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Novel Pathways of Inflammation in Human Fetal Membranes Associated with Preterm Birth and Preterm Pre-labor Rupture of the Membranes

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Abstract

Spontaneous preterm birth (PTB) and preterm pre-labor rupture of the membranes (pPROM) are major pregnancy complications. Although PTB and pPROM have common etiologies, they arise from distinct pathophysiologic pathways. Inflammation is a common underlying mechanism in both conditions. Balanced inflammation is required for fetoplacental growth; however, overwhelming inflammation (physiologic at term and pathologic at preterm) can lead to term and preterm parturition. A lack of effective strategies to control inflammation and reduce the risk of PTB and pPROM suggest that there are several modes of the generation of inflammation which may be dependent on the type of uterine tissue. The avascular fetal membrane (amniochorion), which provides structure, support and protection to the intrauterine cavity, is one of the key contributors of inflammation. Localized membrane inflammation helps tissue remodeling during pregnancy. Two unique mechanisms that generate balanced inflammation are the progressive development of senescence (aging) and cyclic cellular transitions: epithelial to mesenchymal (EMT) and mesenchymal to epithelial (MET). The intrauterine build-up of oxidative stress at term or in response to risk factors (preterm) can accelerate senescence and promote a terminal state of EMT, resulting in the accumulation of inflammation. Inflammation degrades the matrix and destabilizes membrane function. Inflammatory mediators from damaged membranes are

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propagated via extracellular vesicles (EV) to maternal uterine tissues and transition quiescent maternal uterine tissues into an active state of labor. Membrane inflammation and its propagation are fetal signals that may promote parturition. This review summarizes the mechanisms of fetal membrane cellular senescence, transitions, and the generation of inflammation that contributes to term and preterm parturitions.

Keywords

Amniochorion; fetal membranes; EMT; mesenchymal cells; progesterone; cytokines; inflammation; p38MAPK; exosome

1. Background/Introduction

Factors that maintain human pregnancy and signals that can initiate labor and delivery (parturition) are being investigated by many laboratories. Several reports have expanded knowledge on various topics related to pregnancy and childbirth in the past decade but still several ‘unknowns’ exist. These knowledge gaps can be considered one of the major reasons for adverse pregnancy outcomes that complicate pregnancies around the globe, specifically, preterm births (gestation <37 weeks). The World Health Organization estimated the global preterm birth rate for singleton gestation at 10.5% (Beck et al., 2010; Muglia & Katz, 2010; Rubens et al., 2014). A recent report by the March of Dimes, USA showed that the United States experienced an almost 3-decade rise in the preterm birth rate, beginning in 1980 and peaking in 2006 at 12.8%. Although this rate declined in recent years to 11.4%, preterm birth still accounts for 1 in 9 neonates born every year (McCabe, Carrino, Russell, & Howse, 2014). Approximately 65% of preterm births are ‘spontaneous’ (designated as PTB in the rest of this manuscript), while the other 35% are ‘indicated’ preterm births with known maternal or fetal clinically indicated risks that force early delivery (Andrews et al., 2006; Beck et al., 2010; Di Renzo & Roura, 2006; Goldenberg, Culhane, Iams, & Romero, 2008; Kramer et al., 2012; R. Romero, S.K. Dey, & S.J. Fisher, 2014a; Villar et al., 2012). Preeclampsia, gestational diabetes, fetal growth restrictions, multiple fetal pregnancies, and fetal anomalies are some of the indications for early delivery (Ananth & Vintzileos, 2006). PTB has an unknown etiology and presents with two distinct phenotypes: PTB with intact fetal membranes and PTB that follows preterm pre-labor rupture of the membranes (pPROM) (Goldenberg et al., 2008; Villar et al., 2012). A reduction in preterm births, regardless of whether spontaneous or iatrogenic, is extremely important, not only to reduce the rate of neonatal mortality (~1 million/year globally) or morbidity, but also to minimize the societal impact of prematurity (low birth weight) as preterm babies may experience several adult-onset diseases earlier in life (Beck et al., 2010). Adult-onset diseases seen in preterm babies are partly linked to epigenetic changes due to improper programming of the fetus (D.J. Barker et al., 2010; Devaskar & Thamocharan, 2007; Garg et al., 2012; Robins, Marsit, Padbury, & Sharma, 2011). *In utero* fetal growth is a timed event (~40 weeks of gestation) until delivery and epigenetically driven developmental programming occurs at each stage of growth (D.J. Barker et al., 2010). This growth and programming is properly nourished by an *in utero* environment where any disturbances can alter fetal developmental programming impacting organ maturation and predisposing

individuals to adult-onset diseases (D.J. Barker et al., 2010; Barker et al., 2013; Barker, Thornburg, Osmond, Kajantie, & Eriksson, 2010; Thornburg, 2015; Wallack & Thornburg, 2016). Therefore, it is important to know the mechanisms and pathways leading to PTB as it is critical to reduce PTB risk.

Parturition is initiated when fetal growth is completed. Biochemicals released from matured fetal organs indicate the completion of pregnancy and transition quiescent uterine tissues to an active state of labor (Gao et al., 2015; Mendelson, Montalbano, & Gao, 2017). Maternal decidua, myometrium and cervix serve as various pregnancy clocks and their coordinated activation sets off an alarm in response to fetal signals, resulting in parturition (Gonzalez, Dong, Romero, & Girardi, 2011; Menon, Bonney, Condon, Mesiano, & Taylor, 2016; Menon, Mesiano, & Taylor, 2017; Mesiano et al., 2002; Mesiano, Wang, & Norwitz, 2011; Norwitz et al., 2015; Park, Park, Lockwood, & Norwitz, 2005; Smith, Mesiano, & McGrath, 2002). On the fetal side, placental clock signals change in the endocrine milieu as the placenta ages (Smith, 1998; Smith & Nicholson, 2007). Untimely events, initiated either by the mother or the fetus in response to various risk exposures, preset these alarms, leading to premature activation of the labor cascade, ending in PTB. Understanding the mechanisms by which various feto-maternal tissues maintain pregnancy and molecular and physiologic signals that can lead to parturition at term are essential to decipher the mechanisms leading to PTB and pPROM. Based on current knowledge, PTB in a given subject may be considered as a disease of the placenta (Faye-Petersen, 2008), endometrium/decidua (Bukowski et al., 2017; Pavlicev & Norwitz, 2018; Petraglia, Arcuri, de Ziegler, & Chapron, 2012; Rosen, Kuczynski, O'Neill, Funai, & Lockwood, 2001; Sinkey et al., 2020; Snegovskikh et al., 2009), myometrium (Bukowski et al., 2017; Huszar & Naftolin, 1984; Olson et al., 2003; Tattersall et al., 2008) and/or cervix (Keelan, 2018; Mercer et al., 2000; Vink et al., 2016; Yellon, 2017). In a classic review by Brosens et al, placental bed vascular diseases, classification of defective deep placentation associated with different obstetrical syndromes based on salient features of placentation defects have been described (Brosens, Pijnenborg, Vercruysse, & Romero, 2011). Multitudes of risk factors can impact development of fetoplacental units and predispose them to inflammatory activation early on during pregnancy. These classifications are based on clinical and basic research evidences. The majority of current interventions for controlling preterm labor target these organs based on their pathologic involvement. However, systematic reviews and meta-analysis have shown limited success in delaying PTB in specific subsets of subjects by targeting maternal uterine tissues (Conde-Agudelo, Romero, & Nicolaides, 2020; Dodd, Flenady, Cincotta, & Crowther, 2008; Fernandez-Macias, Martinez-Portilla, Cerrillos, Figueras, & Palacio, 2019; Miyazaki et al., 2016; Vogel, Nardin, Dowswell, West, & Oladapo, 2014). This is due to the complexity of pathways leading to PTB and it is also unlikely that PTB is a single organ or single individual (maternal) disorder. Limited clinical and diagnostic success achieved in the past decades has been overshadowed by the high rate of PTB around the globe. Disparity in the PTB rate among different ethnic groups (Manuck et al., 2015; Menon, 2008), and the occurrence of pPROM in a major subset, further complicates our understanding of the pathophysiology of PTB. There are few clinical trials of pPROM leading to PTB (Berghella & Saccone, 2019; Di Sarno, Raffone, & Saccone, 2019; Langen et al., 2018), even though pPROM is associated with an excessive risk of neonatal morbidity and mortality, early

neonatal infections, bronchopulmonary dysplasia, and necrotizing enterocolitis (Schmitz et al., 2019) .

pPROM is associated with 30-35% of all PTB and the number of cases of pPROM in the United States is close to 175,000/year (Menon & Moore, 2020). These numbers exceed all indicated PTBs and have been steady for decades. pPROM is a very poorly studied syndrome. While several tests are available to confirm a diagnosis of pPROM *post facto* (e.g. pooling, fern tests, nitrazine, and Amnisure®), there is no reliable method to predict pPROM *a priori* because the precise pathophysiologic causes or biomarkers for pPROM are unknown (Elci, Gunes Elci, & Sayan, 2020; Kim et al., 2007; Mercer & Lewis, 1997; Simhan & Canavan, 2005). A limited number of studies have been done on pPROM to clearly understand its etiology, pathophysiology, and early diagnostic markers to provide timely interventions to minimize the incidence of rupture and subsequent PTB (Menon & Moore, 2020). PTB and pPROM share similar causal factors, show a redundancy in biomarkers and exhibit similarities in clinical presentations. However, the mechanisms and pathways leading to PTB and pPROM are different and should therefore be considered distinct phenotypes for clinical management (Menon, 2008, 2019; Menon & Fortunato, 2004). Clinicians are often dealt management dilemmas due to similarities in the clinical course of these syndromes and/or due to limited management strategies. Therefore, it will be beneficial to understand the mechanisms leading to these conditions that can provide a better diagnosis and interventions.

The high rate of PTB and pPROM can partly be blamed on the availability of limited or reliable basic science data (mechanisms, pathways, and biomarkers) to synthesize evidence that is needed to conduct clinical trials or provide proper management. As mentioned above, many current interventions target maternal uterine tissues as they are the timekeepers and final promoters of labor and delivery. Fetal signals (endocrine, paracrine, immune etc.) that can offset maternal clocks by transitioning them from a quiescent to an active state are critical contributors of parturition at term and preterm. Therefore, current research has been focused heavily on the fetal side, especially on fetal-derived intrauterine tissues such as the placenta and placental membranes (the amniochorionic membrane or fetal membrane). Fetal signals (their tissue source, pathways of generation, characteristics, propagation and maternal tissue targets) are currently being investigated. Understanding communication signals regarding fetal readiness for delivery and the facilitation of delivery after a normal pregnancy and parturition, will immensely benefit our pursuit in designing strategies to reduce the premature development of fetal signals leading to preterm labor.

This review is focused on certain novel mechanistic aspects of fetal tissue-based signals, their generation and propagation to maternal tissues, and how this indicates parturition. The placenta is a well-studied fetal organ during pregnancy and parturition (Andrews et al., 2006; Cao, Stout, Lee, & Mysorekar, 2014; Challis et al., 2009; Faye-Petersen, 2008; Mor & Kwon, 2015; Seferovic et al., 2019). However, the fetal membranes are often ignored or poorly studied, as their contributions are not regarded as anything beyond the sac that forms the cavity (Menon, 2016). Recent advances in cell biology (Abrahams et al., 2013; Arechavaleta-Velasco, Ogando, Parry, & Vadillo-Ortega, 2002; Canzoneri et al., 2013; Dunand et al., 2014; Feng, Allen, Marinello, & Murtha, 2019; Gomez-Lopez,

Hernandez-Santiago, Lobb, Olson, & Vadillo-Ortega, 2013; C.E. Kendal-Wright, 2007; Lappas, 2013; Malak et al., 1993; Myatt & Sun, 2010; Sato, Collier, Vermudez, Junker, & Kendal-Wright, 2016; Sun, He, & Yang, 2002), organ-on-chip (Gnecco et al., 2017; L. Richardson et al., 2019), and mechanistic models (El et al., 2006; Kumar et al., 2009; Kumar et al., 2011) have made tremendous advances in fetal membrane biology research. There are multiple reasons for using fetal membranes for our studies: 1) they are fetal in origin (Fox, 1981); 2) they are structurally, mechanically and functionally different to the placenta, an organ that has been well studied (Menon, Richardson, & Lappas, 2018); 3) fetal membranes are a rich source of functionally relevant biochemicals that help the pregnancy and promote parturition; 4) membranes start their growth along with the fetus and provide mechanical, structural, immune, antimicrobial and endocrine functions to the uterine cavity that are distinct from the placenta (Fox, 1981; Menon, 2016); 5) membranes are a rich source of stem cells, a property that has been widely used in regenerative medicine (Nogami et al., 2016; Saito, Lin, Murayama, Hashimoto, & Yokoyama, 2012); and 6) pPROM is considered a "disease of the fetal membranes" (Murtha & Menon, 2015). Therefore, we have been using fetal membranes, membrane-derived cells, and the membrane extracellular matrix (ECM) as a model to better understand the contributions to the mechanisms of this tissue, as well as to use it as a proxy to understand fetal contributions to pregnancy and parturition at both term and preterm. We are not attempting to describe the structure and development of fetal membranes here as it has been done in several other recent reviews (Martin, Richardson, da Silva, Sheller-Miller, & Menon, 2019; Menon et al., 2018). We have been studying fetal membranes using an explant culture approach that we developed in the early 1990s (Fortunato, Menon, Swan, & Lyden, 1994), and modified over time or by using cells from the fetal membranes (amnion epithelium, amnion mesenchyme, chorion mesenchyme and chorion trophoblast) (Jin, Richardson, Sheller-Miller, Zhong, & Menon, 2018; L. Richardson & Menon, 2018). By using fetal membrane tissues and cells as a model system, we have developed two independent mechanistic pathways that can contribute to normal term birth and also how membrane cells continue down the PTB and pPROM pathways. The rest of the review will focus on how fetal membrane cells help to maintain pregnancy and will detail two novel mechanistic pathways that send signals to the mother to promote parturition, senescence and cellular transitions.

2. Inflammatory response at term and in PTB

Prior to detailing the mechanisms by which fetal membranes maintain pregnancy and promote parturition by controlling inflammation, we will summarize the overall role of inflammation in pregnancy and parturition. Feto (placenta, membranes, umbilical cord) maternal (decidua, myometrium, and cervix) reproductive tissues maintain immune homeostasis during pregnancy and tolerate the semi-allogeneic fetus until parturition (Alijotas-Reig, Llurba, & Gris, 2014; Chavan, Griffith, & Wagner, 2017; Mor, Cardenas, Abrahams, & Guller, 2011; Negishi, Takahashi, Kuwabara, & Takeshita, 2018b; Schumacher, Sharkey, Robertson, & Zenclussen, 2018; Svensson-Arvelund et al., 2015). Balanced immune interactions by feto-maternal units ensure pregnancy maintenance and feto-placental growth (Gomez-Lopez, StLouis, Lehr, Sanchez-Rodriguez, & Arenas-Hernandez, 2014; Kshirsagar et al., 2012; Negishi et al., 2018b). Pregnancy success

is determined by immune regulatory mechanisms at the feto-maternal interface tissues, ensuring that both innate and adaptive immune cells aptly support feto-placental development by regulating localized inflammation while remodeling uterine tissues (Erlebacher, 2013; Lash, 2015; S. Liu et al., 2017; Nancy & Erlebacher, 2012; Schliefssteiner et al., 2017). On the contrary, parturition in both humans and animals is associated with a physiologic inflammatory process (Dudley, 1999; Goldenberg et al., 2008; Gomez-Lopez et al., 2011; R. Romero, S. K. Dey, & S. J. Fisher, 2014b; Romero, Gotsch, Pineles, & Kusanovic, 2007). This inflammation required to overcome immune balance at the feto-maternal interface tissues and to induce parturition associated changes in maternal tissues is characterized by the infiltration and activation of immune cells (both innate and adaptive immune cells) into the feto-maternal interface, along with the increased production of pro-inflammatory mediators (e.g. IL-1 β , IL-6, IL-8, TNF- α , GM-CSF etc.) and decreased levels of anti-inflammatory mediators (e.g. IL-10, TGF- β etc.) (Boonkasidecha, Kannan, Kallapur, Jobe, & Kemp, 2017; Bukowski et al., 2017; Cappelletti et al., 2017; Edey et al., 2016; Gomez-Lopez et al., 2014; Hamilton et al., 2012; Osman, Young, Jordan, Greer, & Norman, 2006; Peltier, 2003; Rinaldi, Makieva, Saunders, Rossi, & Norman, 2017; Xu et al., 2018; Zhang et al., 2017). Events contributing to physiologic immune imbalances and inflammation at term include, but are not limited to, signals of fetal organ maturation (Mendelson et al., 2017; Montalbano, Hawgood, & Mendelson, 2013), fetal membrane stretch (Mohan, Sooranna, Lindstrom, Johnson, & Bennett, 2007), stress-induced damage to the uterine tissues promoting immune cell chemotaxis (C. S. Buhimschi et al., 2009; Jauniaux, Poston, & Burton, 2006; Sharp, Heazell, Crocker, & Mor, 2010; Wadhwa, Culhane, Rauh, & Barve, 2001), and fetal membrane-placental-decidual senescence (Behnia, Sheller, & Menon, 2016a; Behnia, Taylor, et al., 2015b; Bonney et al., 2016; Hirota et al., 2010a; Menon, Behnia, et al., 2016; Polettini et al., 2015b). The disruption of immune homeostasis leading to parturition is expedited by both (Chen & Chen, 2013; Hinz & Scheidereit, 2014) endocrine and paracrine mediators generated when fetal growth is complete (Buhimschi et al., 2008; Dudley, 1999; Golightly, Jabbour, & Norman, 2011; Iliodromiti et al., 2012; Keelan et al., 2003; Mendelson, 2009; Shynlova, Tsui, Jaffer, & Lye, 2009; Trivedi et al., 2012). Premature disruption of immune homeostasis and an overwhelming host inflammatory response due to infectious or other noninfectious risks can lead to PTB and pPROM (Romero et al., 2014b; Romero et al., 2006; Xu et al., 2018). Histologic chorioamnionitis (HCA), the infiltration of neutrophils into fetal membranes, is often associated with PTB and pPROM and contributes to neonatal morbidity and mortality (Chaiworapongsa et al., 2002; Gomez-Lopez, Romero, Xu, et al., 2017; Menon, Taylor, & Fortunato, 2010). Similarly, decidual cells and immune cells residing in decidua are in an immune harmonious state (McIntire, Ganacias, & Hunt, 2008). Endogenous activation of inflammation due to decidual senescence (Cha, Hirota, & Dey, 2012) or immune activation in response to signals from fetus can disrupt immune homeostasis of this tissue. (Bartmann et al., 2014; Erlebacher, 2013; Gomez-Lopez et al., 2014; J. Liu, Dong, Wang, & Li, 2019; S. Liu et al., 2017; Mor, 2008; Presicce et al., 2015; Rinaldi et al., 2017; Schumacher et al., 2018). In summary, uterine tissue immune balance, a harmonious situation between both maternal and fetal tissues and immune cells harbored by them, is a key factor that maintains pregnancy (Alijotas-Reig et al., 2014; Arenas-Hernandez et al., 2016; Challis et al., 2009; Erlebacher, 2013; Figueiredo & Schumacher, 2016; Gomez-Lopez et al., 2014;

King, Kelly, Sallenave, Bocking, & Challis, 2007; S. Liu et al., 2017; Mor, 2008; Negishi, Takahashi, Kuwabara, & Takeshita, 2018a; Peltier, 2003; Rinaldi et al., 2017; Song & Shi, 2014; Southcombe, Tannetta, Redman, & Sargent, 2011; Szekeres-Bartho, 2002).

Pro-inflammatory activity is one of the key triggers of parturition at term and preterm. One is a physiologic and natural response, while the other is a pathologic and untimely event. Increased pro-inflammation imbalances and collapses all other homeostatic states of various pregnancy-associated tissues to ensure parturition. It is hypothesized that each tissue has its own inflammatory contributions, and we have been investigating how fetal membranes either endogenously, and/or in response to exogenous mediators, lead to an inflammatory state that can cause cellular and fetal membrane ECM degradation and a dysfunctional membrane status. A dysfunctional membrane loses the following capacities: 1) mechanical and structural disruption to support intrauterine cavity; and 2) controlling structural damage that contributes to functional disruptions such as a loss of antimicrobial resistance, immunologic compromises, and a loss of permeability functions resulting in fetal membrane collapse. In the next two segments, we will review how the endogenous activation of inflammatory mediators, primarily inflammatory cytokines, chemokines, and metalloproteinases, causes localized inflammation in the fetal membranes. We will review whether these are capable of disrupting membranes to predispose them to either rupture or generate inflammation to cause labor.

2.1. Fetal membrane stretches and strain and inflammation

The amnion membrane component of the fetal membranes, which is constantly hydrated by the amniotic fluid, is highly elastic and can withstand pressure without undergoing any rupture during normal pregnancy. Pressure on and stretching of the amnion membrane *in utero* are caused by the growing fetus and an increase in amniotic fluid volume as gestation progresses (Adams Waldorf et al., 2015; C. E. Kendal-Wright, 2007). The amnion ECM is comprised of tropoelastins, elastin cross-linking enzymes, lysyl oxidase, and lysyl oxidase-like (LOXL) enzymes that contribute to the amnion's mechanical function (Bryant-Greenwood, 1998; Malak et al., 1993; Poletti et al., 2016; Strauss, 2013). Poletti et al. showed that changes to these enzymes can predispose the membrane to premature rupture, resulting in preterm delivery (Poletti et al., 2016). The membrane undergoes mechanical stretching as it grows and develops alongside the fetus, distends further at term and detaches from the uterine wall (Joyce et al., 2016; Millar, Stollberg, DeBuque, & Bryant-Greenwood, 2000). This has been validated in classical studies utilizing an *in vitro* model of amnion tissue stretch (Buerzle et al., 2013), mimicking fetal descent (Buzle, Mazza, & Moore, 2014; Mauri, Ehret, et al., 2015; Mauri, Perrini, Ehret, De Focatiis, & Mazza, 2015). and a balloon inflation approach to mimic a situation like polyhydramnios in non-human primates. Both studies showed stretching-induced inflammation (Adams Waldorf et al., 2015). Over-distension of the membranes at term or preterm is also mimicked *in vitro* by cyclic stretching of the amnion (Joyce et al., 2016; C. E. Kendal-Wright, 2007; Kendal-Wright, Hubbard, & Bryant-Greenwood, 2008; Kendal-Wright, Hubbard, Gowin-Brown, & Bryant-Greenwood, 2010). Cyclic stretch has been shown to increase cellular stress and pro-inflammatory cytokine production, indicative of a laboring phenotype (C. E. Kendal-Wright, 2007; Kendal-Wright et al., 2010). However, persistent stretch, as reported by

Kendal-Wright et al., demonstrated physical, but not static, strain placed on the amnion cells induces pro-labor chemokines like Interlukin-8 (IL-8) (Kendal-Wright et al., 2008). In support of this finding, we have seen that physiologic stretch experienced by the human amnion membrane does cause stress signals such as p38 mitogen activated protein kinase (MAPK) activation, but this activation does not result in the downstream activation of inflammatory mediators such as NF- κ B (manuscript under review). Therefore, it is likely that stretch-associated inflammation is not sufficient to cause membrane damage but helps to remodel them to withstand sustained stretching throughout pregnancy.

2.2. Fetal organ maturation signals are pro-inflammatory in amniochorion cells

Fetal organ maturation is a key developmental signal from the fetus. Biochemicals such as endothelins (indicating kidney maturation) (Gryglewski, Chlopicki, Uracz, & Marcinkiewicz, 2001; Lockwood, 1994), platelet activation factor (Hoffman, Truong, & Johnston, 1986), surfactant proteins from the fetal lung (Condon, Jeyasuria, Faust, & Mendelson, 2004; Mendelson et al., 2017; Montalbano et al., 2013), epidermal growth factor (Cai, Huang, Leung, & Burd, 2014), and brain-derived neurotrophic factor (Antonakopoulos et al., 2018) are a few of the signals that are considered pro-inflammatory in the amniotic fluid; their presence can increase the overall inflammation. Although these signals are present in the amniotic fluid during fetal growth at low levels, their increased bioavailability may exceed the threshold of a balanced immune status and induce inflammation from fetal membrane cells, specifically amnion epithelial cells that are the innermost lining of the amniotic cavity and constantly bathed in the amniotic fluid and associated biochemicals. Reinl & England summarized a mechanistic process of such signaling that can lead to myometrial smooth muscle cell activation, resulting in parturition (Reinl & England, 2015). Similarly, we further documented in animal model studies that fetal exosomes can travel from the fetal to the maternal side, reaching the uterus and cervix to trigger parturition (Reinl & England, 2015). In summary, fetal biochemical signals of organ maturation have the capacity to cause inflammation in fetal membrane cells that can disrupt membrane functions at term.

One of the above-described mechanisms may contribute to tissue remodeling during pregnancy, whereas others can be a signal of parturition. However, our recent discoveries suggest that there are two other key mechanisms by which fetal membranes generate inflammation that can be detrimental to pregnancy and lead to parturition mechanisms. We will focus the next segments on explaining some of the novel mechanisms involved in fetal membrane maintenance during gestation and how two unique events (senescence (mechanism of aging) and cellular transitions such as epithelial-to-mesenchymal transition (EMT) and mesenchymal-to-epithelial transition (MET)) generate inflammation within the fetal membranes that can promote parturition. These studies are currently focused on amnion membrane cells (amnion epithelial cells [AEC] and amnion mesenchymal cells [AMC]).

3. Fetal membrane senescence, a mechanism of aging, and aging associated inflammation

Fetal membranes start their development at embryogenesis, grow with the fetus, and undergo multiple cycles of developmental changes during gestation. Amnio-chorion fusion happens during the early second trimester, resulting in a single unit structure (Menon & Moore, 2020; Menon et al., 2018). As with any other type of cells, replication of the fetal membrane cells leads to senescence (Campisi, 1997a; von, 2003), a pathway to aging and inflammation (Campisi, 1997b; Chambers & Akbar, 2020). The “Hayflick Phenomenon” explains this process, where cell division is halted after a certain number of divisions (Hayflick, 1961; Hayflick & Moorhead, 1961). Two key function-based definitions proposed by Masoro et al. (Masoro, 1995) and Finch et al. (Finch, 1992) may aid in the understanding of the biologic aging of fetal membranes: 1) fetal membranes are expected to deteriorate during gestation once their maturation is completed during gestation once its maturation is completed around the 12th week of pregnancy, and will be vulnerable to subtle changes in the intrauterine environment, decreasing survival ability; and 2) senescence is a mechanism associated with the deterioration of the membranes, which alters its function and decreases vitality (Menon, 2016). A recent report by Behnia et al. showed an association between amniochorionic membrane senescence and term pregnancies. Additional work determined the mechanism of the development of senescence through various *in vitro* and *in vivo* models (Behnia, Taylor, et al., 2015c). The following features were observed in the development of fetal membrane senescence.

3.1. Amniochorionic membrane senescence shows gestational age dependent telomere attrition

Telomeres are specialized DNA-protein structures located at the ends of eukaryotic chromosomes. Cellular replication leads to their reduction in size and this is considered as one of the biomarkers for aging (Rodier, Kim, Nijjar, Yaswen, & Campisi, 2005; Sanders & Newman, 2013; von & Martin-Ruiz, 2005). The examination of amniochorionic membranes from various gestational ages (starting from 22 weeks) until term (40 weeks) showed a progressive decline in telomere length in both amniochorionic membranes and fetal DNA samples from cord blood (Menon et al., 2012b). We were also able to demonstrate that cell-free fragments of telomeres are shed from the cells and increased in the amniotic fluid of women at term labor compared to term not in labor (Menon, Behnia, et al., 2016). Telomere reduction in fetal membranes is correlated with fetal growth; therefore, the peak of membrane aging is paralleled with the completion of fetal growth and organ maturation, suggesting that senescence is likely a physiologic process preparing membranes for delivery.

3.2. Amniochorion membrane senescence is aided by oxidative stress (OS) build up in the amniotic cavity and activation of p38MAPK

The intrauterine environment undergoes redox changes during pregnancy, as detailed in many reports (G.J. Burton, 2009; Burton & Jauniaux, 2001; Myatt, 2010; Myatt & Cui, 2004). Term pregnancy is characterized by redox imbalance and the accumulation of reactive oxygen species (ROS) in the amniotic fluid and in fetoplacental tissues (Longini et

al., 2007; Menon, Boldogh, et al., 2014; Poletini, Richardson, & Menon, 2018). Multiple reasons can be linked to the build-up of ROS at the feto-maternal interface that can be seen in a previous reference (Menon, 2014) [Figure 1]. The examination of OS-mediated signaling revealed that fetal membrane cells respond with the increased activation of p38MAPK, a responder of stress signals that can determine cell fate (apoptosis, senescence and or other forms of cell deaths) (Choi, Lee, Ha, & Kim, 2011; Jin et al., 2018; Menon et al., 2013; Menon & Papaconstantinou, 2016). p38MAPKs are a family of four stress response signaling isoforms (p38MAPK α , β , γ , δ) that are evolutionarily conserved serine/threonine kinases whose functions differ significantly. We observed activation of p38MAPK α , a form that is reportedly more common in cells (Ruiz-Bonilla et al., 2008). The activation of p38MAPK by phosphorylation results in cell cycle inhibition in amnion cells via the down-regulation of glycogen synthase kinase 3 beta (GSK3 β) in fetal membranes (Lavu et al., 2019; Lavu et al., 2020).

3.3. p38MAPK activation results in cellular senescence

Using multiple models, both *in vitro* and *in situ*, we were able to demonstrate that p38MAPK activation is also a progressive event in fetal membrane cells and maximal activation seen when OS is high (Bonney et al., 2016). As shown in mouse models, p38MAPK activation correlates with gestational age along with the deactivation of GSK3 β (Bonney et al., 2016; Lavu et al., 2020). Senescent cells are characterized by multitudes of changes and produce biochemicals that are specific indicators of senescence and cause injury to cells (Dimri & Campisi, 1994). Senescence-associated galactosidase (SA- β -Gal) is widely used as a biomarker of replicative senescence. As detailed by Kurz et al, SA- β -Gal is a manifestation of residual lysosomal activity at a suboptimal pH, which becomes detectable due to the increased lysosomal content in senescent cells (Kurz, Decary, Hong, & Erusalimsky, 2000). we were able to demonstrate that fetal membrane cells develop senescence-associated β -Galactosidase activity, a key marker of senescence (Behnia, Taylor, et al., 2015c; Coppe et al., 2008; Menon, Behnia, et al., 2016; Poletini et al., 2018). Senescent cells often show injury to organelles and nuclear membranes. Damage to the nuclear lamina leads to nuclear morphology in senescent cells and nuclear membrane Lamin B1 loss is detrimental to cell survival (Freund, Laberge, Demaria, & Campisi, 2012). As a morphological marker of senescent associated nuclear injury, we tested Lamin B1 in fetal membrane cells. In addition to development of SA- β -Gal, we were able to confirm a loss of Lamin B, (Menon, Behnia, et al., 2016), as electron micrographs revealed morphologic changes (enlargement) to organelles such as the mitochondria, ER, and damage to the nuclear membrane (Menon, Boldogh, et al., 2014), DNA fragmentation (Menon et al., 2013; Menon, Poletini, Syed, Saade, & Boldogh, 2014), and the development of unique forms of inflammation (Menon, Behnia, et al., 2016). All of these effects were inhibited by p38MAPK inhibitors, confirming the role of this signal in inducing senescence in fetal membranes.

3.4. Senescent cells release senescence associated secretory phenotype (SASP) and damage associated molecular pattern markers (DAMPs)

Senescence is characterized by inflammation known as SASP as SASP is represented by a unique congregation of various biochemical markers. Although SASP is represented by well

reported cytokines, chemokines, growth factors, metalloproteinases (MMPs), inhibitors and receptors, and vascular growth factors, collective presence of specific markers are unique in senescent cells. Transcriptome analysis of senescent amniochorion cell showed the increased expression of SASP transcripts in term labor and in response to OS *in vitro* compared to their respective controls (Behnia, Taylor, et al., 2015c; Menon, Behnia, et al., 2016). Of note, SASP markers have been reported to be higher in various biological compartments during term labor (although not termed SASP markers) compared to term not in labor conditions. This further associates senescence and SASP with parturition (Behnia, Taylor, et al., 2015c). Most of these SASP markers are known pro-labor factors in fetomaternal tissues (Behnia, Taylor, et al., 2015c). Based on our data, a bioinformatics analysis of SASP markers was performed which revealed that pathological pathways and cellular signaling represented by SASP were derived due to cellular damage resulting from senescence. In our *in vitro* and *in vivo* models, administration of N-acetyl cystine (an anti-oxidant) and p38MAPK inhibitor SB 203580 reversed senescence and SASP, supporting the hypothesis that fetal membrane senescence and inflammation accompanying it are an OS-induced p38MAPK mediated phenomenon. Moreover, cells were further analyzed for degenerative tissue markers. Our studies revealed that OS-induced senescent amnion cells release the high mobility group box (HMGB)1 protein as well as cell free fetal telomere fragments (Bredeson et al., 2014; Menon, Behnia, et al., 2016). These markers belong to a class of Alarmins or damage associated molecular pattern markers (DAMPs) representing cellular injury and or tissue damage. DAMPs are intracellularly sequestered molecules and are hidden from recognition by the immune system under normal physiological conditions. However, under conditions of cellular stress/tissue injury, these molecules can either be actively secreted by stressed immune cells and considered as endogenous danger signals, because they induce potent inflammatory responses by activating the innate immune system during non-infectious inflammation (Roh & Sohn, 2018) (Land, 2015a, 2015b). DAMPs can arise from intracellular proteins, such as HMGB1, histones, IL-33, IL-1 α , S100 proteins, heat-shock proteins (HSPs), DNA and RNA as well as extracellular matrix proteins such as decorin, biglycan, low molecular weight hyaluronan etc.. Besides these, cellular organelles and They can function through a variety of receptors like toll like receptor (TLR) 2, 4, 6 and 9, receptor for advanced glycation endproducts (RAGE) among other on various cell types to cause inflammatory activation in recipient cell (Roh & Sohn, 2018). TLRs expressions are well reported in the fetal membranes. (Abrahams et al., 2013; Agrawal & Hirsch, 2012) and their differential role during infection, parturition at term and preterm are well reported (Hoang et al., 2014; Ilievski, Lu, & Hirsch, 2007; Lim, Barker, & Lappas, 2014; Moco et al., 2013; Sato et al., 2016).

3.5. DAMPs cause feedback activation of senescence

In support of our *in vitro* cell-based models, we conducted an analysis of inflammatory marker concentrations in three distinct biological fluids from the same pregnancy. Our report showed higher concentrations of HMGB1 term labor amniotic fluid and cord blood samples than maternal plasma, supporting the hypothesis that these DAMPs may function as fetal signals of parturition (Menon & Taylor, 2019). DAMPs, Further testing was conducted to determine their functional relevance. Data from our own and other studies using *in vitro* and *in vivo* models revealed the following: 1) both HMGB1 and cell-free fetal telomere DNA

fragments accumulate in the amniotic fluid and they are also powerful stimulants of OS, induce AEC p38MAPK activation and enhance senescence of fetal membranes (Bredeson et al., 2014; Menon, Behnia, et al., 2016; Poletini et al., 2015a); 2) the *in vivo* injection of telomere fragments leads to p38MAPK activation, senescence and PTB in animals (Poletini et al., 2015a); and 3) the intra-amniotic administration of HMGB1 can cause PTB in mouse models (Gomez-Lopez et al., 2016). In addition, our own work has shown senescence in AMCs and chorion trophoblast cells (Feng et al., 2019), which aligns with other labs which have also reported telomere reduction (Phillippe, Sawyer, & Edelson, 2019), senescence of the amniochorion (Gomez-Lopez, Romero, Plazyo, et al., 2017), placenta (Burstein, Frankel, Soule, & Blumenthal, 1973; Cox & Redman, 2017; Ferrari, Facchinetti, Saade, & Menon, 2016; Parmley, 1984), and decidua (Hirota et al., 2010b), along with their association with parturition. In summary, SASP and DAMPs are unique forms of inflammation that are released from senescent fetal membrane cells that can cause parturition.

The above sections summarize that fetal membrane senescence and senescence-associated inflammation can be mechanistically linked to normal term birth. Senescence and SASP/DAMP-mediated inflammation are normal physiologic responses of the fetal membrane as they reach longevity at term, correlating with the completion of fetal growth.

4. Can premature senescence cause preterm birth?

The primary purpose of studying senescence and its association with normal term parturition is to expand our knowledge and understand similar pathways that can contribute to PTB and pPROM. The hypothesis is that PTB and pPROM risk factors accelerate senescence, causing SASP/DAMP-associated inflammation, which induces membrane dysfunction and collapse of the intrauterine structure required to support fetal growth [Figure 2]. All of the above markers of OS and senescence were tested in tissue samples from PTB and pPROM. Also, *in vitro* and *in vivo* models were developed by exposing either membrane explants, primary fetal membrane cells, or animal models to various PTB-inducing stimuli. Clinical sample analysis showed that compared to PTB with intact membranes, pPROM had: 1) shorter fetal membrane and cord blood DNA telomere lengths at delivery (Menon et al., 2012b), 2) reduced levels of antioxidant enzymes (Dutta et al., 2016), 3) the increased evidence of OS, protein peroxidation, lipid peroxidation and DNA damage (Menon, Boldogh, et al., 2014; Menon et al., 2011), and 4) the increased activation of p38 MAPK and tissue senescence (Menon, Boldogh, et al., 2014).

In vitro models showed increased OS, p38MAPK activation, senescence and the generation of SASP in response to various risk factors such as cigarette smoke (Menon et al., 2013; Poletini et al., 2018), environmental pollutants (Behnia, Peltier, et al., 2016; Behnia, Peltier, Saade, & Menon, 2015), noninfectious (sterile) inflammation and infection (Dixon, Richardson, Sheller-Miller, Saade, & Menon, 2018) [Figure 2]. The incidence of senescence in response to infectious stimuli such as lipopolysaccharides (LPS) was much milder than any other risk factors tested in our system (Dixon et al., 2018). This could be due to the fact that all other risk factors than LPS are dominant OS inducers. We do not rule out the possibility that live bacteria or even heat-inactivated bacteria may produce a much more rigorous OS response and hence senescence. *In vitro* models also lack immune cells that are

recruited to control the invasion of pathogens by generating ROS radicals(Menon, Behnia, et al., 2016; Menon, Boldogh, et al., 2014; Menon et al., 2013).

Of note, we did not see a difference in inflammatory markers (mostly cytokines and chemokines) in the amniotic fluid, cord blood, or maternal plasma between infection and OS, suggesting that inflammation is a key trigger in both conditions, regardless of the mechanism or pathway (Menon & Taylor, 2019). Interestingly, molecular markers, cellular level changes in senescence and inflammatory mediators exhibited tremendous similarities between normal term birth and pPROM (specifically early pPROM <34 weeks), whereas they were different in PTB. These include increased reactive oxygen species, telomere length reduction, increase in markers of lipid and protein peroxidation, imbalance in MMPs and TIMPs, p38MAPK, senescence associated cellular organelle morphology, localized MMP9 activation, and loss of Lamin B1(Dutta et al., 2016; Menon, Boldogh, et al., 2014; Menon, McIntyre, Matrisian, & Fortunato, 2006; Menon et al., 2012a). This suggests that fetal membrane aging and aging-induced inflammation are a (patho)physiologic requirement for term labor and pPROM, causing membrane weakening and rupture prior to labor. Although inflammatory marker concentrations and their profiles are similar between PTB and pPROM, it is likely that mechanisms leading to the increase in fetal membrane inflammation is independent of senescence and SASP in PTB (Dutta et al., 2016). Conversely, we would like to emphasize that pPROM is a disease associated with fetal membrane senescence. OS induced by various pPROM risk factors prematurely ages fetal membranes, making them dysfunctional leading to mechanical and structural weakening and rupture (Behnia, Peltier, et al., 2015; Behnia, Sheller, & Menon, 2016b; Menon et al., 2013).

To note, DAMPs can be generated independently of senescence of cells and they are primarily produced in response to infection, specifically in response to Pathogen-associated molecular patterns (PAMPs)(Santoni et al., 2015). Therefore, infection associated adverse pregnancy events can produce DAMPS via PAMPs as well as by inducing senescence of cells(Elovitz, Wang, Chien, Rychlik, & Phillippe, 2003; Hoang et al., 2014; Jaiswal et al., 2013; Padron, Saito Reis, & Kendal-Wright, 2020).

5. Cellular transitions and generation of inflammation

The above section detailed how inflammation is generated in response to senescence. In this section we describe yet another mechanism by which fetal membrane cells generate local inflammation that can be beneficial during gestation for tissue remodeling or the overwhelming presence of which can lead to membrane degeneration. Fetal membrane cells have stem cell-like properties, as they can proliferate, migrate, express stem cell markers and are capable of transitioning into other cell types (L. Richardson & Menon, 2018). These properties are essential for fetal membrane remodeling and to maintain its integrity, as membranes are constantly under shear stress and stretch from the fluid and fetus, respectively, during pregnancy. Studies conducted using fetal membrane cells to examine cellular transition mechanisms and generation of inflammation are listed below.

5.1. Fetal membrane microfractures

The remodeling of membrane cells includes the shedding of cells, leading to the development of microfractures in the membrane structure (L. S. Richardson et al., 2017). Microfractures are not just gaps created by cellular shedding or puckering, they also show degradation of the basement membrane and matrix collagen as well as function as a passage for shed cells (L. S. Richardson et al., 2017). This process generates localized inflammation, which is required for membrane matrix remodeling. Although intrauterine OS levels fluctuates during gestation (G. J. Burton, 2009; Jauniaux et al., 2006; Myatt, 2010), redox balance sustains the remodeling process (Agarwal, Gupta, & Sharma, 2005). However, the process is stalled as the membrane reaches longevity at term and demonstrates structural, functional, and biomolecular changes that are characteristic of aging (Behnia, Taylor, et al., 2015a). The number of microfractures and their morphometry (width and length) were higher in term laboring membranes than term not in labor membranes (L. S. Richardson et al., 2017). Similarly, pPROM had a higher number of microfractures than PTB with intact membranes (L. Richardson & Menon, 2018).

5.2. Microfracture healing involves cellular transitions and inflammation

In a recent study, Richardson et al. created artificial scratches (representing microfractures) using AECs. The following observations were made during microfracture healing (L. Richardson & Menon, 2018): 1) epithelial cells proliferate and transition into mesenchymal cells (EMT) in the early stages of healing with the expression of cytoskeletal and cell adhesion markers; 2) the transition to mesenchymal cells increases the migratory capacity of AECs; 3) healing of wounds/microfractures is accompanied by cell transitioning back to the epithelium (MET); 4) amniotic fluid accelerated the healing process, whereas OS stopped healing; 5) antioxidants reversed OS effect and augmented healing, suggesting that OS can cause a terminal or static state of EMT and prevent membrane microfracture healing; and 6) migration and healing was associated with localized inflammation.

5.3. EMT is associated with term parturition

In addition to wound healing, as described above, the transdifferentiation of EMT is seen in embryogenesis and required for embryonic stem cell differentiation (DaSilva-Arnold, James, Al-Khan, Zamudio, & Illsley, 2015). Furthermore, EMT is also pathologically associated with fibrosis and cancer metastasis (Hay, 1995; Lamouille, Xu, & Derynck, 2014). In reproductive tissues, EMT has been linked to various stages of placental development including the differentiation of cytotrophoblasts to extravillous trophoblast cells (DaSilva-Arnold et al., 2015), as well as trophoblast differentiation (Vicovac & Aplin, 1996). EMT is characterized by the repression of epithelial cell-associated genes and the concomitant activation of genes that transition them into a mesenchymal phenotype (Lamouille et al., 2014). The reverse of EMT is seen with MET, which generates epithelial cells during various developmental stages. During the early stages of development, MET facilitates the embryo's engagement in gastrulation and organogenesis (Pei, Shu, Gassama-Diagne, & Thiery, 2019). Based on our observations during microfracture healing, we hypothesized that the amnion layer fetal membrane may undergo EMT to generate inflammation that will weaken this layer. This hypothesis was aided by a couple of supportive reports suggesting

EMT associated with membrane rupture and healing (Janzen et al., 2017; Mogami, Hari Kishore, Akgul, & Word, 2017). We examined human and mouse amnion membranes and noted that membranes from term parturition have a substantially higher number of AMCs than epithelial AECs (L. S. Richardson, Taylor, & Menon, 2020). This helped to expand our hypothesis that term labor may be associated with a terminal state of EMT due to the accumulation of highly inflammatory AMC in the ECM, as previously described (Sato et al., 2016). Descriptive data determined the following: 1) compared to term not in labor, term labor is associated with amnion membrane EMT in both mice and humans (L. S. Richardson et al., 2020); 2) AECs during normal gestation exhibit a 'metastate' expressing both epithelial (cytokeratin) and mesenchymal marker (vimentin) (L. S. Richardson et al., 2020); 3) classic markers (cytoskeletal, adhesion, and transcription factors) associated with mesenchymal transitions were expressed in amnion layer; and 4) EMT is associated with MMP9 induction and collagen degradation that can cause basement membrane degradation to structurally and functionally weaken the membrane, cause dysfunction and predispose them to rupture. In summary, term labor is associated with EMT and localized inflammation [Figure 3].

5.4. Mechanisms of EMT mediated by transforming growth factor (TGF)- β

Using molecular and cell biological approaches, Richardson et al. reported that EMT in amnion cells are mediated by TGF- β . TGF- β is seen in the amniotic fluid during gestation, but its concentrations are higher at term labor compared to term not in labor samples (L. S. Richardson et al., 2020). The exposure of cells to OS experienced at term labor increased TGF- β release from AECs, an increase that was reduced by antioxidant NAC treatment (L. S. Richardson et al., 2020). Blocking OS induced TGF- β -mediated signaling through gene silencing of TGF- β -activated kinase 1 binding protein 1 (TAB1) reduced EMT transcription factors and mesenchymal junction markers, maintaining epithelial characteristics. The silencing of TGF- β reduced p38MAPK activation. The inhibition of EMT in AECs by treatment with p38MAPK inhibitors further supports the role of p38MAPK in TGF- β -TAB1-mediated EMT (L. Richardson, Dixon, Aguilera-Aguirre, & Menon, 2018). EMT accumulates AMCs in the matrix and prompt inflammatory activation in the membranes (Sato et al., 2016; Whittle, Gibb, & Challis, 2000). Therefore, OS-induced p38MAPK activation can cause both senescence and EMT in cells. It is still unclear whether these two processes are interdependent or independent; however, both of these processes co-exist at term labor membranes in both humans and mouse models of normal gestation and parturition [Figure 3].

5.5. Mechanism of MET mediated by progesterone

Mesenchymal cells perform endocrine functions during gestation; however, these are tightly regulated and require a limited number of cells. As mesenchymal cells are highly susceptible to inflammation and ROS (Sato et al., 2016), their numbers need to be tightly regulated, which is achieved by reprogramming them back to epithelial cells through MET to maintain membrane integrity. MET will reestablish epithelium cell-to-cell contact and increase nascent collagen production to remodel the degraded matrix. The pregnancy maintenance hormone progesterone, an anti-inflammatory hormone, was tested to determine whether it mediated the reversal of EMT by promoting MET (L. S. Richardson et al.,

2020). This mechanism is also thought to regulate local inflammation. Progesterone, through the progesterone receptor membrane component 2 (PGRMC2), induces MET via the proto-oncogene c-MYC. The silencing of PGRMC2 using siRNA and/or reducing c-MYC using its pharmacologic inhibitor increased mesenchymal transcription factors and cellular junction markers indicative of the persistence of the mesenchymal fibroblastoid phenotype (L. S. Richardson et al., 2020) [Figure 3].

5.6. Cyclic EMT-MET maintains membrane integrity during gestation and a terminal state of EMT at term increases inflammation

Based on the mechanisms described above, fetal membranes maintain their integrity through a cyclic EMT-MET process. This helps to maintain the 10:1 epithelial to mesenchymal ratio during gestation (Myatt & Sun, 2010). TGF- β in the local cellular and amniotic fluid environment can promote EMT and progesterone-mediated MET can revert them back to AEC. This cyclic process balances the number of AMCs in the ECM and allows microfractures to reseal and the membrane to remodel. At term, increased OS promotes two key events that lead to a static state of EMT and the accumulation of AMC: 1) OS causes TGF- β levels to increase and promote EMT; and 2) the OS-mediated reduction in progesterone receptor expression leads to membrane functional progesterone withdrawal, resulting in a lack of MET. This will result in localized inflammation and matrix degradation, predisposing membrane weakening and preparing them for labor [Figure 3].

6. Can premature EMT activation and lack of MET cause preterm birth?

Like senescence, EMT at term is a normal and physiologic process to ensure inflammation and membrane dysfunction and prepare the fetal tissues for parturition. The goal of understanding these mechanisms is to see whether the premature activation of EMT can cause pathways leading to PTB and pPROM. Unpublished reports from our lab have shown that EMT markers and inflammation are evident in PTB, but not in pPROM when tissues are gestationally age-matched. These markers include decreased E-cadherin (epithelial marker) and increased vimentin and N-cadherin (both mesenchymal markers). This contrasts with senescence, which was dominant in pPROM, but not in PTB when membranes were intact. We are not elaborating this section as it is still under investigation. In summary, PTB and pPROM may have similar risk factors but the cellular biologic mechanisms that generate inflammation in fetal membranes are different. pPROM involves an accelerated senescence, generating inflammation that collapses the structural and functional integrity of the fetal membrane, whereas PTB has a terminal state of EMT, forcing inflammation.

7. How do membrane senescence and EMT associated inflammation promote parturition?

Multiple mechanisms have been proposed to destabilize membranes. Collagenolytic processes by endogenously generated MMPs and an imbalance in MMP/inhibitor (tissue inhibitor of matrix metalloproteinase -TIMP) have been well reported. It was unclear, however, how these MMPs are activated. In this review, we introduced two new mechanistic events that can generate localized inflammation. Senescence and EMT-

associated Inflammation of the membranes generate an array of cytokine mediators that belong to different classes with distinct functions that imbalance membrane function and cause them to collapse. These inflammatory mediators are disseminated to the maternal side via senescent and EMT impacted cell-derived extracellular vesicles (EV; exosomes of 50-150 nm). Our studies have shown that EV carrying this inflammatory load at term can reach the myometrium, decidua, and cervix and cause parturition-associated inflammatory activation (Hadley et al., 2018; Jin & Menon, 2017; Menon, 2019; Menon et al., 2017; Sheller-Miller, Trivedi, Yellon, & Menon, 2019). These EV-mediated mechanisms can also be considered as “fetal signals of parturition”, indicating fetal readiness for delivery either at term or preterm. In preterm birth, maternal plasma contains fetal exosomes that are different to those in normal term birth. Fetal exosomes carrying inflammatory pathway mediators can be identified in maternal plasma as early as the late first trimester, indicating underlying pathophysiological conditions (Menon et al., 2020; Menon et al., 2019). Sheller-Miller et al. showed that exosomes carrying inflammatory mediators can cause preterm parturition in animal models (Sheller-Miller et al., 2019). As shown in Figure 4, senescent cells generate inflammatory mediator-enriched exosomes that can reach maternal uterine tissues and cause inflammatory activation (Hadley et al., 2018), and transition them into an active state of labor. Inflammatory mediators are not just cytokines, chemokines or immune cells, but also include SASP and DAMP-enriched exosomes (Sheller-Miller, Urrabaz-Garza, Saade, & Menon, 2017). In summary, we demonstrated two unique forms of inflammatory activation in fetal membranes and how they can mechanistically mediate parturition.

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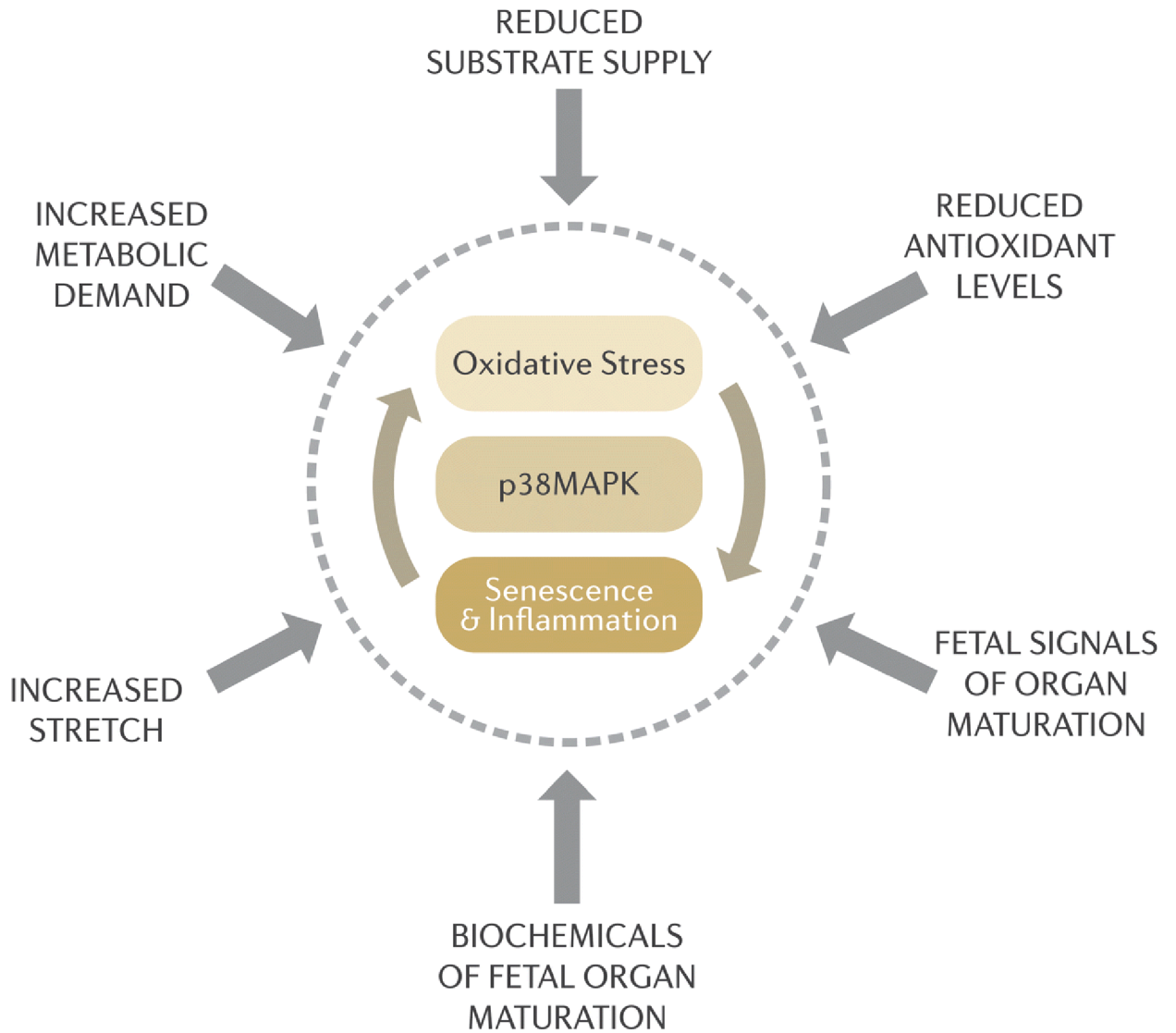


Figure 1: Oxidative stress induces p38MAPK mediated activation of senescence and inflammation in fetal membranes at term.

Various factors as shown in the figure can contribute to excessive reactive oxygen species (ROS) build-up in the intra-amniotic cavity. This ROS can accelerate fetal membrane senescence and senescence-associated secretory phenotype (SASP). Senescence and SASP factors, in a feedback loop, can cause further damage to non-senescent and neighboring tissues to cause further enhanced inflammation. This is a natural and physiological process during normal parturition.

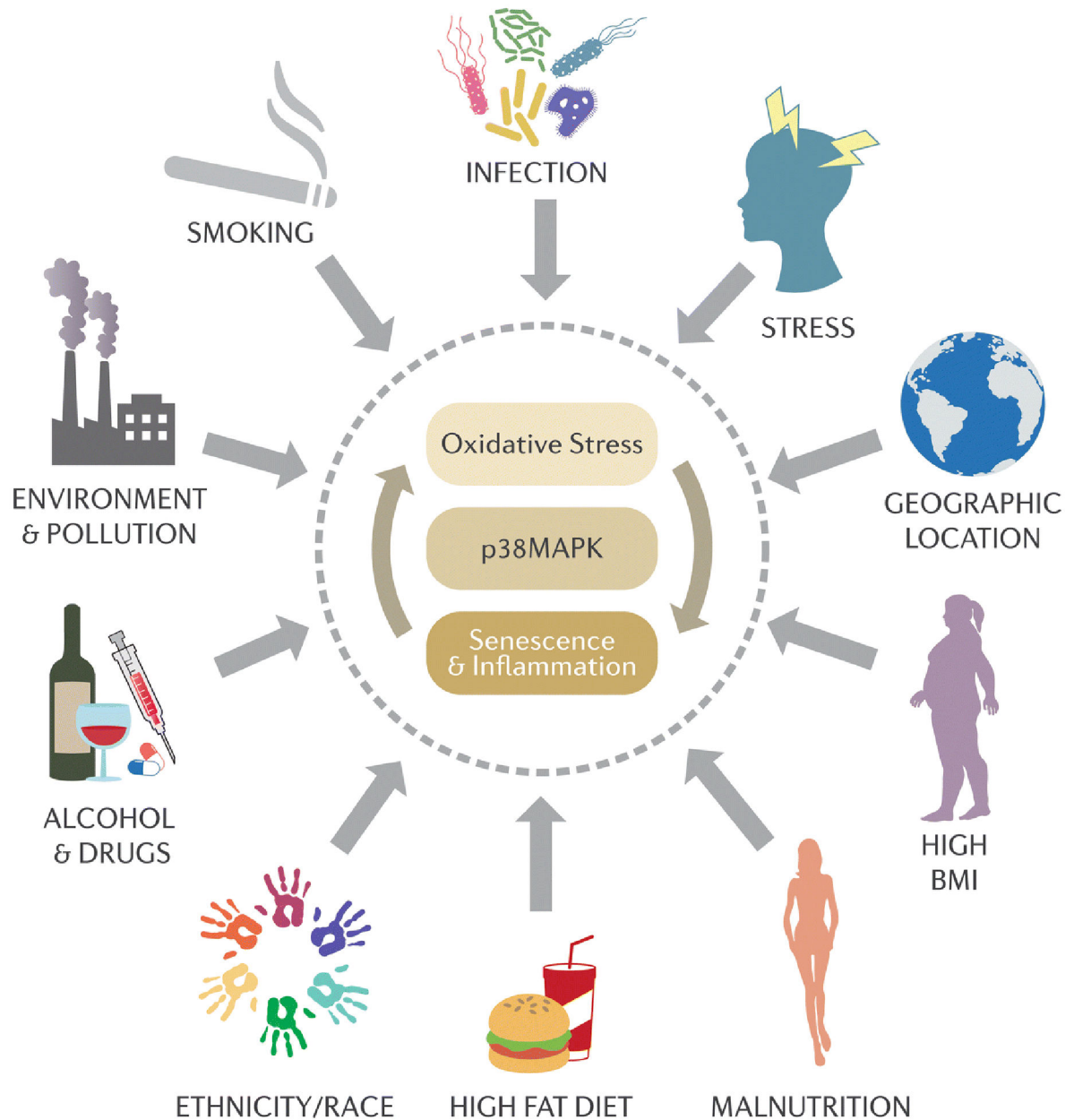


Figure 2: Pregnancy risk factors increase oxidative stress and cause increased activation of p38MAPK, senescence and inflammation in fetal membranes in preterm pregnancies.

Multiple risk factors can cause an increase in ROS in the intra-amniotic cavity. Pathways of ROS generation, characteristics of oxidative stress and p38MAPK activation may not be the same for all risk factors. Regardless, many of these factors can increase p38MAPK activation pathologically prior to term. Senescence and SASP factors can cause preterm labor and or pPROM. Premature activation of p38MAPK is the pathological activation of senescence leading to preterm parturition.

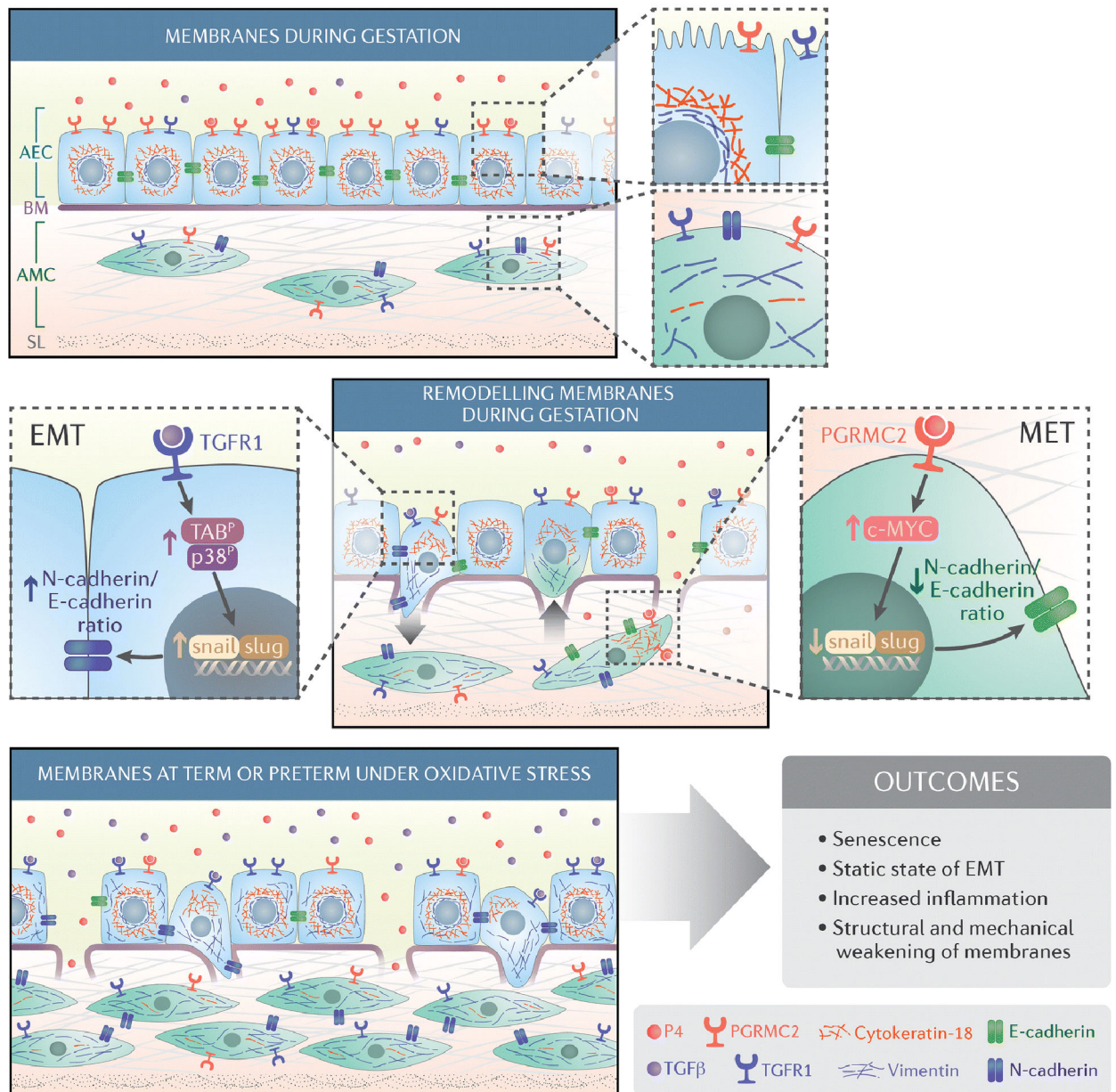


Figure 3: Schematics of changes to membrane structure during normal gestation and parturition.

Top panel: Membranes during gestation: Fetal membranes, specifically amnion layer undergoes a cyclic transition of EMT \leftrightarrow MET that maintains membrane homeostasis and a 10:1 ratio between AEC and AMC.

Middle panel: During gestation, the TGF- β /TAB/p38MAPK-mediated pathway forces the EMT to shed AECs which are transformed into AMC. Since the accumulation of AMC is an unstable state, these cells are transitioned back to AMC by the P4/PGRMC2/c-MYC pathway.

Bottom panel: At term, ROS buildup and p38MAPK activation (see Figure 1), can lead to a terminal state of EMT with the accumulation of AMC and no MET to balance the cell

ratio between AEC and AMC. This is an unstable state of inflammation and cause local inflammatory build-up, matrix degradation and membrane weakening.

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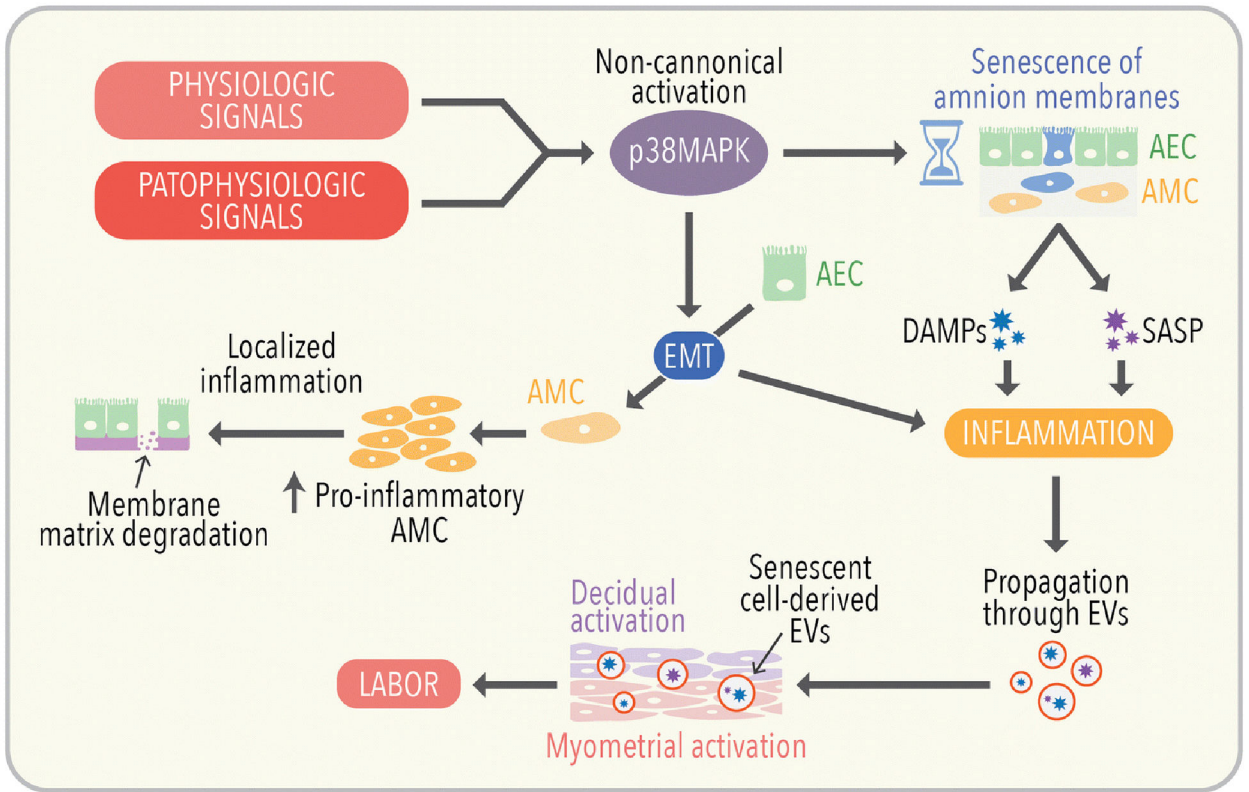


Figure 4: Two distinct mechanisms of inflammatory activation in fetal membranes and fetal inflammatory signaling for parturition.

Physiologic (Fig 1) or pathologic (Fig 2) signals increase ROS in the intra-amniotic cavity and cause non-canonical activation of p38MAPK in human fetal membrane cells. p38MAPK activation can lead to:

1. Senescence of the fetal membrane cells, the production of SASP and the generation of DAMPS.
2. p38MAPK forces a terminal state of EMT and the accumulation of AMCs, causing membrane matrix damage and weakening along with increased localized inflammation. Inflammatory mediators generated are packaged into extracellular vesicles released by fetal membrane cells – fetal signals – that can reach the myometrium and decidua and cause their activation (inflammation) which can transition these tissues from their quiescent state to an active state of labor.