Research Article





Tocotrienols activate diacylglycerol kinase a via 67 KDa laminin receptor

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ABSTRACT

Background: Diabetes is a significant social issue. Controlling diabetic complications such as nephropathy is crucial for the quality of life (QOL) of diabetic patients. The abnormal activation of protein kinase C (PKC) through increased diacylglycerol levels (DG) due to hyperglycemia is a common mechanism that causes diabetic complications. Diacylglycerol kinase (DGK) can attenuate PKC activity by converting DG to phosphatidic acid. Thus far, d- α -tocopherol (α Toc) treatment has been shown to prevent early changes of diabetic renal dysfunctions by activating DGK α via the 67KDa laminin receptor.

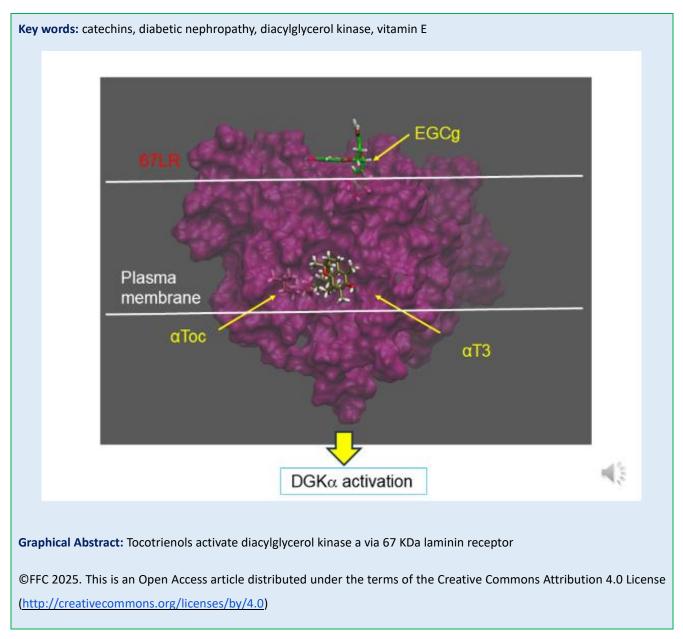
Vitamin E is classified into tocopherols (Toc) and tocotrienols (T3) with four derivatives: α , β , $\gamma \alpha$, β , γ , and δ . We have found that all Tocs induced the translocation of DGK α . However, it remains unclear if T3s also induced translocation of DGK α .

Objective: The objective of the study was to investigate the relationship between tocotrienols (T3s) and induced translocation of DGK α , while exploring the mechanism underlying T3s-induced translocation of DGK α .

Results: Similarly to a-tocopherol, α -T3 binds to 67LR, resulting in the induction of a translocation of DGK α . As well as α -T3, β , γ , and δ Tocs induced translocation of DGK α .

Conclusion: α T3, like α Toc, activates DGK α by binding to Leu58 and Trp176 in 67LR. It means 67LR may be important for the functions of T3s, especially the non-oxidant effects of T3s.

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INTRODUCTION

In recent years, diabetes has become a significant health concern with an estimated 400 million patients worldwide [1-2]. Prolonged diabetes may lead to complications such as diabetic retinopathy, diabetic neuropathy, and diabetic nephropathy (DN). One-third of diabetic patients suffer from DN, which is characterized by proteinuria, decreased renal glomerular filtration, hypertension, and edema. One of the causes of DN is an increase in diacylglycerol (DG), which activates protein kinase C (PKC), specifically the β subtype [3-7]. However, DG kinase phosphorylates DG to produce phosphatidic acid, decreasing DG [8-11]. Therefore, activation of DGK may improve diabetic nephropathy indirectly by suppressing the abnormal hyperactivity of PKC. Previous data have shown that oral administration of epigallocatechin gallate (EGCG), a type of catechin, and d- α -tocopherol (α Toc), a type of vitamin E, increased DGK α activity in the renal glomeruli of diabetic mice, which improved DN [12-17]. The activation of DGK α was also achieved by direct binding of EGCG and α Toc to the 67 kDa laminin receptor (67LR) [18-19]. However, the binding sites of EGCG and α Toc were different. EGCG binds around the 166th lysine of the 67LR, whereas α Toc

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binds to the pocket created by the 67LR [18].

Vitamin E is the most abundant fatsoluble antioxidant in vivo and is classified into tocopherols (Toc) and tocotrienols (T3) with four derivatives, γ , β , α , and δ , respectively. The former has saturated phytyl groups and the latter has unsaturated isoprenoid side chains (Fig. 1) [20]. Toc is found in soybeans, rapeseeds, and sunflower seeds, while T3 is abundant in oil and cereals such as palm oil, wheat, and barley. α T3 has about 50 times higher antioxidant capacity than α Toc, and its unique physiological functions, such as inhibition of atherosclerosis, cell proliferation, and nerve cell protection, have been reported [21-22]. However, the detailed mechanism of action is still unknown except for its antioxidant capacity.

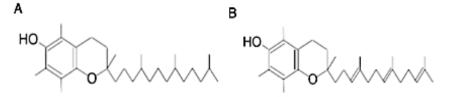


Figure 1. Basic structure of tocopherol (A) and tocotrienol (B)

Numerous functions of α -tocopherol have been studied, including ischemic protection by preventing LDL oxidation, vascular protection, prevention of diabetic complications, and prevention of skin blemishes. Most of these functions are thought to be due to anti-oxidation by oxidizing itself. The antioxidant alone cannot explain many reported functions, such as anticancer effects. Azzi et al. have evaluated these non-antioxidant functions since around 1990 [23]. Yet, this study found that α Toc activates DGK α by direct binding to 67LR, which may account for the non-oxidative functions of α Toc. T3 also has many functions, and its molecular structure is similar. This suggests that 67LR is also involved in some functions of T3s. Therefore, this study determined if T3 can bind to 67LR and activate DGK α .

MATERIALS & METHODS

Materials: Dulbecco's Modified Eagle medium (DMEM), trypsin-EDTA, fetal Bovine Serum (FBS), Opti-MEM, and penicillin-streptomycin were obtained from Gibco. Fugene HD Transfection Reagent was from Promega. Anti-FLAG mouse antibody was purchased from Santa Cruz Biotechnology. Peroxidase-conjugated affiniPure goat anti-Mouse IgG and peroxidase-conjugated affiniPure goat anti-Rabbit IgG were from Jackson Immuno Research Laboratories. Plasmids encoding FLAG-67LR, GFP-DGK α , and anti-GFP antibody were generated in our laboratory [12]. Anti-67LR antibody was from GeneTex. T3s and α Toc were from Cayman Chemical and FUJIFILM Wako Pure Chemicals. EGCG was from LKT Laboratory.

Cell culture: DDT1-MF2 cell, a smooth muscle cell line derived from a leiomyosarcoma of the ductus deferens of a Syrian hamster, was cultured in DMEM-high glucose medium supplemented with penicillin (100 units/ml) and streptomycin (100 mg/ml), 5% FBS without heat treatment, and 20 mM L-glutamine. All cells were cultured at 37°C and 5% CO₂.

Lipofection: DDT1-MF2 cells (1.0×10^5) were seeded in a polylysine-coated glass-bottomed petri dish (MATSUNAMI, 3.5 cm diameter) and cultured for 24 hours. Three hundred µl of OPTI-MEM and 5 µl of Fugene HD transfection reagent (Promega) were mixed well for each glass-bottomed petri dish. Two µg each of plasmid encoding the cDNA of both GFP-DGK α and FLAG-67LR (or mutants) were added to this solution, mixed well, and allowed to stand at room temperature for 10 minutes.

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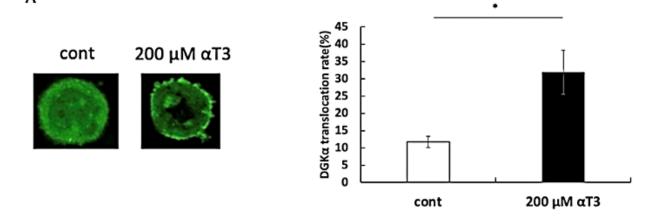
The entire volume of the prepared gene transfer solution was slowly added to the center of a glassbottomed petri dish seeded the previous day and incubated for 24 hours.

Detection of DGK α **activation:** Twenty-four hours after the lipofection, cells were stimulated with several concentrations of T3s or α Toc for 15 minutes and fixed with 4% paraformaldehyde. Localization of GFP-DGK α was then observed based on the green fluorescent under confocal laser microscopy. DGK α translocation rate (%), which is the number of cells showing membrane localization of GFP-DGK α /total cells observed expressing GFP-DGK α , was used for the activation because DGK α is translocated from the cytoplasm to the plasma membrane when activated.

Molecular docking: Since the α Toc binding site is closed in the crystal structure of 67LR (PDB ID: 3BCH) [24], the 3D structure of α Toc-bound form of 67LR in the previously performed MD simulation [18] was employed as the receptor. The receptor was prepared using the Protein Preparation Wizard [25, 26]. **A** αToc and EGCG were prepared using the LigPrep function [27, 28]. The box center for the docking calculations was defined using the centroid of selected residues that were found to constitute the EGCG and αToc binding site [29, 18]. The grid required for the docking procedure was generated using a scaling factor of 1.0 and a partial charge cutoff of 0.25, while the X, Y, Z dimensions of the inner box were set to cover the whole binding site. For the ligand docking, a scaling factor of 0.8 and a partial charge cutoff of 0.15 were used to allow complete flexibility of the structures. The poses were selected according to the binding mode and the XP GScore. The Glide Extra-Precision (XP) scoring function was used for the calculations [30]. The binding free energy was calculated by the MM-GBSA method using Prime-MM-GBSA [31].

RESULTS & DISCUSSION

To investigate whether α T3 activates DGK α , we first stimulated DDT1-MF1 cells with 200 μ M α T3, which significantly increased the rate of DGK α translocation compared to untreated cells. (Fig. 2A, B).



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Figure 2. Effect of aT3 on DGK α activation (A) Typical image of GFP-DGK α before (control) and after DDT-MF2 cells expressing both FLAG-67LR and GFP-DGK α by 200 mM α T3 (experiments). (B) The translocation rate was calculated as described in Materials and methods, * shows p<0.05 by student's T-test.

Next, the translocation rate of DGK α was examined by varying the concentration of α T3. The results showed that α T3 above 50 μ M increased the translocation rate of DGK α , which is a percentage of the cells showing membrane localization of GFP-DGK α in a concentrationdependent manner (Fig. 3A, B).

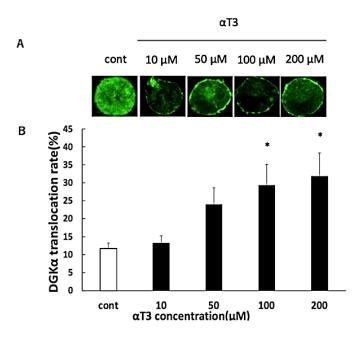


Figure 3. Dose-dependent effect of aT3 on DGK α activation. DDT-MF2 cells expressing both FLAG-67LR and GFP-DGK α were stimulated by 10, 50, 100 or 200 mM α T3. We performed the experiments more than 3 times and observed more than 30 cells in each experiment. Pictures above the graph are typical images of the cell showing the membrane localization of GFP-DGK α , so-called translocation. The percentage of the cells showing the membrane localization of GFP-DGK α , so-called translocation. The percentage of the cells showing the membrane localization of α T3. * Shows p<0.05 by Dunnett test.

Next, the translocation rate of DGK α was examined by pretreatment with 67LR antibody prior to α T3 stimulation, to determine whether 67LR, as well as α Toc, mediates DGK α activation by α T3. The results showed that stimulation of α T3 after pretreatment with 67LR antibody was significantly reduced, compared to the groups that were not treated with the antibody or treated with the control (Fig. 4). This indicates that the activation of DGK α by α T3 was mediated by 67LR.

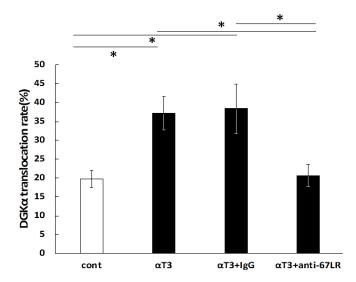


Figure 4. Involvement of 67LR in the aT3-induced activation of DGKα. After preincubation without or with anti-67LR or control antibody (IgG), DDT-MF2 cells expressing both FLAG-67LR and GFP-DGKα were stimulated by 200 mM αT3. We performed the experiments more than 3 times and observed more than 30 cells in each experiment. The translocation rate was calculated as described in Materials and methods, * shows p<0.05 by student's T-test.

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To examine the binding sites of α T3 and 67LR, we overexpressed FLAG-67LR with mutations on Leu58 and Trp176, which are the α Toc binding sites, or on Lys166, which is the EGCG binding site, and examined their effects on the translocation of DGK α . Figure 5 showed that the translocation rate of DGK α was not significantly changed with the use of the Lys166 mutant. However, the translocation rate significantly decreased in the group mutated in the α Toc binding site.

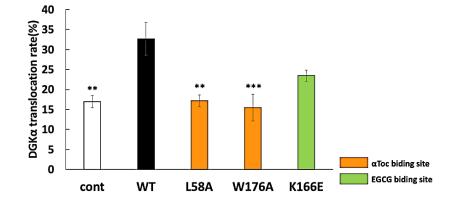


Figure 5. The binding site of α T3 in 67LR. To determine the binding site of α T3, DDT-MF2 cells expressing both GFP-DGK α and FLAG-67LR (WT) or its mutant were stimulated by 200 mM α T3. We performed the experiments more than 4 times and observed more than 30 cells in each experiment. * shows p<0.05 by Student's T-test vs WT.

The results indicated that α T3, like α Toc, activates DGK α via binding to Leu58 and Trp176 in 67LR. Since α Toc and α T3 have very similar structures (Fig. 1), this finding aligned with expectations. Molecular docking simulation supports the hypothesis that both bind to the same pocket with similar strength; binding free energy of α Toc and α T3 was -94.3 and -75.3, respectively (Fig. 6). However, binding free energy of EGCG to the site was -

58.01, which was very low compared with α Toc and α T3 (data not shown). These results indicate that α Toc and α T3 bind to the same site in 67LR, activating DGK α . According to the analysis, 67LR is most often expressed in cancer cells [32]. However, both α Toc and α T3 have anticancer effects. So, the binding of both α Toc and α T3 may be involved in this phenomenon.

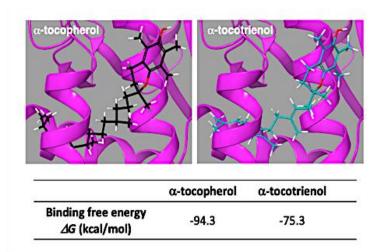
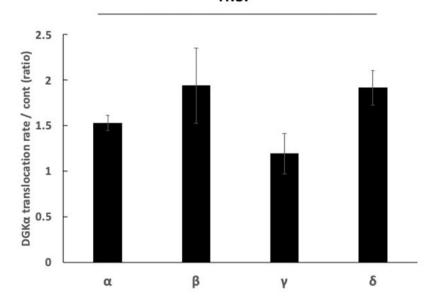


Figure 6. Comparison of the binding free energy of α T3 and α Toc in 67LR by docking simulation. The docking simulations were performed using 3D structures of 67LR as a receptor and α Toc and α T3 as ligands. The docking poses that gave the best binding score were shown, and the calculated binding free energy was calculated by the MM-GBSA method.

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T3 has four isoforms (α -, β -, γ -, and δ -), characterized by a Croman ring attached to the isoprenoid side chain at the C2 position. T3 also has anticancer properties, inducing growth arrest and cell death in several cancer cells derived from breast, gastrointestinal, liver, prostate, lung, and bone cancer [33-36]. However, anticancer effects vary among the T3

isoforms. It was hypothesized that 67LR would show a different affinity to the T3 isoforms. Thus, we examined whether there were differences in DGK α translocation rate among the T3 isoforms. All T3 isoforms significantly activated DGK α , and the strengths to activate DGK α were β -, δ -, α -, and γ -T3, respectively (Fig. 7). However, no significant differences were observed (Fig. 7).

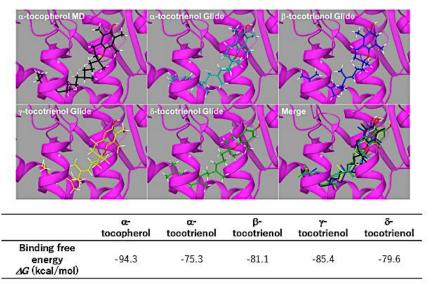


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Figure 7. Comparison of the ability of α , β , γ , and dT3 to activate DGK α . To compare the activity to induce DGK α translocation (activation), DDT-MF2 cells expressing both GFP-DGK α and FLAG-67LR (WT) were stimulated by 100 mM α , β , γ , or δ T3. We performed the experiments 4 times and observed more than 30 cells in each experiment to calculate the translocation ratio. Each column shows a ratio to control (no stimulation). * Shows p<0.05 by the Turky-Kramer test.

Figure 8 shows no difference between binding free energies of T3 isoforms in α Toc binding sites of 67LRs, which supports the results in Fig. 7. This finding suggests that all T3s activate DGK α via 67LR. A difference in translocation of DGK α in each T3 may be observed when there is a concentration dependence. However, a significant correlation between increased 67LR expression and tumor progression has been recently reported. The overexpression of 67LR has been observed in many cancers such as chronic lymphocytic leukemia, acute myeloid leukemia, multiple myeloma, and prostate cancer [37-40]. Yet, it has been reported that EGCG binding to 67LR may induce apoptosis via protein kinase B, nitric oxide, and cyclic AMP34, 35. Therefore, the mechanism of 67LR-mediated DGK α activation by T3 revealed in this study may explain the anticancer effects of various T3, but further studies are needed to conclude.

As described above, DGK α activation via 67LR is crucial to ameliorating DN by EGCG and VtE. Therefore, T3 may be effective on DN. Similarly, Naigenin may activate DGK α via 67LR [41]. Yet, we cannot rule out the possible involvement of an anti-oxidative effect in the VtE or EGCG-mediated amelioration of DN because anti-oxidative impact is vital for the amelioration of DN [15, 42].



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Figure 8. Comparison of binding free energy of α , β , γ , δ T3, and α Toc to 67LR by docking simulation. The docking simulations were performed using 3D structures of 67LR as a receptor and α Toc, α , β , γ , and δ T3 as ligands. The docking poses that gave the best binding score were shown, and the binding free energy was calculated using the MM-GBSA method.

CONCLUSION

 α T3, like α Toc, activates DGK α by binding to Leu58 and Trp176 in 67LR. Therefore, 67LR may be important for the functions of T3s, especially the non-oxidant effects of T3s, such as the anti-cancer function.

Abbreviations: d-α-tocopherol (αToc), diacylglycerol (DG), diacylglycerol kinase (DGK), Dulbecco's Modified Eagle medium (DMEM), diabetic nephropathy (DN), ethylenediamine tetraacetic acid (EDTA), epigallocatechin gallate (EGCG), fetal bovine serum (FBS), green fluorescent protein (GFP), knock out (KO), protein kinase C (PKC). tocopherol (Toc), tocotrienol (T3), 67 KDa laminin receptor (67LR)

Competing interests: The Authors have no conflict of interest.

Author's contribution: Tomoko Namba performed almost all the experiments, Dr. Daiki Hayashi performed the Glide simulation, Dr. Yasuhito Shirai designed and conducted the experiments, and Drs. Itsuko Fukuda and Shuji Ueda helped design and financially supported a series of experiments. Acknowledgements: Dr. Edward Dennis at UCSD kindly supervised our molecular docking experiments. This work was partly supported by the Daiichi Sankyo Foundation of Life Science and the Grant–in–Aid for scientific research (C) of Japan's Ministry of Education, Culture, Sports, Science and Technology.

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