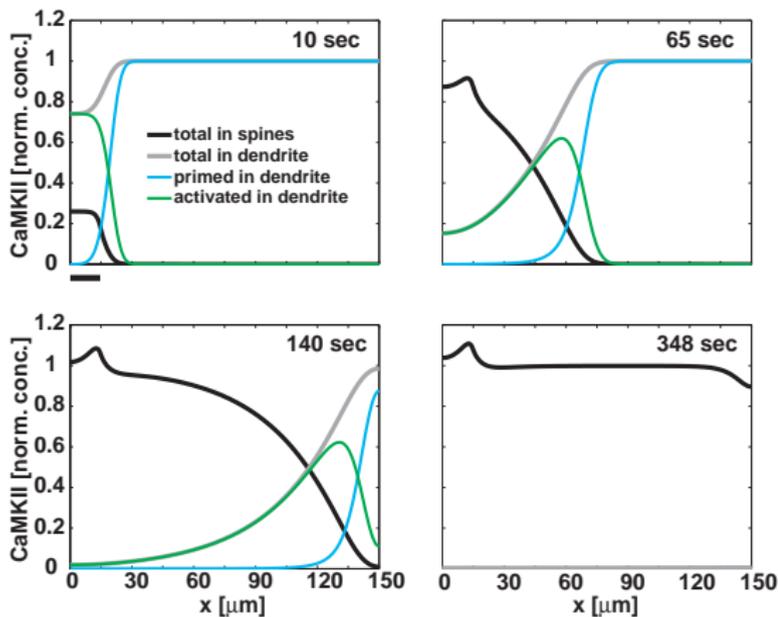
Future directions – trafficking of other proteins

$$\frac{\partial p}{\partial t} = D \frac{\partial^2 p}{\partial x^2} - kap$$

$$\frac{\partial a}{\partial t} = D \frac{\partial^2 a}{\partial x^2} + kap - ha$$

$$\frac{\partial s}{\partial t} = ha$$

BAE & Bressloff, *J. Comput. Neurosci.*



Thank you!

Thanks to

- Paul Bressloff (Oxford)
- Michael Ward (UBC)
- National Science Foundation

