

Induction of VEGF Expression by Alpha-Tocopherol and Alpha-Tocopheryl Phosphate via PI3Kγ/PKB and hTAP1/SEC14L2-Mediated Lipid Exchange

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ABSTRACT

In several studies, vitamin E has been observed to influence angiogenesis and vasculogenesis. We recently showed that the phosphorylated form of α -tocopherol (α T), α -tocopheryl phosphate (α TP), increases the expression of the vascular endothelial growth factor (VEGF). Thus, α TP may act as an active lipid mediator increasing VEGF expression, angiogenesis, and vasculogenesis. Here, we investigated the molecular signaling mechanisms by which α TP induces VEGF expression using cultured HEK293 cells as model system. α T and more so α TP increased VEGF-promoter activity in a phosphatidylinositol-3-kinase gamma (PI3K γ)-dependent manner. In contrast, after overexpression of PI3K γ and/or protein kinase B (PKB), VEGF promoter activity was inhibited by α T and more so by α TP. Inhibition by α T and α TP was dependent on the lipid kinase activity of PI3K γ , whereas an induction was seen with the protein kinase activity, consistent with a model in which PKB inhibition by α T or α TP occurs only when activated at the plasma membrane and possibly involves a phosphatase such as PHLPP1. PI3K γ -induced VEGF expression was reduced when the human tocopherol-associated protein 1 (hTAP1/SEC14L2) was overexpressed suggesting formation of an inactive PI3K γ /hTAP1 heterodimer, that could be reactivated by α T and more so by α TP. We suggest a novel signaling mechanism by which α TP stimulates PI3K γ activity by stimulating hTAP-mediated phosphatidylinositol exchange and presentation to the enzyme and/or dissociation of an inactive heterodimer. At cellular level, hTAP may act as sensor for intracellular lipid information (location, type, and amount of lipid) and translate it into responses of PI3K-mediated signaling and gene expression. J. Cell. Biochem. 116: 398–407, 2015. © 2014 Wiley Periodicals, Inc.

KEY WORDS: VEGF; SEC14-LIKE PROTEINS; TOCOPHEROL; PHOSPHOLIPIDS; VITAMIN E; KINASE; TOCOPHERYL PHOSPHATE; hTAP; SUPERNATANT PROTEIN FACTOR; SPF; PKB; PI3K; PI3K; PI3Kγ

V itamin E was discovered as a dietary factor essential for reproduction in rats [Evans and Bishop, 1922]. Since then, vitamin E has revealed many important molecular properties such as the scavenging of reactive oxygen and nitrogen species, and the modulation signal transduction and gene expression (reviewed in [Zingg, 2007]). A congenital disease, ataxia with vitamin E deficiency (AVED), which is characterized by low levels of α tocopherol (α T) in plasma due to mutations in the α -tocopherol transfer protein gene (α -TTP), has been described [Ben Hamida et al., 1993]. An effect of vitamin E on angiogenesis and vasculogenesis has been the subject of several studies but the molecular mechanisms involved are not clear (reviewed in [Zingg et al., 2012]). Using THP-1 monocytes, we recently observed that the phosphorylated form of α -tocopherol (α T), α -tocopheryl phosphate

(α TP), induced VEGF expression leading to increased angiogenesis in human umbilical vascular endothelial cell (HUVEC) in culture, whereas α T was not effective [Zingg et al., 2010a].

VEGF is an endothelial cell-specific mitogen that promotes angiogenesis and mediates, among other actions, a successful pregnancy to the final stage by inducing angiogenesis and vasculogenesis during placenta and embryonic development [Zygmunt et al., 2003; Haigh, 2008]. We proposed that fetal resorption in the vitamin E-deficient state may be the consequence of decreased expression of VEGF as a result of insufficient production of VEGF in the absence of vitamin E, followed by impaired formation of an extensive vascular net in the placenta leading to placental ischemia, and inadequate nutrient supply to the fetus [Zingg et al., 2012]. In support of this concept, increased placental angiogenesis and vascular

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network formation was recently detected in pregnant ewes supplemented with vitamin E, possibly resulting from stimulation of VEGF expression [Kasimanickam et al., 2010]. Accordingly, in mice deficient in vitamin E as result of a knockout of the α -TTP gene (α - $TTP^{-/-}$), the absence of embryonic blood vessels in the trophoblast was observed, in addition to a failure of trophoblasts to survive and an abnormally forming small labyrinth [Jishage et al., 2001]. A lower level of VEGF in VEGF knockout mice (homozygous and even heterozygous) leads to abnormal embryonic blood vessel formation and impairs the production of a viable offspring [Carmeliet et al., 1996; Ferrara et al., 1996]. Likewise, successful blastocyst implantation for pregnancy establishment is critically dependent on the presence of sufficient expression of VEGF [Sengupta et al., 2007]. In the brain, VEGF in the Purkinje cell layer of cerebellum is necessary for proper granule cell migration [Ruiz de Almodovar et al., 2010], and mice not able to upregulate VEGF during hypoxia develop motor neuron degenerations that are symptoms related to vitamin Edeficient mice [Oosthuyse et al., 2001; Ulatowski et al., 2014].

Regulation of VEGF expression by vitamin E has been reported in several in vitro and in vivo experimental systems, in which either activation [Zhang et al., 2004; Daghini et al., 2007] or inhibition of VEGF [Tang and Meydani, 2001; Nespereira et al., 2003; Schindler and Mentlein, 2006] has been observed, but the molecular mechanisms of modulation of VEGF by vitamin E are yet to be understood (reviewed in [Zingg et al., 2012]). We reported that the induction of VEGF by α TP was due to upregulation of the phosphatidylinositol-3-kinase (PI3K) and/or protein kinase B (PKB/Akt) signaling pathway, a pathway that is downregulated by αT [Munteanu et al., 2006]. We suggested that αTP acts as an active "lipid mediator" and activates the PI3K/PKB pathway and that the un-phosphorylated aT inhibits its activation in THP-1 cells [Kempna et al., 2004; Munteanu et al., 2006; Numakawa et al., 2006]. Thus, the cellular response to αT may depend on the ratio of $\alpha T/\alpha TP$ and thus on degree of their interconversion in different tissues or cell types. Based on these results we proposed that in certain tissues or conditions, αT is converted locally to αTP , which activates the PI3K/PKB signaling pathway and ultimately stimulates VEGF production leading to increased angiogenesis and vasculogenesis (reviewed in [Zingg et al., 2010b]).

In a recent study, we have shown that αT and more so αTP activate in vitro phosphatidylinositol-3-kinase gamma (PI3K γ) activity by inducing tocopherol-associated protein 1 (hTAP1/SEC14L2)-mediated lipid exchange [Zingg et al., 2014]. By using HEK293 cells as a model system, we propose here novel molecular signaling mechanisms by which αT and more so αTP can both stimulate and inhibit the VEGF promoter depending on the activation state of the PI3K γ /PKB pathway.

MATERIALS AND METHODS

MATERIALS

RRR- α -tocopherol (α T) (from Cognis, Cincinnati, OH) was dissolved in ethanol as 50 mM stock solutions and the concentrations confirmed spectrophotometrically. Stock solutions (50 mM) of α -tocopheryl phosphate (α TP), D- α -[5-methyl-14C]-tocopheryl phosphate (14C- α TP) (0.13 μ Ci/mg) (provided by Phosphagenics Ltd (Melbourne, Australia)) and D- α -[5-methyl-14C]-tocopherol $(14C-\alpha T)$ (57 mCi/mmole) (Amersham Pharmacia Biotech) were prepared in ethanol or water [Munteanu et al., 2004; Negis et al., 2007]. The specific inhibitor of PI3K γ , AS-605240 (5-(Quinoxalin-6ylmethylene)thiazolidine-2,4-dione) (Alexis Biochemicals, San Diego, CA) was dissolved in DMSO.

CELL CULTURE

HEK293 cells (ATCC, CRL-1573) were grown in Dulbecco's modified Eagle's medium, 10% fetal calf serum, and 2 mmol/L L-glutamine containing 100 µg/mL streptomycin and 100 U penicillin.

TRANSFECTION

HEK293 cells (70% confluent) were transfected with pCGCG-luc, a reporter plasmid containing 3169bp of the human VEGF promoter in front of the *firefly* luciferase gene (kindly provided by S. J. Prior, University of Maryland, Baltimore, MD [Prior et al., 2006]), with expression vectors for phTAP1, phTAP2, phTAP3, or phTTP [Kempna et al., 2003; Zingg et al., 2008], PI3Ky-wt, PI3Ky-mut, PI3Ky-L, PI3Ky-P, PI3Ky-LP (all kindly provided by Dr. H. A. Rockman, Duke University, Durham, NC [Ma et al., 1998; Naga Prasad et al., 2005]), pPKBwt, pPKB(R25C), and pPKB(K179M) (kindly provided by Dr. J. Downward, Imperial Cancer Research Fund, London, UK [Watton and Downward, 1999; Munteanu et al., 2006]), together with the Renilla internal control plasmid pRL-TK (Promega, Madison, WI), for 3 h using Fugene (Promega) as transfection reagent, and then treated with αT (40 μ M) or αTP (40 μ M) for additional 21 h. Extracts were prepared, and promoter activities were measured using the Dual-Luciferase assay kit (Promega) by means of a GLOmax luminometer (Promega). The VEGF promoter-firefly luciferase activities were normalized to the thymidine kinase promoter-Renilla luciferase activities, and the activities of the control transfections were set to 100%.

UPTAKE OF αT AND αTP

HEK293 cells were plated in 24 wells plates at 70% confluency overnight, transfected for 24 h with expression vectors for hTAP1, hTAP2, hTAP3, or hTTP, or empty control vector pMH [Kempna et al., 2003], and after changing medium incubated with ¹⁴C- α T (40 μ M) or ¹⁴C- α TP (40 μ M) for 5 h. Cells were washed with PBS, trypsinized, removed to eppendorf tubes and washed two times with PBS. The radioactivity associated with the cell pellet was measured using a scintillation counter and the data plotted as % of total input.

STATISTICAL ANALYSIS

All values are expressed as the mean \pm standard error of the mean (SEM) as explained in the figure legends. Student's *t*-test was used to determine the significant differences between two conditions. A P < 0.05 was considered as significant and indicated by * or # in the graphs.

RESULTS

αT and αTP regulate VEGF promoter activity VIA modulation of PI3K γ and PKB

Using gene expression arrays and THP-1 monocytes, we recently detected a number of genes, such as VEGF, that are upregulated specifically by α TP [Zingg et al., 2010a]. α TP stimulated PKB (Ser473)

phosphorylation and free radicals production in a wortmannin and AS-605240-sensitive manner, suggesting the involvement of phosphatidylinositol-3-kinases (PI3K) [Zingg et al., 2010a, 2014], whereas α T antagonized these reactions. Interestingly, VEGF [Dutra et al., 2011], angiogenesis [Madeddu et al., 2008; Siragusa et al., 2010], as well as the production of free radicals [Lehmann et al., 2009] are all regulated by PI3K γ suggesting that α TP may indeed activate these events via PI3K γ .

To assess further whether α TP activates VEGF expression via PI3K γ , expression vectors for wild-type or inactive mutant PI3K γ (pPI3K γ -wt or pPI3K γ -mut, respectively [Ma et al., 1998; Naga Prasad et al., 2005]) were transfected together with a human VEGF-promoter-luciferase reporter vector (pCGCG-Luc [Prior et al., 2006]) into HEK293 cells. These cells were chosen since unlike THP-1 monocytes, transfection of HEK293 is highly efficient allowing co-transfection of multiple vectors. Moreover, HEK293 cells may represent a suitable model since they express low levels of VEGF [Liang et al., 2002] and PI3K γ [Hirsch et al., 2000], yet VEGF can be activated [Kurig et al., 2009; Zingg et al., 2012].

VEGF-promoter activity was increased with α TP (Fig. 1A), and in contrast to THP-1 cells, an intermediate activation was also observed by aT possibly resulting from its conversion to aTP in these cells [Zingg et al., 2010a, 2012, 2014]. Overexpression of pPI3Ky-wt strongly activated VEGF-promoter activity, but in this case aT and α TP effects were inhibitory (Fig. 1A). These results suggest that α T and more so α TP can increase the endogenous PI3K γ activity, which is similar to the previously described in vitro results [Zingg et al., 2014]. However, since after PI3K γ overexpression, the effects of αT and α TP on PI3K γ /PKB/VEGF signaling were inhibitory (Fig. 1A), it is plausible that additional regulation may occur when the levels of PI3Ky are high in cells. In these situations, PKB is most likely membrane-attached and fully activated by phosphorylation as in HMC-1 mastocytoma cells or other cancer cells [Munteanu et al., 2006], and αT and αTP may either interfere with PKB membrane translocation and/or activate phosphatases able to de-phosphorylate and inactivate PKB, such as PP2A [Ricciarelli et al., 1998; Wei and Xia, 2006] or PH domain and leucine-rich repeat protein phosphatase 1 (PHLPP1) [Huang et al., 2013] (reviewed in [Zingg, 2007]). In fact, when pPKBwt was overexpressed, αT and more so α TP-inhibited VEGF-promoter activity, but not when the mutants pPKB(R25C) (mutated in the PH-domain required for membrane translocation) and pPKB(K179M) (mutated in the catalytic domain) were overexpressed, both without (Fig. 1B) and with (Fig. 1C) additional overexpression of PI3Ky-wt. These results are in line with a model in which the inhibitory effect occurs only with membraneassociated activated PKB which becomes de-phosphorylated as result of activation of PHLPP1 by aT, aTP or PI3P [Huang et al., 2013].

αT and αTP increase VEGF promoter activity in an htap1-dependent manner via modulation of Pi3K γ

In previous studies, the recombinant human tocopherol-associated protein 1 (hTAP1/SEC14L2) reduced the in vitro activity of the PI3K γ possibly by sequestering the substrate phosphatidylinositol (PI) and/or forming an inactive PI3K γ /hTAP1 heterodimer; addition α T or α TP stimulated PI3K γ , possibly by facilitating egress of PI from



Fig. 1. α T and α TP increase VEGF expression in HEK293 cells, but inhibit it when activated by PI3K γ or PKB. (A) The treatment with α TP, and less with α T, induces VEGF promoter activity in human embryonic kidney 293 (HEK293) cells; when PI3K γ is overexpressed, α T and more so α TP reduce PI3K γ -induced VEGF promoter activity (\pm SEM, n = 4, *P<0.05 relative to untreated control (c)). (B) α T and more so α TP inhibit VEGF-promoter activity after overexpression of pPKBwt, but not when the mutants pPKB(R25C) (mutated in the PH-domain) and pPKB(K179M) (mutated in the catalytic domain) are overexpressed (n = 4, *P<0.05 relative to untreated control (c), both in the absence (B) and presence (C) of PI3K γ overexpression (n = 4, *P<0.05 relative to untreated control).

hTAP1 to the enzyme [Kempna et al., 2004; Zingg et al., 2014]. In view of these in vitro results it appeared that in HEK293 cells PI3K γ is only activated by α TP when hTAP1 is overexpressed. Therefore, it was interesting to investigate, whether overexpression of hTAP would alter the response of PI3K γ /PKB/VEGF to α T or α TP. To assess whether the ability of α T or α TP to activate PI3K γ and the VEGF-promoter is regulated by hTAP1, the above experiments (Fig. 1A) were performed in the presence of an expression vector for hTAP1 (phTAP1) or of an empty control vector (pMH), both in the presence of wild-type or inactive mutant PI3K γ (pPI3K γ -wt or pPI3K γ -mut,

respectively). Similar to the experiments done in vitro [Zingg et al., 2014], hTAP1 expression reduced the ability of PI3Ky-wt to activate the VEGF-promoter, and the addition of αT and more so αTP reactivated hTAP1-inhibited PI3Ky and VEGF-promoter activity (Fig. 2A). When PI3Ky-mut was used, these regulatory effects of hTAP1 and α T or α TP were absent or much weaker (Fig. 2B). VEGFpromoter activity was only induced by aT and aTP after overexpression of hTAP1, but not of the related proteins hTAP2/ SEC14L3, hTAP3/SEC14L4, or hTTP (Fig. 2C). Inhibition of PI3Ky by the specific inhibitor AS-605240 (1 µM) prevented induction of VEGF promoter activity, suggesting that in these cells PI3K γ is the isoform responsive to αT and αTP (Fig. 2D). Interestingly, activation of the VEGF promoter by αT and αTP still occurred in the presence of the src inhibitor PP2, indicating that this tyrosine kinase, which acts upstream of the PI3K, is not involved in the observed effects (Fig. 2D). These results suggest that increase of VEGF promoter activity by αT and more so αTP involves PI3K γ and hTAP1, whereas, in the absence of hTAP1 when only PI3K γ is overexpressed α T and α TP suppress VEGF promoter activity (Figs. 1A and 2A).

DIFFERENTIAL REGULATORY EFFECTS OF αT and αTP on PI3K γ LIPID and PROTEIN KINASE ACTIVITY

PI3K γ can either phosphorylate PI or several proteins including tropomyosin, MAPK, Ras, 4EBP1, or PKC α [Bondeva et al., 1998;

Naga Prasad et al., 2005]. Protein phosphorylation activity has been first characterized as auto-phosphorylation and has been measured only with the cytosolic form of PI3K γ [Bondeva et al., 1998; Dolle et al., 2011]. Moreover, PI3Ky can serve as scaffold for proteins such as β-ARK, PDE3B, and PDE4, for kinase-independent regulation of cellular processes [Dolle et al., 2011]. To evaluate whether the regulatory effects of αT and αTP on VEGF expression occur as a result of phosphorylation of protein or lipid (PI), PI3Ky mutant forms [Naga Prasad et al., 2005], with either only lipid kinase activity (PI3K γ -L), only protein kinase activity (PI3K γ -P), or both reconstituted lipid and protein kinase activity (PI3Ky-LP) were transfected into HEK293 cells with pMH (empty control vector) or with hTAP1 co-transfection, and the effects of αT and αTP treatment on VEGF expression measured. Interestingly, VEGF expression mediated by lipid kinase activity of PI3K γ -L was inhibited by α T and α TP both in the presence and absence of hTAP1 (Fig. 3A). Whereas, VEGF expression mediated by PI3Ky-P protein kinase activity was stimulated (Fig. 3B). In contrast, for the reconstituted PI3Ky-LP lipid and protein kinase the two responses were apparently almost neutralized by each other (Fig. 3C). Overexpression of hTAP1 mainly increased VEGF expression when PI3Ky protein kinase was also expressed (Fig. 3B). These results suggest that in cells the stimulatory effects of αT and more so αTP on VEGF expression may involve the cytosolic protein kinase activity of PI3Ky, possibly requiring hTAP1-



Fig. 2. Modulation of VEGF expression by hTAPs, α -TTP, α T, or α TP. (A) Overexpression of hTAP1 reduces PI3K γ -induced VEGF promoter activity; in this case α T and more so α TP reactivate VEGF promoter activity (\pm SEM, n = 4, **P* < 0.05, relative to empty control vector (pMH), "*P* < 0.05 relative to untreated control), most likely by enhancing lipid exchange and presentation of phosphatidyl inositol to PI3K γ by means of the nanoreactor formed by hTAP1 similar to previously published in vitro results with recombinant enzymes [Zingg et al., 2014]. With empty control vector (pMH), inhibition of VEGF expression was seen. (B) VEGF promoter activity is only weakly changed by α T and α TP after overexpression of a mutant form of PI3K γ ; in this case overexpression of hTAP1 only weakly reduces VEGF promoter activity that is restored by α T and α TP (\pm SEM, n = 4). (C) Overexpression of hTAP2, hTAP3, or hTTP did not change the response of the VEGF-promoter to α T or α TP (\pm SEM, n = 4, **P* < 0.05, relative to untreated control, "*P* < 0.05 relative to untreated empty control vector (pMH)). (D) Induction of VEGF promoter activity by α TP is inhibited by the specific PI3K γ inhibitor AS-605240 (1 μ M), but not by the c-src tyrosine kinase inhibitor PP2 (10 μ M) (\pm SEM, n = 4, **P* < 0.05 relative to control (c), #*P* < 0.05 relative to control (c1)).



Fig. 3. Modulation of VEGF expression by PI3K γ lipid kinase (PI3K γ -L), PI3K γ protein kinase (PI3K γ -P), or reconstituted PI3K γ lipid and protein kinase (PI3K γ -LP) activity and hTAP1. (A) α T and more so α TP reduced VEGF promoter activity in PI3K γ -L overexpressing HEK293 cells, both in the presence of empty control vector (pMH)) or of hTAP1 (n = 6, *P < 0.05 relative to untreated control (c)). (B) α T and more so α TP increased VEGF promoter activity in PI3K γ -P overexpressing HEK293 cells, both in the presence of empty control vector (pMH) or of hTAP1 (n = 6, *P < 0.05 relative to untreated control (c)). (C) α T and α TP weakly increased VEGF promoter activity in reconstituted PI3K γ -LP overexpressing HEK293 cells, both in the presence of empty control vector (pMH) or of hTAP1 (n = 6, *P < 0.05 relative to untreated control (c)). (D) α T and α TP weakly increased VEGF promoter activity in reconstituted PI3K γ -LP overexpressing HEK293 cells, both in the presence of empty control vector (pMH) or of hTAP1 (n = 6, *P < 0.05 relative to untreated control (c)).

mediated transport of αT , αTP , or PI to cytosolic PI3K γ and/or allosteric activation of its kinase activity by these lipids. Whereas the inhibitory effects of αT and αTP on VEGF expression may occur when the lipid kinase activity of PI3K γ generates sufficient PI3P to trigger PKB translocation to the plasma membrane and then becomes inactivated by the PKB phosphatase PHLPP1 [Huang et al., 2013].

OVEREXPRESSION OF hTAPs AND α -TTP INFLUENCES CELLULAR α T AND α TP UPTAKE

It is plausible that the higher stimulation of VEGF promoter activity by αTP in the presence of hTAP1 is the result of increased transport and cellular uptake of α TP. Thus, hTAP1/2/3, and as control α -TTP, were overexpressed in HEK293 cells and the uptake of radioactive D- α -[5-methyl-14C]-tocopherol (14C- α T) or D- α -[5-methyl-14C]-tocopheryl phosphate (14C- α TP) measured. Within the time of the experiment (5 h), about half of α TP was taken up when compared to αT (Fig. 4). Thus, since αTP despite a reduced uptake was more active than αT , αTP may act as an intact and more active molecule. Cellular αT and αTP uptakes were differently regulated, in that hTAP1, hTAP3, and hTTP-inhibited α T uptake (Fig. 4A), whereas they increased it for α TP (Fig. 4B). Thus, since the three proteins hTAP1, hTAP3, and hTTP induced an increased aTP uptake, but only one, hTAP1, increased VEGF expression (Fig. 2C), the effects seen on VEGF expression cannot be solely due to increased aTP uptake. Most likely increased VEGF expression is the consequence of enhanced lipid/ protein catalytic activity of PI3K γ , e.g., as result of increased lipid presentation and exchange by hTAP1 and/or dissociation of an inactive heterodimer [Ile et al., 2006; Zingg et al., 2014]. However, it remains to be determined to what degree binding and activation of a receptor and/or transporter by αTP at the plasma membrane such as CD36 [Zingg et al., 2014] contributes to enhanced activation of endogenous PI3Ky activity by aTP in certain cell types.



Fig. 4. Uptake of α TP into HEK293 cells by hTAPs and hTTP proteins. HEK293 were transfected with expression vectors for hTAP1, hTAP2, hTAP3, hTTP, or empty control vector pMH [Kempna et al., 2003], and then treated with (A) ¹⁴C- α T (40 μ M) or (B) ¹⁴C- α TP (40 μ M) for 5 h. Cellular uptake of α T or α TP was measured as described in materials and method. The radioactivity associated with the cell pellet was measured using a scintillation counter, and the data expressed as % of total input (\pm SEM, n = 4, *P < 0.05).

DISCUSSION

In previous studies, we observed increased expression of VEGF and of angiogenic activity secreted from THP-1 monocytes after in vitro treatment with α TP [Zingg et al., 2010a, 2012, 2014]. Here, we analyzed in detail the signaling mechanisms by which α TP induces VEGF by using a human embryonic kidney cell line (HEK293) as a model system. Although the relevance of embryonic kidney cells for angiogenesis is to date unclear, supplementation of normal pigs with vitamin E and C increased VEGF expression and angiogenesis in the kidney suggesting a regulatory role of α T and/or α TP for angiogenesis in this tissue [Daghini et al., 2007].

We find that αTP and to a lesser degree also αT increases VEGF expression. Some activation by aT suggests either an intrinsic lower ability of this compound to activate PI3Ky or some conversion to α TP by α T kinase activity in these cells [Zingg et al., 2014]. Using a specific inhibitor of PI3Ky, AS-605240, we show that VEGF induction by αTP is mediated by PI3K γ , which is expressed in HEK293 at low levels [Hirsch et al., 2000]. Interestingly, when PI3Ky and/or PKB were overexpressed, αT and more so αTP inhibited VEGF expression, suggesting that after activation and translocation to the plasma membrane, these enzymes become inhibitable by aT and aTP, most likely by increasing PKB dephosphorylation by PHLPP1 [Huang et al., 2013]. Moreover, overexpression of hTAP1 with and without PI3Ky inhibited VEGF expression, but in this situation αT and more so αTP could stimulate it. These results suggest that in the presence of hTAP1, αT and more so αTP increased PI3K γ activity either by disrupting an inactive

PI3K γ /hTAP1 heterodimer or by facilitating lipid exchange and presentation, as supported also by our earlier in vitro data [Zingg et al., 2012].

As outlined in the reaction scheme (Fig. 5), the ratio between hTAPs, enzymes and different lipid ligands may be an important factor in determining the cellular response to αT and αTP and activation of PI3Ky and VEGF expression. The source of αT and αTP (intracellular, extracellular), the efficiency of transport and the conversion of αT to αTP by αT kinase or vice versa of αTP to αT by a phosphatase in a given tissue and cell type [Zingg et al., 2010a] may explain that in response to aT, both induction [Numakawa et al., 2006] and inhibition [Munteanu et al., 2006] of the PI3K/ PKB pathway has been observed. Additional regulatory mechanisms of PI3K γ regulation by α TP may occur, as suggested by experiments with mutant PI3K γ enzymes either having only lipid kinase, protein kinase, or both reconstituted lipid and protein kinase activities. In these experiments, aT and more so aTP inhibited VEGF expression when the lipid kinase was active triggering PKB presence at the plasma membrane, whereas they stimulated VEGF expression via protein kinase activation. However, when both kinase activities were present an intermediate response was obtained. Since protein kinase activity was mainly observed with cytosolic PI3Ky [Bondeva et al., 1998; Dolle et al., 2011], hTAP1 may play a double role by transporting activating lipids to cytosolic locations, and/or by stimulating catalytic turnovers by increasing lipid exchange and presentation to enzymes.

In this study we focused only on the regulation of the VEGF promoter activity by αT and αTP in a cell culture system, without an





in depth analysis of the transcription factors activated by PI3K γ /PKB stimulation. The in vivo regulation of the VEGF expression is very complex and several transcription factors that are regulated by PI3K/PKB such as Hif1 α and CREB are involved [Pages and Pouyssegur, 2005]. Moreover, additional post-transcriptional regulatory mechanisms of VEGF mRNA and protein induction by α TP may be involved that may need to be investigated further, such as alternative splicing and mRNA stability, miRNAs, and proteasome inhibition [Munteanu et al., 2007; Arcondeguy et al., 2013].

In monocytes/macrophages, the observed stimulation of VEGF production may enhance reparative angiogenesis and tissue remodeling as observed after experimental myocardial infarction occurring in a PI3K γ /PKB-dependent manner [Siragusa et al., 2010]. In the placenta, the stimulation of VEGF expression and vasculogenesis potentially by αTP may explain the essentiality of vitamin E against fetal resorption (reviewed in [Zingg et al., 2012]). Moreover, induction of VEGF expression may explain other effects of vitamin E, such as the prevention of ischemia/reperfusion injury in the cardiovascular and nervous system (Lambrechts et al., 2003; Zhang et al., 2004; Mukherjee et al., 2008). Additionally, αT, after conversion to aTP, may possibly promote survival of muscle cells and neurons, stimulate neurite outgrowth, and prevent neurodegenerative processes [Jin et al., 2006; Sakowski et al., 2009; Ulatowski et al., 2014]. A rapid generation of a functional vascular system triggered by α TP and VEGF may be particularly required in newly formed tissues and organs such as the placenta, embryo, and possibly needed during repair after tissue injury, thus avoiding nutrient/oxygen deprivation and ischemia.

It should be emphasized that in this study a relatively high concentration of αT and αTP was used for cell treatments; such high concentrations may occur only for aT in plasma and tissues after supplementation. The tissue levels of α TP are generally low since foods contain only low amounts of a TP [Ogru et al., 2003] and since the uptake into the body as intact molecule is not efficient [Libinaki et al., 2005; Gianello et al., 2005, 2007; Mustacich et al., 2007; Zingg et al., 2010a]. A local synthesis of αTP in specific cells may thus be necessary to reach high enough concentrations for its function at precise subcellular sites relevant for signaling. In our previous experiments, no direct effect was detected when endothelial cells were exposed to αTP [Zingg et al., 2010a], suggesting that αTP may orchestrate angiogenesis in endothelial cells by triggering VEGF production and secretion from neighboring cells in the vascular system (e.g., from vascular smooth muscle cells (VSMC), monocytes/ macrophages, kidney cells, or trophoblasts [Pennington et al., 2012]).

We hypothesize that in the normal physiological situation, the pro-angiogenic signal coming from α TP can be switched off by dephosphorylation to α T [Zingg et al., 2010a]. It is possible that chronic activation of PI3K/PKB/VEGF by α TP in a pathological situation (e.g., by aberrant activation of α T kinase or inactivation of α TP phosphatase in tumor tissues) may facilitate the development, growth, and migration of neoplastic cells by increasing VEGF expression, angiogenesis, and tumor growth. On the other hand, increased expression of alkaline phosphatase in certain cancer cells may prevent the stimulatory effects of α TP. These possible correlations may be at the basis of the finding that no adverse

effects have been reported during dietary supplementation with α TP in animal models [Libinaki et al., 2005; Gianello et al., 2007], and high concentrations of α TP (possibly beyond the hydrolytic capacity of the phosphatases) inhibited cell proliferation and migration, and induced apoptosis in cancer cell lines [Munteanu et al., 2004; Rezk et al., 2007; Saitoh et al., 2009; Zingg et al., 2010a]. In fact, our present results suggest that α TP rather reduces VEGF expression when the PI3K/PKB signaling pathway is activated, as it often occurs in cancer cells.

So far no genetic disease has been linked to hTAPs [Nile et al., 2010], but specific polymorphisms in hTAP1 have been linked with prostate cancer risk [Wright et al., 2009; Zingg and Azzi, 2009]. Reduced expression of hTAP1 in proliferating prostate and breast cancer cells indicates that it interferes with cell proliferation [Ni et al., 2005; Wen et al., 2007; Wang et al., 2009; Johnykutty et al., 2009], possibly as shown here as a result of hTAP-mediated decreased signaling to PI3Ky/PKB/VEGF, Ras/Erk, and Raf kinase [Johnson and Kornfeld, 2010]. The binding of the synthetic vitamin E analog α -tocopheryl succinate (α TS) to hTAPs inhibits cell proliferation and induces apoptosis by affecting the Ras, Mek/Erk and PI3K/PKB pathways, and αT or αTP may act in a similar manner [Ni et al., 2005; Donapaty et al., 2006; Neuzil et al., 2006]. Moreover, downregulation of hTAP1 in tumor tissues may remove its inhibitory effect on PI3K/PKB/VEGF and facilitate the development, proliferation, and migration of neoplastic cells by increasing angiogenesis and tumor growth (reviewed in [Zingg et al., 2012]). In this situation, excess vitamin E may stimulate tumor angiogenesis and e.g., contribute to a higher risk for prostate cancer observed in some studies with vitamin E supplementation [Lippman et al., 2009; Klein et al., 2011].

Taken together, we describe novel signaling pathways by which α TP increases VEGF expression involving PI3K γ and hTAP1mediated lipid exchange, whereas VEGF expression is decreased when PI3K γ /PKB are overexpressed. The enhanced expression of VEGF induced by α TP may explain not only the essential roles of vitamin E on reproduction, but also its effects against ischemia/ reperfusion injury and during wound healing. It may also serve as a survival factor for brain and muscle cells.

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