DermaPep™ A440

Innovative Anti-Aging Tetrapeptide



Represented in the USA by TRI-K Industries, Inc.

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Anti-aging

DermaPep[™] A₃₅o

DermaPep™ A420

DermaPep[™] A440

DermaPep[™] A530

Whitening

DermaPep[™] A₃₅o

DermaPep[™] UL

DermaPep™ W220

DermaPep[™] W411

Anti-inflammatory

DermaPep[™] A₃₅o

DermaPep[™] A440

DermaPep[™] A530

DermaPep[™] UL

DermaPep™ W220

DermaPep™ W411

DermaPep™ A440 Anti-Aging Peptide

Introduction

Collagen represents the main component of the extracellular matrix of the dermal connective tissue and its concentration decreases as we age. It is known that the loss of collagen, especially in photoaged skin, is not only due to a reduction of the synthesis but also due to its enhanced degradation. The enzyme mainly responsible for collagen breakdown and inhibition of new collagen formation in the skin is MMP-1 which cleaves collagen type I, III, VII, VIII. Exposure of the skin to UV radiation results in the upregulation of several different MMPs and the repeated upregulation of these collagen-degrading enzymes over time is thought to underlie the collagen damage that is one of the hallmarks of photoaging.

In a healthy skin, the synthesis and degradation of the matrix are well in balance. However, this intricate balance gets disrupted as we age, in which too little of the matrix is synthesized and too much is degraded. Many of well-known skin aging treatments are aimed at replenishing skin matrix by stimulating the synthesis of the matrix. Unfortunately, this approach fails in some people, especially old people, whose ability to respond to matrix synthesis boosters is considerably declined.

DermaPepTM A440, innovative antiaging tetrapeptide, has been developed to maximally and effectively inhibit these matrix degrading MMPs and to restore collagenlevels back to normal. In vitro results show that DermaPepTM A440 greatly inhibited the expression and production of MMP-1 under both UV-stimulated and non-stimulated condition and restored collagen levels significantly. In addition, DermaPepTM A440 considerably lowered the MMP levels that have been produced by ROS or RNS. Finally, in vivo clinical study results prove that DermaPepTM A440 is surely one of the best antiaging agents in the market.

Function

- Anti-wrinkle effect for face and body
- Reduction of MMP1 synthesis
- Restoration of procollagen type 1
- Providing exceptional anti-aging benefits
- Anti-aging effect against environmental stimulus such as UV irradiation

Applications

DermaPep™ A44o can be incorporated in cosmetic formulations such as emulsions, oily sera, gels and creams for anti-aging and anti-wrinkle purpose.

Formulation

- Dermatological tolerance: Standard testing has been performed on DermaPep™ A440 which has showed neither cytotoxic effects nor any irritation or sensitization reaction in healthy volunteers with an occlusive single patch test.
- Recommended concentration: 1-3 %

Product Information

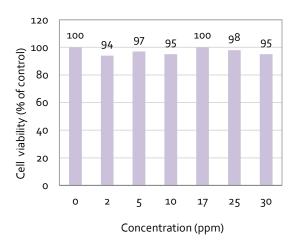
- Appearance : Transparent solution
- INCI/CTFA-Declaration : Myristoyl Tetrapeptide-34 (and) Butylene Glycol
- Active ingredient content
- : 0.15% Myristoyl Tetrapeptide-34
- Powder purity: 95 % up
- Preservative : None

DermaPep™ A440 Safety

Cell Cytotoxicity Test

Hs68 cells were seeded into 96-well plates (2 x 10³) and treated with different concentrations of Myristoyl Tetrapeptide-34 for 24 hours. Cell viability was assessed by MTT assay and was shown relative to untreated control. Absorbance was measured by an ELISA reader at 540 nm.

MTT Assay



Skin Irritation Test

Skin irritation test was performed on 31 healthy Asian volunteers (Spincontrol Asia (study report number IR-6Qo1-MW-DEo8), Bangkok, Thailand).

Evaluation and grading of skin irritation

- Erythema (no redness = 0 to very strong redness=5)
 After 24 hrs, mean score = 0.00 (max = 0)
 After 48 hrs, mean score = 0.00 (max = 0)
- Oedema (no oedema = o to very strong oedema = 5)
 After 24 hrs, mean score = 0.00 (max = 0)
 After 48 hrs, mean score = 0.00 (max = 0)
- Scaling (no scaling = 0 to very strong scaling = 5)
 After 24 hrs, mean score = 0.00 (max = 0)
 After 48 hrs, mean score = 0.00 (max = 0)

Mutagenicity (Ames Test)

Myristoyl tetrapeptide-34 at 5 different concentrations (30,60,90,180, 360 ppm) were evaluated to predict its potential to cause mutagenicity. Five different auxotrophic mutants were used to assesse directly or in the presence of liver S9 fractions.

No mutagenicity was observed.

In vitro Ocular Irritation Test

DermaPep[™] A440 at 3 different concentrations (2, 4, 8 %) and butylene glycol were evaluated with the ocular irritection assay system (CA, USA).

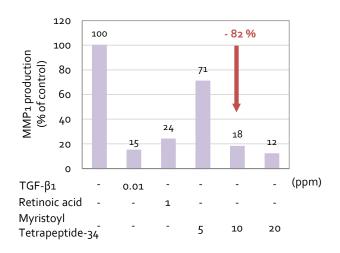
The ocular results demonstrated that the DermaPepTM A440 was classified as mild irritants.

Inhibition of MMP-1 Production

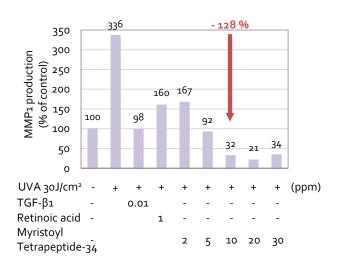
Hs68 cells were seeded into 6-well plates and treated with TGF- β , retinoic acid or Myristoyl Tetrapeptide-34 followed by UV radiation or not for 48 hours. Equal amounts of media were analyzed by ELISA assay for MMP-I. Cell viability was used for sample standardization.

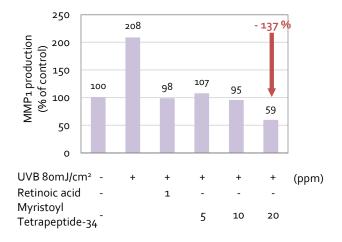
Myristoyl Tetrapeptide-34 significantly decreased MMP-1 production.

Inhibition of MMP-1 Production



Inhibition of MMP-1 production after UVA or UVB irradiation





Effect on Procollagen 1 Production

Hs68 cells were seeded into 6-well plates and treated with retinoic acid or Myristoyl Tetrapeptide-34 followed by 3oJ/cm² UVA. After 48 hours equal amounts of cell lysates were subjected to electrophoresis and analyzed by western blot for procollagen type I.

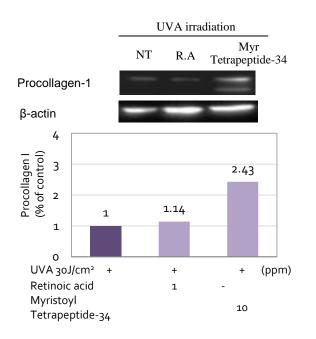
Myristoyl Tetrapeptide-34 significantly stimulated the production of type I procollagen.

Changes in MMP-1 and ECM gene expression

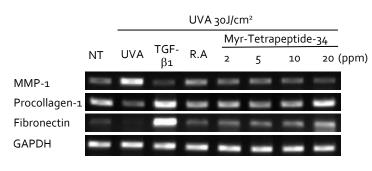
Hs68 cells were seeded into 6-well plates and treated with TGF- β , retinoic acid or Myristoyl terapeptide-34 followed by 30J/cm² UVA. After 48 hours, total cellular RNA was extracted and reverse-transcription PCR was performed to determine MMP-1, procollagen 1, fibronectin. GAPDH mRNA level was used for sample standardization.

Myristoyl Tetrapeptide-34 significantly inhibited UVA stimulated MMP-1 expression and stimulated the procollagen-1 and fibronectin expression.

Stimulation of Procollagen I Production



Changes in MMP-1 and ECM gene expression

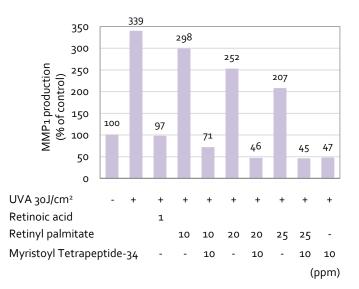


Comparison with Retinoids

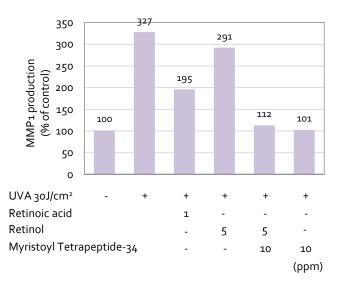
Hs68 cells were seeded into 6-well plates and then treated with retinyl palmitate, Myristoyl tetrapeptide-34 or both (left figure) or retinol, Myristoyl Tetrapeptide-34 or both(right figure) followed by 3oJ/cm² UVA . Equal amounts of media were analyzed by ELISA assay for MMP-I. Cell viability was used for sample standardization

Myristoyl Tetrapeptide-34 showed greater effect on reducing MMP-1 than retinyl palmitate or retinol alone. In addition, there was no synergy between retinyl palmitate and Myristoyl Tetrapeptide-34 or retinol and Myristoyl Tetrapeptide-34, respectively.

Comparison with Retinyl Palmitate on MMP-1 Inhibition



Comparison with Retinol on MMP-1 Inhibition

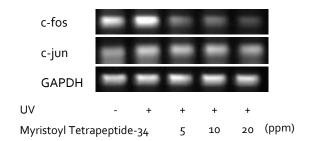


Signaling of MMP-1 expression

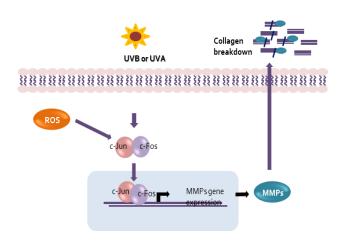
Hs68 cells were seeded into 6-well plates and pretreated with the retinoic acid or Myristoyl Tetrapeptide-34 for 24 and then stimulated by UV irradiation. After 30 minutes, total cellular RNA was extracted and reverse-transcription PCR was performed to determine c-jun and c-fos mRNA level. β -actin was used for sample standardization.

Myristoyl Tetrapeptide-34 significantly decreased UVB induced c-fos mRNA expression

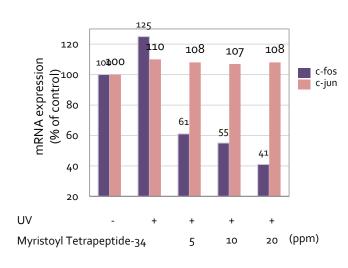
AP-1 component gene expression(PCR)



Mechanism of MMP-1 expression



c-fos/c-Jun gene expression

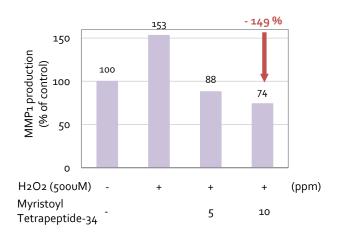


Inhibition of ROS Induced MMP-1 Production and AP-1 Signaling

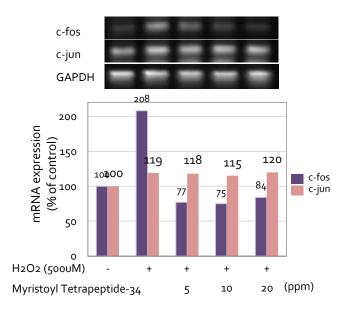
Hs68 cells were seeded into 6-well plates and treated with Myristoyl Tetrapeptide-34 followed by or H2O2. Equal amounts of media were analyzed by ELISA assay for MMP-I. Cell viability was used for sample standardization. For AP-1 signaling, total cellular RNA was extracted and reverse-transcription PCR was performed to determine c-jun and c-fos mRNA level. GAPDH was used for sample standardization.

Myristoyl Tetrapeptide-34 significantly decreased ROS radical stimulated MMP-1 production and c-fos mRNA expression.

Inhibition of MMP-1 production against H2O2



AP-1 component gene expression(PCR)

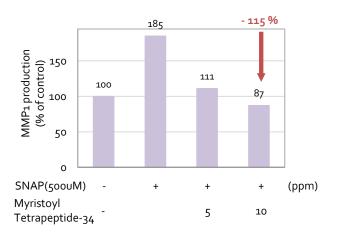


Inhibition of RNS Induced MMP-1 Production and AP-1 Signaling

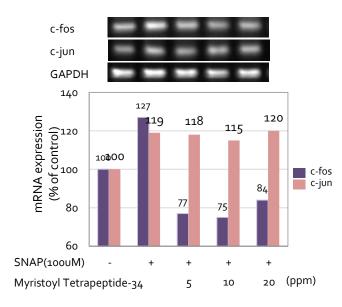
Hs68 cells were seeded into 6-well plates and treated with the Myristoyl Tetrapeptide-34 followed by SNAP(S-nitroso-N-acetylpenicillamine. Equal amounts of media were analyzed by ELISA assay for MMP-I. Cell viability was used for sample standardization. For AP-1 signaling, total cellular RNA was extracted and reverse-transcription PCR was performed to determine c-jun and c-fos mRNA level. GAPDH was used for sample standardization.

Myristoyl Tetrapeptide-34 significantly decreased RNS radical stimulated MMP-1 production and c-fos mRNA expression.

Inhibition of MMP-1 production against SNAP



AP-1 component gene expression(PCR)



Anti-inflammatory Effects of Myristoyl Tetrapeptide-34 on Human Fibroblasts

Hs68 cells were seeded into 6-well plates and pretreated with TGF- β , retinoic acid or Myristoyl Tetrapeptide-34 followed by 30J/cm² UVA . After 24 hours, total cellular RNA was extracted and reverse-transcription PCR was performed to determine IL-1 β , IL-6, IL-8 . GAPDH mRNA level was used for sample standardization.

Myristoyl Tetrapeptide-34 greatly inhibited UVA-induced expression of pro-inflammatory cytokines in human skin fibroblasts.

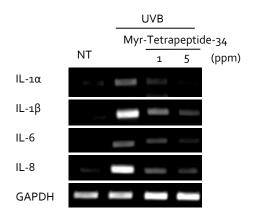
Anti-inflammatory Effects of Myristoyl Tetrapeptide-34 on Keratinocytes

Human keratinocyte HaCaT cells were treated with Myristoyl Tetrapeptide-34 and then stimulated with UVB (100 mJ/cm²). After 24 hours, total cellular RNA was extracted and reverse-transcription PCR was performed to determine IL-1 α , IL-1 β , IL-6. GAPDH mRNA level was used for sample standardization.

Myristoyl Tetrapeptide-34 significantly inhibited UVB-induced expression of pro-inflammatory cytokines in human keratinocytes.

Pro-inflammatory cytokines Level (Hs68)

Pro-inflammatory cytokines Level (HaCaT)



DNA Chip Assay

CHIP data of pro-inflammatory related gene

| Index | ProbeName | Description | Gene Symbol | Genbank Accession | UniGeneID | A440/UV intensity |
|-------|---------------|--|----------------|----------------------|-----------|----------------------|
| 1078 | A_23_P121253 | tumor necrosis factor (ligand) superfamily, member 10 (TNFSF10), transcript variant 1, mRNA $$ | TNFSF10 | NM_003810 | Hs.478275 | 0.712051591 |
| 2139 | | tumor necrosis factor (ligand) superfamily, member 13b (TNFSF13B), transcript variant 1, mRNA $$ | TNFSF13B | NM_006573 | Hs.525157 | 0.444733865 |
| 7521 | A_23_P376488 | tumor necrosis factor (TNF), mRNA | TNF | NM_000594 | Hs.241570 | 0.679159864 |
| 3365 | A_23_P165624 | tumor necrosis factor, alpha-induced protein 6 (TNFAIP6), mRNA | TNFAIP6 | NM_007115 | Hs.437322 | 0.542426693 |
| 10479 | A_23_P71530 | tumor necrosis factor receptor superfamily, member 11b (TNFRSF11B), mRNA | TNFRSF11B | NM_002546 | Hs.81791 | 0.50666134 |
| 4776 | A_23_P218646 | tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF 6B), mRNA $$ | TNFRSF6B | NM_003823 | Hs.434878 | 0.587013243 |
| 25168 | | tumor necrosis factor receptor superfamily, member 6b, decoy (TNFRSF 6B), mRNA $$ | TNFRSF6B | NM_003823 | Hs.434878 | 0.589897775 |
| 1337 | A_23_P126836 | tumor necrosis factor (ligand) superfamily, member 4 (TNFSF4), mRNA | TNFSF4 | NM_003326 | Hs.181097 | 0.858187786 |
| 21238 | A_33_P3246833 | interleukin 1 receptor antagonist (IL1RN), transcript variant 4, mRNA | IL1RN | NM_173843 | Hs.81134 | 0.398674612 |
| 12985 | A_24_P200023 | interleukin 1 receptor, type I (IL1R1), mRNA | IL1R1 | NM_000877 | Hs.701982 | 0.547208757 |
| 31154 | A_33_P3396389 | interleukin 1 receptor, type I (IL1R1), mRNA | IL1R1 | NM_000877 | Hs.701982 | 0.654729398 |
| 10576 | A_23_P73780 | interleukin 1 receptor-associated kinase 1 (IRAK1), transcript variant 1, mRNA | IRAK1 | NM_001569 | Hs.522819 | 0.826588326 |
| 11156 | A_23_P85209 | interleukin 13 receptor, alpha 2 (IL13RA2), mRNA | IL13RA2 | NM_000640 | Hs.336046 | 0.454015552 |
| 1964 | A_23_P138680 | interleukin 15 receptor, alpha (IL15RA), transcript variant 2, mRNA | IL15RA | NM_172200 | Hs.524117 | 0.696403722 |
| 5072 | A_23_P252062 | ariant 3, MKNA | PPARG | NM_138711 | Hs.162646 | 0.620602594 |
| 28110 | A_33_P3350726 | peroxisome proliferator-activated receptor gamma (PPARG), transcript v ariant 3, mRNA | PPARG | NM_138711 | Hs.162646 | 0.587618207 |

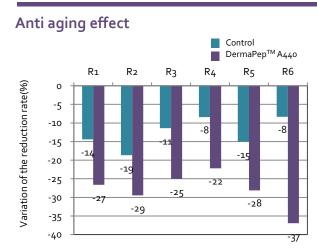
CHIP data of ECM protein related gene

| Index | ProbeName | Description | Gene Symbol | Genbank Accession | UniGeneID | UV/NT intensity | A440/UV intensity |
|-------|---------------|--|----------------|----------------------|-----------|-----------------|-------------------|
| 25105 | A_33_P3304668 | collagen, type I, alpha 1 (COL1A1), mRNA | COL1A1 | NM_000088 | Hs.172928 | 0.445770446 | 1.932809889 |
| 11097 | A_23_P83818 | collagen, type V, alpha 1 (COL5A1), mRNA | COL5A1 | NM_000093 | Hs.210283 | 0.749845964 | 1.621188441 |
| 33475 | A_33_P3629678 | collagen, type V, alpha 1 (COL5A1), mRNA | COL5A1 | NM_000093 | Hs.210283 | 0.559161349 | 1.618872875 |
| 31278 | A_33_P3398236 | collagen, type XXVIII, alpha 1 | COL28A1 | AJ890452 | Hs.491104 | 0.934087883 | 3.457301057 |
| 14419 | A_24_P334130 | fibronectin 1 (FN1), transcript variant 7, mRNA | FN1 | NM_054034 | Hs.203717 | 0.850157729 | 2.353647741 |
| 2910 | A_23_P156327 | transforming growth factor, beta-induced, 68kDa (TGFBI), mRNA | TGFBI | NM_000358 | Hs.369397 | 0.682829566 | 2.148446928 |
| 25707 | A_33_P3313825 | transforming growth factor, beta receptor II (70/80kDa) | TGFBR2 | AJ786388 | Hs.82028 | | 1.62420191 |
| 15179 | A_24_P402438 | transforming growth factor, beta 2 (TGFB2), transcript variant 2, mRNA | TGFB2 | NM_003238 | Hs.133379 | 0.971181556 | 1.69905287 |
| 8176 | A_23_P405129 | latent transforming growth factor beta binding protein 2 (LTBP2) , mRNA $$ | LTBP2 | NM_000428 | Hs.512776 | 1.024822695 | 2.42692819 |
| 3781 | A_23_P19663 | connective tissue growth factor (CTGF), mRNA | CTGF | NM_001901 | Hs.410037 | 0.332717584 | 3.151554741 |
| 30141 | A_33_P3380797 | fibroblast growth factor 3 (FGF3), mRNA | FGF3 | NM_005247 | Hs.37092 | 0.965838192 | 2.016988657 |
| 3838 | A_23_P200741 | dermatopontin (DPT), mRNA [NM_001937] | DPT | NM_001937 | Hs.80552 | 0.762254902 | 1.327178047 |
| 13766 | A_24_P274270 | signal transducer and activator of transcription 1, 91kDa (STAT1), transcript variant beta, mRNA | STAT1 | NM_139266 | Hs.642990 | 2.028586279 | 0.346110016 |
| 9975 | A_23_P62115 | TIMP metallopeptidase inhibitor 1 (TIMP1), mRNA | TIMP1 | NM_003254 | Hs.522632 | 0.850732951 | 1.517430948 |
| 3547 | A_23_P1691 | matrix metallopeptidase 1 (interstitial collagenase) (MMP1), transcript variant 1, mRNA $$ | MINIPI | NM_002421 | Hs.83169 | 1.564405238 | 0.568818868 |
| 3181 | A_23_P161698 | matrix metallopeptidase 3 (stromelysin 1, progelatinase) (MMP3), mRNA | | NM_002422 | Hs.375129 | 2.803817166 | 0.82164818 |
| 16187 | A_24_P82106 | matrix metallopeptidase 14 (membrane-inserted) (MMP14), mRN A $$ | MMP14 | NM_004995 | Hs.2399 | 1.135234216 | 0.695241922 |

Anti-Aging Efficacy on Asian Skin Type

In a clinical study with 21 Asian healthy female volunteers (Korea), aged 30 to 65, 2 % formula containing DermaPep™A440 had been applied twice daily during 12 weeks on the face area. Evaluation was performed by Transparancy profilometry analysis using visiometer SV600(Courage-Khazaka electronic GmbH, Germany) and digital photography.

DermaPep[™]A440 showed significant improvement in all the parameters measured and proved to be the powerful anti-wrinkle agent.



R1: Skin roughness R2: Maximum roughness R3: Average roughness R4: Smoothness depth

R5: Arithmetic average roughness

R6: Expert's assessment

Illustrative Picture_Code No.11915-K1-12



| 2% DERMAPEP™ A440 | | | | | | | |
|---------------------------------|-------|--------------|--------------|---------------|--|--|--|
| Code 11915-K1-12 | T0 | T+4 weeks | T+8 weeks | T+12 weeks | | | |
| Skin Roughness | 134.2 | 93.08 | 99.35 | 63.18 | | | |
| Maxium Roughness | 90.13 | 66.23 | 68.86 | 49.11 | | | |
| Average Roughness | 53 | 48.03 | 45.6 | 35.5 | | | |
| Smoothness Depth | 62.31 | 36.36 | 38.47 | 26.45 | | | |
| Arithmetic average Roughness | 31.24 | 13.62 | 17.04 | 9.74 | | | |
| Experts assessment | 3 | 3 | 2 | 1 | | | |



Last revised on Apr 17h, 2013