

REVIEW

Placental membrane-type metalloproteinases (MT-MMPs): Key players in pregnancy

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ABSTRACT

Membrane-type matrix metalloproteinases (MT-MMPs) are a sub-family of zinc-dependent endopeptidases involved in the degradation of the extracellular matrix. Although MT-MMPs have been mainly characterized in tumor biology, they also play a relevant role during pregnancy. Placental MT-MMPs are required for cytotrophoblast migration and invasion of the uterine wall and in the remodeling of the spiral arteries. They are involved in the fusion of cytotrophoblasts to form the syncytiotrophoblast as well as in angiogenesis. All these processes are crucial for establishing and maintaining a successful pregnancy and, thus, MT-MMP activity has to be tightly regulated in time and space. Indeed, a de-regulation of MT-MMP expression has been linked with pregnancy complications such as preeclampsia (PE), fetal growth restriction (FGR), gestational diabetes mellitus (GDM) and was also found in maternal obesity. Here we review what is currently known about MT-MMPs in the placenta, with a focus on their general features, their localization and their involvement in pregnancy disorders.

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Introduction

The extracellular matrix (ECM) is a plastic matrix giving structure and grounding for the 3 dimensional organization of tissues. ECM is involved in multiple aspects of cell function including cell proliferation, differentiation, adhesion, migration and invasion.¹ Its degradation and remodeling is central for structural and developmental changes. Thus, the establishment of pregnancy as well as embryonic and fetal development requires ECM degradation to allow implantation, placental development, angiogenesis and parturition.² Vice versa, ECM composition modulates these processes. ECM degradation is tightly regulated since imbalances lead to pregnancy complications.³

Several proteases are involved in ECM degradation during pregnancy. These include serine proteases, cathepsins and matrix metalloproteinases (MMPs).⁴ MMPs are a family of 24 zinc dependent endopeptidases capable of degrading virtually all ECM components. They have been classified into 5 groups: collagenases, gelatinases, stromelysins, membrane-type MMPs (MT-MMPs) and other MMPs.^{5,6} This review will focus on MT-MMPs, a subgroup of 6 membrane anchored MMPs: MT1-MMP (MMP14), MT2-MMP (MMP15), MT3-MMP

(MMP16), MT4-MMP (MMP17), MT5-MMP (MMP24) and MT6-MMP (MMP25).

Because of their key role in ECM degradation, various MMPs have been studied in depth regarding their function in pregnancy, but the majority of this work has focused on MMP2 and MMP9.^{7–9} In contrast to secreted MMPs, MT-MMPs are membrane anchored and thus, allow a directed and spatially regulated mode of action. MT-MMPs have been shown to be required for tumor proliferation, invasion and angiogenesis,¹⁰ processes that are also taking place in pregnancy, although tightly regulated. However, despite their function in ECM degradation, little is known about the role of MT-MMPs during pregnancy. This review summarizes the function of placental MT-MMPs, focusing on their localization, regulation and their involvement in the pathophysiology of pregnancy.

MT-MMPs: General features

MT-MMP structure

All members of the MMP family share a similar structure differing mainly in their domain organization. These

domains include a pro-domain, a catalytic domain, a hinge region and a hemopexin domain.¹¹

MT-MMPs, which are inserted in the membrane, can be further classified into 2 groups: i) MT-MMPs that are anchored by a transmembrane domain followed by a cytoplasmic domain (MT1-, MT2-, MT3- and MT5-MMP), and ii) MT-MMPs that are anchored by a glycosylphosphatidylinositol (GPI)-anchor and which lack a cytoplasmic domain (MT4- and MT6-MMP).¹² Figure 1 shows the domain assembly of MT-MMPs.

MT-MMPs can function both as monomers or homodimers.^{13,14} Interestingly, homodimerization has been shown to play an important role in the regulation of MT-MMP activity, e.g. in MMP-2 activation mediated by MT1- and MT2-MMP (section 2.3.).

Although ECM degradation by the catalytic domain is the primary function of MMPs, other functions have been attributed to the rest of the MT-MMP domains. For instance, the hemopexin domain is involved in protein-protein interactions, allowing MT1-MMP oligomerization.¹⁵ The cytoplasmic domain can interact with other intracellular proteins, transducing information from the extracellular environment.¹⁶ Furthermore, it is thought to regulate MT-MMP intracellular trafficking, degradation and surface distribution.¹⁰ The GPI-anchor in MT4- and MT6-MMP is involved in signal transduction as well as in lipid raft localization, and confers their sensitivity to phospholipases.¹⁷

MT-MMP activation

MMPs are secreted as inactive zymogens with the protease inhibiting pro-domain located at the C-terminus. The pro-domain contains a cysteine residue that interacts with a zinc ion in the catalytic domain. Disruption

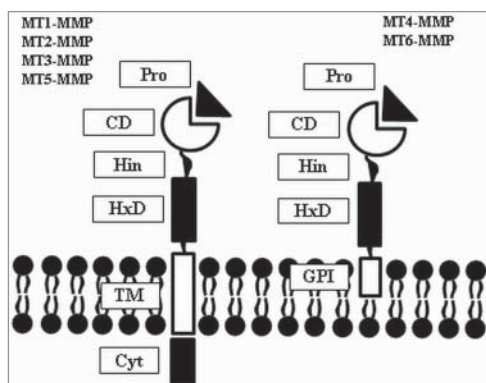


Figure 1. MT-MMP structure. MT-MMPs are anchored to the membrane either via transmembrane domain (MT1-, MT2-, MT3- and MT5-MMP) or via GPI-anchor (MT4- and MT6-MMP). Cyt, cytoplasmic domain; TM, transmembrane domain; GPI, glycosylphosphatidylinositol; HxD, hemopexin domain; Hin, hinge region; CD, catalytic domain; Pro, pro-domain.

of this interaction is required for MMP activation. This occurs in a 2-step process. First, the pro-domain is cleaved in a sequence-specific manner. Then, the intermediate MMP product as well as other proteases can completely remove the pro-domain, resulting in a fully-activated MMP.¹⁸ Additionally, MMP activation can also result from a conformational change of the pro-domain, or by activation by reactive oxygen species, which interact with the free cysteine in the pro-domain.¹⁹

In contrast to the general activation process of MMPs, MT-MMPs and MMP11, 21 and 28 contain a recognition sequence for Golgi-associated pro-protein convertases, i.e. RXXKR, with R=Arg, K=Lys and X=non basic amino acid, and are thus activated in the Golgi network prior to secretion.¹⁹⁻²¹ MT-MMP activation might also occur in 2 steps, since MT1-MMP activation requires a first cleavage in the pro-domain prior to the second cleavage mediated by pro-protein convertases.²² Once in the plasma membrane, active-MT1-MMP can undergo autocatalysis generating an inactive cleavage product, which regulates the pool of active-MT1-MMP present on the cell surface by inhibiting active-MT1-MMP endocytosis.^{23,24} The membrane anchored MT1-MMP can also be cleaved and shed. Non-autocatalytical shedding results in a soluble active MT1-MMP.²⁵ This shedding of an active MT1-MMP species produces a soluble enzyme and allows MT1-MMP action beyond the surface of the MT1-MMP producing cells.²⁶

MT-MMP activity

MT-MMPs accept a wide array of substrates (Table 1). They degrade various ECM components, including those found in the uterine wall and the spiral arteries, such as fibronectin, vitronectin, collagen IV and laminin.²⁷ Furthermore, MT-MMPs can cleave pro-forms of cytokines including TGF- β and TNF- α , resulting in their activation.²⁸⁻³⁰ In addition to their regulation of the immune response, these cytokines also modulate some placental functions. TGF- β has been shown to inhibit trophoblast invasion.³¹ TNF- α was able to induce MT1- and MT2-MMP expression in the human first trimester trophoblast cell line ACH-3P and in placental explants from the first trimester.³² However, several studies have also shown that TNF- α limits trophoblast invasion by increased secretion of plasminogen activator inhibitor (PAI1). PAI1 is the principal inhibitor of plasminogen activator (tPA) and urokinase (uPA), both activators of fibrinolysis.^{33,34} This mechanism would reduce ECM degradation independent of the MMP/TIMP system.

Besides ECM degradation and cytokine activation, MT-MMPs can also inactivate cytokines,³⁵ activate other MMPs such as MMP2, 8, 9 and 13^{30,36} as well as other

Table 1. Major substrates of MT-MMP (adapted from ref. 18 with additional data from refs. 10,12,16,17,36,50. *Contradictory data showing that MT4-MMP and MT6-MMP are able/not able to activate pro-MMP2.¹⁷

MT-MMPs	ECM substrates	Non-ECM substrates
MT1-MMP (MMP14)	Aggrecan, collagen I, II, III, dermatan sulfate proteoglycan, entactin, fibrin, fibrillin, fibrinogen, fibronectin, gelatin, laminin-1, -2/4, -5, lumican, nidogen, perlecan, tenascin, vitronectin	α_2 -Macroglobulin, α_1 -PI, $\alpha_v\beta_3$ integrin, β -glycan δ -like 1, ADAM9, ApoE, CD44, CXCL12, death-receptor 6, EMPRIN, factor XII, ICAM-1, IL-8, KiSS-1, LRP1, MCP-3, myelin-inhibitory protein, progranulin, pro-MMP2, pro-MMP8, pro-MMP13, pro-TGF- β , pro-TNF- α , RANKL, syndecan, tissue transglutaminase
MT2-MMP (MMP15)	Aggrecan, collagen I, IV, entactin, gelatin, fibrin, fibronectin, laminin-1, nidogen, perlecan, tenascin	pro-MMP2, pro-MMP13, pro-TNF- α , tissue transglutaminase
MT3-MMP (MMP16)	Aggrecan, collagen III, fibronectin, gelatin, laminin-1, vitronectin	α_2 -Macroglobulin, α_1 -PtdIns, casein, pro-MMP2, pro-MMP13, syndecan, tissue transglutaminase
MT4-MMP (MMP17)	Fibrin, fibrinogen, gelatin	ADAMTS4, KiSS-1, pro-TNF- α , pro-MMP2*
MT5-MMP (MMP24)	Chondroitin sulfate proteoglycan, dermatan sulfate proteoglycan, gelatin, fibronectin, plasminogen	pro-MMP2, pro-MMP13, tissue transglutaminase
MT6-MMP (MMP25)	Chondroitin sulfate proteoglycan, collagen IV, dermatan sulfate proteoglycan, fibrin, fibrinogen, fibronectin, gelatin, laminin	α_1 -PI, CCL2, CCL7, CCL13, CCL15, CCL23, CXCL2, CXCL5, CXCL12, pro-MMP2*, pro-MMP9, urokinase plasminogen activator receptor, vimentin

proteases such as ADAM9 (a disintegrin and metalloproteinase 9)¹² and ADAMTS4 (ADAM with thrombospondin motifs 4).³⁷

Pro-MMP2 activation by MT1-MMP has been thoroughly studied, revealing a complex mechanism where TIMP2 (tissue inhibitor of MMPs 2), a natural MMP inhibitor, is also required. Once located on the membrane, MT1-MMP forms a homodimer. One MT1-MMP of this dimer binds TIMP2, which will subsequently bind pro-MMP2.²³ Then, the other MT1-MMP molecule of the dimer can cleave pro-MMP2 and produce active MMP2. This mechanism is dependent on TIMP2 concentration and occurs only at basal TIMP2 levels. High TIMP2 levels inhibit both, MMP2 and MT1-MMP.³⁸ Interestingly, MT2-MMP can activate pro-MMP2 without interaction with TIMP2. This mechanism also involves an MT2-MMP dimer, of which one MT2-MMP molecule binds pro-MMP2 directly, followed by activation by the other MT2-MMP molecule.³⁹ Contradictory data regarding MMP2 activation have been reported for the other 3 MT-MMP members.¹⁷

MT-MMP regulation

It has been hypothesized that the different members of the MMP family originate from gene duplication, as they are widely spread across several chromosomes.²¹ However, promoter architecture highly differs among distinct MMPs, which allows discriminative regulation at the level of gene expression.

Based on basic promoter conformation, i.e., the distribution of the cis-elements along the promoter, MMPs are classified into 3 different groups: i) MMPs containing a TATA box at -30 bp and a proximal binding site for the transcription factor activator protein-1 (AP-1); ii) MMPs containing a TATA box but no proximal AP-1 binding

site; and iii) MMPs lacking both, TATA box and AP-1 binding site.^{40,41}

MT-MMPs belong to the third group, i.e., they lack a TATA-box and an AP-1 binding site, with the exception of MT2-MMP, which contains a TATA box. Thus, lack of common MMP regulatory elements in the promoter regions of MT-MMPs might allow their regulation different from other MMPs. Moreover, the basic promoter conformation represents only one determining aspect of the regulation of MMP expression. The majority of the regulatory elements in MMP promoters are non-canonical, which might also explain the finely tuned regulation with a different set of MMPs being activated in various cell types and at different developmental stages.

Finally, MMP expression is also regulated by cytokines, hormones and growth factors, adding a new level of complexity. Several interleukins such as IL-1, IL-6 and IL-15 have been shown to promote trophoblast invasion, whereas IL-10 inhibits this process.⁴ The role of placental hormones, e.g. human chorionic gonadotropin, in this regard remains still contradictory.⁴² We previously reported that insulin-like growth factors (IGFs) as well as insulin increase the expression of MT1-MMP in isolated first trimester trophoblast.⁴³ Whether this entails invasion promotion remains to be studied.

Inhibition of MT-MMP activity could be considered as the last level of regulation of ECM degradation. Plasma proteins such as α_2 -macroglobulin are thought to be broad-range inhibitors of endopeptidases, including MMPs. However, α_2 -macroglobulin action has been mainly found in tissue fluids. Thus, its relevance for MT-MMP inhibition remains unclear.⁴⁴

By contrast, tissue inhibitors of MMPs (TIMPs) are proteins specifically targeting MMPs. Four members of the TIMP family have been described, which bind to the catalytic domain of MMPs in a 1:1 molar ratio. The zinc ion in the MMP catalytic domain is then chelated, which

results in MMP inhibition.⁴⁵ TIMPs are broad-range MMP inhibitors affecting MMPs in general, but with different preference and efficiency. For instance, TIMP1 is a weak inhibitor of MT1-, MT2-, MT3- and MT5-MMP.⁴⁶ Furthermore, different TIMP members can fulfill different functions: As mentioned above, low levels of TIMP2 are even required for MMP2 activation by MT1-MMP. TIMP4 also interacts with MT1-MMP and MMP2, but this interaction results in inhibition of both MMPs.^{47,48} Thus, the balance between the different TIMP members can also fine tune MMP activity.

Placental MT-MMPs and their relevance during pregnancy

Table 2 summarizes the cellular and tissue distribution of MT-MMPs in human placenta and decidua in the course of pregnancy.

MT-MMPs during the first trimester of pregnancy

The placenta is a fast developing and growing villous organ. Specialized placental cells, the cytotrophoblasts,

undertake various processes essential for placental function and pregnancy success. Three different cytotrophoblast subpopulations can be distinguished: the villous cytotrophoblasts (VTs), which can fuse and form the syncytiotrophoblast (ST), a syncytium that represents the classical placental barrier, transporting maternal nutrients to the fetal circulation and producing pregnancy hormones. Finally, the extravillous cytotrophoblasts (EVTs), which migrate from tips of some specialized placental villi, invade the maternal decidua and anchor the placenta in the uterus. This process predominantly takes place in the first trimester of pregnancy.⁴ Some EVT's also reach the uterine spiral arteries and remodel them into wide, low resistance vessels, allowing an adequate blood supply to the fetus.⁴⁹ All these processes, i.e. trophoblast fusion, invasion and the remodeling of the spiral arteries, require ECM degradation, pointing out once more the relevance of MMPs in pregnancy. However, due to their involvement in the process of trophoblast invasion, MMP expression has been mainly characterized in first trimester trophoblasts.

Because of their membrane anchor, MT-MMPs can be located to specific membrane regions, where they

Table 2. MT-MMP localization in human placenta and decidua.

MT-MMPs	Cell type/Tissue	Trimester	References
MT1-MMP (MMP14)	Feto-placental endothelial cells	1 st	59
	Syncytiotrophoblast (ST)	1 st	58,59
	Villous cytotrophoblasts (VTs)	1 st	43,52,58,59,92
	Extravillous cytotrophoblasts (EVTs)	1 st	53-56,58-60
	Cell columns	1 st	57,60
	Perivascular EVT's	1 st	57,58
	Decidua	1 st	60,64,65,91
	Decidua	2 nd	64
	Feto-placental endothelial cells	3 rd	59,62
	ST	3 rd	59,62,66
	VTs	3 rd	59,62
	Decidua	3 rd	64
	Amniochorion	3 rd	69
MT2-MMP (MMP15)	VTs	1 st	43
	EVTs	1 st	53,54,60,61
	Decidua	1 st	64,65,91
	Decidua	2 nd	64
	ST	3 rd	3,67
	Decidua	3 rd	64
	Amniochorion	3 rd	70
MT3-MMP (MMP16)	Decidua	1 st	64,65,91
	Decidua	2 nd	64
	ST	3 rd	3,68
MT4-MMP (MMP17)	Decidua	3 rd	64
	Decidua	1 st	64
	Decidua	2 nd	64
MT5-MMP (MMP24)	Feto-placental endothelial cells	3 rd	68
	Decidua	3 rd	64
	Amniochorion	3 rd	70
	Decidua	1 st	64,65,91
	Decidua	2 nd	64
MT6-MMP (MMP25)	ST	3 rd	93
	Decidua	3 rd	64
	Amniochorion	3 rd	70
	ST	3 rd	68
	Amniochorion	3 rd	70

play an active role in pericellular proteolysis,⁵⁰ a feature required in the spatially directed process of trophoblast invasion. Indeed, in first trimester trophoblasts MT1-MMP is localized in invadopodia, a structure at the cellular edge of ECM degradation and invasion progression.⁵¹ MT1-MMP expression in cytotrophoblasts is maintained through the first trimester of pregnancy.⁵² This suggests that, together with its role as pro-MMP2 activator, MT1-MMP may be involved in trophoblast invasion. In fact, MT1-MMP expression has been observed in EVT^s⁵³⁻⁵⁶ and in perivascular EVT^s, i.e. EVT^s migrating into the spiral arteries.^{57,58} Blocking of MT1-MMP with neutralizing antibodies reduced migration of a first trimester trophoblast cell line.⁵⁹

Among the other members of the MT-MMP family, only MT2-MMP is expressed in isolated first trimester trophoblasts at levels similar to MT1-MMP.^{43,60} MT2-MMP is predominantly expressed in EVT^s, with weaker expression in VT^s.^{54,61} Thus, a role of MT2-MMP in trophoblast invasion is also likely.

However, besides trophoblast invasion, also other functions of placental development may involve MT-MMPs in the first trimester of pregnancy. MT1-MMP is produced in fetoplacental endothelial cells forming the first placental vessels.⁶² In fact, MT1-MMP is a key player in matrix degradation during angiogenesis,⁶³ and blocking of MT1-MMP reduces the network formation potential of human fetoplacental endothelial cells isolated from full term placentas.⁶² This suggests a role of MT1-MMP in neovascularisation or vascular development of the early placenta.

In addition to its location in EVT and endothelial cells with both cell types performing directed migration during trophoblast invasion and angiogenesis, respectively, MT1-MMP is also expressed in VT^s and ST.^{58,59} Here, MT1-MMP has been observed to promote VT proliferation as well as their fusion to ST *in vitro*.⁵⁹

All MT-MMP members, except MT6-MMP, are expressed in first trimester decidua. MT1- and MT4-MMP are expressed in EVT^s reaching the decidua, but also in decidual fibroblasts and uterine natural killer cells, 2 cell types controlling the depth of trophoblast invasion. MT2-MMP is predominantly produced in cytotrophoblasts, whereas MT3- and MT5-MMP are mainly located in decidual stromal cells.⁶⁴ When MT1-, MT2-, MT3- and MT5-MMP expression was studied in the decidual secretory endometrium, in decidua parietalis and in decidua basalis, all 4 MT-MMPs were detected in all parts of the decidua, as well as in the syncytiotrophoblast and EVT^s reaching the decidua basalis.⁶⁵

Placental MT-MMPs during the second trimester of pregnancy

For obvious reasons, sampling during the second trimester of pregnancy is mainly based on non-invasive techniques such as serum screening and ultrasound. Therefore, in this period placental MT-MMPs remain uncharacterized. All MT-MMP members except MT6-MMP were detected in second trimester decidua.⁶⁴

Placental MT-MMPs during the third trimester of pregnancy

All MT-MMP members are present in the third trimester placenta.^{3,66-68} This is especially surprising for MT6-MMP, which is not expressed in the placenta during the first trimester of pregnancy. In the third trimester, MT1-MMP is expressed in several placental cell types including the syncytiotrophoblast, the underlying VT^s and the endothelium of the fetoplacental vessels.^{59,62,66} As mentioned above, *in vitro* experiments suggest a role of MT1-MMP in proliferation and fusion of VT^s as well as in fetoplacental angiogenesis.^{59,62} MT2-, MT3-, MT5- and MT6-MMP have been localized in the syncytiotrophoblast,^{67,68} with MT2-, MT3- and MT5-MMP being also present in VT^s.³ By contrast, MT4-MMP expression was predominantly observed in fetal vessels.⁶⁸

Similar to their expression in the first and the second trimester of pregnancy, all MT-MMP members except for MT6-MMP are expressed in the decidua at term.⁶⁴ MT-MMPs have also been characterized in fetal membranes after delivery. With the only exception of MT3-MMP, all MT-MMPs are expressed in the amniochorion.^{69,70}

Changes in placental MT-MMP expression during pregnancy

Another aspect regarding placental MT-MMPs that has been insufficiently documented is the dynamic in their expression during the course of pregnancy. In a microarray study⁷¹ several genes involved in different biological processes such as cell proliferation, differentiation and angiogenesis were differentially regulated in first and third trimester human placenta, with most genes involved in these processes, including MT1-MMP but not MT2-MMP, being higher expressed in first trimester placenta.

In another microarray study we have previously analyzed MT-MMP gene expression in cytotrophoblasts from first trimester placenta (FT) and showed that only MT1- and MT2-MMP could be detected in FT.⁴³ Here we present novel RT-PCR data comparing MT-MMP

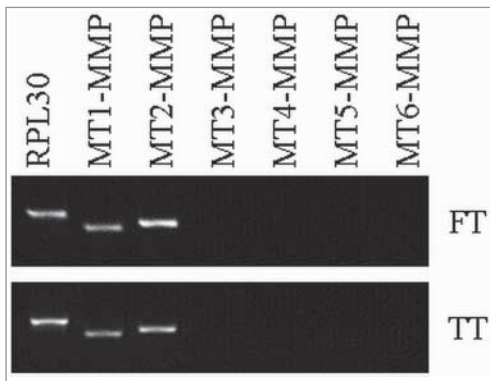


Figure 2. RT-PCR analysis of MT-MMP expression in primary trophoblasts isolated from first trimester placenta (FT) vs. third trimester placenta (TT). For the housekeeping gene RPL30 24 cycles were used, for all MT-MMP 27 cycles.

expression between cytotrophoblasts isolated from first (FT) and third trimester placenta (TT) by tissue digestion, Percoll gradient centrifugation and immunopurification^{72,73} (Fig. 2). Of all MT-MMPs, only MT1-MMP and MT2-MMP were expressed in both FT and TT, with MT2-MMP expression being higher in FT (Fig. 2). This expression change underlines the functional difference of FT and TT, since in contrast to FT, TT cannot fulfill endovascular remodeling both *in vivo* and *in vitro*.⁷⁴ Interestingly, we were not able to detect MT3- and MT5-MMP in TT, despite its previously reported presence³ (Table 2). This may in part be due to differences between mRNA and protein, or due to differences in the trophoblast subpopulations investigated. While freshly isolated trophoblasts mainly resemble cytotrophoblast cells, Zhu *et al.* investigated whole placental tissue and identified the syncytiotrophoblast as major MT3- and MT5-MMP producing site.

Placental MT-MMPs in pregnancy complications

It is well established that severe pregnancy complications associated with shallow or impaired trophoblast invasion and spiral artery remodeling, such as preeclampsia (PE) and fetal growth restriction (FGR), have their origin in early pregnancy.⁴⁹ However, PE and FGR clinically manifest only during the second trimester.⁷⁵ Therefore, the relationship of MMPs with PE and FGR has been mainly analyzed in third trimester placentas from preterm and term pregnancies.

Investigating chorionic villous biopsies provides the possibility to study placental tissue in the period of potential functional failure. Huisman *et al.*⁷⁶ found no differences in MMP2 and MMP9 activity in first trimester placenta from pregnancy subsequently developing PE

or FGR, but the study was limited by the small sample size of the groups.

Also from the studies investigating placenta from third trimester, the majority of analyses focused on MMP2 and MMP9.⁷⁷⁻⁷⁹ Few publications have studied the role of MT-MMPs in FGR. We have previously shown a decrease in MT1-MMP mRNA and protein levels in FGR placentas when compared to gestational age matched healthy controls.⁵⁹ This change in FGR may be limited to MT1-MMP, because MT2-, MT3- and MT5-MMP are unaltered in FGR.³ However, more data are required to draw firm conclusions.

In PE placental MT6-MMP is down-regulated,⁶⁸ whereas MT2-MMP^{67,80} and MT4-MMP⁶⁸ are up-regulated. Since MT4-MMP is involved in angiogenesis, its up-regulation might be a compensatory mechanism to overcome poor blood supply to the fetus with placental hypervascularisation.⁶⁸ The role of MT-MMPs in the pathology of PE may involve their function in cleavage and thus activation or inactivation of circulating substrates. For instance, soluble endoglin (sEng), a soluble form of the TGF- β receptor, is increased in PE.⁸¹ Both sEng and MT1-MMP have been localized in the ST of third trimester placenta. Indeed, although its expression is not upregulated in pre-eclamptic placentas, MT1-MMP has been reported to cleave endoglin.⁶⁶

Thus, contrasting the concept that poor placentation may relate to lower MMP expression, some MT-MMP members are up-regulated in PE and FGR during the third trimester of pregnancy. This might represent a compensatory mechanism to overcome poor placentation due to MT-MMP down-regulation in the first trimester of pregnancy.⁸²

Diabetes in pregnancy is associated with both an increased risk for PE⁸³ and FGR, suggesting placental invasion failure, as well as excessive fetal fat accretion and fetal overgrowth. Furthermore, the placenta in diabetes is characterized by hypervascularization.⁸⁴ Accordingly, MT1-MMP imbalances have been described in diabetes. MT1-MMP is increased in first trimester placentas of Type 1 diabetic women, and upregulation of trophoblast MT1-MMP by insulin, IGF-1, IGF-2 and TNF- α may underlie this observation.⁴³ Gestational diabetes mellitus (GDM) is a glucose intolerance with first clinical manifestation in the second trimester. We showed an up-regulation of MT1-MMP protein levels in third trimester placenta as a result of GDM in lean women. The upregulation of MT1-MMP by insulin and IGF-2 in endothelial cells accounts for the increase.⁶² In contrast, MT1-MMP is downregulated in GDM placentas of obese women.⁸⁵ This suggests a complex interplay between diabetes-associated and obesity-associated

factors to ultimately determine placental MT1-MMP levels in the third trimester of pregnancy.

Indeed, maternal obesity is associated with an increased risk of preterm delivery, PE and fetal death⁸⁶ and might also play a role in the regulation of placental MT-MMPs. In a rat model mimicking maternal obesity, trophoblast invasion and blood vessel remodeling is altered.⁸⁷ Leptin, a classical hormone upregulated in obesity and involved in placental lipid metabolism,⁸⁸ can also modulate placental ECM molecule expression⁸⁹ and up-regulates MT1-MMP expression in human EVT's.⁹⁰ However, more studies are required to identify the obesity-related factors regulating MT-MMPs.

MT-MMP expression is also altered in pregnancy disorders leading to miscarriage. Decidual MT2- and MT5-MMP expression is increased in first trimester decidua of spontaneous abortions,⁹¹ whereas absence of MT4-MMP in amniochorion has been described in preterm premature rupture of the membranes (pPROM) when compared with term labor placenta.⁷⁰

Conclusion

In the present review we have summarized current knowledge about MT-MMPs with a focus on the human placenta. Their location, functions and involvement in pregnancy complications highlight the crucial role of MT-MMPs for a successful pregnancy. However, it is obvious that MT-MMPs are understudied and more research is required to completely understand their role in placental implantation and trophoblast invasion. Additional studies addressing MT-MMP activity and function in PE and FGR could help in finding diagnostic markers for these pregnancy complications.

Disclosure of potential conflicts of interest

No potential conflicts of interest were disclosed.

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Authors contributions

AM reviewed the literature and wrote the first draft. UH modified the first draft and critically discussed and reviewed the manuscript. NG, UL, GD and MD critically discussed and reviewed the manuscript.

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