



Revealing SNAI2 as a Pseudo-Primed Substrate of GSK3

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Abstract

❖ SNAI2, a labile protein, functions as a **pseudo-primed substrate**, that is **phosphorylated by glycogen synthase kinase 3 (GSK3)** and **targeted to degradation**

Introduction

• **EMT (Epithelial-Mesenchymal Transition)** is **crucial** in various biological processes, including **neural crest migration**, **wound healing**, **proliferation**, and notably, **cancer metastasis**^{1,2}

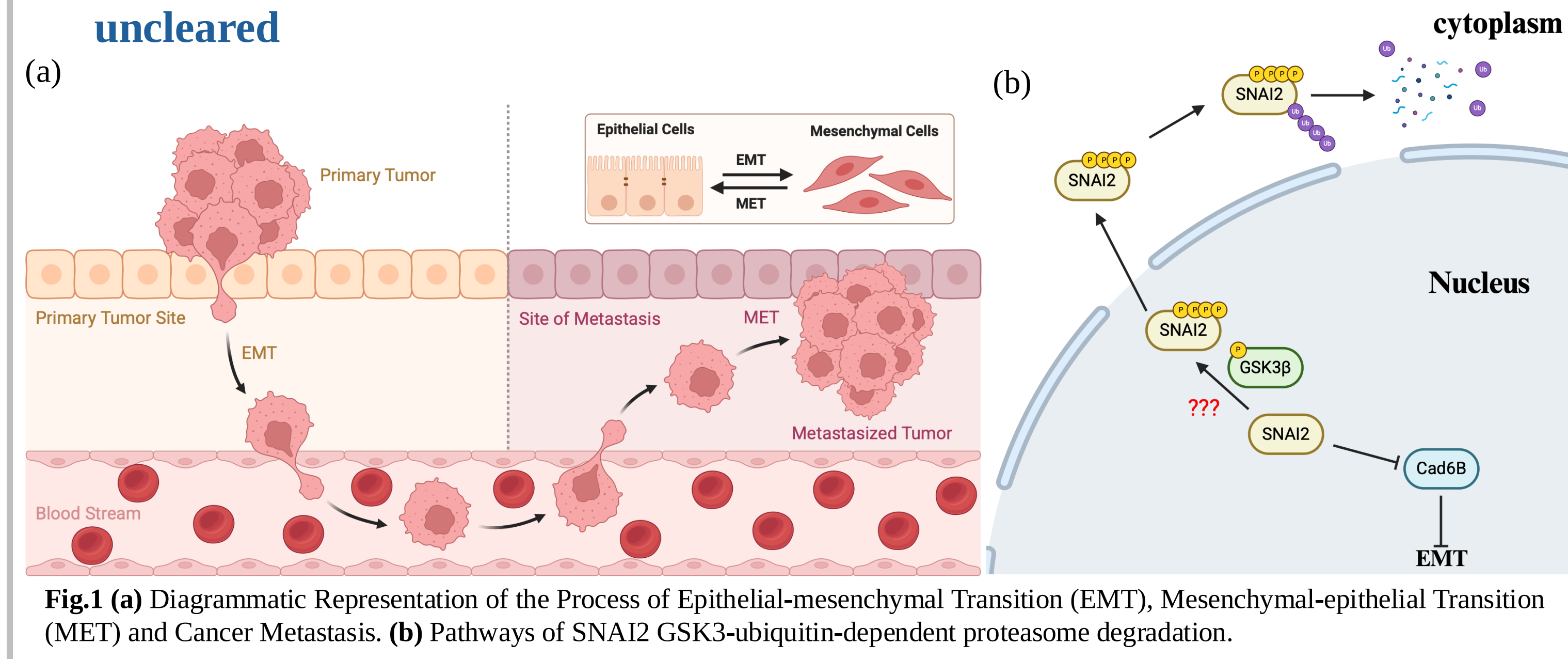
• SNAI2 is an **EMT transcription factor (TF)** with a **repressor activity**³

• SNAI2 is crucial for the **initiation of neural crest EMT** by **repressing cadherin6B**³
➢ **Neural crest cells** undergo **EMT** during **delamination** and **migrate** to various regions of the body to **differentiate** into **diverse cell types**⁴
➢ The **EMT** in **cancer cells** that **facilitates metastasis** also **aligns** with the **gene-regulatory module of neural crest EMT**⁵⁻⁶

• SNAI2 can be phosphorylated by GSK3 (Glycogen synthase kinase-3)
➢ Promotes SNAI2 to undergo **ubiquitin-dependent proteasomal degradation**⁷
➢ **Important regulation** of SNAI2 expression levels during EMT⁷

• **GSK3 substrates** can be either **primed (pre-phosphorylated)** or **non-primed (non-phosphorylated)**⁸

• However, the **mechanism** by which **SNAI2 is recognized by GSK3** is still **uncleared**

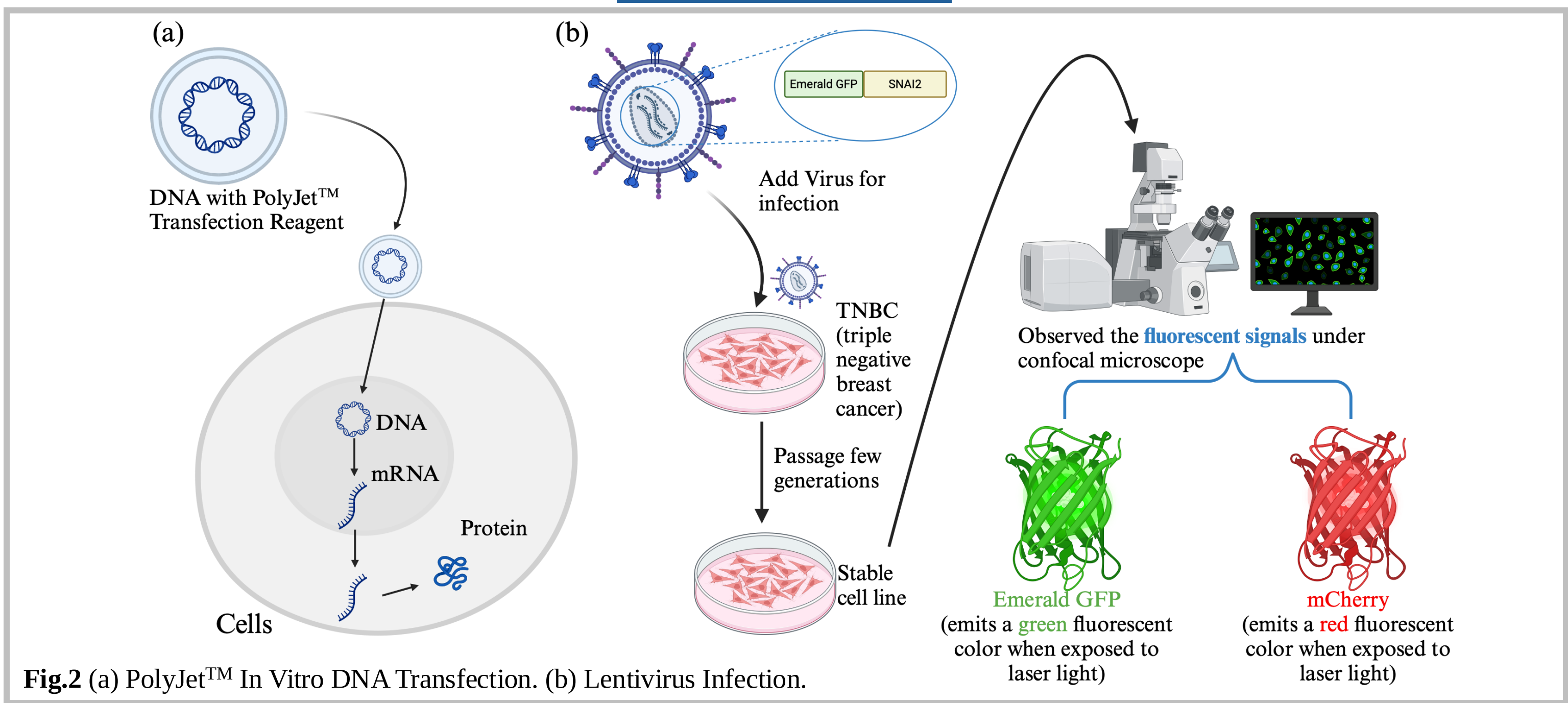


Objectives

❖ Elucidate the role of a **negatively charged region C-terminal** to the GSK3 phosphodegron to **mediate the phosphorylation and degradation** of SNAI2 by GSK3

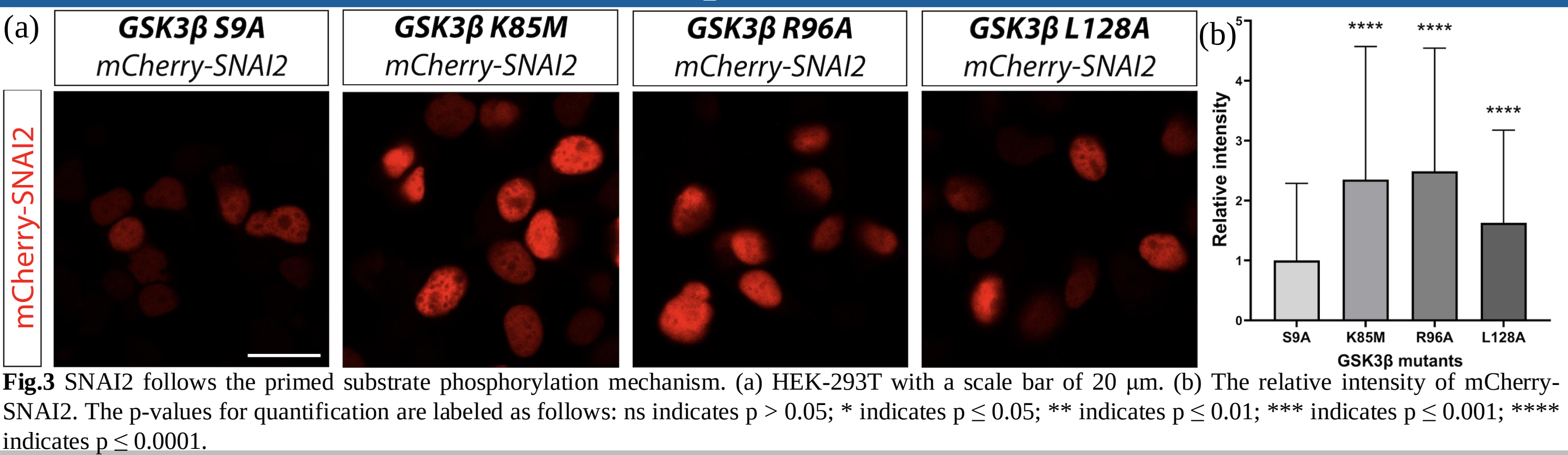
❖ Generating **constructs encoding SNAI2 mutants** fused with **EGFP (emerald)**

Methods



Results

1. SNAI2 acts as a primed substrate of GSK3

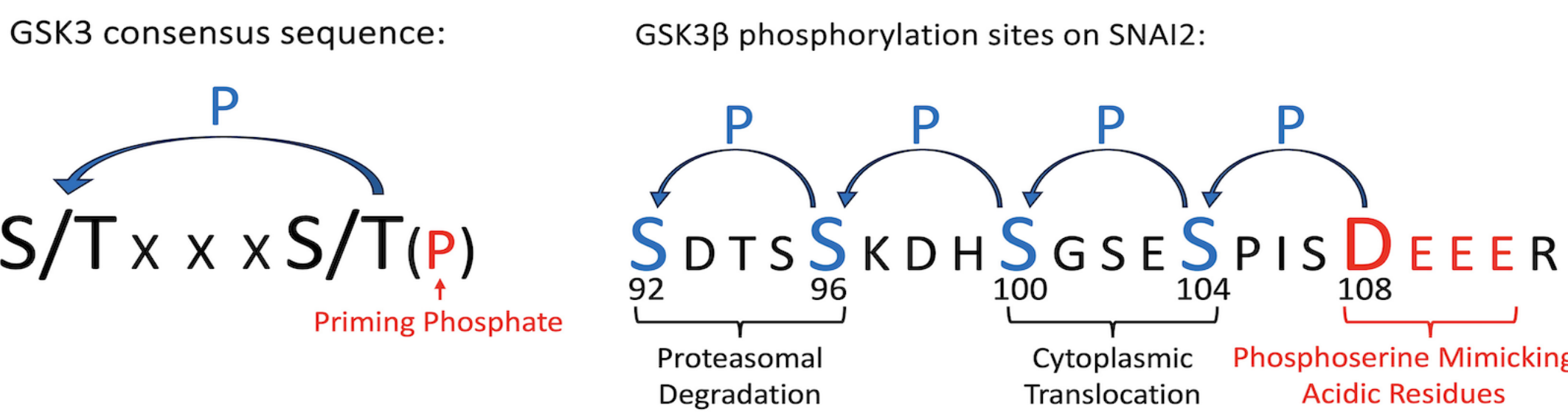


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• Although SNAI2 follows the **primed substrate phosphorylation mechanism**, it **does not contain any primed (pre-phosphorylated) sites** for GSK3 to **recognize**

2. Negatively charged DEEE amino acid sequence is found in SNAI2



Conservation of phosphoserine mimicking acidic residues and phosphodegron on SNAI2:

	100	110	120
Consensus	PXSDTSSKDHS	SGSESPIS	DEEERLQXKLS
Homo sapiens	PPSDTSSKDHS	SGSESPIS	DEEERLQSKLS
Gallus gallus	PPSDTSSKDHS	SGSESPIS	DEEERIQSKLS
Mus musculus	PSDTSSKDHS	SGSESPIS	DEEERLQPKLS
Xenopus tropicalis	PQSDTSSKDHS	SGSESPIS	DEEERLQTKLS
Rattus norvegicus	PSSDTSSKDHS	SGSESPIS	DEEERLQPKLS

Fig.4 Negatively charged DEEE amino acid sequences in SNAI2 across different vertebrates and the proposed mechanism of GSK3β-dependent phosphorylation on SNAI2.

• However, it **remains unknown** whether **DEEE** on SNAI2 functions as a **pseudo-priming site** for GSK3 **phosphorylation** and **degradation**

3. Generating SNAI2 mutants and EGFP (emerald)-SNAI2 fusion protein

(a) SNAI2 phosphodegron⁸⁰
Wild Type (WT): PSSSLGRVSPPPPDTSSKDHS⁹²SGSESPIS¹⁰⁰DEEERIQSKLS¹⁰⁸
4A: PSSSLGRVSPPPADTSSKDHS⁹²AGSEAPIS¹⁰⁰DEEERIQSKLS¹⁰⁸
D108A: PSSSLGRVSPPPPDTSSKDHS⁹²SGSESPIS¹⁰⁰AEEERIQSKLS¹⁰⁸
D108E: PSSSLGRVSPPPPDTSSKDHS⁹²SGSESPIS¹⁰⁰EEERIQSKLS¹⁰⁸
ΔDEEE: PSSSLGRVSPPPPDTSSKDHS⁹²SGSESPIS¹⁰⁰RIRIQSKLSDPHAI¹⁰⁸
DEEE/A: PSSSLGRVSPPPPDTSSKDHS⁹²SGSESPIS¹⁰⁰AAARIQSKLS¹⁰⁸

(b) N- Emerald-GFP SNAI2 C-

Fig.5 (a) The protein sequences of different SNAI2 mutants. (b) The Emerald-SNAI2 fusion protein.

4. The DEEE site functions as the pseudo-priming site on SNAI2

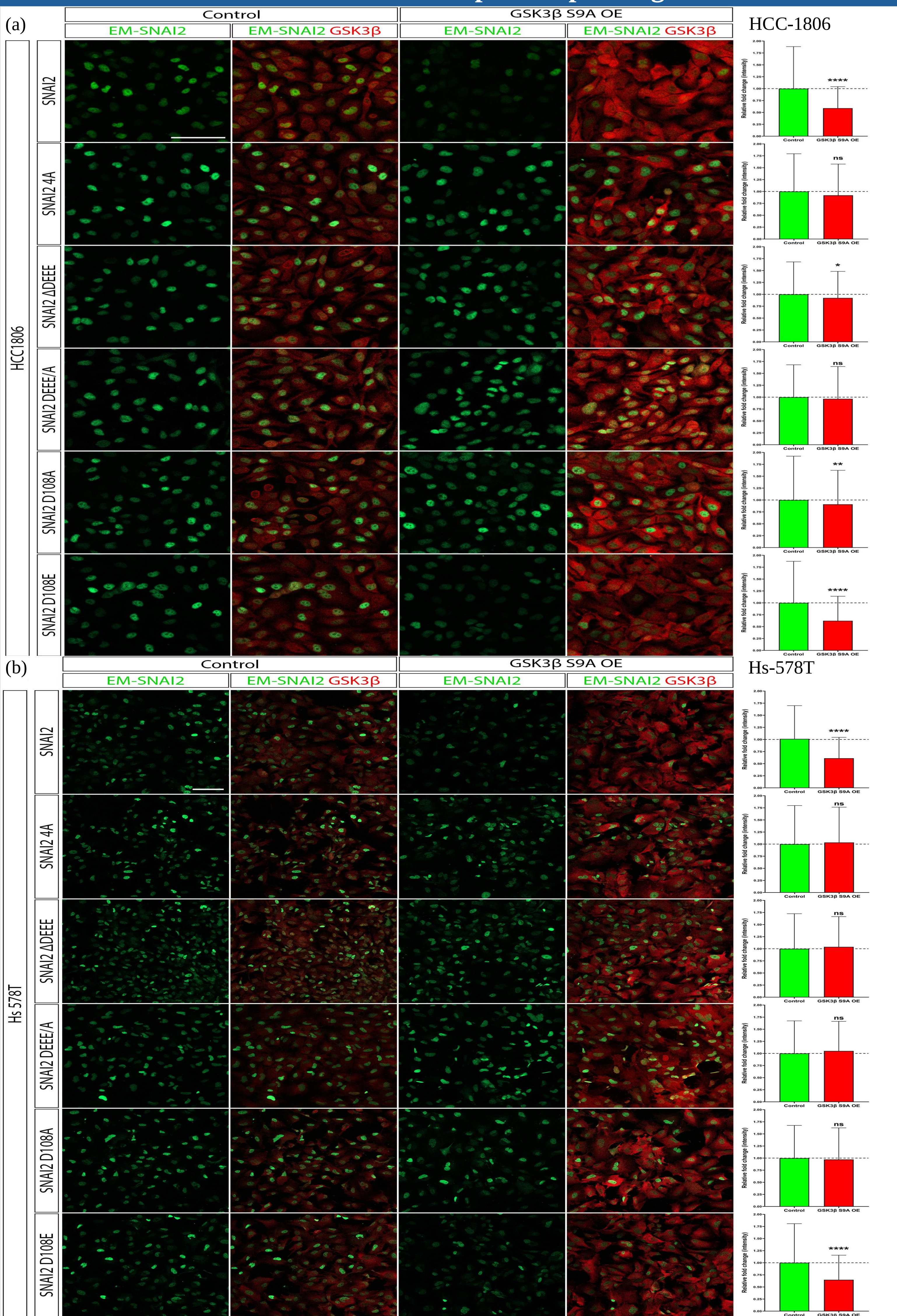


Fig.6 The DEEE site functions as the pseudo-priming site on SNAI2. The p-values for quantification are labeled as follows: ns indicates $p > 0.05$; * indicates $p \leq 0.05$; ** indicates $p \leq 0.01$; *** indicates $p \leq 0.001$; **** indicates $p \leq 0.0001$. (a) HCC-1806 with a scale bar of 50 μ m. (b) Hs-578T with a scale bar of 100 μ m.

Discussion

• The **DEEE sites** can serve as the **pseudo-priming site** on SNAI2 for GSK3 to **recognize**

• **Pre-phosphorylation** by another kinase, including other components such as **CK1**, in the **Wnt complex** is **not necessary** for GSK3 to bind to SNAI2

Future Direction:

• The experiment will be **repeated** in the **chick cranial neural crest cells** for further **validation**