

Characterization of a Novel Function of G α i as an Intracellular pH Sensor

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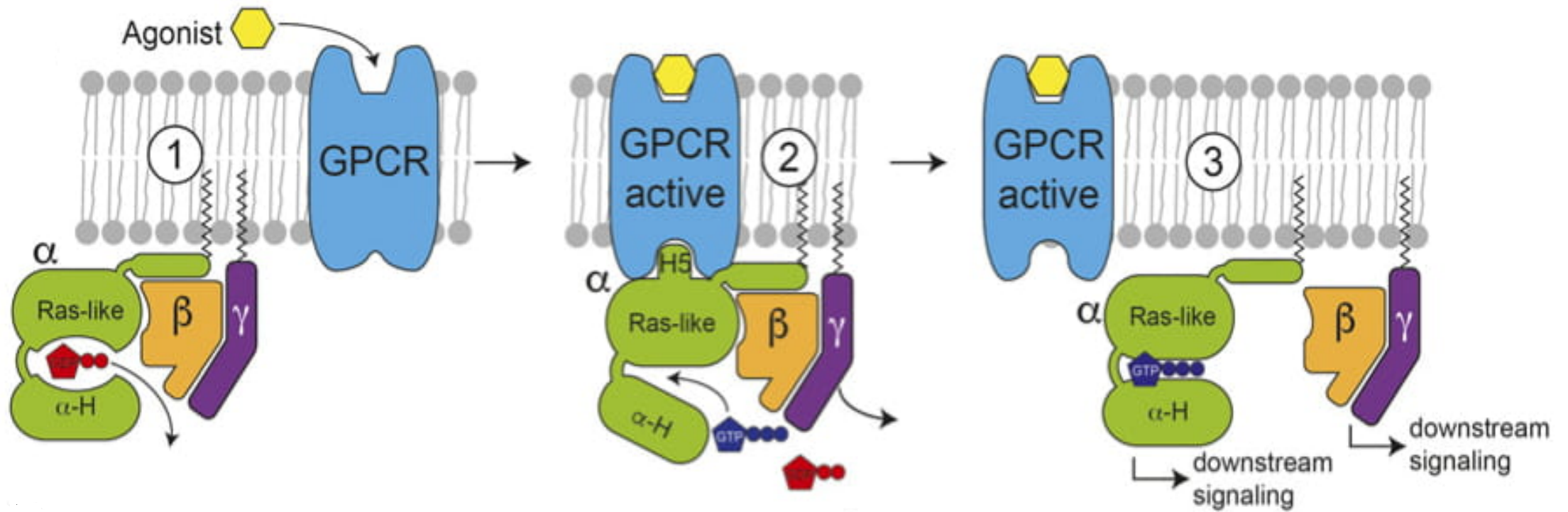
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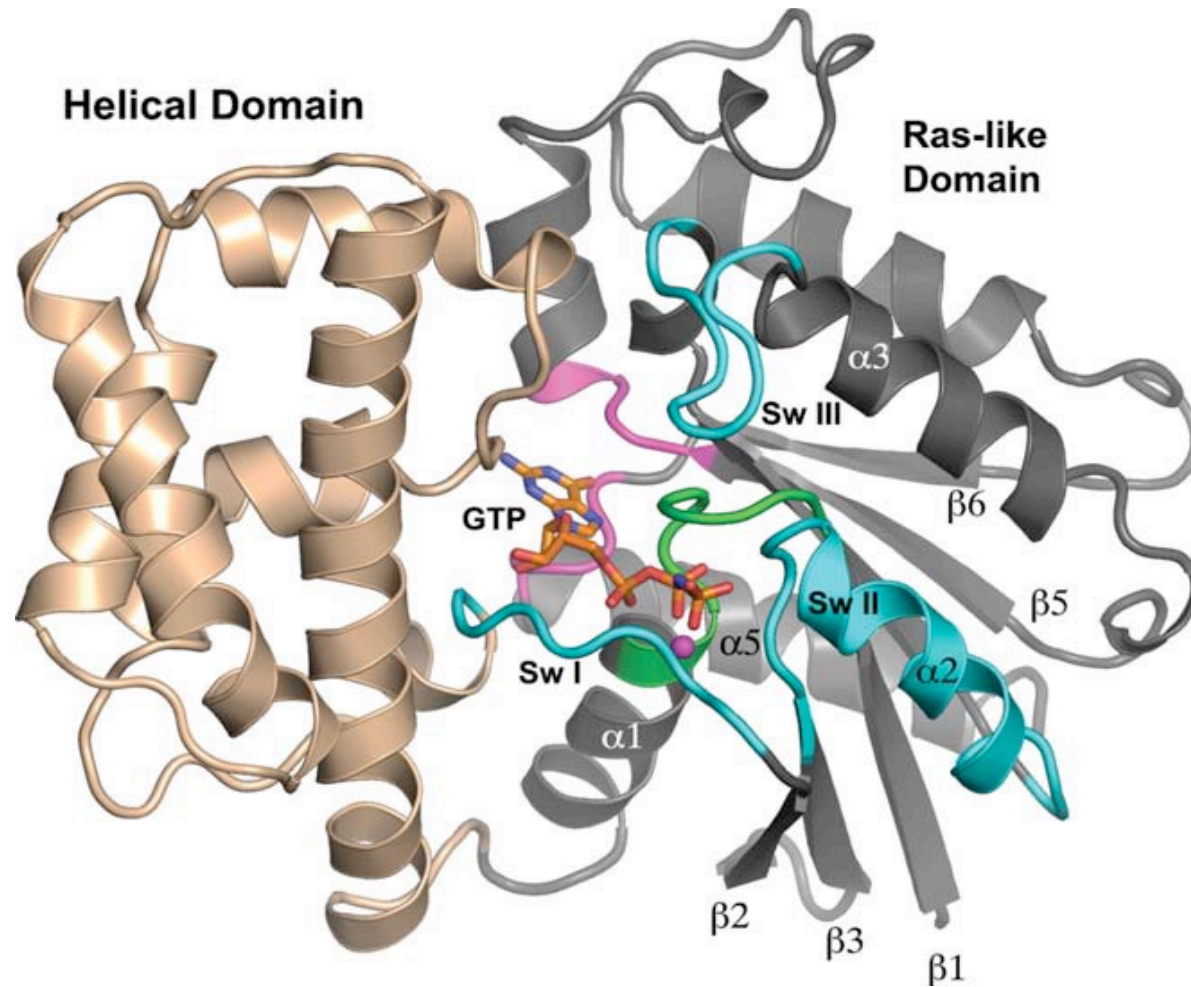


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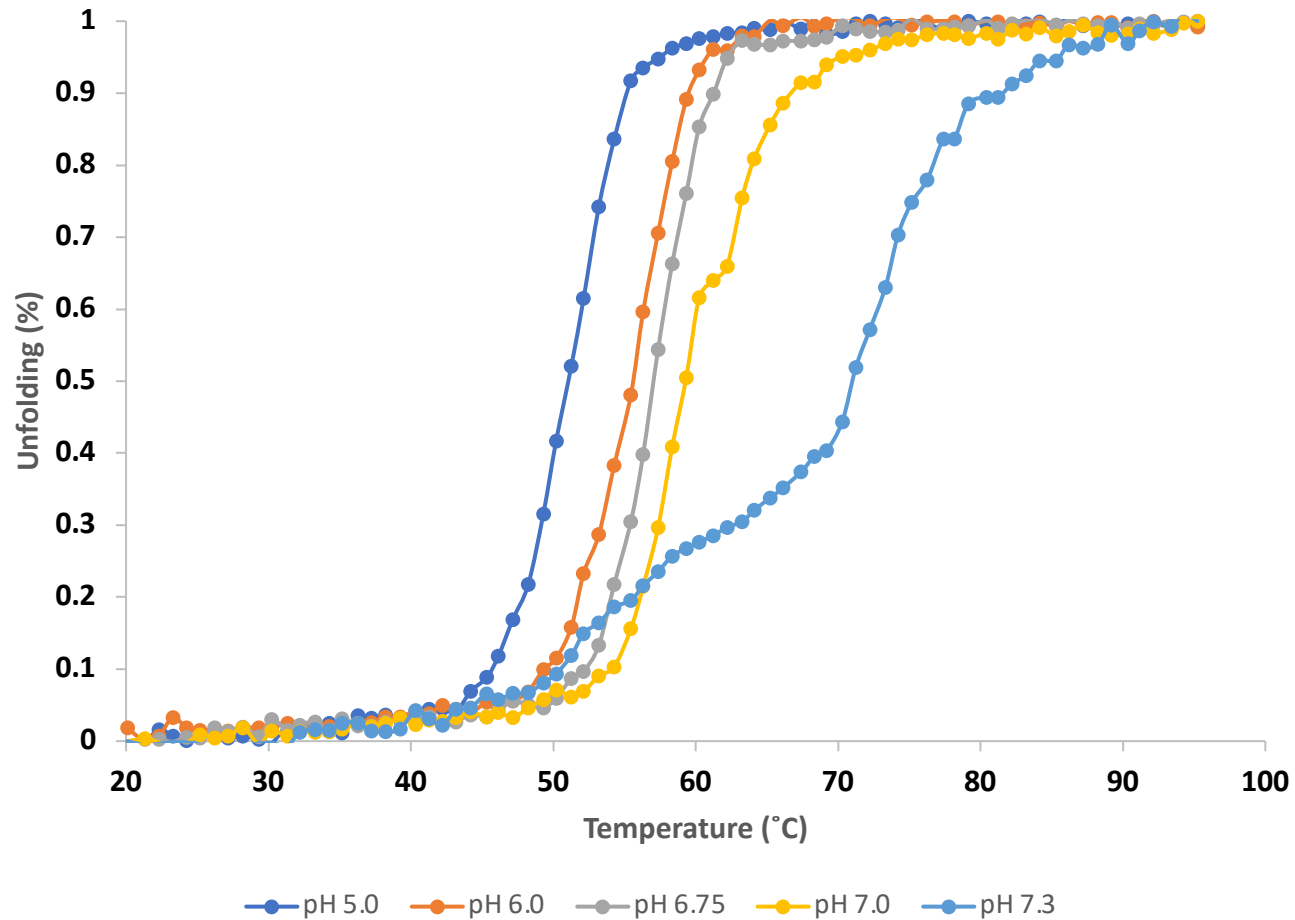
G α is a downstream component of GPCR



$G\alpha$ is a molecular switch

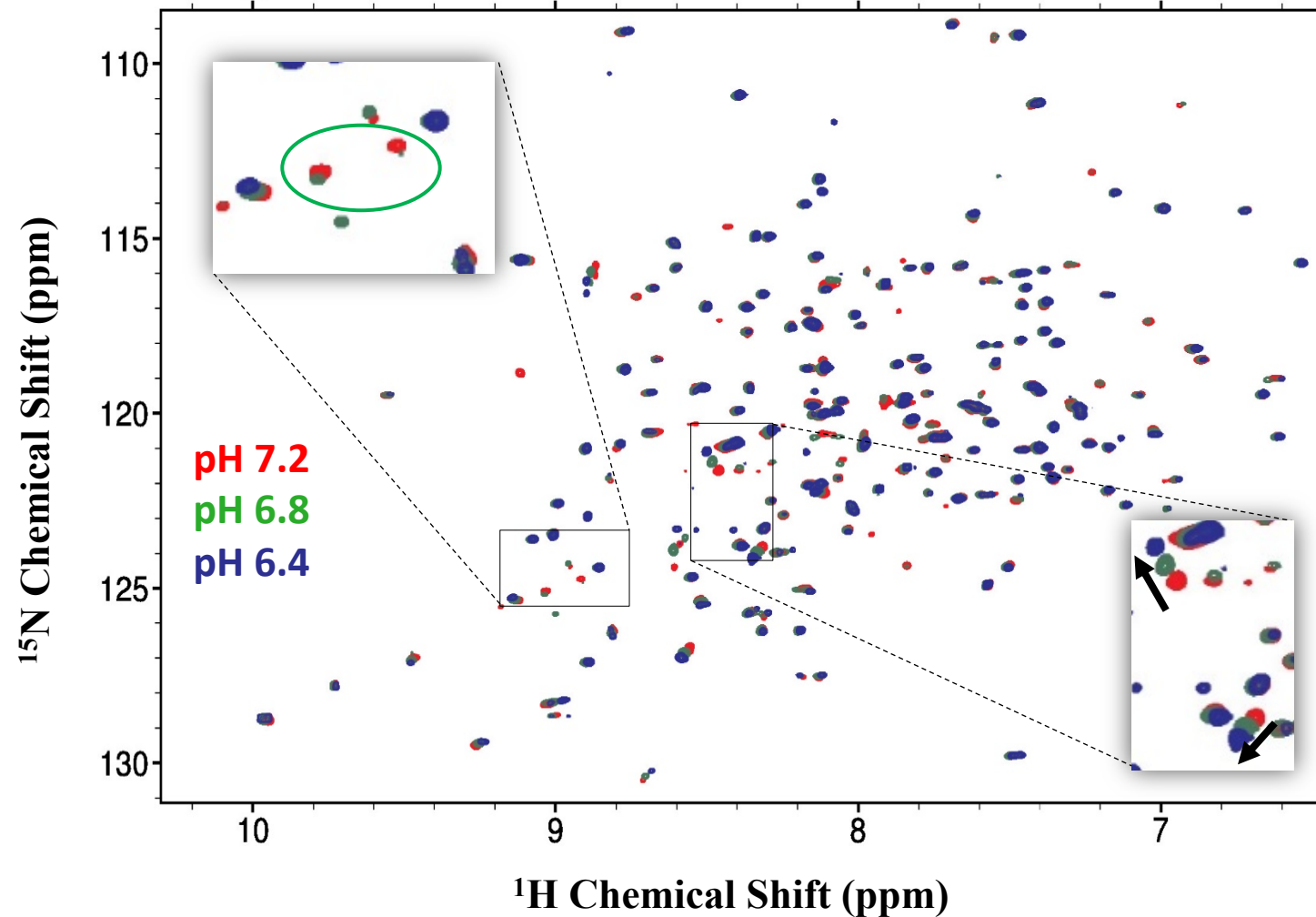


Thermal stability of G α i-GDP is pH-dependent

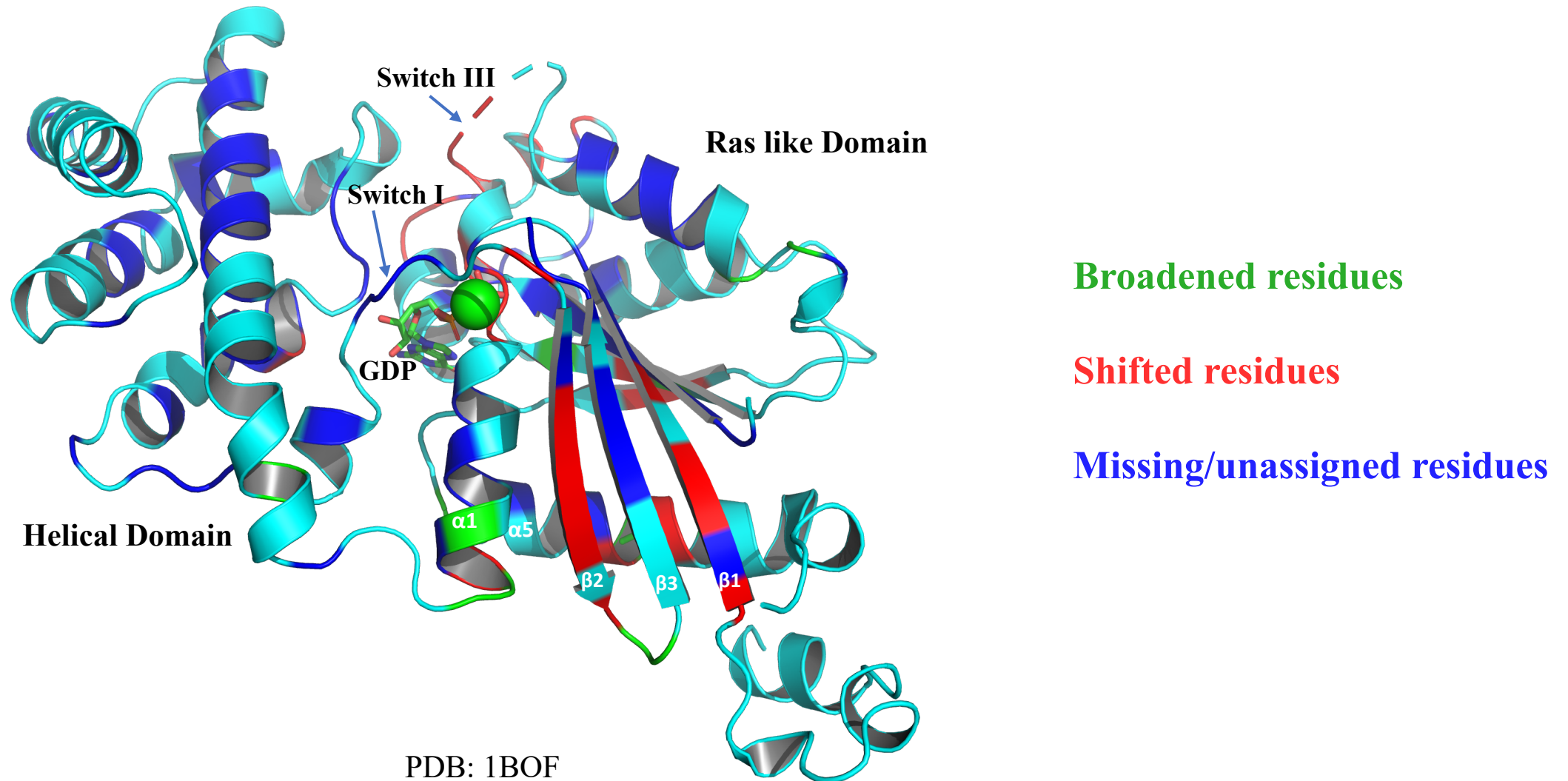


pH	T _m
5.0	50.8
6.0	55.6
6.75	57.2
7.0	58.3, 64
7.3	56.3, 74.2

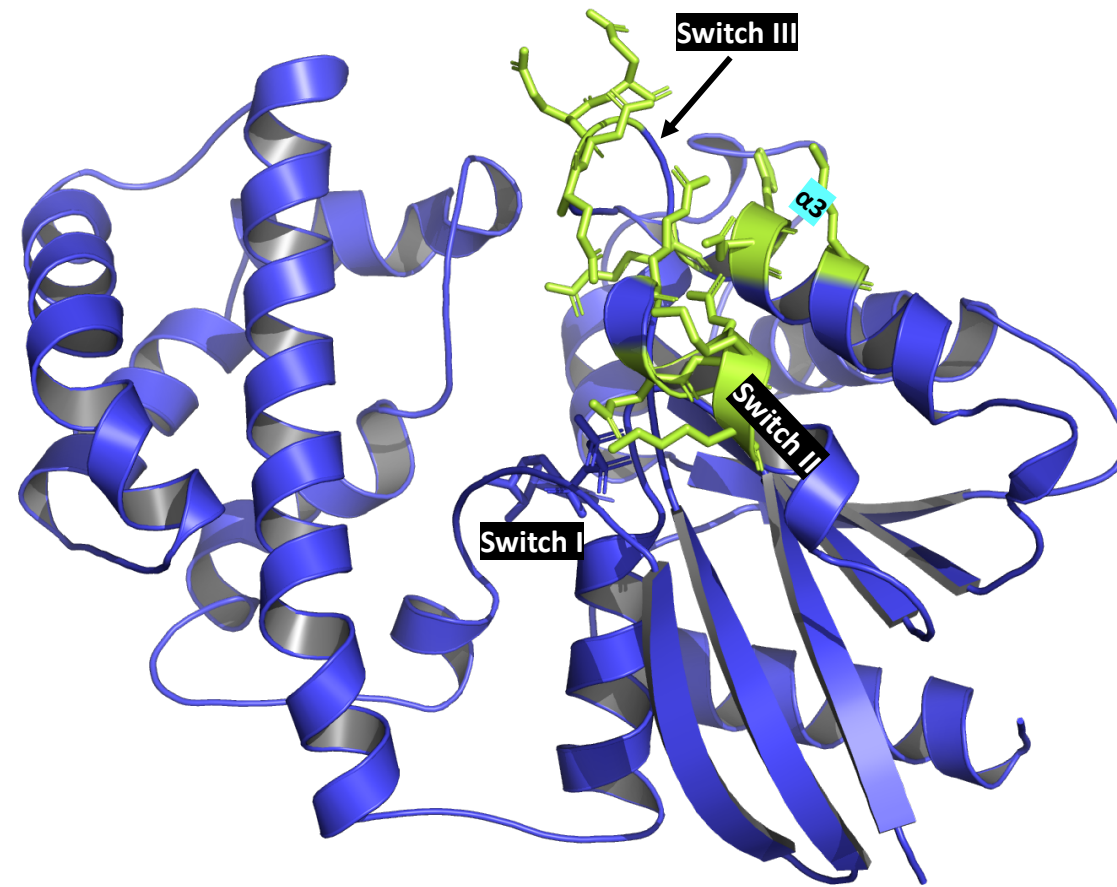
Chemical shift and broadening of G α i-GDP is pH-dependent



Residues in switch regions show NMR chemical shift and broadening for pH 6.4 to 7.2

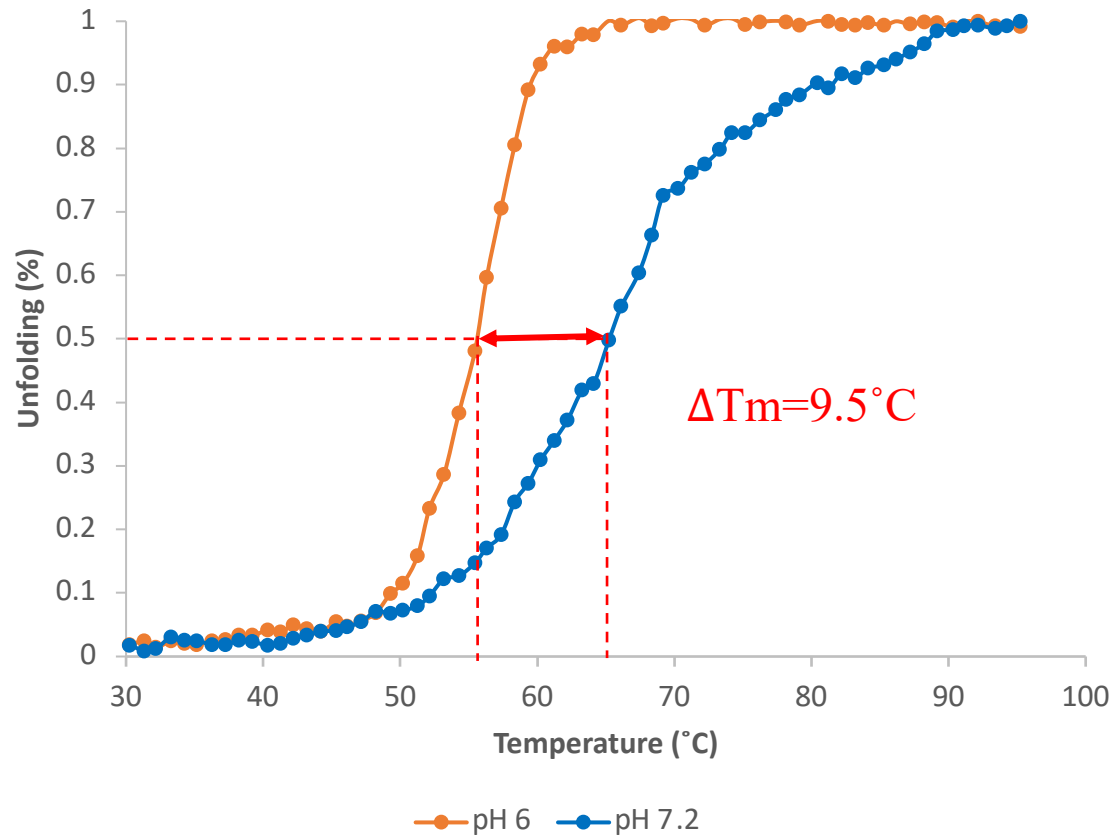


Switch regions of $G\alpha_i$ contain charged amino acids forming interactive network

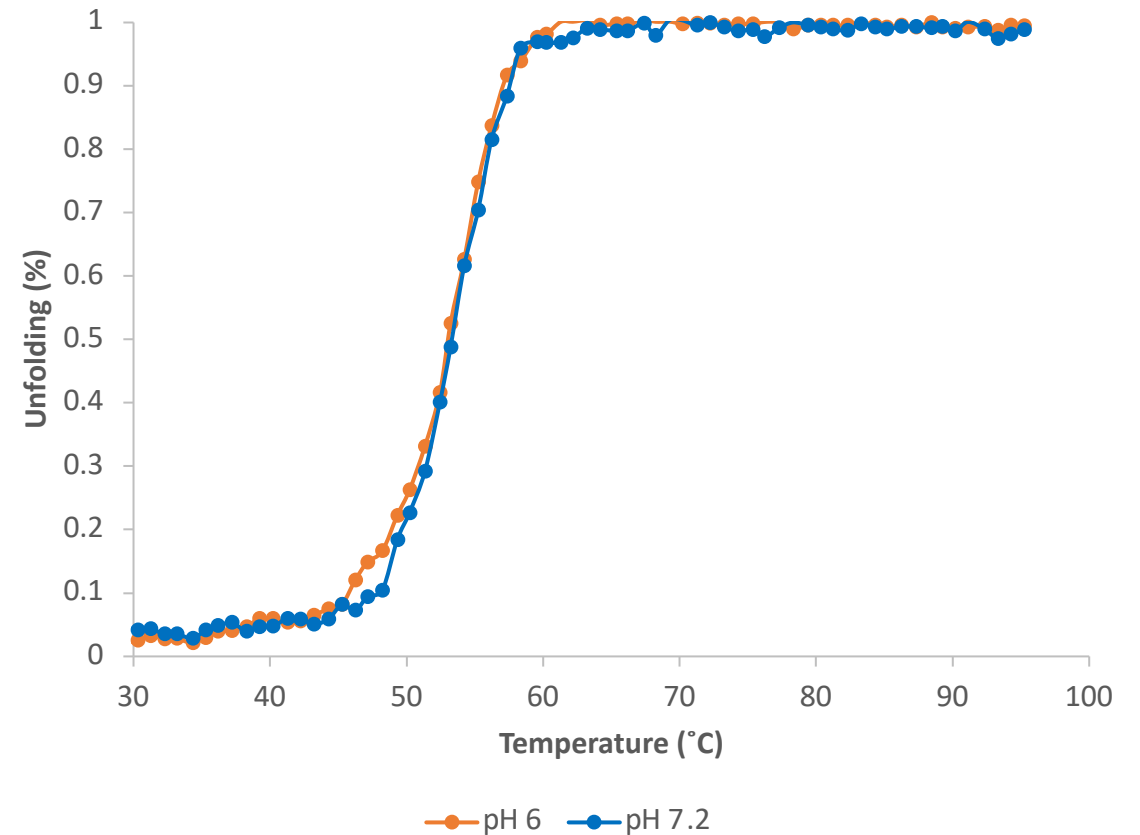


PDB: 1CIP

Gai-GDP Triple mutant from the switch network abolishes pH-dependent thermostability

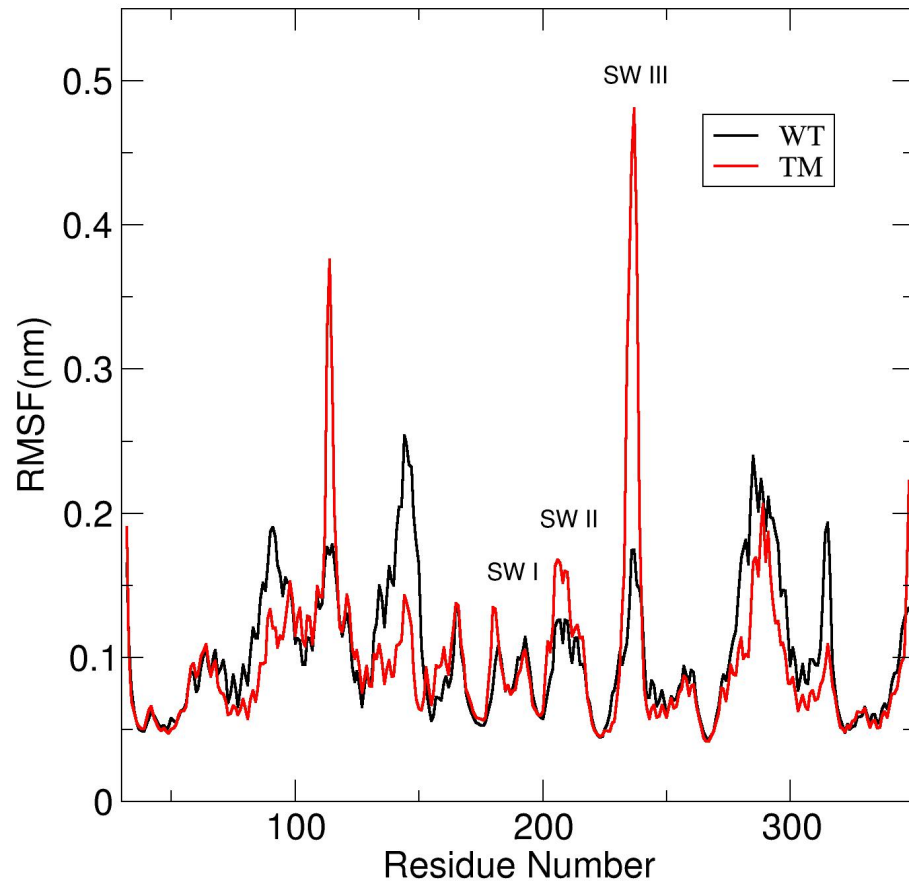


WT Gai

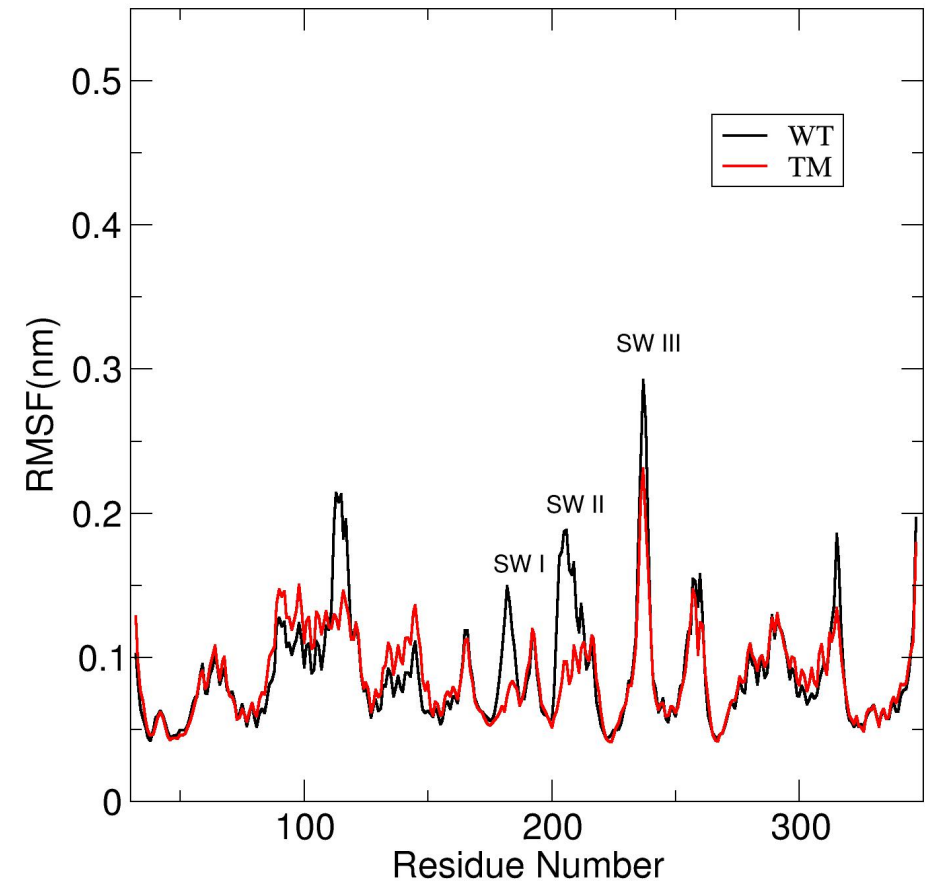


Triple mutant

Triple mutant makes switches more dynamic in $G\alpha i$ -GDP but not in $G\alpha i$ -GTP

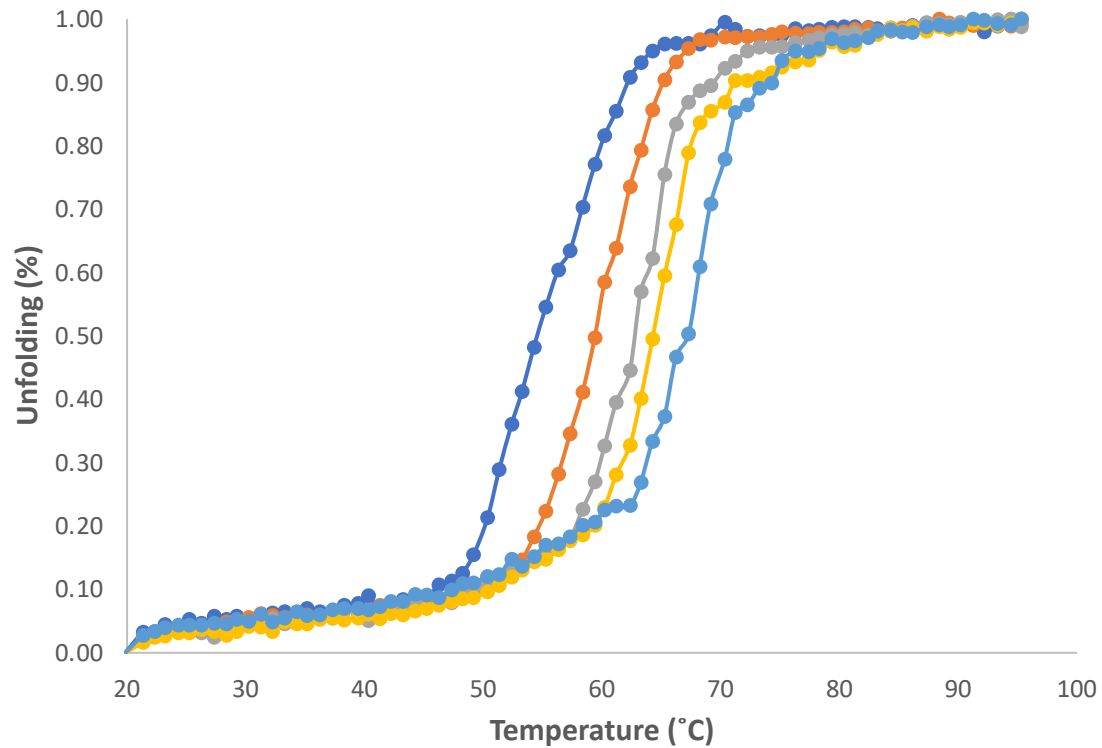


Gai-GDP



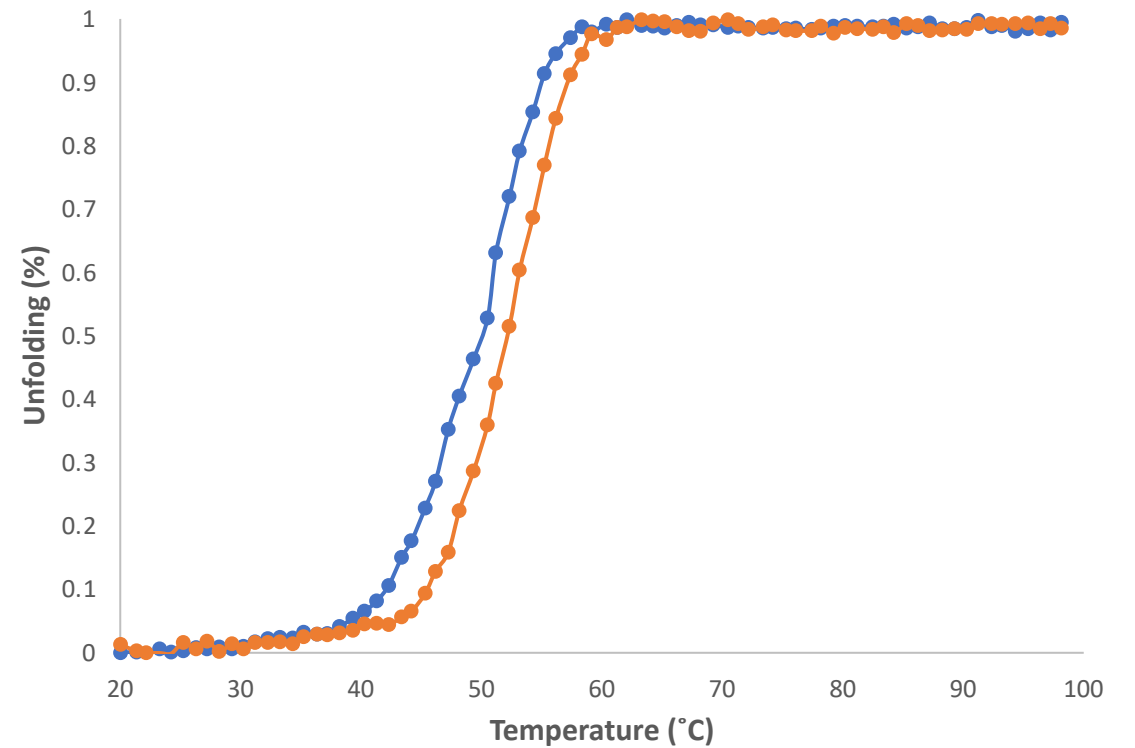
Gai-GTP

$G\alpha i$ -GMPPCP Triple mutant does not completely abolish pH-dependent thermostability



● pH 6.0 ● pH 6.4 ● pH 6.8 ● pH 7.2 ● pH 7.6

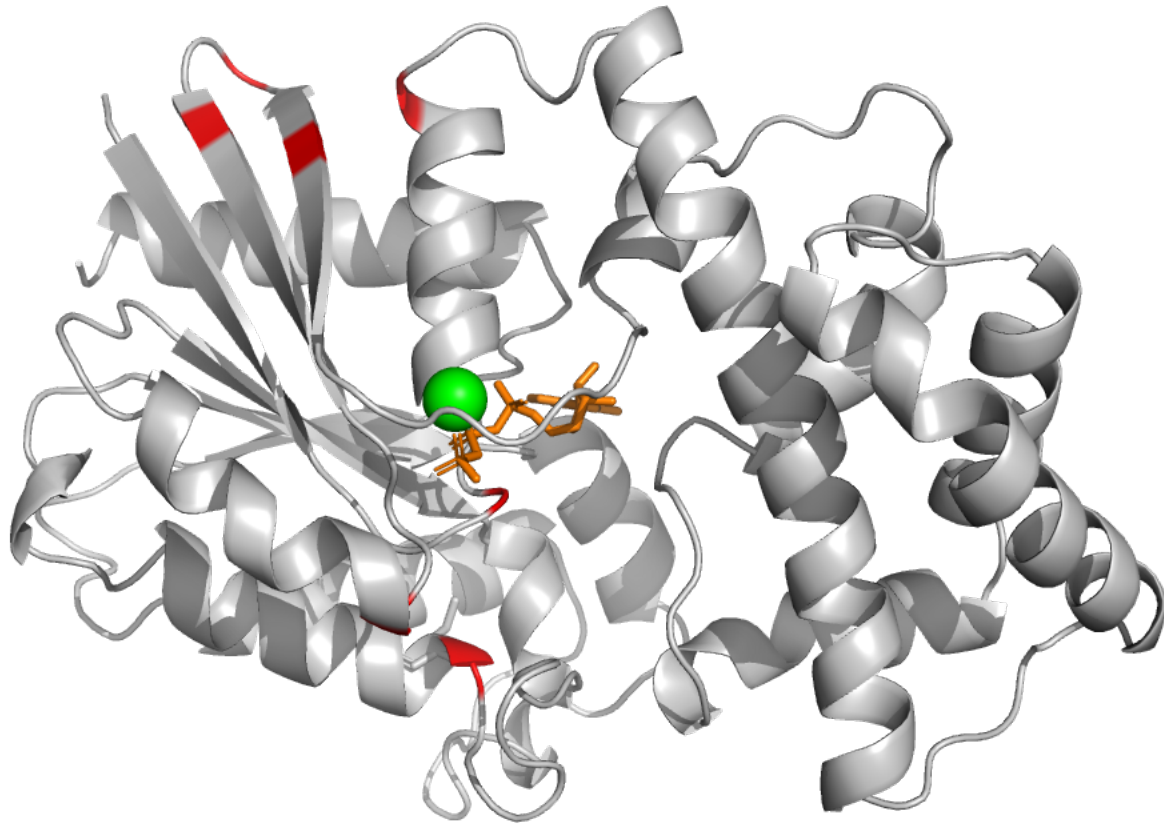
WT $G\alpha i$



● pH 6.4 ● pH 7.2

Triple mutant

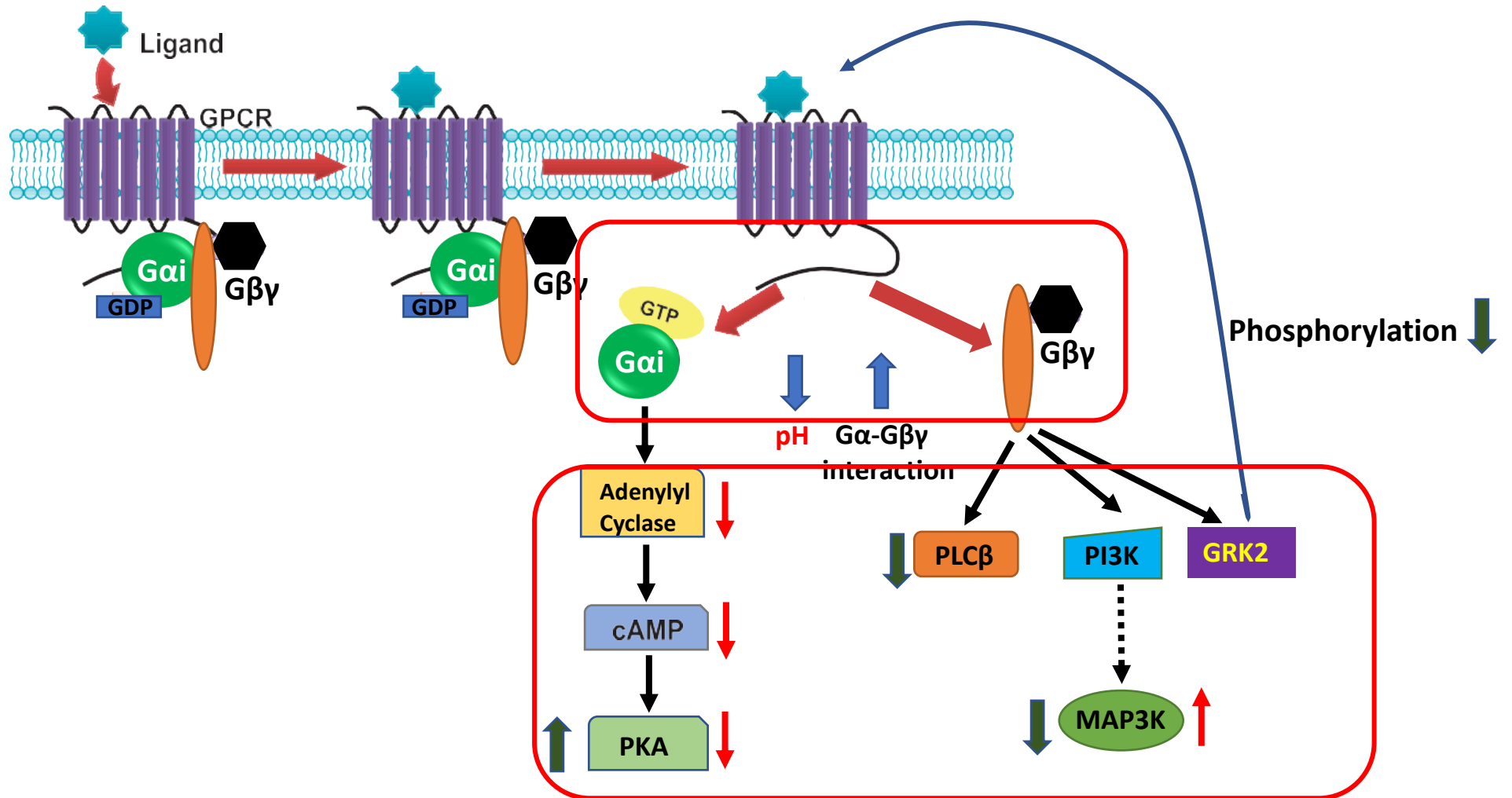
Some residues in the GDP release network may affect pH-dependent thermostability in G α i-GMPPCP



GDP release network residues

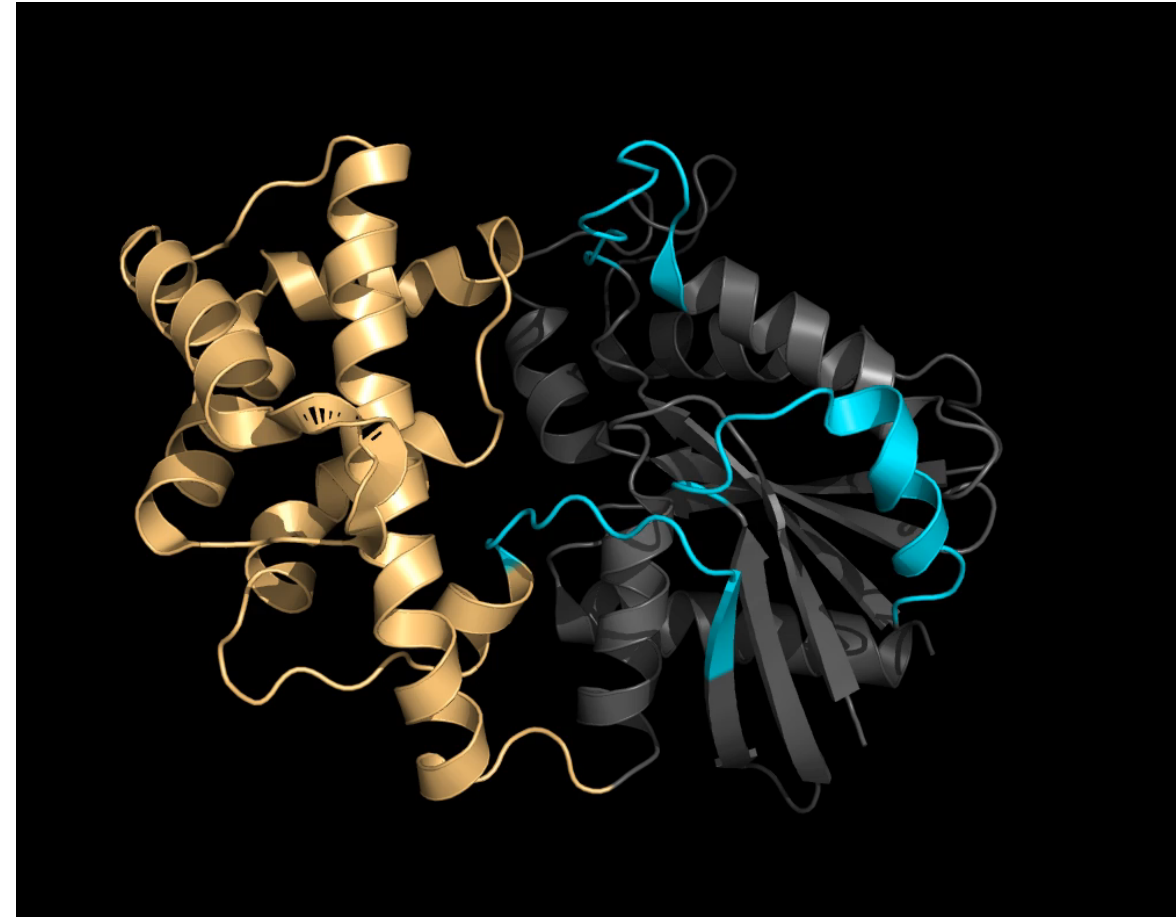
Residue	T _m (pH 6.4)	T _m (pH 7.2)	Δ
WT	59.5	64.5	5
E43	60	64	4
H57	59	64	5
H188	60	66.5	6.5
K192	56	63	7
H195	59	64.5	5.5
R205	54.5	63	8.5
R242	56.5	65.5	9
TM	50	52	2

Future Perspective



Conclusion: $G\alpha$ is an intracellular pH sensor

- Thermal stability of $G\alpha_i$ is pH-dependent in both active and inactive forms
- Low pH alters $G\alpha$ structure and destabilizes $G\alpha$ demonstrated by lower overall T_m
- Mutant $G\alpha$ acts as low-pH mimetic suggesting the identification of key pH-sensing residues
- Active $G\alpha$ may have additional pH-sensing network



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Thank you!