

Identifying a Gene Orchestrating Skin Regeneration via Tissue Rebuilding ~ Inspiration from Aesthetic Treatments ~

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Introduction



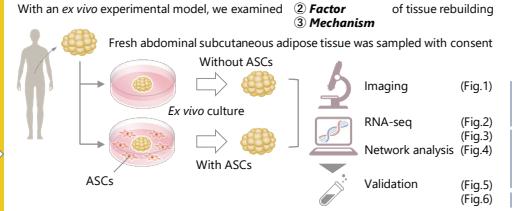
Hypothesis

Subcutaneous injection of adipose-derived stem cells (ASCs) occur biological responses that trigger tissue remodeling

Aim

Clarify "Rebuilding Induction"
- What key factor induces rebuilding?
- What is the mechanism?

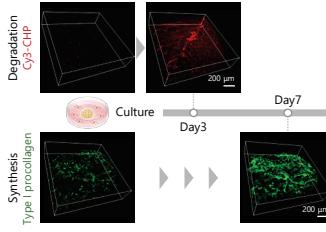
Study flow



Result

① Phenomenon: In an ex vivo co-cultured experimental model, not only rebuilding but also wound healing was observed

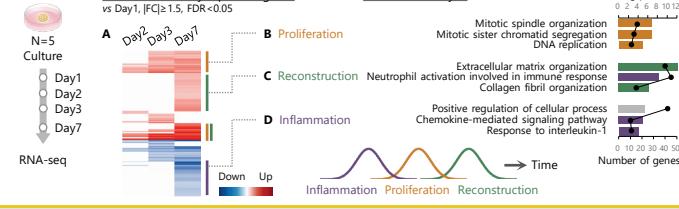
1. Rebuilding, sequential collagen degradation and synthesis, occurs in the presence of ASCs



2. Alongside rebuilding, wound healing processes, inflammation, proliferation and reconstruction, occurred

Purpose: To understand the basic phenomena that occur in the ex vivo subcutaneous adipose tissue

Method: RNA-seq, analysis of changes in differentially expressed genes compared to baseline (Day1) for Days2, 3, and 7

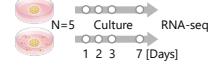


② Factor: TSG-6 was identified as a key factor in triggering tissue rebuilding instead of scarring

3. ASCs mediate suppression of pro-fibrotic signals and early termination of inflammation

Purpose:
To understand the influence of ASCs co-culture in the ex vivo subcutaneous adipose tissue

Method: RNA-seq, analysis of changes in differentially expressed genes compared to ASC(-) for each culture day



4. TSG-6 was selected as a key paracrine factor involved in rebuilding based on the high degree of centrality in the network analysis

Purpose: To identify the genes involved in tissue rebuilding

Methods: A protein-protein interaction network was generated

- A protein from differentially expressed gene analysis, ASC(+) vs ASC(-) at day1
- A protein directly interacting with red/blue from the database
- Direct interaction

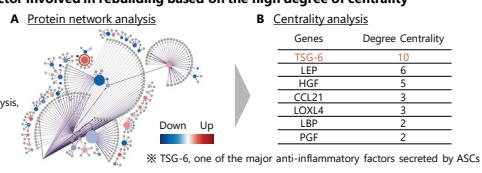
5. TSG-6 released by ASCs mediates the effects of ASCs

Purpose: To reveal the role of TSG-6 in inducing ASC's biological responses

Method: After treatment with an TSG-6 neutralizing antibody, either qPCR or imaging was performed.

(A) qPCR for α-SMA, a marker of myofibroblast. (B,C) Imaging analysis of (B) degradation and (C) synthesis of fibrous structures.

N=4, Mean ± SE, Tukey HSD test, *p < 0.05, **p < 0.01

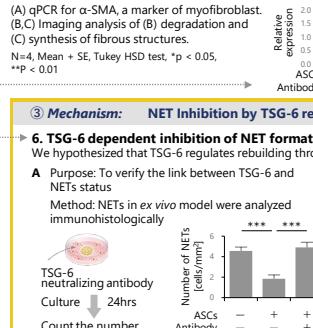


6. TSG-6-dependent inhibition of NET formation suppresses fibrosis

We hypothesized that TSG-6 regulates rebuilding through neutrophil extracellular traps (NETs).

A Purpose: To verify the link between TSG-6 and NETs status

Method: NETs in ex vivo model were analyzed immunohistologically



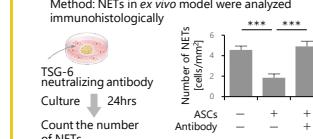
③ Mechanism: NET Inhibition by TSG-6 reduces fibrosis, shifts to rebuilding

7. TSG-6-dependent inhibition of NET formation suppresses fibrosis

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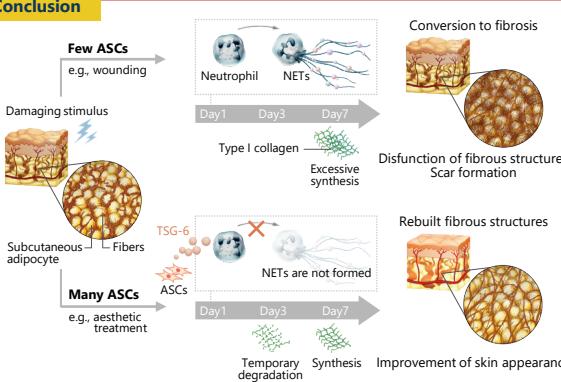
Method: NETs in ex vivo model were analyzed immunohistologically



B Purpose: To investigate the effects of NETs on the fibrous structures

Method: By qPCR, gene expression changes in subcutaneous adipose tissue exposed to NETs were examined

Conclusion



Impact

Transformation of approach methods:

- Current paradigm
Elucidation of the causes of skin problems

↓ "Stop the aging process"

"Restore the skin to its younger state"

- New paradigm
Elucidation of the mechanisms underlying skin improvement

↓

"Cosmetics"

to enhance efficacy
to expand applications for fibrotic skin problems
- Aesthetic treatments
- Medical fields
to reduce downtime/to maximize efficacy
to reduce scarring

In the future: "Rebuilding Induction" by ASC injection in a bottle

