

# Insulin Signaling Pathway

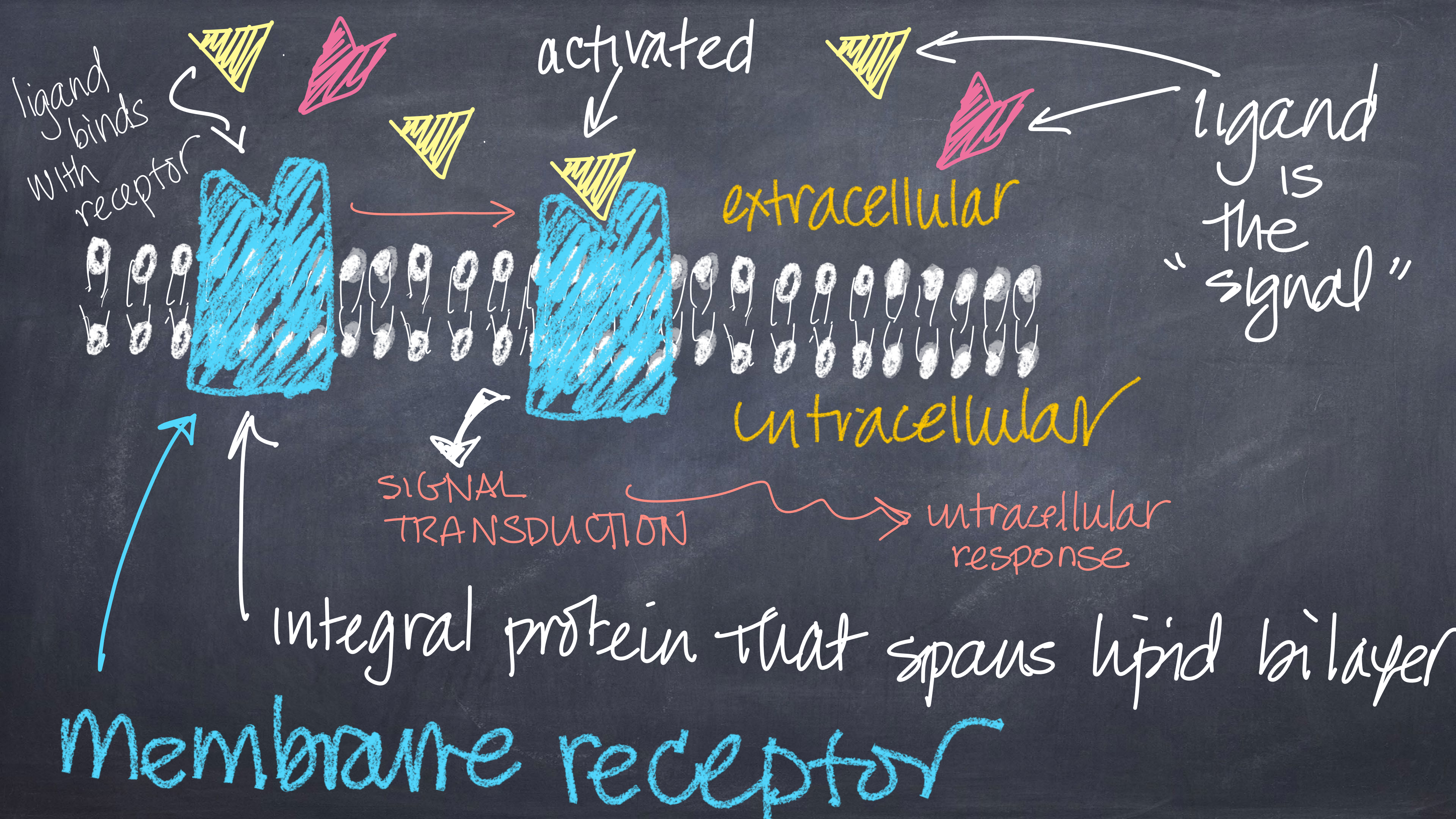
Boyle - Metabolic Brain Disorders

# Insulin Signal Transduction Pathway

- Primary messenger: insulin
- The release of insulin is induced by having specific levels of glucose in the blood.
- Insulin binds to the insulin receptor.

# The Basics of Signaling

- 1. primary messenger → attaches to a receptor
- 2. receptors are a way to mediate information from the outside of the cell to the inside of the cell
- 3. once information is communicated or transduced into the cell there may be a secondary messenger (for example, cAMP or  $Ca^{++}$ ).
- 4. There needs to be a mechanism to terminate the signal.



ligand binds with receptor

activated

ligand is the "signal"

extracellular

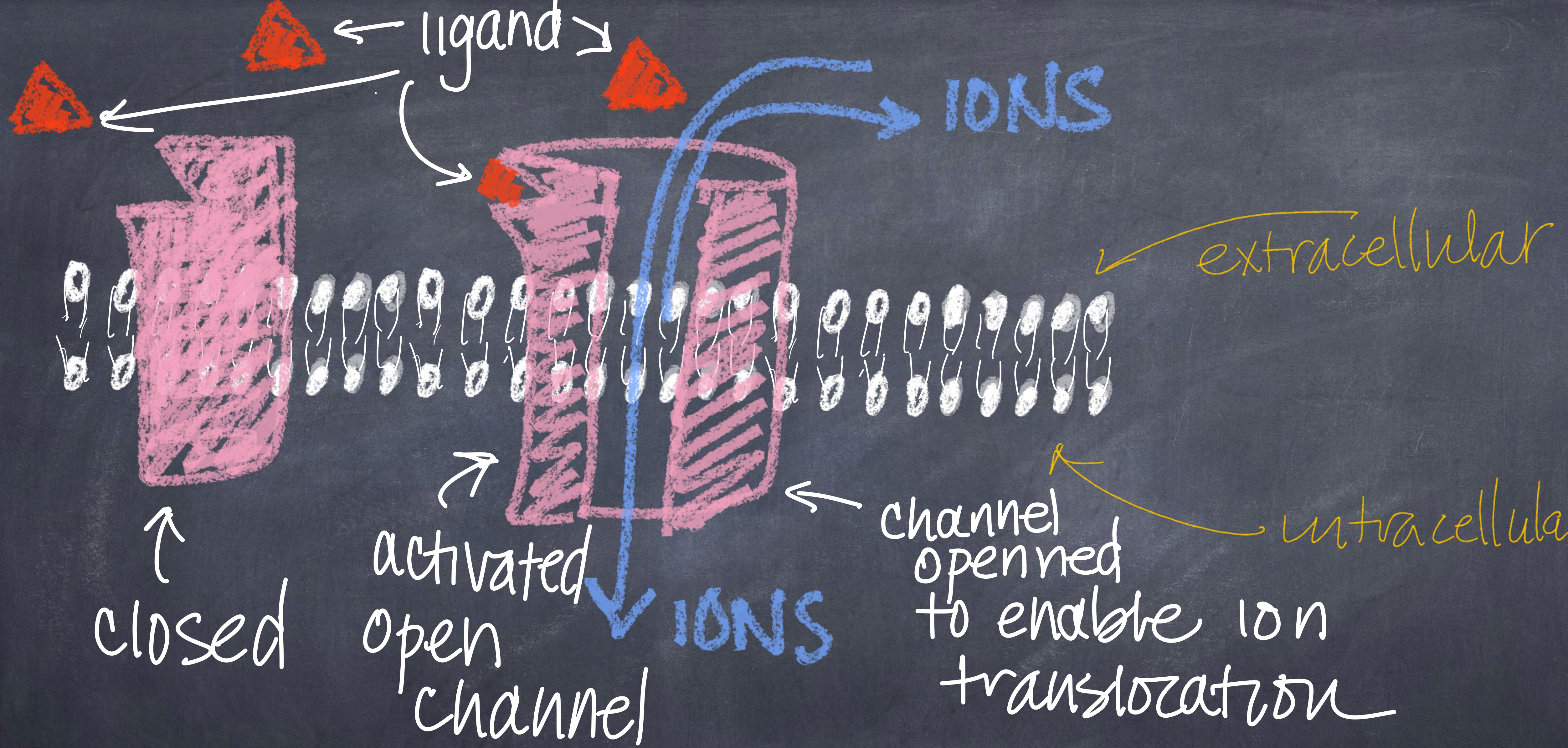
intracellular

SIGNAL TRANSDUCTION

intracellular response

integral protein that spans lipid bilayer

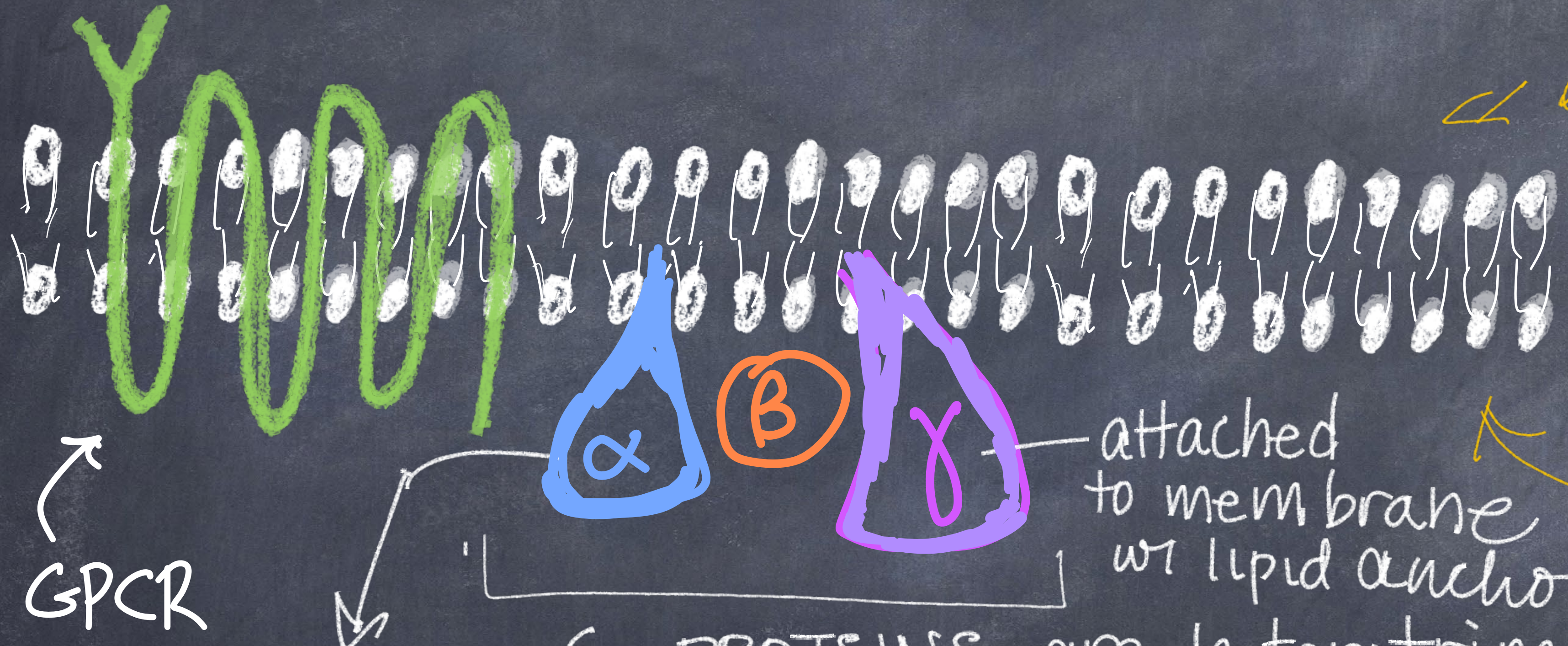
membrane receptor



# LIGAND GATED ION CHANNELS

G-PROTEINS CAN BIND GTP &/OR GDP

GPCR



extracellular

GPCR

attached to membrane w/ lipid anchor

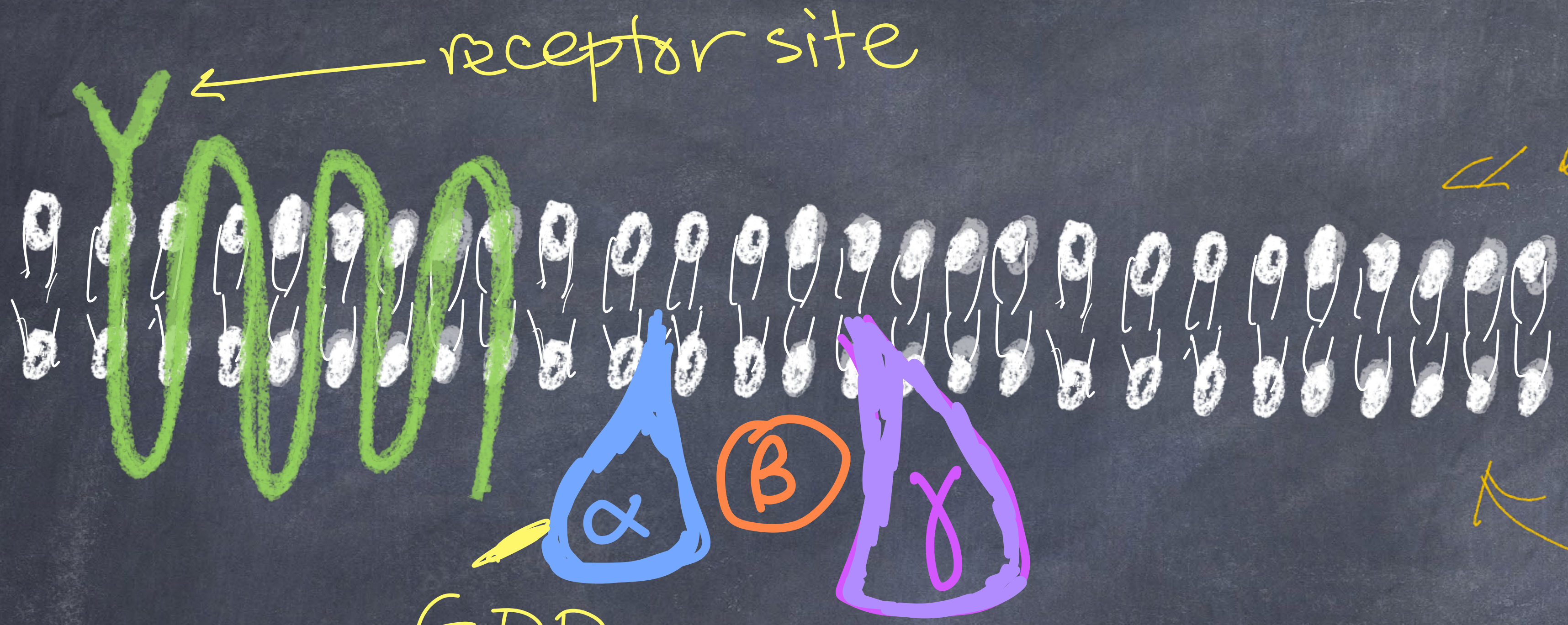
intracellular

G-PROTEINS are heterotrimeric attached to membrane with lipid anchors

G-PROTEIN COUPLED RECEPTORS

G-PROTEINS CAN BIND GTP &/OR GDP

GPCR

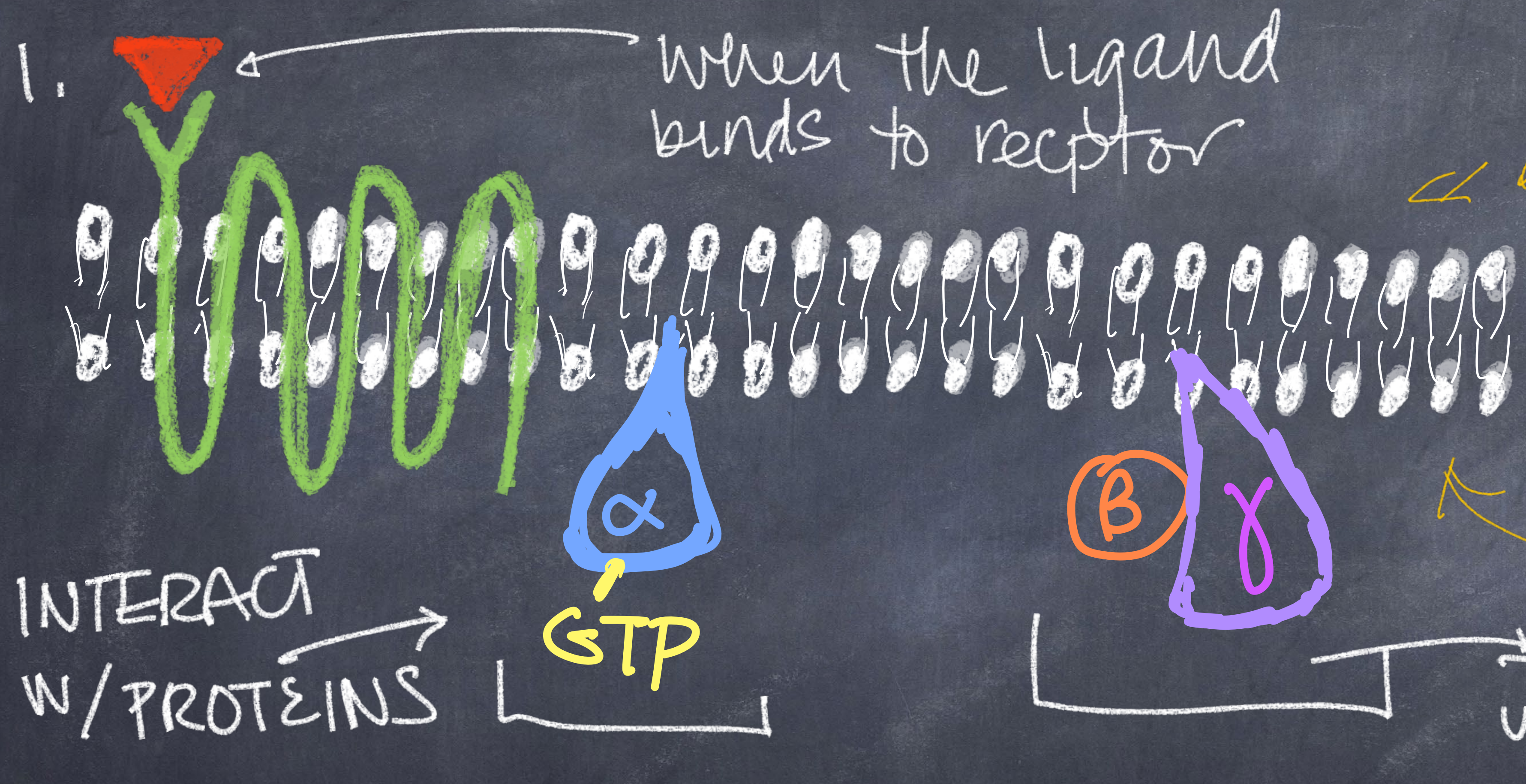


GDP binds to  $\alpha$  subunit

inactive state (GDP)

G-PROTEINS CAN BIND GTP &/OR GDP

GPCR



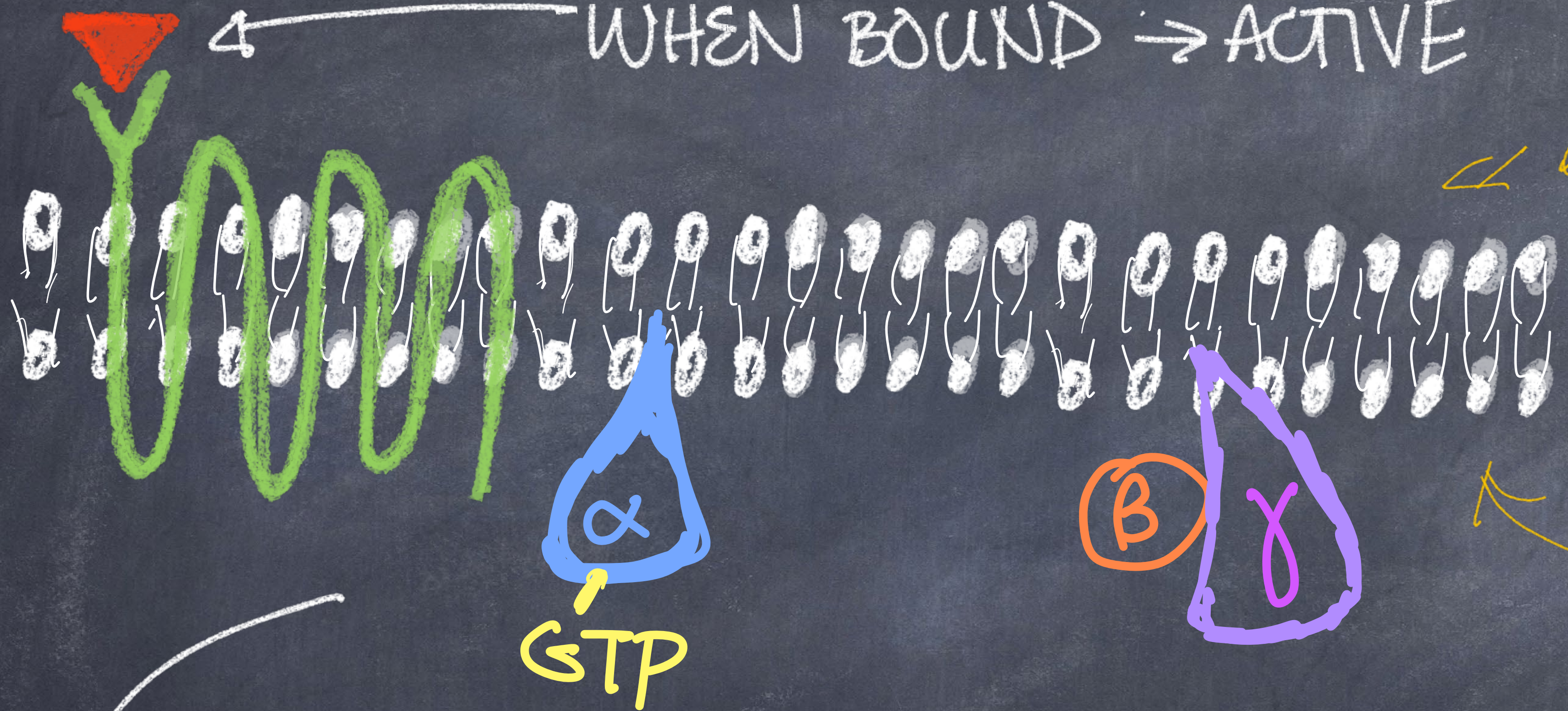
- 2. CONFORMATIONAL CHANGE →  $\alpha$  BINDS GTP
- 3.  $\beta \gamma$  SUBUNIT DISSOCIATE → TARGET PROTEINS



LONG LASTING ...

GPCR

← WHEN BOUND → ACTIVE



extracellular

intracellular

→ the signal stays "active" for as long as the ligand is bound to the receptor.

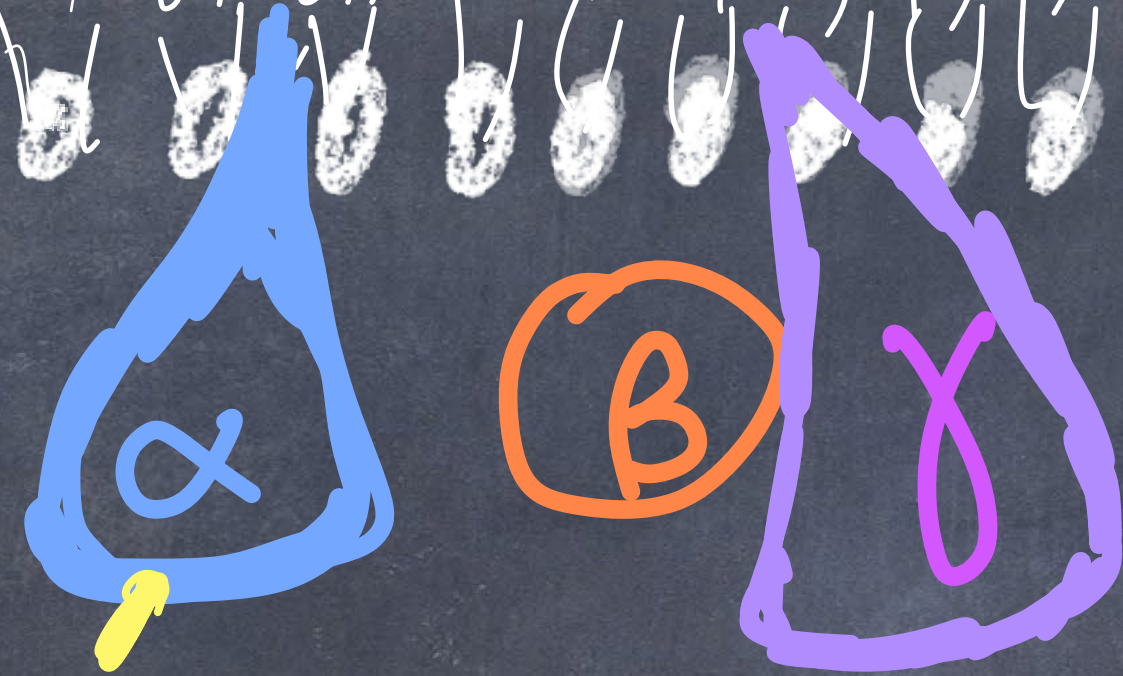
THE END ...

GPCR

LIGAND IS NO LONGER BOUND



extracellular

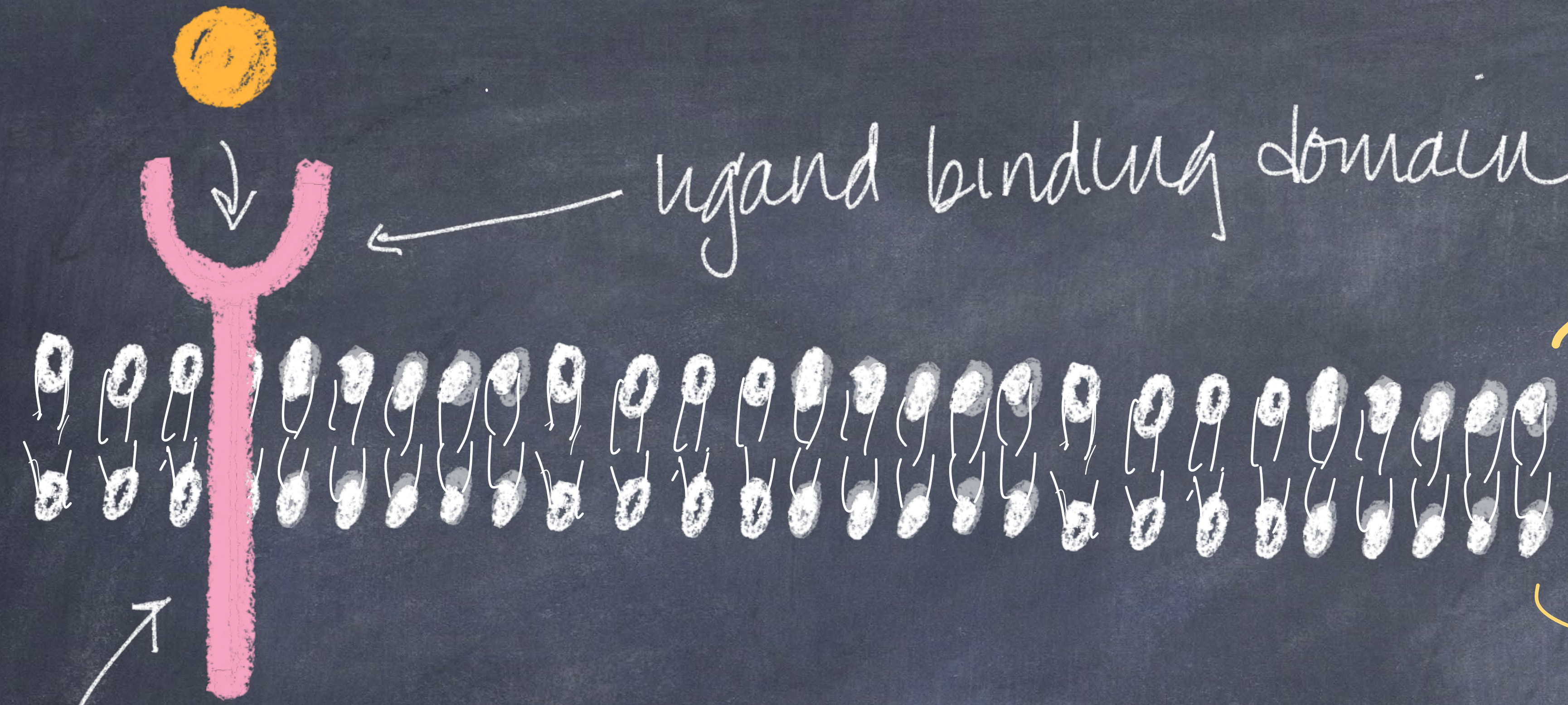


intracellular

GDP

GTP is hydrolyzed to GDP

G. PROTEINS RETURN TO ORIGINAL CONFORMATIONAL STATE



ligand binding domain

extracellular

intracellular

functional domain  
enzyme

catalytic

# ENZYME LINKED RECEPTORS

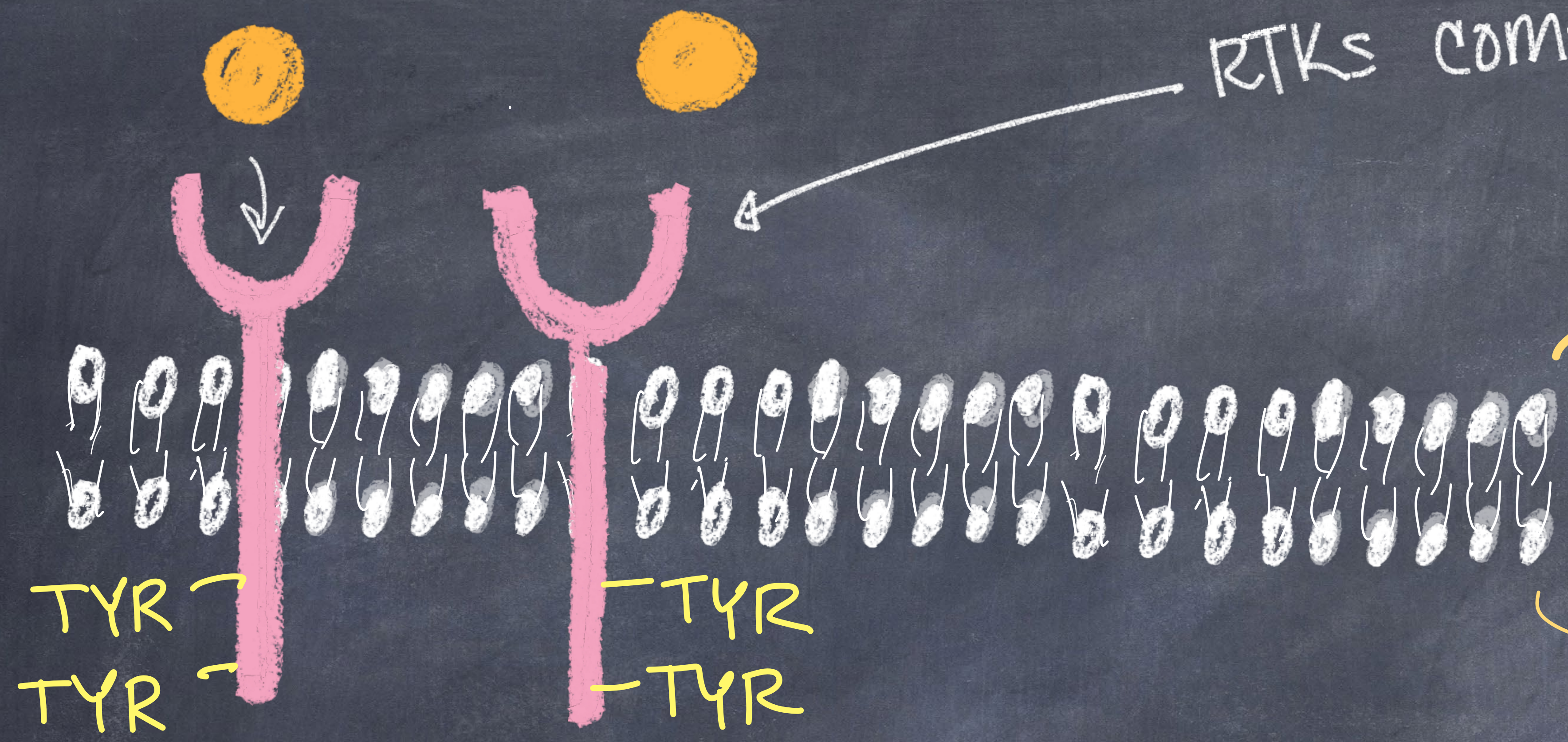
RTKs come in pairs

extracellular

intracellular

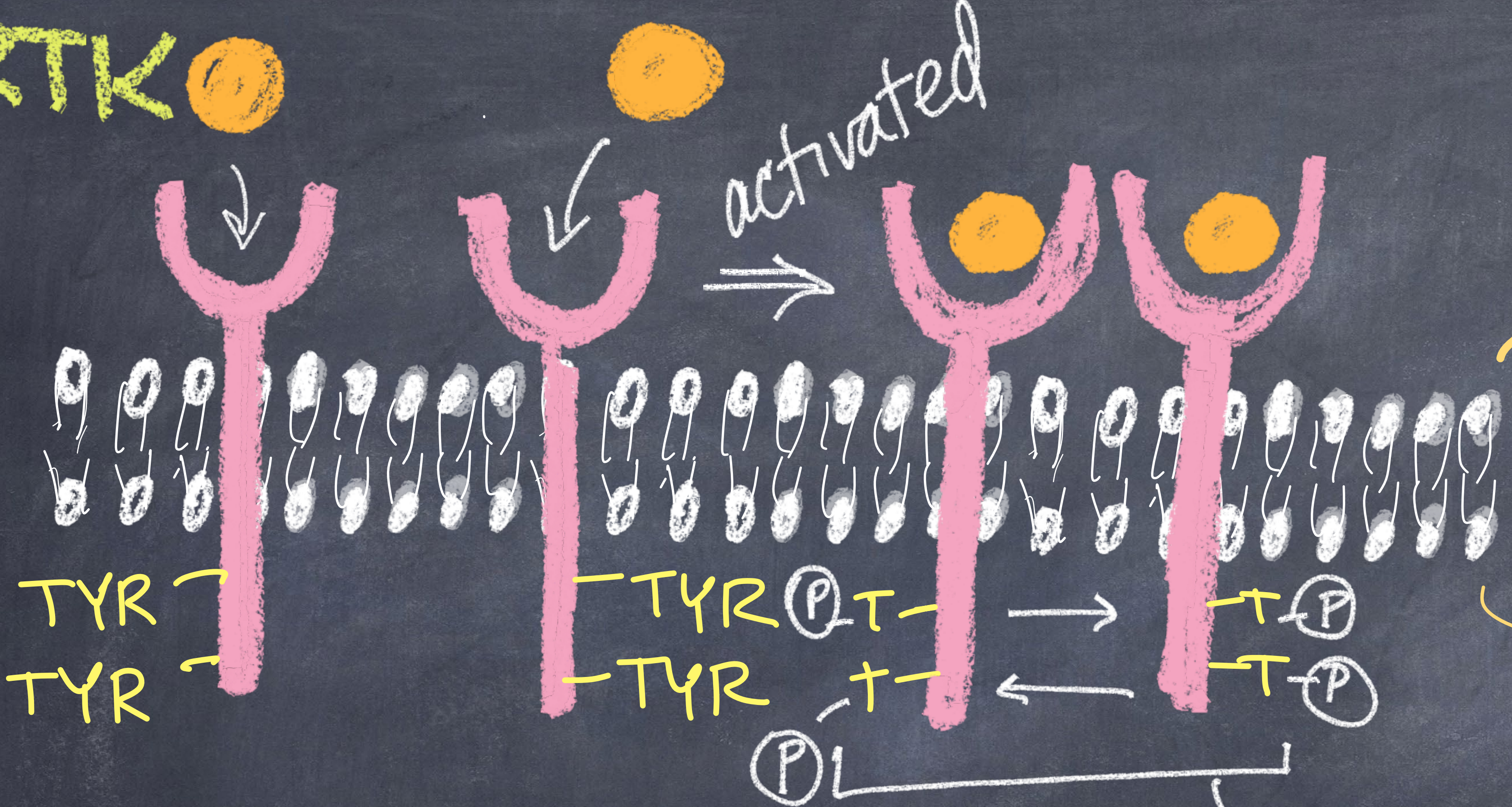
most common!  
RTK

# RECEPTOR TYROSINE KINASE



it has tyrosines here

RTK



extracellular

intracellular

TYR  
TYR

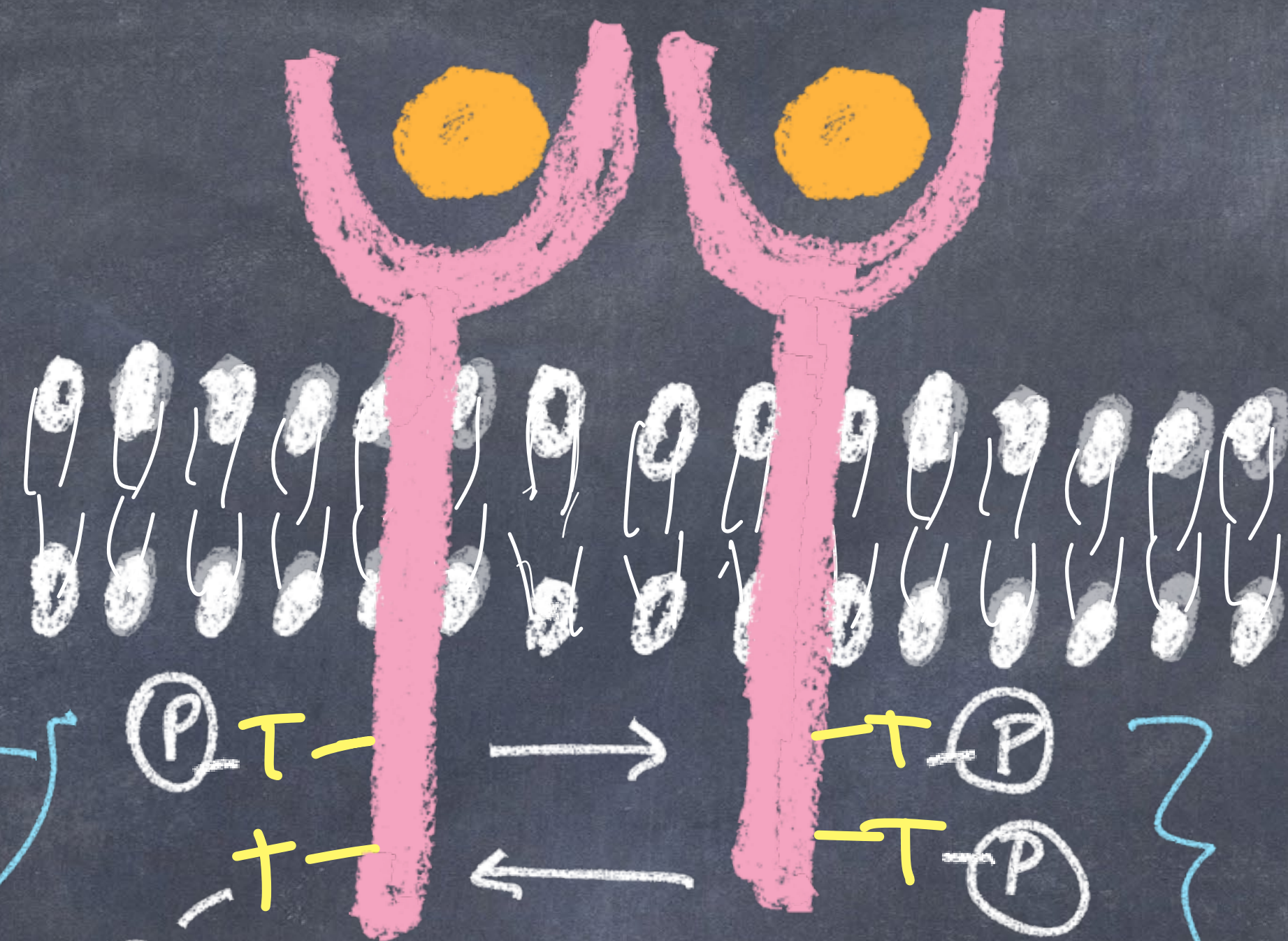
TYR-P  
TYR-P  
P-L

become a  
"cross linked"  
dimer

ONE RTK WILL P THE OTHER

ATP → ADP + P

# RTK



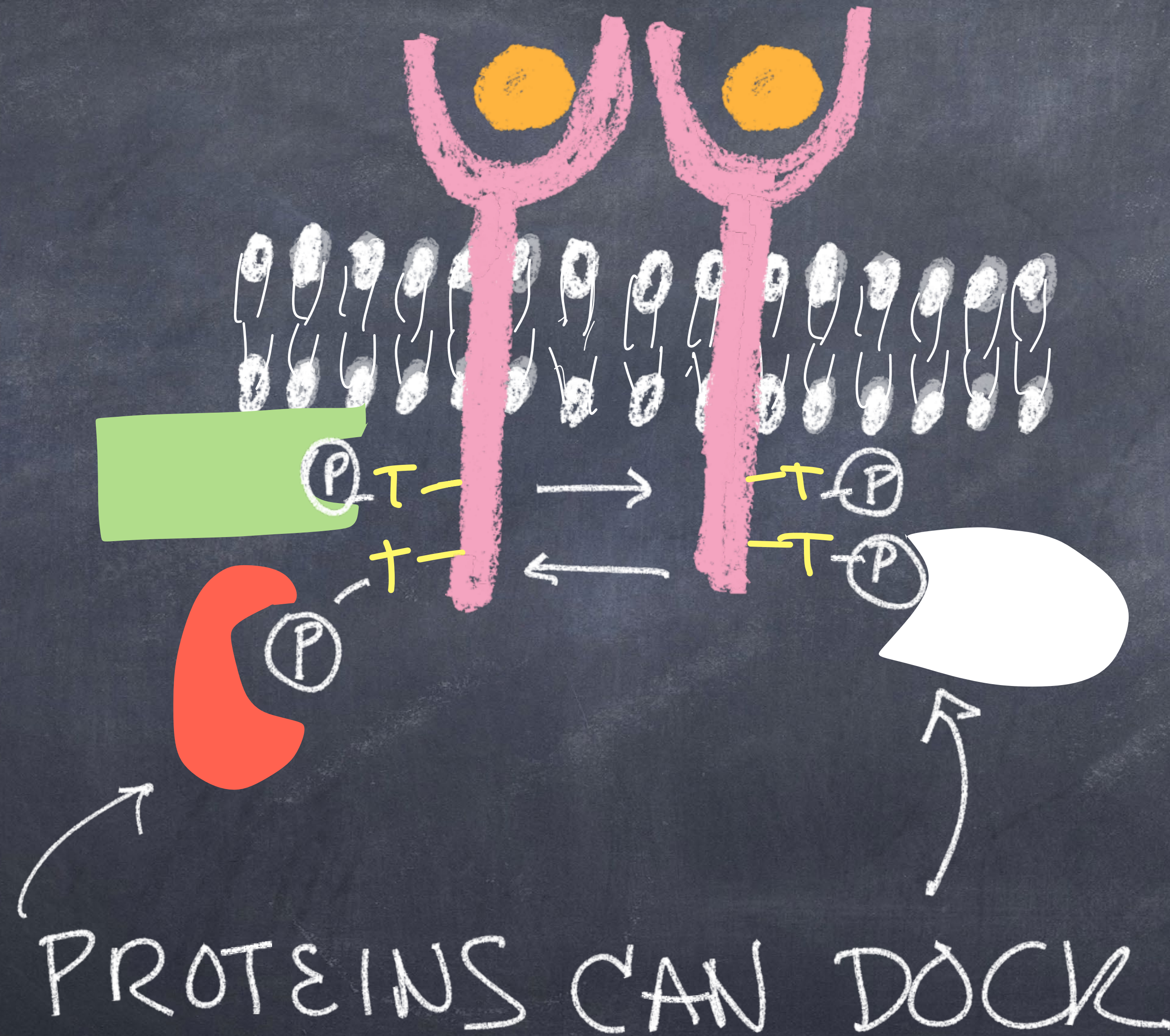
extracellular

intracellular

← THESE WILL SERVE AS DOCKING PLATFORMS

FOR DIFFERENT INTRACELLULAR PROTEINS

# RTK



extracellular

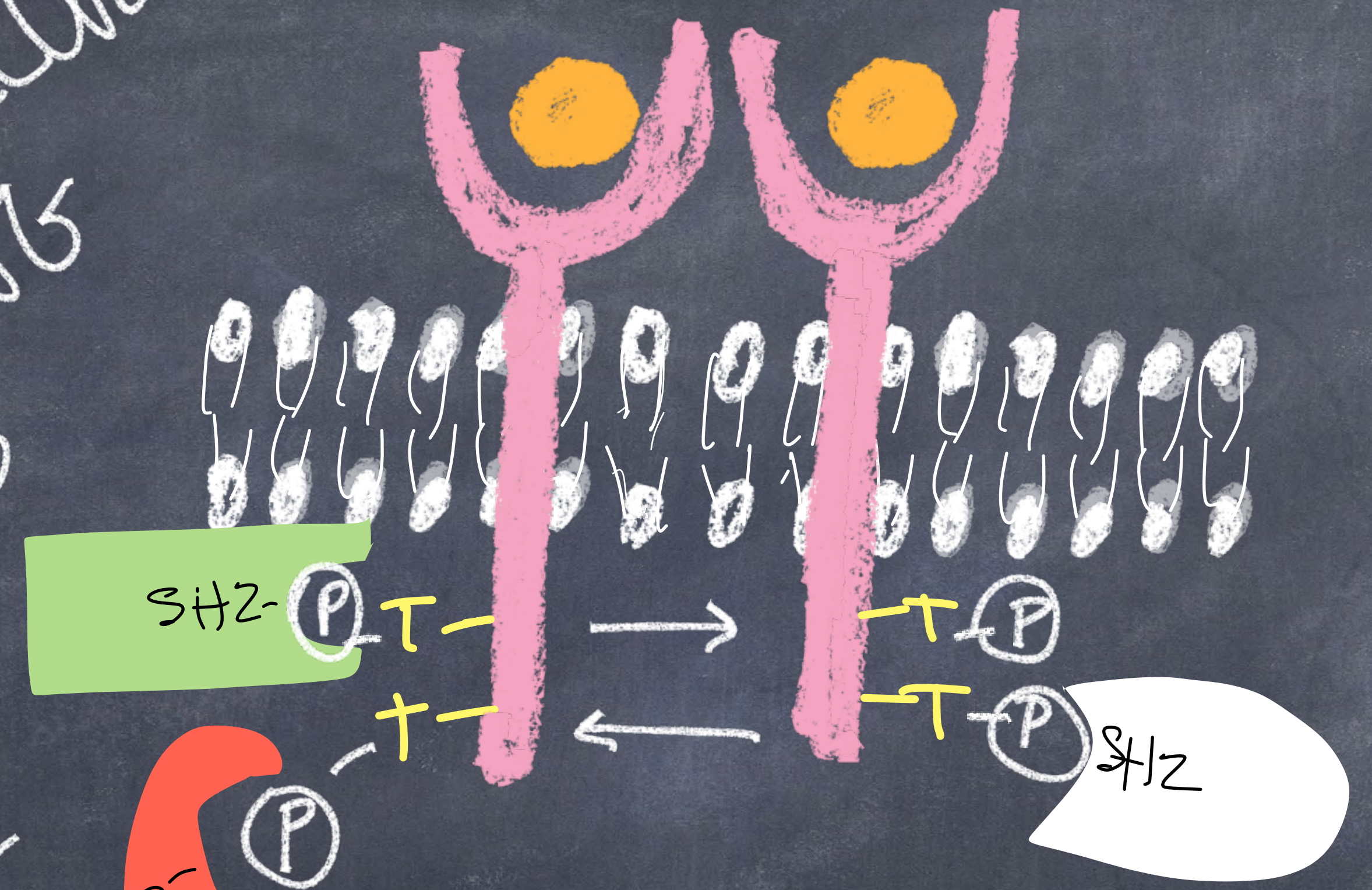
intracellular

PROTEINS CAN DOCK

# RTK

MULTIPLE  
INTRACELLULAR  
SIGNALING  
PATHWAYS  
IN  
PARALLEL

NOW!!



extracellular  
intracellular

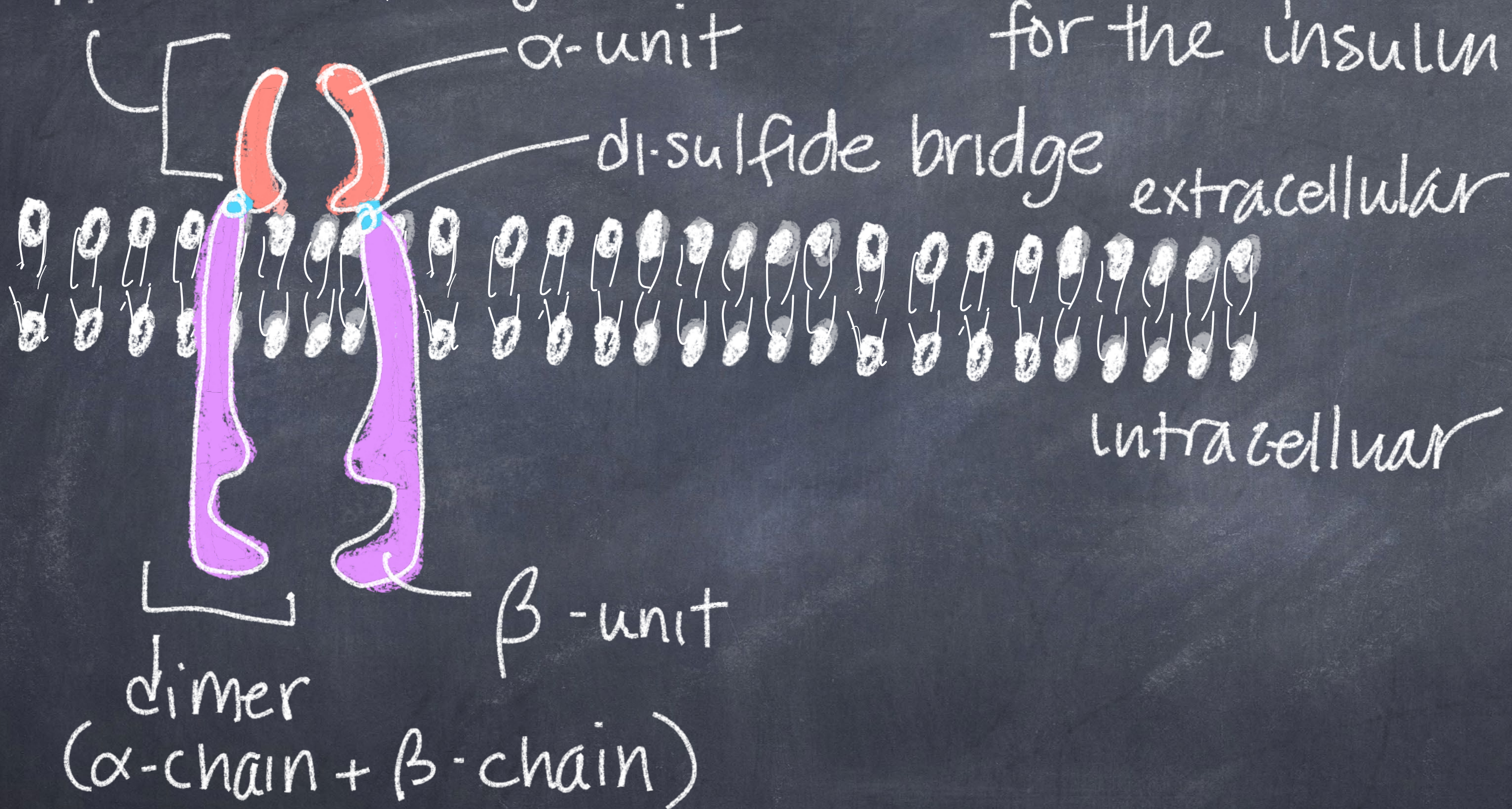


PROTEINS THAT HAVE SH2  
DOMAINS CAN DOCK ON  
T-P

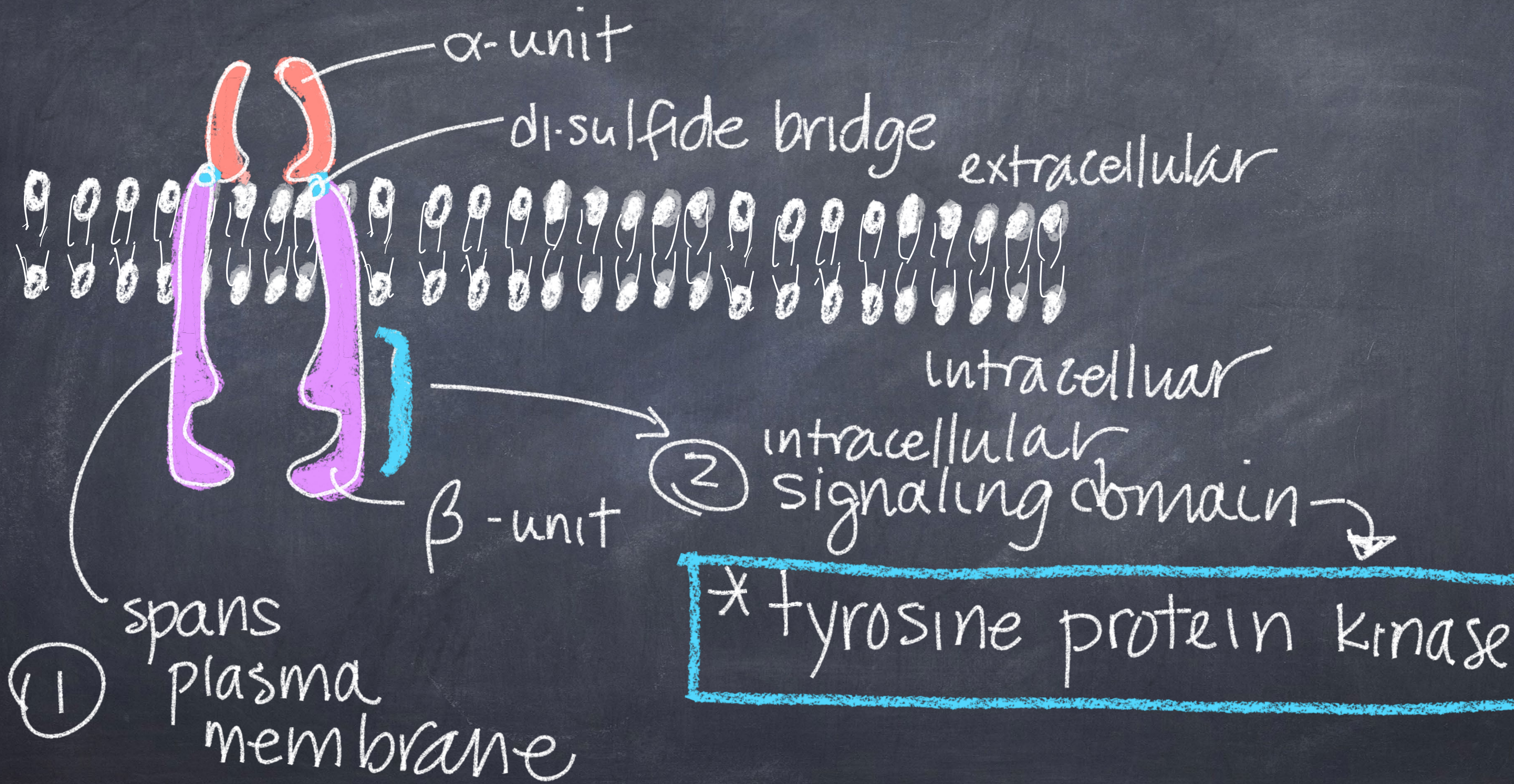


extracellular region

2  $\alpha$ -units form a "pocket" for the insulin



INSULIN RECEPTOR (IR)



Recall: tyrosine protein kinase

① is an enzyme, the Ⓟ

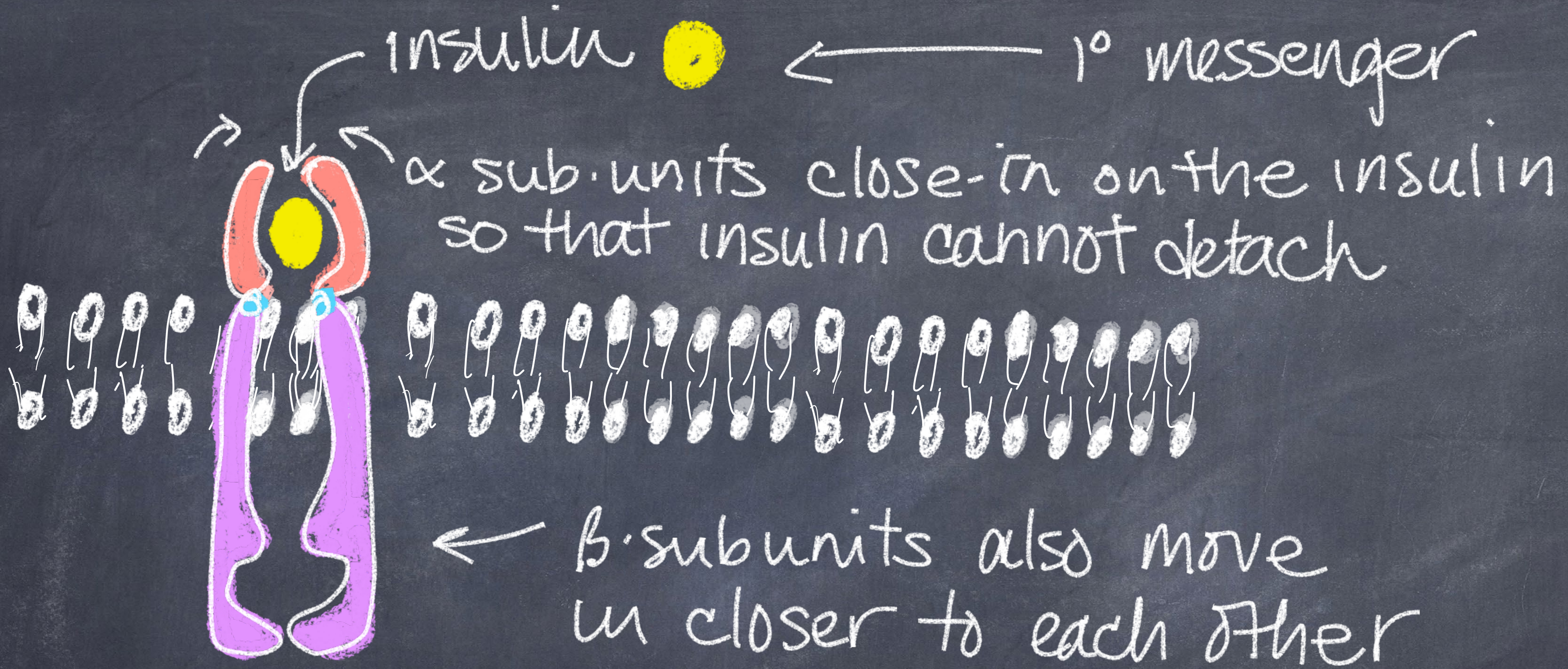
tyrosine  
amino  
acids



note:

\* the protein kinase  
is found in the structure

\* tyrosine protein kinase

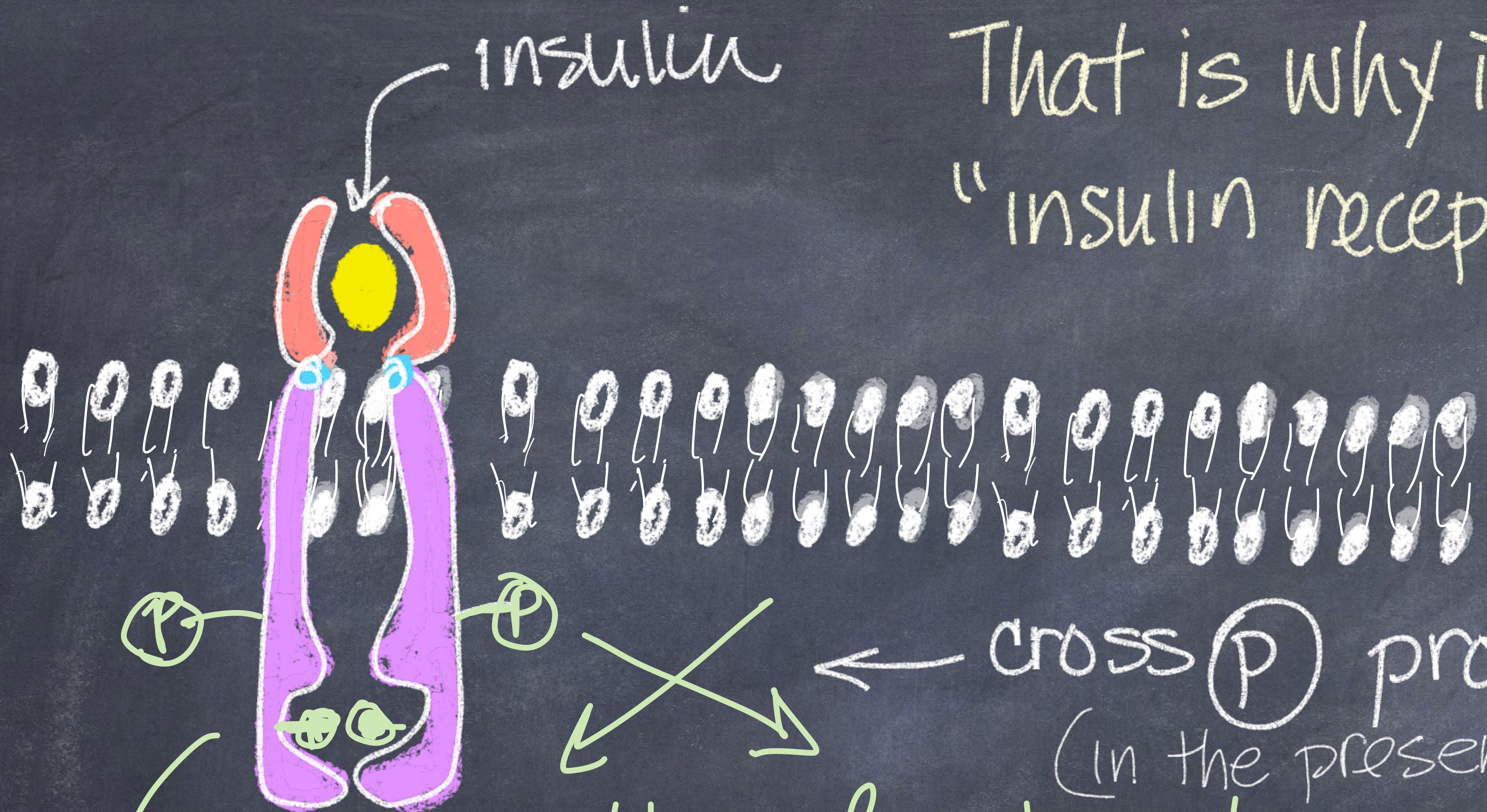


insulin ● ← 1<sup>o</sup> messenger

α sub-units close-in on the insulin so that insulin cannot detach

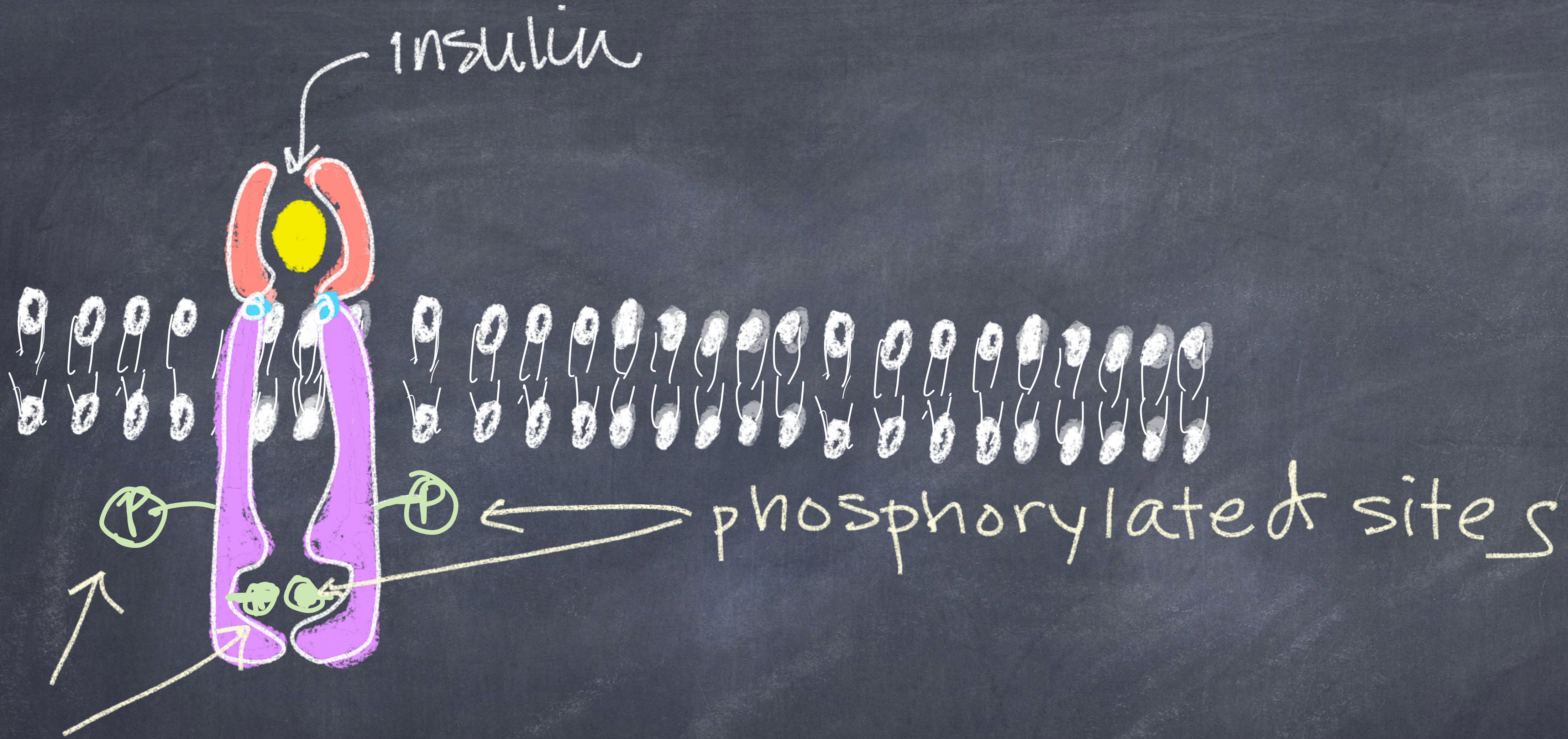
← β-subunits also move in closer to each other

That is why it is called  
"insulin receptor protein  
kinase"

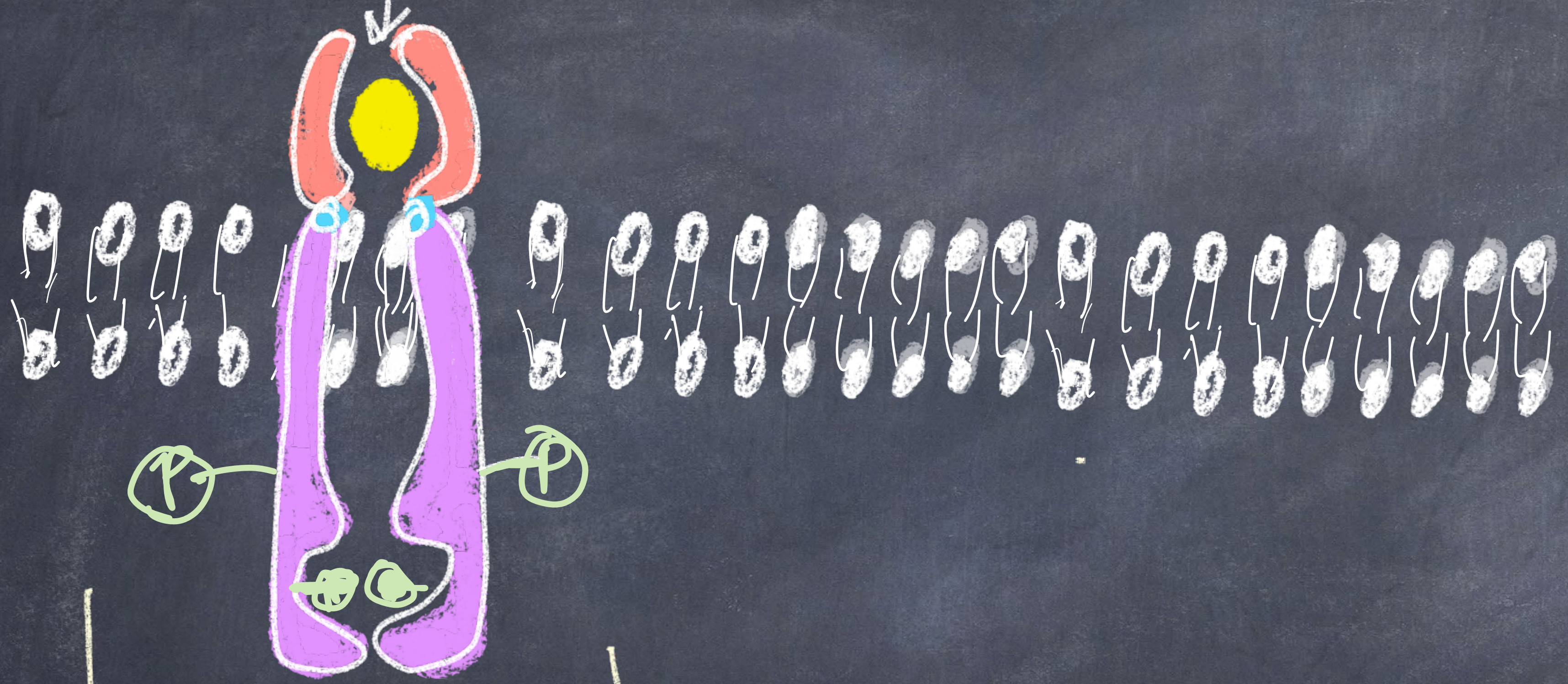


cross (P) process  
(in the presence of ATP)

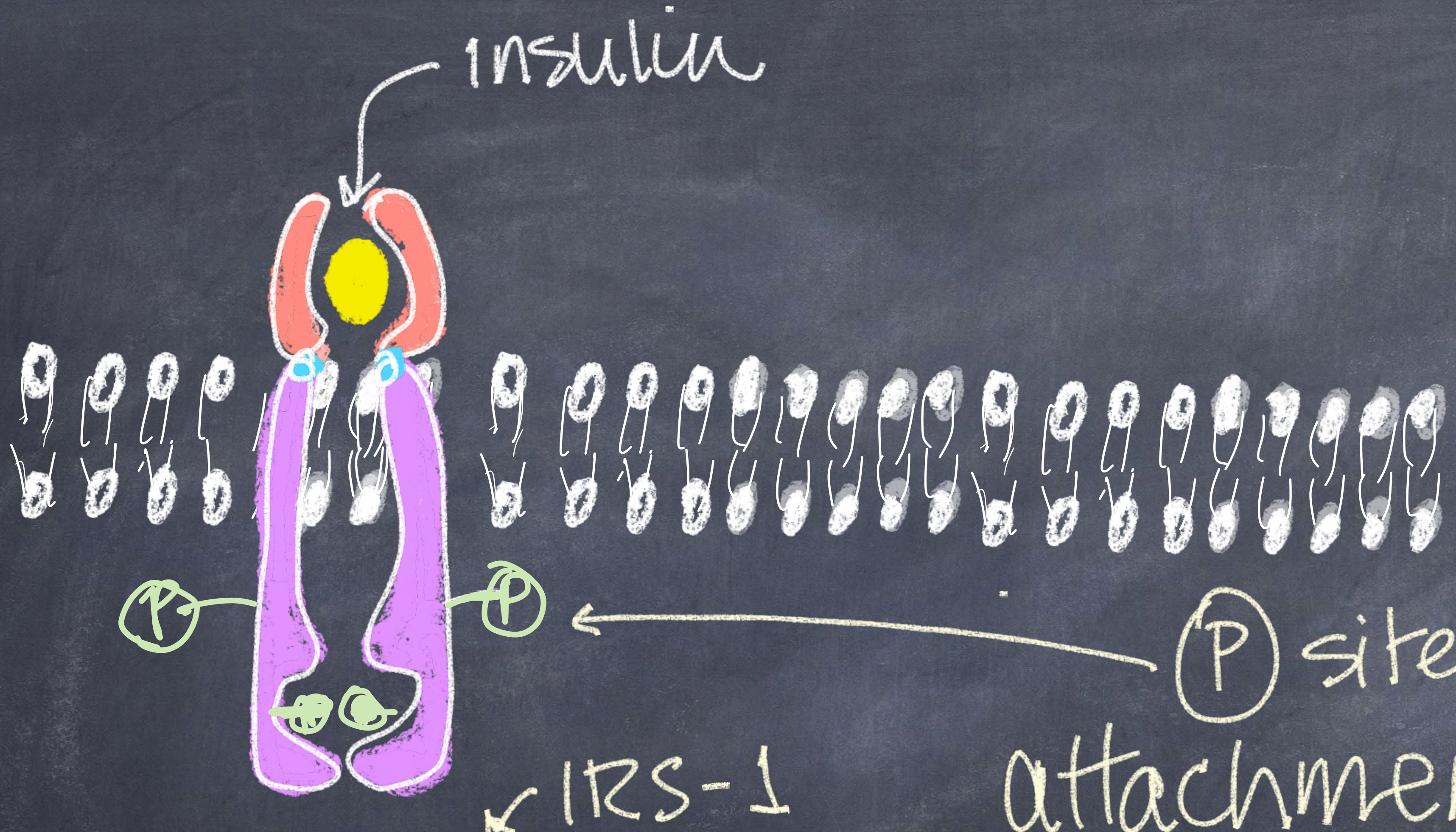
as the  $\beta$ -subunits move closer  
together one subunit activates  
the other subunit.



insulin



activated.



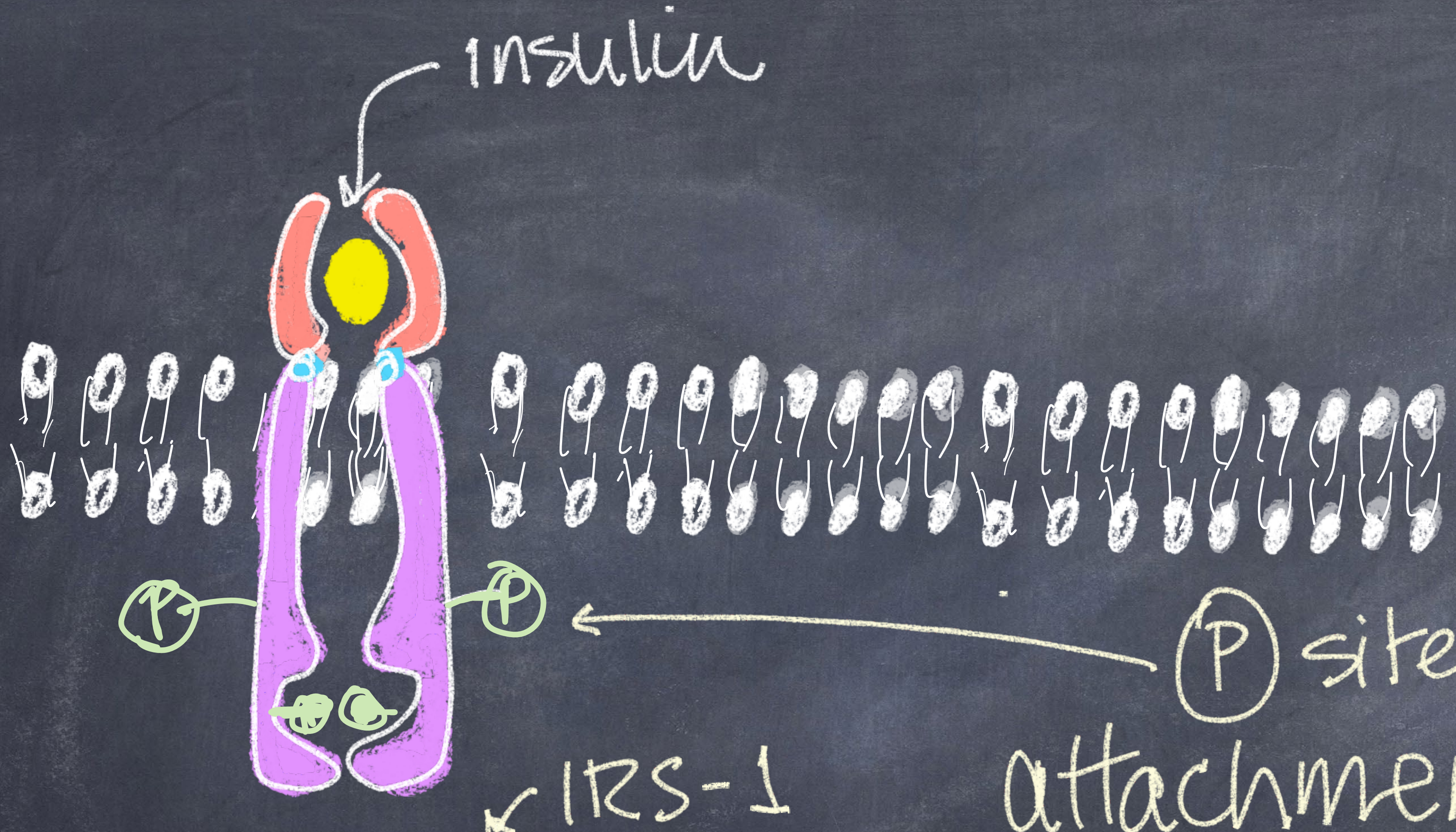
insulin

IRS-1

(P) sites act as attachment sites for other proteins  
→ e.g. IRS-1

Insulin Receptor Substrate



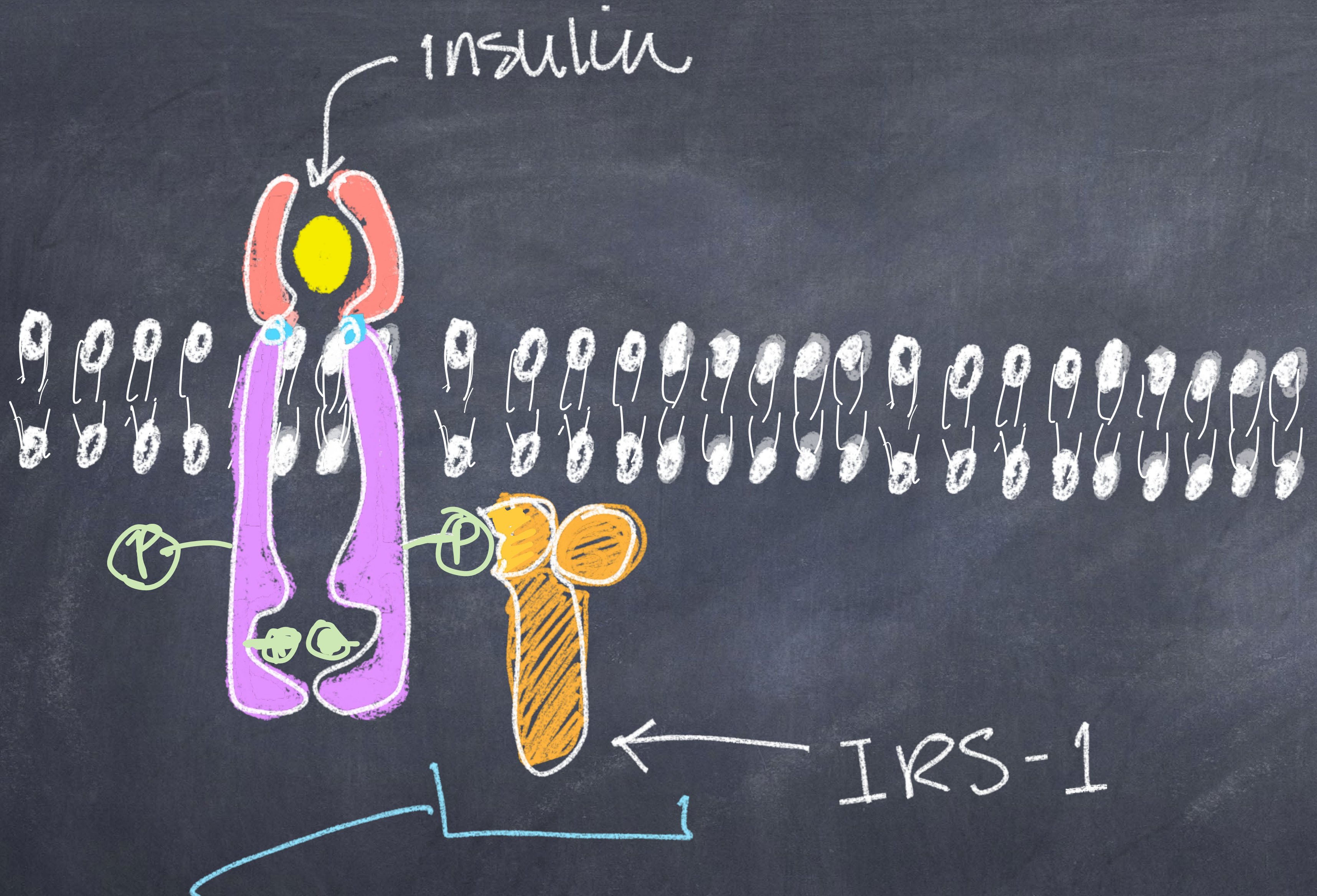


insulin

IRS-1

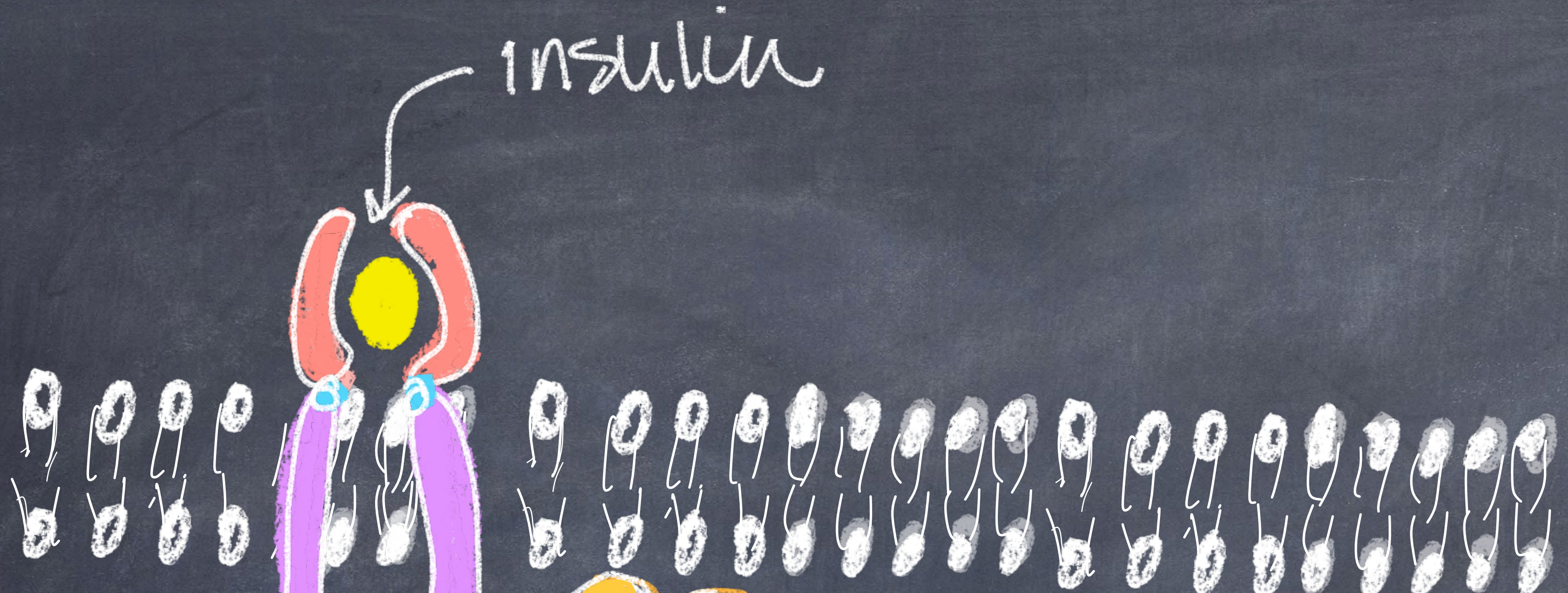
(P) sites act as attachment sites for other proteins  
→ e.g. IRS-1

Insulin Receptor Substrate



IRS molecules are called adaptor proteins

IRS = insulin receptor substrate



insulin



IRS-1

← upon binding, IRS-1 is phosphorylated by the insulin receptor kinase