

Arterial calcifications and increased expression of vitamin D receptor targets in mice lacking TIF1 α

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Calcification of arteries is a major risk factor for cardiovascular mortality in humans. Using genetic approaches, we demonstrate here that the transcriptional intermediary factor 1 α (TIF1 α), recently shown to function as a tumor suppressor in murine hepatocytes, also participates in a molecular cascade that prevents calcifications in arterioles and medium-sized arteries. We further provide genetic evidence that this function of TIF1 α is not exerted in hepatocytes. The sites of ectopic calcifications in mutant mice lacking TIF1 α resemble those seen in mice carrying an activating mutation of the calcium sensor receptor (*Casr*) gene and, in TIF1 α -deficient kidneys, *Casr* expression is increased together with that of many other vitamin D receptor (VDR) direct target genes, namely *Car2*, *Cyp24a1*, *Trpv5*, *Trpv6*, *Calb1*, *S100g*, *Pthlh*, and *Spp1*. Thus, our data indicate that TIF1 α represses the VDR pathway in kidney and suggest that an up-regulation of *Casr* expression in this organ could account for ectopic calcifications generated upon TIF1 α deficiency. Interestingly, the calcifying arteriopathy of TIF1 α -null mutant mice shares features with the human age-related Mönckeborg's disease and, overall, the TIF1 α -null mutant pathological phenotype supports the hypothesis that aging is promoted by increased activity of the vitamin D signaling pathway.

aging | ectopic calcification | mouse knockout | transcriptional regulation | vitamin D signaling

Initially identified through its ability to interact directly with nuclear receptors in a ligand-dependent manner, transcriptional intermediary factor 1 (TIF1 α), also known as tripartite motif (TRIM24) protein, was subsequently described as both a negative and positive regulator of ligand-induced transactivation acting through chromatin modification (1–6).

To address the physiological functions of TIF1 α , we monitored large cohorts of TIF1 α -deficient (TIF1 α ^{-/-}) mice. Hepatic tumors were detected at necropsy in a vast majority of TIF1 α ^{-/-} mutants, thus indicating that TIF1 α deficiency predisposes to liver tumor formation. Tumor predisposition was not observed in TIF1 α ^{-/-} mutants heterozygous for the retinoic acid receptor α (*Rara*) gene, thereby providing genetic proof that TIF1 α and *Rara* act in opposition to each other in liver carcinogenesis (6).

In the present study, a systematic histological analysis of TIF1 α ^{-/-} mutants at different ages was carried out to examine the effect of TIF1 α gene deletion on a wide range of tissues. This analysis revealed that, aside from occasional metastases from liver tumors (6), TIF1 α ^{-/-} mutants displayed calcifications, increasing with age, in extra-hepatic connective tissues, namely arterioles, medium-sized arteries, lungs, and vibrissae. These ectopic calcifications were correlated with an increase in expression of several vitamin D direct targets, therefore raising the possibility that TIF1 α could repress the vitamin D receptor (VDR) signaling pathway *in vivo*.

Results

Calcifications of Renal Arteries and Arterioles in Young TIF1 α ^{-/-} Mutants. Tissues of 2-month-old TIF1 α ^{-/-} mutants were indistinguishable from those of age-matched WT littermates (data

not shown). In contrast, hematoxylin/eosin staining of kidney sections at 3 months of age revealed deposits of an acellular, basophilic (i.e., taking up hematoxylin), material (C; compare Fig. 1B with A) in about three-fourths of the TIF1 α ^{-/-} males and females, but never in age-matched WT littermates. These abnormal deposits were restricted to glomerular arterioles and medium-sized (i.e., hilar and arcuate) arteries and spared the veins that often bordered an affected artery. They contained calcium, stained in brown by the von Kossa method, (C; compare Fig. 1D and F with C and E), but did not take up the lipid-soluble dye oil red O and thus did not represent atherosclerotic plaques. No extrarenal ectopic calcification was detected in 3-month-old TIF1 α ^{-/-} mutants.

Extensive Calcifications of Arterioles, Arteries, Vibrissae, and Lungs in Old TIF1 α ^{-/-} Mutants. Histological sections of 8- to 10-month-old TIF1 α ^{-/-} mutants showed calcifications in arterioles and medium-sized (i.e., muscular) arteries with an incidence of 100% in kidney (Fig. 1I and J) and tongue (Fig. 1G), 6% in brown fat (Fig. 1H), 13% in snout dermis, 20% in heart, 30% in retina, and 65% in thyroid. It is noteworthy that these percentages underscore the actual extent of arterial calcifications as they were established through analysis of a single section of each organ. The fact that calcifications were detected in only 30% of histological sections through limb muscles, but were consistently observed in alizarin red-stained whole-mount preparations (C in Fig. 1M–O), supports this statement. We did not detect arterial calcifications in muscular arteries supplying liver, spleen, lung, and testis. Importantly, no calcification was ever detected in large (i.e., elastic) arteries including aorta and its major branches (data not shown), whereas calcified foci were scarce in major arteries supplying limbs (i.e., humeral and femoral arteries; e.g., Ah; Fig. 1N), in sharp contrast with their branches of smaller caliber (A in Fig. 1M–O). Altogether, these observations indicate that the extent of the pathological mineralization process is negatively correlated to the caliber of arteries and, consequently, that it spares the large vessels forming the bed of atherosclerotic plaques.

Calcifications in lungs and vibrissae affected 100% and 88% of the 8- to 10-month-old TIF1 α ^{-/-} mutants, respectively. In lungs, calcifications were confined to alveolar walls and displayed features characteristic of pulmonary alveolar microlithiasis (C in Fig. 2B and refs. 7 and 8). In vibrissae, mineralization was restricted to the capsules (CA in Fig. 2D, F, and H). Age-matched WT littermates did not show evidence of ectopic calcified deposit at any sites (Figs. 1L and 2A, C, E, and G),

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Table 1. Serum biochemical profiles in WT (*TIF1 α ^{+/+}*) and *TIF1 α ^{-/-}* mice (means \pm SD) at the age of 8 months

Factor	Female mice		Male mice	
	WT	<i>TIF1α</i>	WT	<i>TIF1α</i>
Glucose, mmol/liter	4.79 \pm 1.09	3.87 \pm 0.73	6.53 \pm 1.30	6.23 \pm 1.85
Phosphorus, mmol/liter	2.64 \pm 0.64	2.18 \pm 0.19	2.72 \pm 0.16	2.62 \pm 0.22
Calcium, mmol/liter	2.43 \pm 0.17	2.40 \pm 0.15	2.45 \pm 0.06	2.51 \pm 0.18
Urea, mmol/liter	9.84 \pm 2.34	8.27 \pm 1.54	10.30 \pm 2.39	10.10 \pm 1.19
Creatinine, μ mol/liter	43.50 \pm 3.78	39.67 \pm 3.27	43.25 \pm 7.59	44.33 \pm 4.84
Cholesterol, mmol/liter	2.61 \pm 0.65	2.14 \pm 0.37	3.13 \pm 0.50	3.45 \pm 0.35
Triglyceride, mmol/liter	1.52 \pm 0.37	1.65 \pm 0.64	2.16 \pm 0.88	2.31 \pm 0.25
Protein, g/liter	57.33 \pm 3.79	56.75 \pm 6.45	58.50 \pm 2.12	65.00 \pm 4.24
Free fatty acids, mEq/liter	1.54 \pm 0.39	1.43 \pm 0.1	1.01 \pm 0.18	1.92 \pm 0.13
ASAT, units/liter	192.67 \pm 90.43	148.83 \pm 49.81	164.50 \pm 67.27	182.83 \pm 131.69
ALAT, units/liter	51.00 \pm 20.07	62.67 \pm 31.62	83.50 \pm 64.42	76.33 \pm 37.33
LDH, units/liter	1,051.17 \pm 448.16	896.00 \pm 368.75	1,054.50 \pm 632.40	1,183.83 \pm 544.27

n = 5 mice of each genotype and gender. ASAT, aspartate aminotransferase; ALAT, alanine aminotransferases; LDH, lactate dehydrogenase.

suffering from glomerulosclerosis had normal circulating creatinine levels (Table 1) proved that their rate of glomerular filtration was not significantly decreased.

The Function of *TIF1 α* in Preventing Arterial Calcifications Is Not Exerted in Hepatocytes. Hepatocytes, which represent major targets of the *TIF1 α* -null mutation (6), are known to secrete inhibitors of calcium deposition in soft tissues (11–13). Thus, we investigated whether inactivating *TIF1 α* solely in hepatocytes would induce calcification of kidney arteries. To this end, mice harboring floxed alleles of *TIF1 α* (defined as *TIF1 α ^{L2/L2}* mice) were crossed with *Alb-Cre* transgenic mice that express Cre exclusively in hepatocytes to generate *TIF1 α ^{L2/L2} Alb-Cre* mice, in which *TIF1 α* is ablated in all hepatocytes at 6 weeks of age (*TIF1 α ^{hep-/-}* mice; ref. 6 and references therein). All 10 *TIF1 α ^{hep-/-}* mice analyzed at 3 months of age displayed the liver abnormalities characteristic of the *TIF1 α ^{-/-}* mutation in the germ line (i.e., in the whole organism) (6). In contrast, none of these *TIF1 α ^{hep-/-}* mutant mice displayed calcifications of kidney vessels (data not shown). This observation rules out the possibility that the ectopic calcification phenotype of *TIF1 α ^{-/-}* mutants is a systemic disease of liver origin.

***TIF1 α* Deficiency Increases the Expression of Vitamin D Target Genes Involved in Calcium Homeostasis in the Kidney.** As the kidney arterial bed is the earliest target organ of *TIF1 α* deficiency-induced calcifications, we next focused our investigations on expression, in kidney, of genes involved in calcium homeostasis. To this end, total kidney RNA of 3-month-old *TIF1 α ^{-/-}* mutant and WT littermates was analyzed by quantitative RT-PCR. In *TIF1 α ^{-/-}* mutants, mRNA levels of *Casr* (encoding a calcium-sensing receptor), *Cyp24a1* [encoding the 25-hydroxyvitamin D-24-hydroxylase that controls intracellular levels of 1,25(OH)₂D₃], *Trpv5* and *Trpv6* (encoding calcium-regulating ion channels), *Calb1* and *S100g* (encoding calcium-binding proteins), *Spp1* (encoding the secreted phosphoprotein 1, osteopontin), *Abcc6* (encoding a transmembrane transporter of the multidrug resistance protein family), *Pthlh* (encoding the parathyroid hormone-like peptide), and *Car2* (encoding carbonic anhydrase type 2) all were significantly up-regulated (Fig. 3A). On the other hand, genes expressed in WT kidneys and not implicated in calcium homeostasis, namely *Aldh1a3*, *Wt1*, and *Epo*, were not significantly altered in their expression levels in kidneys lacking *TIF1 α* (*P* > 0.05; Fig. 3A). Altogether, these results indicate that the *TIF1 α* -null mutation selectively increases, in kidney, the expression levels of numerous genes involved in calcium homeostasis. Among these, *Car2*, *Cyp24a1*,

Casr, *Trpv5*, *Trpv6*, *Calb1*, *S100g*, *Spp1*, and *Pthlh* are known to be direct vitamin D targets (refs. 14–18 and references therein). Therefore, their increased expression in *TIF1 α* -deficient kidneys indicates that *TIF1 α* down-regulates the vitamin D pathway in kidney.

***TIF1 α* Is Expressed in Tissues Exerting Crucial Functions in Calcium Homeostasis, but Is Undetectable in Endothelial Cells.** In *TIF1 α ^{-/-}* mutants, ectopic calcifications developed in contact with endothelial cells lining the lumen of arteries and arterioles, making up the bulk of the lung alveolar wall or forming the interface between capsules and blood sinuses of vibrissae. Endothelial cells therefore represented good candidates as primary targets of the *TIF1 α* null mutation. However, *TIF1 α* could not be detected in endothelial cells or in vascular smooth muscle cells by immunohistochemistry on sections of WT mice, whereas it was present in a variety of epithelial cells types (Fig. 3, Table 2, and SI Fig. 4). These data indicate that *TIF1 α* may act in tissues known to play critical roles in calcium homeostasis and expressing the VDR, namely renal tubules and glomeruli (T and G in Fig. 3B and C), intestinal epithelium, and parathyroid gland parenchyma (Table 2, SI Fig. 4, and refs. 19 and 20).

Discussion

In Addition to Its Role as a Liver Tumor Suppressor, *TIF1 α* Is Instrumental in Preventing Ectopic Calcifications and Other Features of Premature Aging. The present results show that, in addition to hepatic tumors (6), *TIF1 α ^{-/-}* mice spontaneously develop pathologic calcifications in arterial vessels, lungs, and vibrissae. Arterial calcifications first appear in the walls of medium-sized arteries and arterioles in the kidney by the age of 3 months and, with aging, they extend to other muscular arteries and become associated with calcifications of lung alveoli and vibrissae capsules. Arterial calcifications vary in prevalence among different organs of *TIF1 α ^{-/-}* mutants. Liver, spleen, lung, and testis appear relatively resistant to the process, whereas kidney is most vulnerable. Such variation argues against a change in serum pH or ion content as solely responsible for the pathological calcified deposits and favors at least a partial influence from local factors such as vascular dynamics or local differences in endothelial cell metabolic activity.

In mice, arterial calcification is a classical age-related lesion (21), which affects *TIF1 α ^{-/-}* mutants already 3 months old of age. In this context, it is noteworthy that *TIF1 α ^{-/-}* mice exhibit other features of premature aging. Among these, lung microolithiasis is present in all *TIF1 α ^{-/-}* mice as early as 8 months of age while affecting a maximum of 17% WT individuals >25 months

Table 2. TIF1 α is detected in a large variety of epithelia

Epithelia	Signal	Figure
Cardiovascular system		
Myocardium	0	
Arteries and veins (endothelium and smooth muscles)	0	Fig. 3C
Capillaries (endothelium)	0	
Respiratory system		
Tracheal epithelium and cartilage	0	
Alveolar epithelium	0	
Digestive system		
Esophagus (epithelium)	+	
Stomach (epithelium of fundic glands)	+	
Ileum (epithelium)	+	
Colon (epithelium)	+	SI Fig. 4D
Liver parenchyma	+	
Pancreas (exocrine)	+	SI Fig. 4E
Urinary system		
Kidney glomeruli (Bowman's capsules)	+	Fig. 3C
Kidney tubules	+	Fig. 3B and C
Urinary bladder epithelium	+	
Genital system		
Testis		
Sertoli cells	++	SI Fig. 4A
Spermatocytes	0 (PR) to ++ (D)	
Round spermatids	+++	
Leydig cells	0	
Epididymis (epithelium of the head)	++	SI Fig. 4C
Epididymis (epithelium of the tail)	0	
Vas deferens (epithelium)	++	
Cranial prostate (epithelium)	++	
Granulosa cells	+	SI Fig. 4I
Oocytes	0	
Oviduct (epithelium)	++	SI Fig. 4K
Uterus (epithelium and glands)	++	SI Fig. 4J
Uterus (stroma)	+	SI Fig. 4J
Endocrine glands		
Pancreas (Langerhans cells)	+	SI Fig. 4E
Parathyroid glands	+	SI Fig. 4G
Thyroid gland	0	
Adrenal gland (cortex and medulla)	+	
Miscellaneous connective tissues		
White and brown adipose tissue	0	
Striated muscle (oesophagus)	0	
Skin (epidermis and dermis)	0	

Connective tissues are not stained with the anti-TIF1 α antibody with the notable exception of the uterine stroma. Note that tissues of age-matched TIF1 α ^{-/-} mutants were used as negative controls. PR and D, preleptotene and diplotene spermatocytes, respectively. 0, absence of nuclear immunofluorescent signal. +, ++, and +++ indicate increasing intensities of nuclear immunofluorescent signals.

Kidney is a key target organ of the vitamin D endocrine system, and both vitamin D deficiency and VDR ablation lead to impaired renal functions (refs. 43 and 45 and references therein). In this context, it is noteworthy that (i) the *Casr* promoter is known to contain functional vitamin D responsive elements, (ii) *Casr* mRNA levels in kidney are increased upon vitamin D administration and decreased in VDR-null mutant mice (15, 45), and (iii) TIF1 α , VDR, and *Casr* all are expressed in distal convoluted tubules (refs. 20 and 46 and Fig. 3B and C). We also show that, in addition to *Casr*, expression of other direct vitamin D target genes, including *Car2*, *Cyp24a1*, *Trpv5*, *Trpv6*, *Calb1*, *S100g*, *Spp1*, and *Pthlh*, is up-regulated in kidney upon TIF1 α ablation. Altogether, these data indicate that TIF1 α normally functions to repress the vitamin D endocrine system and additionally suggest that ectopic calcifications in TIF1 α ^{-/-} mice are causally related to a disturbed VDR signaling pathway. Interestingly, the fact that these metabolic disturbances correlate with

a calcifying arteriopathy and other features of premature aging in TIF1 α ^{-/-} mice provides support for the hypothesis that aging is promoted by an increased activity of the vitamin D signaling pathway (26, 27).

In conclusion, the present data demonstrate that TIF1 α plays an indispensable role in regulating a molecular pathway involving VDR and *Casr*, which functions to prevent ectopic calcifications *in vivo*.

Materials and Methods

Mice. All mice were on a mixed C57BL/6–129/Sv genetic background and housed in an animal facility licensed by the French Ministry of Agriculture (agreement B67-218-5, 1999-02-09). Animal experiments were supervised by M.M. who is qualified for experimenting with mice (French Ministry of Agriculture agreement 67-62, 2003-05-30).

Histology, Alizarin Red-Stained Whole-Mount Preparations, and Immunohistochemistry. Tissues from TIF1 α ^{-/-} mutants and WT littermates were collected at 2, 3, and 8–10 months of age. Tissues were fixed in 4%

(wt/vol) buffered formalin for 24 h, then either embedded in paraffin or frozen in liquid nitrogen vapors (47). Histological sections from paraffin-embedded tissues were stained either with hematoxylin/eosin or processed using the von Kossa method for localizing calcium histochemically (48). In the latter staining method, nuclei were counterstained either with hematoxylin/eosin, safranin O, or DAPI, and the DAPI fluorescence was then converted into a bright-field image with Photoshop (Adobe). Organs that were systematically analyzed included: heart, aorta white fat and interscapular brown fat, striated muscle, liver, trachea, lung, kidney, testis, eye and adnexia, thyroid gland tongue spleen, and skin. Oil red O staining of frozen histological sections was performed as described (47). Histological observations were repeated on at least eight males and eight females per genotype and age group. For alizarin red staining of mouse carcasses (comprising bones, striated muscles, and muzzle skin), mice were killed, eviscerated, fixed in ethanol, cleared in KOH, and stained with alizarin red (see *SI Text*). Immunohistochemistry with the monoclonal anti-TIF1 α antibody 5T1E8 (1) was performed according to standard protocols (see *SI Text*).

RNA Isolation, Reverse Transcription, and Real-Time PCR Analysis. Total RNA from whole kidneys was isolated by using Rneasy (Qiagen), and 5 μ g was used

for cDNA synthesis primed with oligo(dT)₂₄ (Invitrogen). The final product was then diluted 80 times, and 4 μ l was mixed with the forward and reverse primers listed in *SI Table 4* (250 nM of each primer at final concentration) and 5 μ l of SYBR Green master mix. Real-time PCR was performed with the LightCycler 1.5 system (Roche). Each cDNA sample was tested in triplicate, and the expression level of each gene was normalized to the hypoxanthine-guanine phosphoribosyltransferase level.

Blood Sample Analyses. Blood collection and establishment of biochemical and hematological parameters were as described (49).

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