Addressing skin firmness through DNA methylation

ADEBIOTECH - Romainville | March 13, 2018 Reymermier Corinne - Debret Romain



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The fibroblast submitted to the proof of time and lifestyle



Our Challenge: Wake up our skin cells to reactivate the optimum fibroblast machinery



Epigenetic mechanism through promoter methylation





Relationship DNMT3A / LOXL1 promoter proven in aging



LOXL1 mRNA expression is decreased with aging due to a decrease of its promoter activity which is correlated with an increase of DNMT3A driven methylation **BASE** hare Dreations,

We create chemistry

Epigenetic regulation: Screening strategy **To reverse methylation and restore** *LOXL1* **expression**



Preselection in BBCS library

Cell viability and organoleptic assessment of ingredients

Q-PCR screening: Induction of gene expression of 4 genes related to ECM synthesis and organization (*col1a, loxl1, fbln5 and magp1*)

HRM-PCR selection: Decrease of lox/1 promoter methylation

Confirmation: Decrease of recombinant DNMT3A in transitory transfection system

Validation: Induction of LoxI1 and Col1 protein increase in ECM in aged fibroblasts

Origanum majorana extract (Dermagenist™) was selected



Dermagenist[™] reverses methylation and restores *LOXL1* expression



Dermagenist[™] prevents the methylation of LOXL1 promoter driven by DNMT3 suggesting a rejuvenation effect of cell's epigenetic pattern □BASF

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Dermagenist[™] induces LOXL1 protein involved in collagen fibers organization



Dermagenist[™] reactivates COL I proteins to restore skin firmness



