

# Fertilization triggers early proteomic symmetry breaking in mammalian embryos

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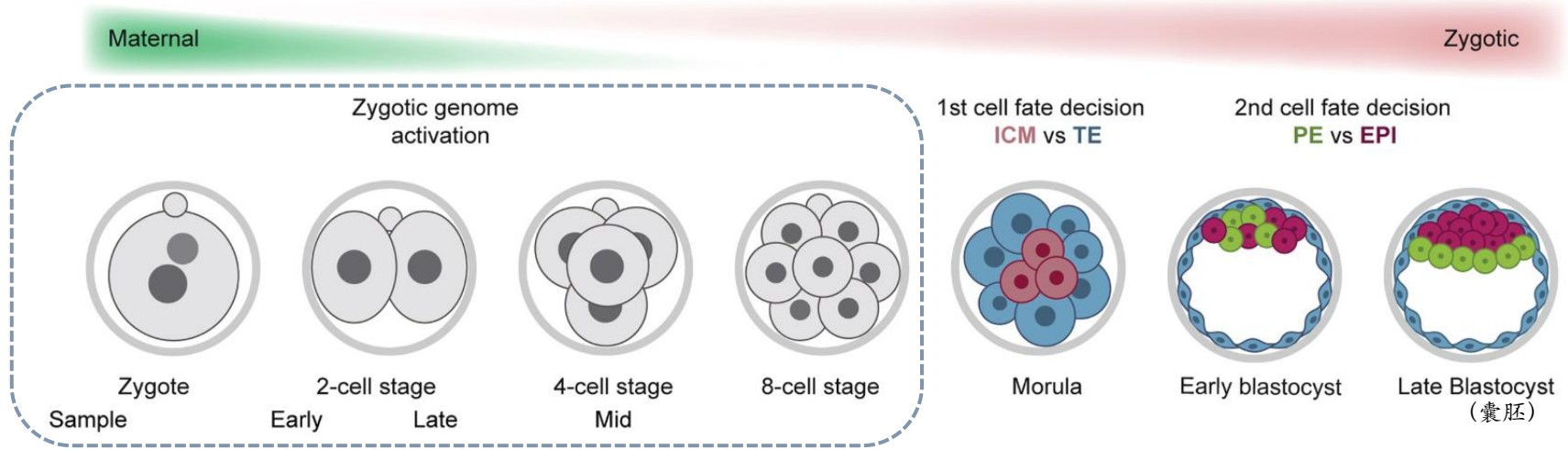
**Reporter:** Jingrui Lu

**Date:** 2026.3.6



# Introduction

# Blastocyst formation



**Do all blastomeres remain developmentally equivalent?**

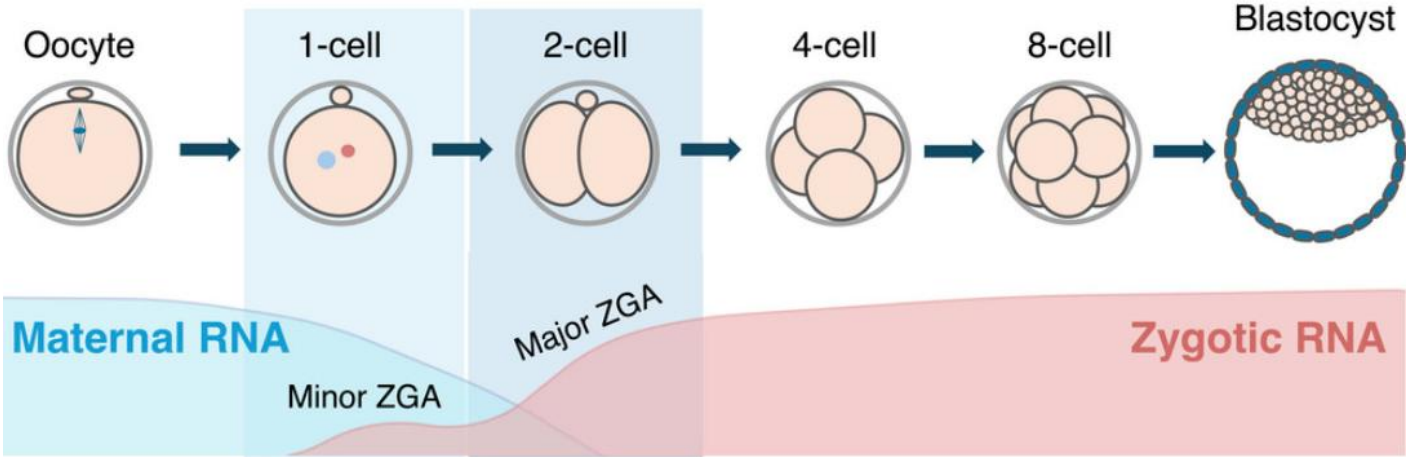
## 1st decision

Outer cells differentiate into **trophoblast** (TE);  
Inner cells form the **inner cell mass** (ICM)

## 2nd decision

ICM subdivides into **epiblast** (EPI) and **primitive endoderm** (PE)

# Zygotic genome activation (ZGA)



Solberg *et al.*, 2024

Rely on **maternal** ribosomes, mRNAs and proteins

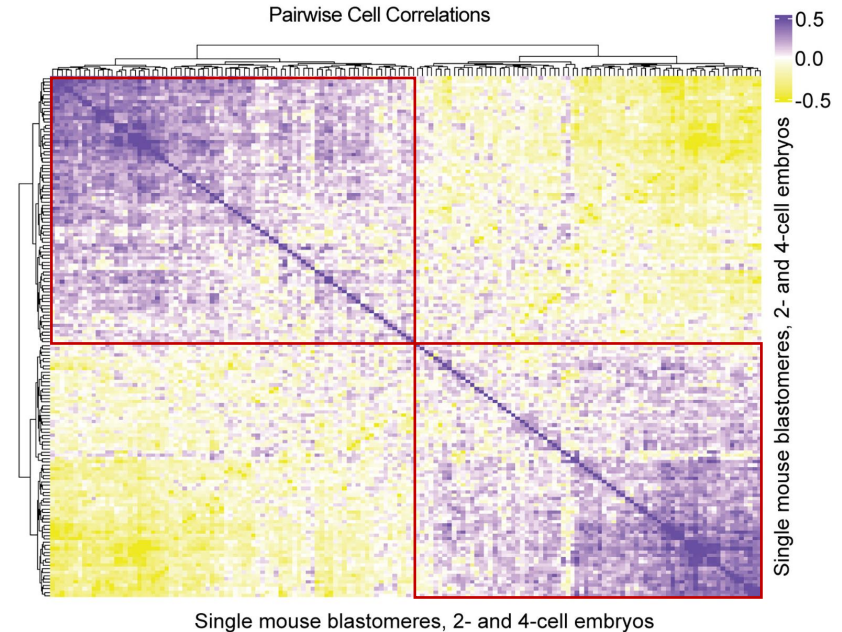
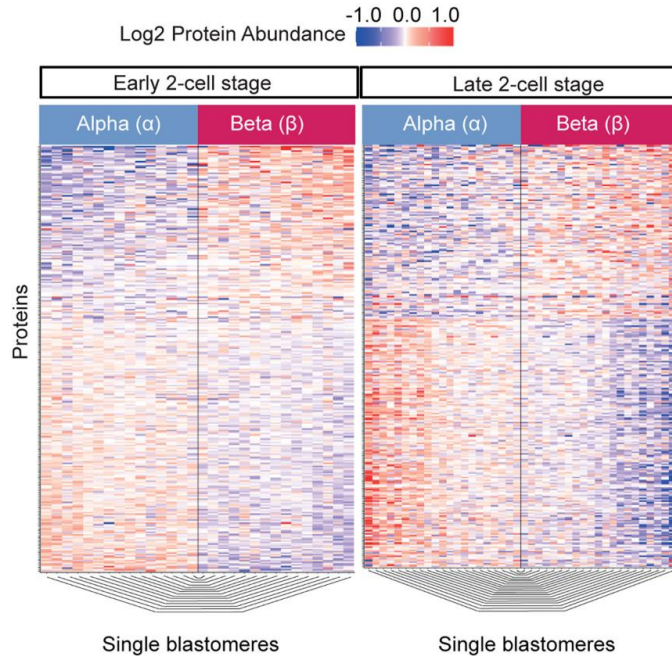


Activate and transcribe its **own genome**

**Proteomics change**

- **Maternal proteins** are actively degraded
- **Zygotic proteins** are actively produced

# Proteomic asymmetry in blastomeres of 2-cell- and 4-cell-stage embryos

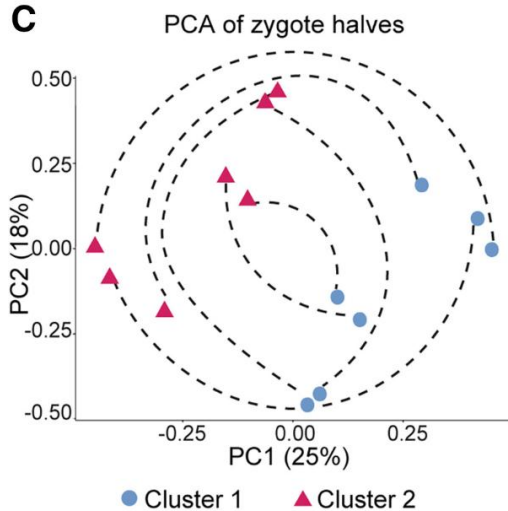
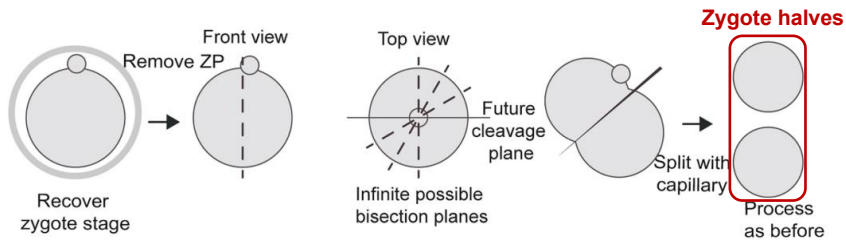


- In both early and late 2-cell stage, each embryo contains two types of blastomere (alpha and beta).

- Proteomic asymmetry persisted across 4-cell embryos.

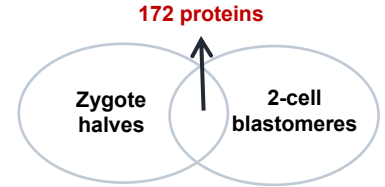
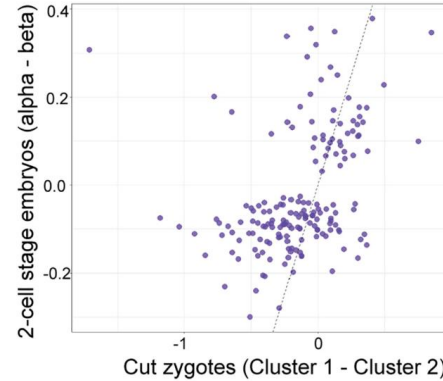
Do such asymmetries originate earlier?

# Proteomic asymmetry at zygote stage

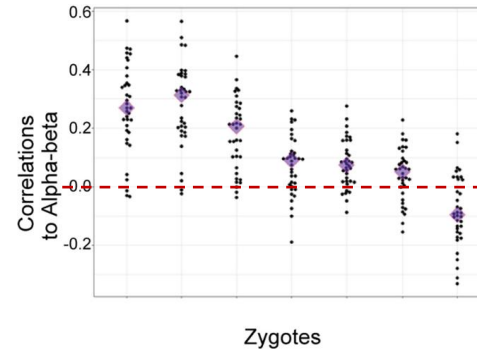


➤ Proteomic asymmetry at the zygote stage.

## Correlation of protein fold change



Cluster 1 ➡ Alpha-like  
Cluster 2 ➡ Beta-like



➤ Most zygote halves show alpha-beta-like differences.



How alpha and beta differences evolve over time?

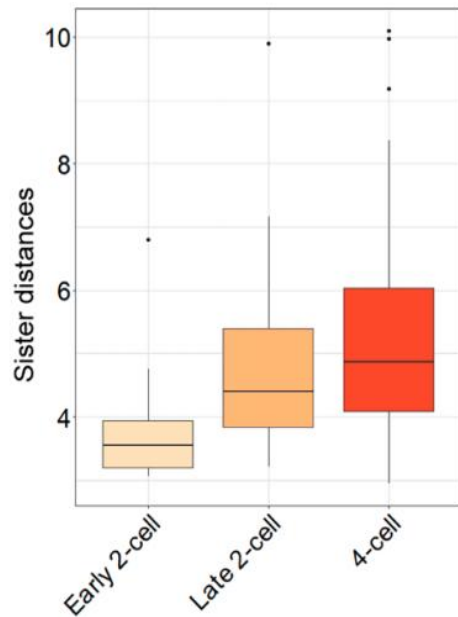
# Dynamics of alpha and beta differences

Alpha identity    Beta identity

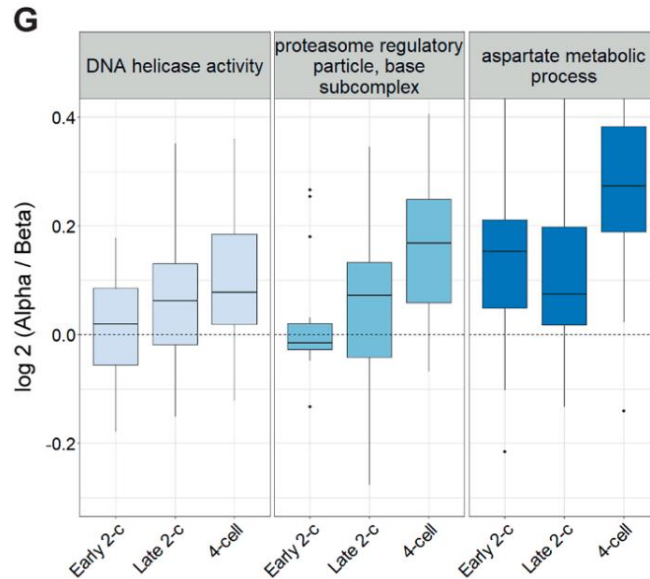
Psmc4

Gps1  
Nedd8

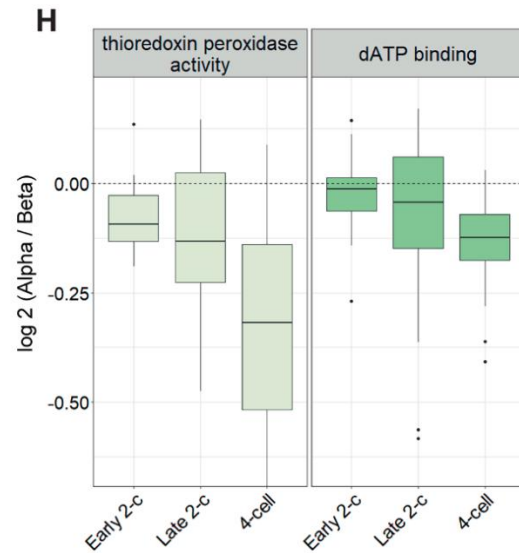
## Proteomics difference



## Alpha identity

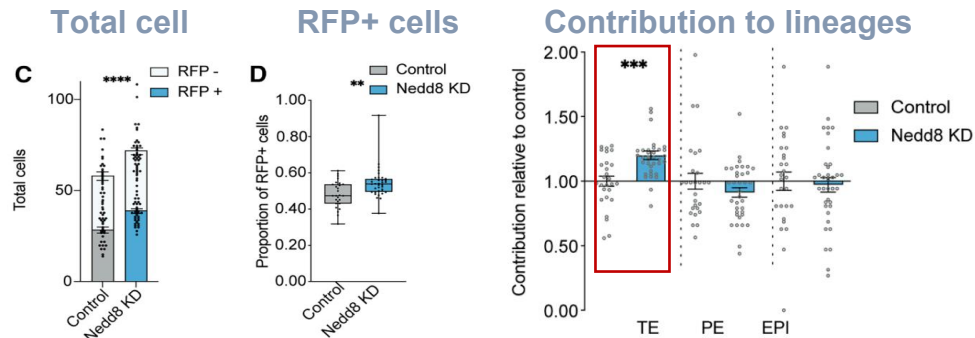
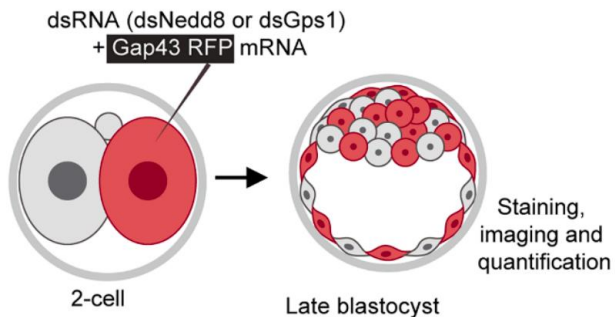


## Beta identity



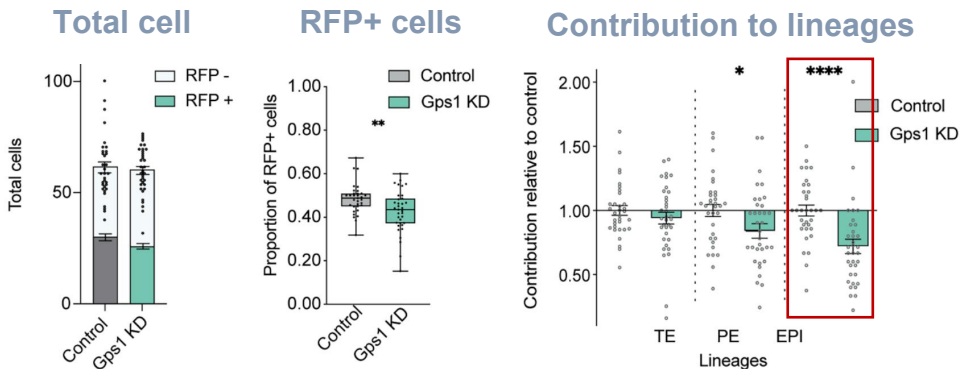
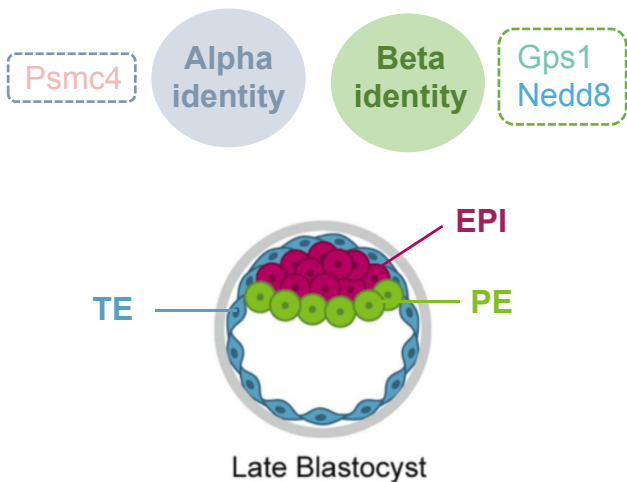
# The role of beta-enriched proteins in lineage fate

## Nedd8



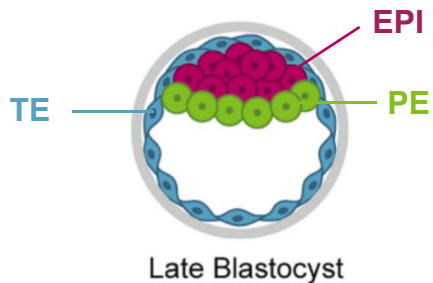
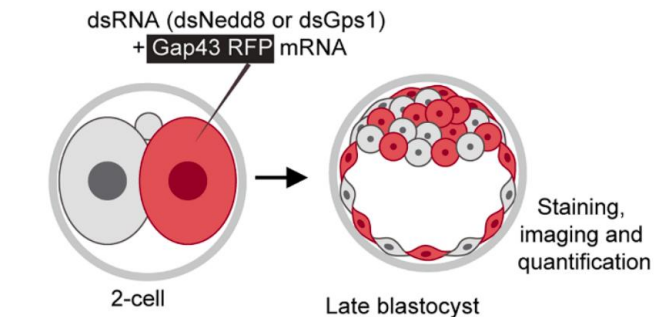
➤ **Nedd8 restricts proliferation and TE differentiation**

## Gps1



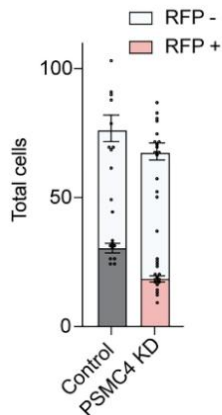
➤ **GPS1 promotes proliferation and EPI differentiation**

# The role of alpha-enriched proteins in lineage fate

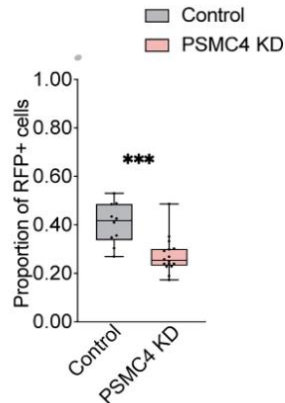


## Psmc4

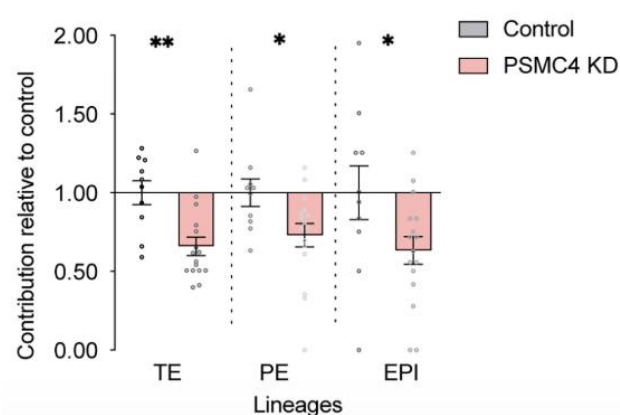
### Total cell



### RFP+ cells

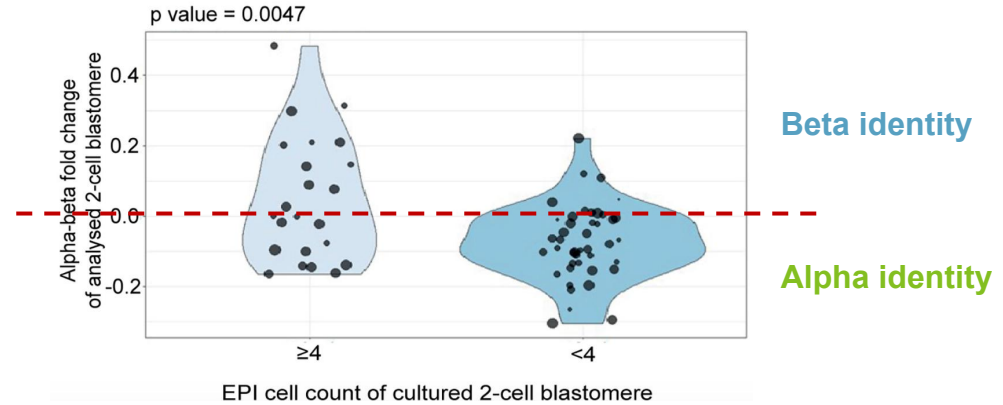
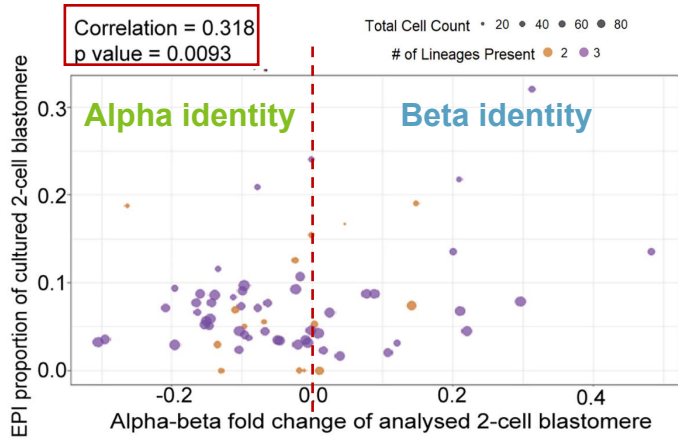
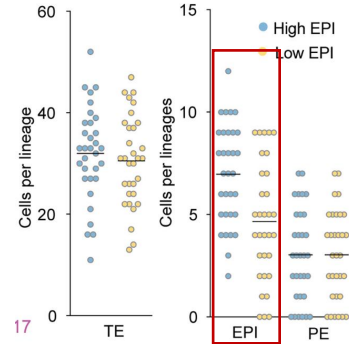
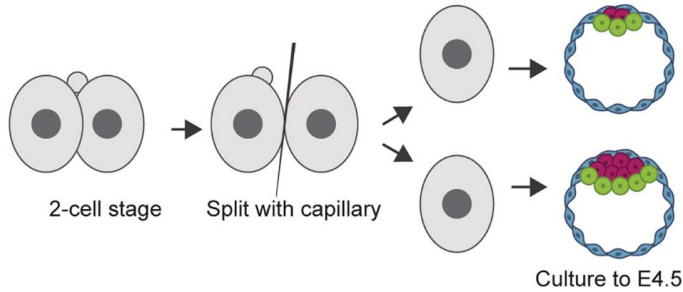


### Contribution to lineages



➤ Psmc4 promotes **proliferation** and support **all lineage** specialization

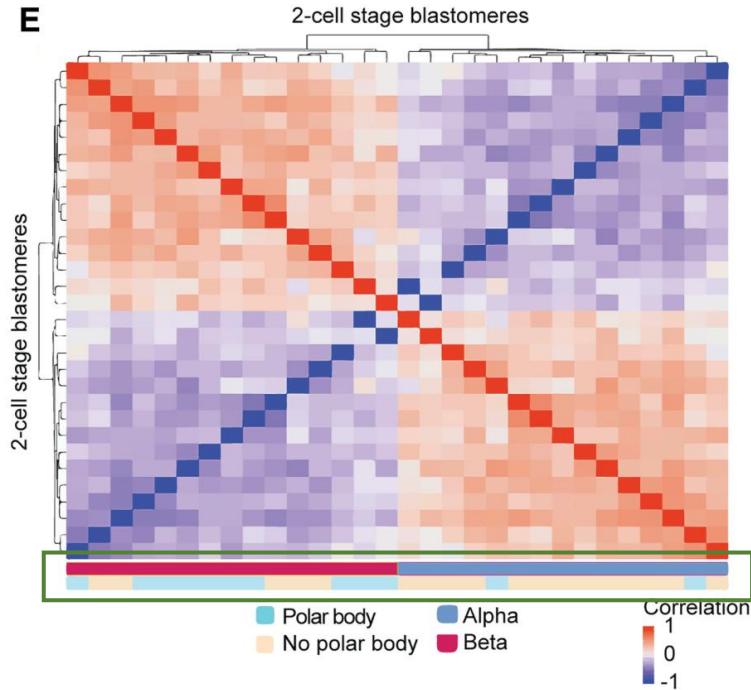
# Different developmental potential of alpha and beta blastomeres



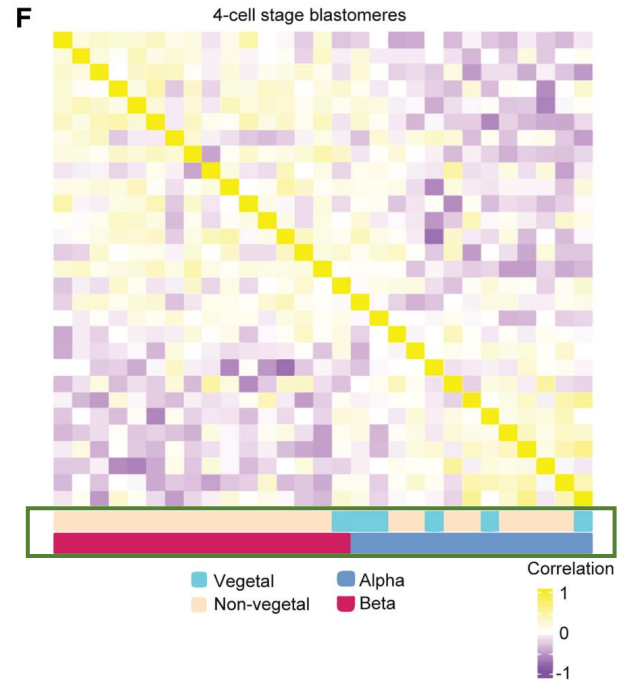
Beta blastomeres → Higher proportion of EPI  
Alpha blastomeres → Lower proportion of EPI

➤ Beta blastomeres promote epiblast formation

# Different developmental potential of alpha and beta blastomeres



- Blastomere **inheriting the second polar body** was more likely to be classified as **beta**

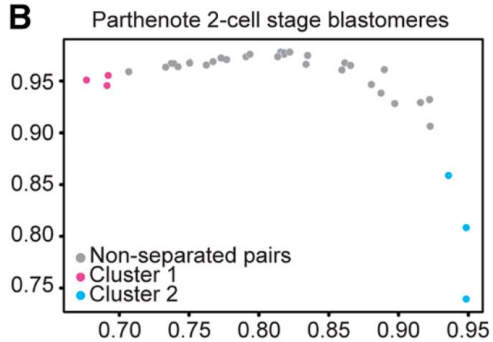


- **Vegetal blastomeres** were significantly more likely to be classified as **alpha**

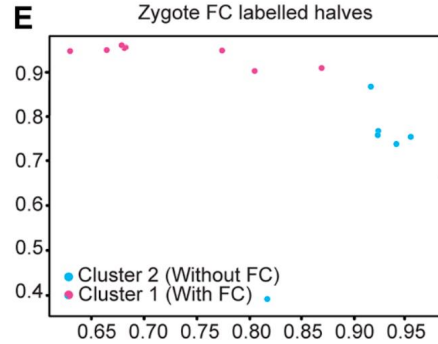
How to trigger such proteomics asymmetry?

# Fertilization triggers the breaking of symmetry

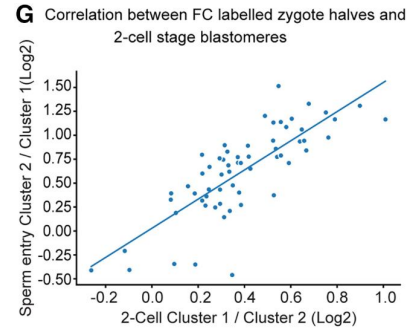
## Parthenogenetic embryos (孤雌胚胎)



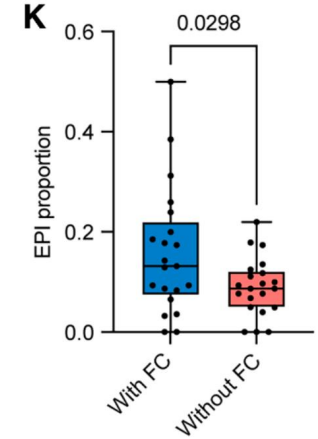
## Fertilized embryos



FC: Fertilization cone (受精锥)



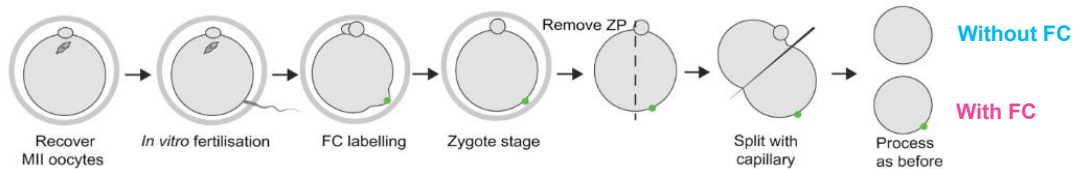
## Lineage development



➤ **Fertilization triggers proteome asymmetry in the zygote.**

➤ **Fertilization leads to higher EPI proportion.**

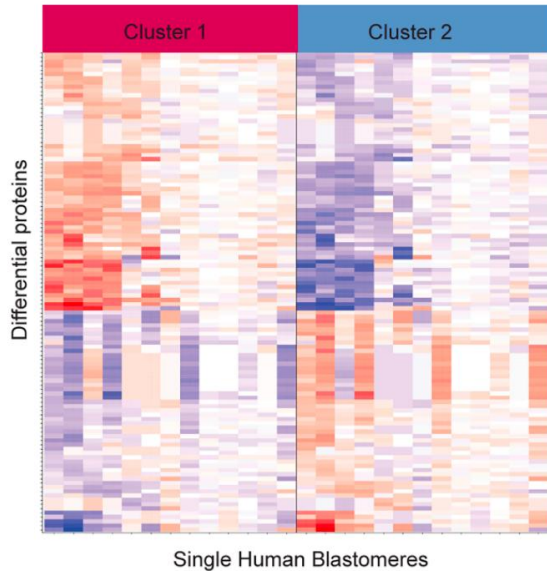
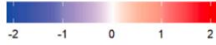
## Fertilized embryos processing procedure



Whether the proteomic asymmetry is **conserved** in **human** embryos?

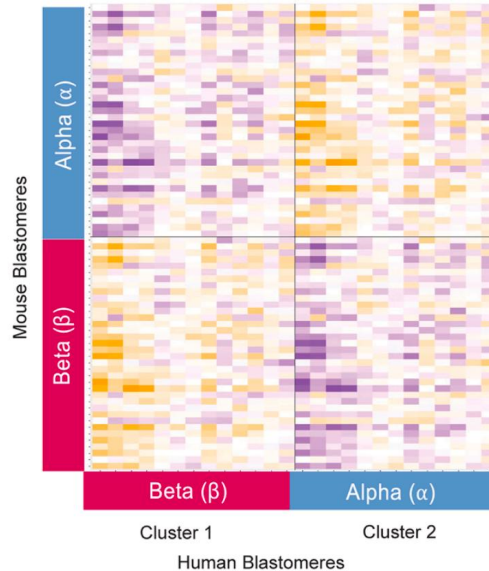
# Proteome asymmetry is conserved in human 2-cell\_x0002\_stage embryos

C Log2 Protein Abundance

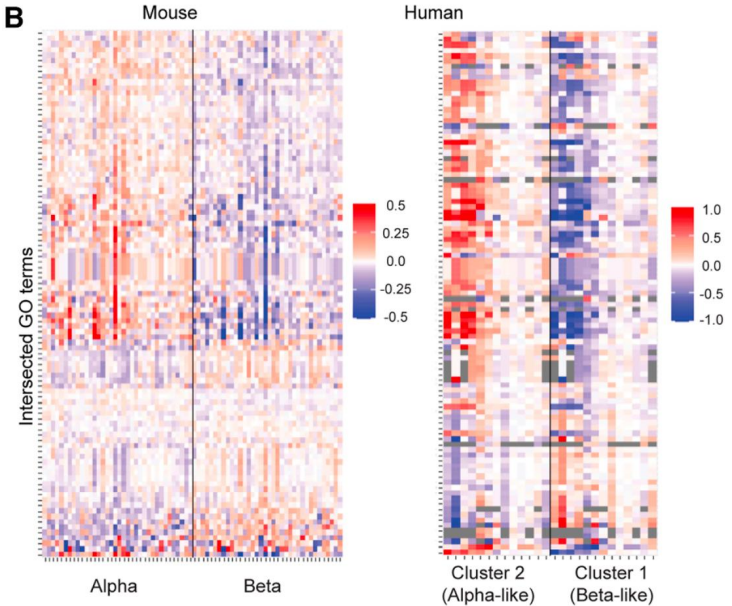


## Cross-species annotation

A Correlations based on intersected proteins



B



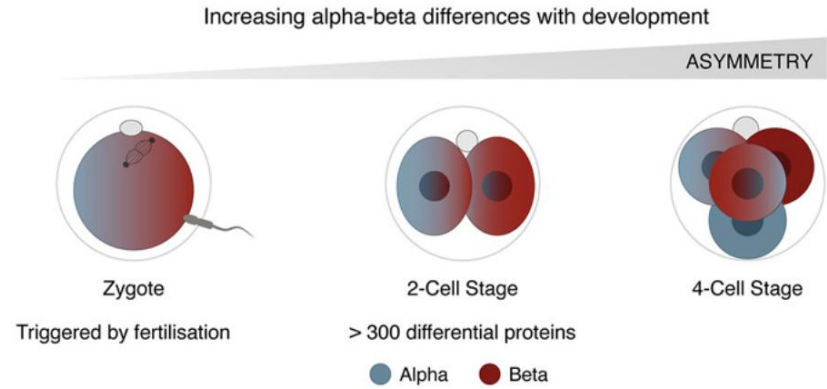
➤ **Strong directional concordance** between mouse and human.



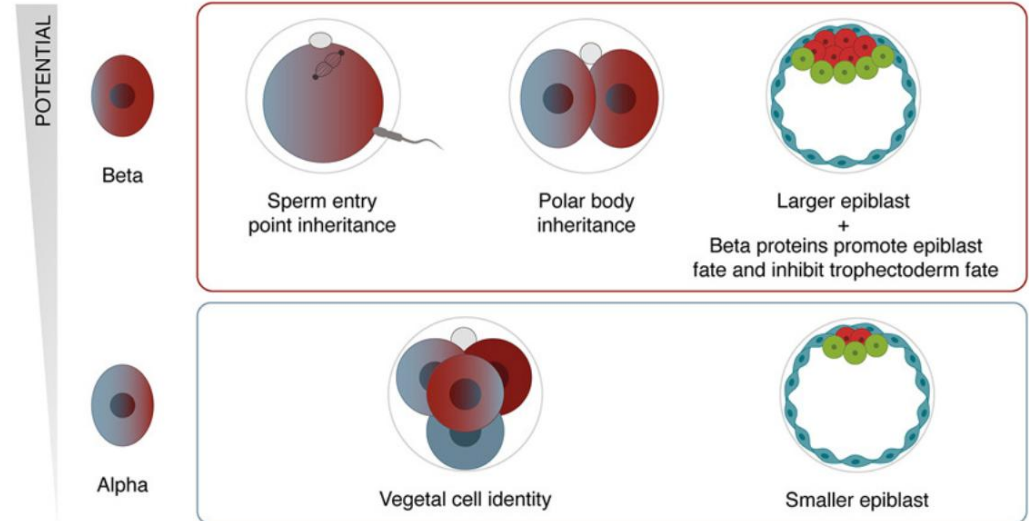
# Summary

# Summary

- Single-cell proteomics reveals early **proteome asymmetry** in human and mouse.



## ALPHA-BETA ASYMMETRY IS LINKED WITH DEVELOPMENTAL POTENTIAL





**Thanks for your listening!**

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