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Complement Component 3 expressed by the endometrial ectopic tissue is involved in the endometriotic lesion formation through mast cell activation

Short Title:

Role of complement component 3 in endometriosis

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Summary: C3 produced by the endometriotic tissue is involved in the lesion development 31 through mast cell activation 32

34 Abstract

The pathophysiology of endometriosis (EM) is an excellent example of immune 35 dysfunction, reminiscent of tumor microenvironment as well. Here, we report that an interplay 36 between C3 and mast cells (MCs) is involved in the pathogenesis of ectopic EM. C3 is at the 37 epicenter of the regulatory feed forward loop, amplifying the inflammatory microenvironment, in 38 which the MCs are protagonists. Thus, C3 can be considered a marker of EM and its local 39 synthesis can promote the engraftment of the endometriotic cysts. We generated a murine model 40 41 of EM via injection of minced uterine tissue from a donor mouse, into the peritoneum of the recipient mice. The wild type mice showed greater amount of cyst formation in the peritoneum 42 compared to C3 knock-out mice. This study offers an opportunity for novel therapeutic 43 44 intervention in EM, a difficult to treat gynecological condition.

46 MAIN TEXT

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48 Introduction

Endometriosis (EM) is a chronic condition that affects about 5-10% of women in 49 50 reproductive age. EM is characterized by pain and infertility as a consequence of the presence of functional endometrial tissue outside the uterine cavity (1). The most common locations for the 51 ectopic implants are the ovaries, peritoneum, and the utero-sacral ligaments. The presence of 52 ectopic tissue in these areas induces a condition of chronic inflammation. Current evidence 53 suggests that immune dysfunction is the most likely causative factor for the EM pathogenesis. In 54 particular, the pathways involved in immune-cell recruitment, cell-adhesion, and inflammatory 55 processes encourage the implantation and survival of endometriotic lesions. Recently, it has been 56 57 shown that the complement system is one of the most predominant pathways altered in EM (2).

The complement system is an important part of the innate immunity and acts as a bridge 58 between innate and acquired immune system. It is involved in host defense against infectious 59 agents and altered self. Three different molecules are responsible for the recognition phase of 60 complement: C1q, Mannose-Binding Lectin (MBL), and C3. The binding of the recognition 61 molecules to the target ligands initiate the three different complement pathways: the classical (via 62 C1q), alternative (via C3) and lectin (via MBL) pathway (3, 4). The three pathways eventually 63 converge in the formation of the C3 and C5 convertases. This then results in the generation of the 64 main effector molecules of the complement system: the opsonins C3b and C4b, the 65 anaphylatoxins C3a, C4a and C5a, and the Membrane Attack Complex (MAC) that causes the 66 target cell lysis. The small complement fragments, anaphylatoxins, cause local inflammatory 67 responses by acting, for instance, on mast cells (MCs), inducing an increase in blood flow, 68 vascular permeability, and leukocyte recruitment (5, 6). 69

The current consensus is that EM involves a local pelvic inflammatory process with 70 altered function of immune-related cells in the peritoneal environment. Recent studies suggest 71 72 that the peritoneal fluid (PF) of women with EM contains an increased number of activated 73 macrophages that secrete various humoral mediators locally, including growth factors, cytokines 74 and possibly, oxidative products (7), leading to the development and progression of EM and EMassociated infertility. Although the contributions of specific immune cell subsets and their 75 mediators to the onset and the course of the inflammatory process in endometrial lesions are still 76 poorly understood, evidence suggests that MCs are crucially involved in the inflammatory process 77 associated with EM. High numbers of degranulated MCs have been found in endometriotic 78 79 lesions (8-13).

Complement component C3 is expressed by the epithelial cells found in endometriotic tissue (14-16). In this study, we examined differential expression of C3 in eutopic and ectopic tissues

derived from EM lesions and its consequences in the pathogenesis of EM using human EM tissues and C3 gene-deficient mice.

85 **Results**

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C3 is present in ectopic, but not in eutopic, endometrium and is locally expressed by endometriotic cells

We initially confirmed the presence of C3 in human endometriotic tissue sections by 88 immunofluorescence (IF, fig. S1) and by immunohistochemical assays (Fig. 1), in uterine and 89 ectopic endometrium. IHC, using anti-human C3 polyclonal antibody, showed moderate 90 91 cytoplasmic expression of C3 by endometrial stromal cells, and rarely in some glandular 92 epithelial cells in proliferative as well as secretory endometrium (Fig.1, A and B). A strong signal was present in adnexal and peritoneal EM (Fig. 1, C to G). C3 was found to be widely distributed 93 in the EM tissue, with variable intensity; it was mostly localized in the glandular-like structures 94 95 and in the vessels.

In order to investigate the local synthesis of C3, total RNA was isolated from EM cysts 96 and C3 gene expression was analyzed by RT-qPCR. Ectopic EM tissues expressed high levels of 97 C3 transcripts compared to the eutopic normal endometrium (Fig. 2A). To assess the contribution 98 of EM epithelial cells to the local C3 production, we isolated EM epithelial/stromal cells from 99 human EM cysts, which showed the characteristic epithelial phenotype (Fig. 2B); ~98% of the 100 101 cells were positive for the epithelial marker, Cytokeratin 8/18 (fig. S2). We analyzed their expression of C3 by RT-qPCR using HepG2 (hepatocyte cell line) as a calibration control. 102 AN3CA cells were used to assess the expression of C3 by normal epithelial endometrial cells. 103 Because the IHC analysis revealed a marked positivity in the EM vessels, the expression of C3 104 was also investigated in endometriotic endothelial cells (EEC) and corresponding eutopic cells 105 106 (uterine microvascular endothelial cells; UtMEC). We observed a very low expression of C3 transcript by both endothelial cell types (fig. S3). Interestingly, EM epithelial/stromal cells 107 expressed a higher amount of C3 compared to AN3CA and HepG2 cells (Fig. 2C). In addition, 108 these cells secreted the C3 protein in the culture supernatant, as measured via ELISA (Fig. 2D). 109 To establish the specificity and exclusivity of C3 expression, we also measured C5 mRNA level; 110 we could not detect C5 transcripts either in the EM tissue or cultured EM epithelial cells (data not 111 shown). 112

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Pro-inflammatory stimuli induce C3 expression by normal endometrial cells, in an in vitro model of EM

Because the ectopic endometrial microenvironment is characterized by inflammation, we 116 set up an in vitro model of EM, stimulating the endometrial cell line, AN3CA, with pro-117 inflammatory cytokines. Our results demonstrated that normal endometrial cells under resting 118 conditions were unable to produce C3; however, when stimulated with TNF- α , and to a lesser 119 extent with IL-1 β , cells started to express C3 mRNA (Fig. 3A), and secrete an increased level of 120 121 C3 protein, as measured by ELISA (Fig. 3B) as well as by western blot analysis (Fig. 3, C and D). We demonstrated also that the stimulation of C3 transcription induced by TNF-a was dose-122 dependent (fig. S4). 123

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125 C3^{-/-} mice are refractory to developing EM cysts in a syngeneic in vivo model of EM

To further investigate the role of C3 in EM pathogenesis, we set up a syngeneic model of EM in C57BL/6 WT or C3 gene-deficient (C3^{-/-}) mice. Estrus was induced in donor animals via

administration of estradiol. Then, the minced uterus of the donor mouse was injected in the peritoneum of 2 recipient mice (Fig. 4A). After 3 weeks, the animals were sacrificed, and the peritoneal cysts were counted. WT mice developed a higher number of cysts compared to C3 deficient mice (Fig. 4B), indicating that lack of C3 could prevent the EM cyst formation.

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Peritoneal washing isolated from WT mice with EM present more degranulated MCs compared to C3^{-/-} mice

We then analyzed the peritoneal washing of WT and C3^{-/-} mice for the presence of 135 infiltrating leukocytes. The samples were cytocentrifuged and the cells were stained with Giemsa. 136 A high number of leukocytes was observed in the peritoneal washing of both WT and C3^{-/-} treated 137 mice. The number of infiltrating leukocytes was comparable with those observed in WT untreated 138 139 (CTRL Untr) mice. Surprisingly, the fluids collected from the WT animals contained a lower number of MCs, compared to the C3^{-/-} mice (Fig. 4, C and D). Next, we stained the 140 cytocentrifuged leucocytes with a monoclonal antibody to CD117 (a marker of mature MCs). The 141 peritoneal washing collected from the WT mice contained CD117 positive cells as well (Fig. 4E). 142 We wanted to confirm if the MCs present in WT peritoneal washing were degranulated. We thus 143 measured the levels of MC enzyme, tryptase, in the murine peritoneal washing (Fig. 4F). The 144 ELISA for tryptase demonstrated that WT mice with EM had a higher amount tryptase compared 145 to C3^{-/-} or CTRL mice. 146

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C3a level is higher in the PF of EM patients, which can act on EM MCs

C3a is one of the most important stimuli for MC activation. Thus, we measured the levels of the C3a in the PF of EM patients, and compared them with those obtained from myoma and fibroma patients (women undergoing laparoscopy, without alterations of peritoneal cavity environment). Our results showed that the PF of EM patients had significantly higher level of C3a compared to the control patients (Fig. 5A).

We next considered that the C3a present in the peritoneal cavity of EM patients could stimulate the MCs within the EM lesions. We therefore stained human EM tissue sections with toluidine blue to highlight MCs present in the tissue. The histochemical analysis confirmed that EM lesions were rich in MCs (Fig. 5, B to D), compared to normal endometrium (fig. S5); the IHC for C3a on the same sections corroborated the presence of this anaphylatoxin as well (Fig. 5, E and F).

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161 C3a is involved in an auto-amplifying loop of inflammation between MCs and EM 162 cells

In order to reconstruct the cross-talk between EM cells and MCs, we set up a co-culture 163 assay, seeding endometrial cell line (AN3CA) at the bottom of a 24-well plate and placing the 164 HMC-1 (a MC cell line) onto the upper chamber of a transwell system (1 µm diameter pores). To 165 understand the role of C3a in this cross-talk, the co-culture was also performed in the presence of 166 a pool of EM-PF and/or a blocking anti-C3a antibody. AN3CA cells, cultured with HMC-1 167 without any other stimulus, expressed low levels of C3. When HMC-1 cells were stimulated with 168 EM-PF in the co-culture, AN3CA cells began to express higher quantities of C3 (Fig. 6A). 169 Surprisingly, AN3CA cells, stimulated with only EM-PF (in absence of HMC-1), were not able to 170 express C3; they produced C3 only in response to EM-PF stimulated HMC-1, or to TNF-α (used 171 as a positive control). Pre-incubation of EM-PF with a blocking monoclonal anti-C3a antibody 172 173 completely abrogated the effect of EM-PF stimulation, bringing it to resting values (Fig. 6B).

174 With a view to gauge a potential novel immunotherapeutic approach for EM, we 175 investigated in our co-culture model the effect of anti-TNF- α antibodies. Infliximab and 176 Adalimumab (both in clinical use) showed a strong effect in blocking C3 expression by AN3CA, 177 whereas Certolizumab did not block significantly the effects induced by the EM-PF on HMC-1 178 cells (Fig. 6C).

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181 Discussion

There is sufficient evidence for the expression of complement components and regulators 182 in human endometrium (17-20). Dysregulation of the complement system is largely considered to 183 play an important role in the pathogenesis of EM (2, 21, 22). The importance of the complement 184 proteins in EM was highlighted recently by high-throughput studies (2, 21-23). The complement 185 system appears to dominate the chronic inflammation in EM and remains a precipitating factor in 186 EM-associated ovarian cancer. Complement factors C3, C4A, C7, factor D, factor B, factor H, 187 and mannose-associated serine protease 1 (MASP1), are differentially expressed in EM compared 188 to normal uterine tissues (2). Furthermore, single-nucleotide polymorphisms in the C3 gene show 189 association with an increased risk of EM and EM-associated infertility (15). 190

The presence of C3 in EM tissue was first highlighted in 1980 (24). Subsequent studies confirmed the presence of higher levels of C3 in the EM lesions (2, 14-16, 21, 23, 25-28). However, how C3 contributes to the pathogenic mechanisms in EM has not been elucidated yet. We show, in this study, the presence of C3 in eutopic and ectopic endometrium (Fig. 1); a strong signal was observed in EM tissue (Fig 1, C to F). C3 was found to be widely distributed in the EM lesions with variable intensity, being mostly localized in the glandular-like structures and in the vessels.

C3 appeared to be produced locally and in higher amounts in the ectopic tissues compared to the normal uterus (Fig. 2A). EM cells, isolated from ovarian cysts of women of fertile age, produced C3 in culture (Fig. 2, C and D) (22, 26, 27). We also sought to investigate the factors produced by endometrial implants located in the abdominal cavity, which influenced the C3 production, and consequent inflammatory microenvironment in the uterine endometrium (25).

A potential candidate for the upregulation of C3 expression by endometrial cells is the PF 203 rich in pro-inflammatory factors (29). PF TNF- α concentrations are elevated in EM patients; its 204 higher concentration correlates with the severity and stage of the disease (30-32). Furthermore, 205 TNF- α and IL-1 β are known to be elevated in EM milieu (29, 33). The stimulation of endometrial 206 207 cells with TNF- α and IL-1 β induced the production of C3. Surprisingly, the stimulation of endometrial stromal cells with EM-PF did not cause increased C3 gene expression, indicating that 208 the concentration of TNF- α in the PF (around 50 pg/ml) was not sufficient for a direct 209 endometrial cell activation (31). We wondered what the source of higher concentration of TNF- α 210 in the endometrial tissue could be. A partial explanation came from the study of $C3^{-1}$ mice. We 211 observed that WT mice developed a higher number of cysts and presented degranulated MCs in 212 PF, as demonstrated by the presence of a considerably higher amount of tryptase, compared to C3⁻ 213 ¹⁻ mice (Fig. 3). Thus, endometrial cells under pro-inflammatory stimuli express C3; the pro-214 inflammatory milieu can induce the generation of C3a, which in turn, can activate the MCs. 215 Activated MCs can release histamine, in addition to secreting TNF- α (and a host of inflammatory 216 factors). 217

One of the possible causes of complement activation in the EM microenvironment is the 218 setting off of the coagulation cascade, which is caused by the typical periodic bleeding in the EM 219 tissue. Thrombin cleaves C3 to C3a and C3b; activated platelets are also involved in C3 cleavage 220 (34). Another activator of the C3 is heme that is released from hemoglobin during hemolysis; 221 222 heme induces deposition of C3 fragments on the erythrocytes (35). Alternative pathway activation can occur through its up-regulator, properdin, binding to activated platelets promoting $C_3(H_2O)$ 223 recruitment and complement activation (36). In addition, stimulation of endothelial cells by C3a 224 or other factors promptly induces expression of P-selectin, which by binding to C3b, induces the 225 formation of C3 convertases (37). 226

We confirmed the presence of higher levels of C3a in the EM-PF compared to control non-EM healthy women (Fig. 5A). The presence of complement component (C4) and complement activation products (C3c SC5b-9) in the PF and in the sera of EM women have been
previously described (*38*). Significantly higher PF levels of C1q, MBL and C1-INH in women
with EM compared to control group has also been reported (*39*).

C3a present in the EM-PF can act, through C3aR interaction, which is abundantly 232 expressed on the MCs present in the EM tissue (9, 25). The involvement of MCs in the EM lesion 233 formation was previously investigated. A recent study showed that numbers of MCs in total as 234 well as activated MCs were clearly increased in the EM lesions in both animals and humans. 235 Thus, the use of MC stabilizers and inhibitors may prove to be effective in treating EM and its 236 associated pain (40). An increased presence of activated and degranulated MCs in deeply 237 infiltrating EM and its close histological relationship with nerves strongly suggests that MCs 238 contribute to the development of pain and hyperalgesia in EM, possibly by a direct effect on the 239 240 nerve structures (10). A diffuse infiltration of numerous MCs was observed throughout the stromal lesions, which exhibited degranulation; scattered granules were also observed. In the 241 eutopic endometrium and normal uterine serosa of the EM patients and the controls, MCs were 242 rarely detected (11). We identified MCs in the EM cyst tissues (Fig. 5, B to D). Very few cells 243 were localized in the endometrial stroma, which is characteristic of endometrial gland-like 244 regions. A good proportion of MCs were localized around blood vessels and the interstitium with 245 fibrosis, that is, the fibrotic interstitium of endometrial cysts. MCs thus are likely to be involved 246 in the development and progression of EM. Localization of MCs suggests its particularly close 247 relationship with fibrosis and adhesion (41, 42). 248

In conclusion, we demonstrated that the complement component C3 is locally synthesized by 249 250 ectopic endometrial tissue, while the normal uterine endometrium does not express C3. Normal endometrial cells, under pro-inflammatory stimuli, produce C3. C3 gene-deficient mice present 251 less endometriotic lesions in a syngeneic EM mouse model. C3a seems to recruit massive MCs in 252 the EM lesions, which can have a damaging pathogenic consequence in the ectopic EM. C3 seems 253 central to a regulatory feed forward loop, and is able to amplify the inflammatory 254 255 microenvironment, in which the MCs are protagonists. This study opens up a new window for the identification of new targets for treating EM. 256

258 Materials and Methods

Patients.

Patients and control women were enrolled at the Institute for Maternal and Child Health, IRCCS Burlo Garofolo, Trieste, Italy. The study group (EM) consisted of a total of 7 women, diagnosed with moderate/severe and minimal/mild EM, according to the revised criteria of the American Society for Reproductive Medicine ASRM (*43*).

Primary cell isolation and culture

Endometriotic cells and endothelial microvascular uterine cells (UtMEC) were isolated, as described earlier (*44*).

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Murine model of EM

C3^{-/-} mice were kindly provided by Prof. Marina Botto, Centre for Complement and Inflammation Research, Department of Medicine, Imperial College, London (UK) and generated as described previously (45). The mouse model of EM was adapted from Somigliana and Mariani (46, 47).

273 Gene expression analysis

RNA was extracted from cells using the kit supplied by Norgen Biotek Corp. (Aurogene, 274 Rome, Italy) according to the manifacturer's instructions and reverse transcripted to cDNA 275 through SuperMix kit (Bioline). qPCR was carried out on a Rotor-Gene 6000 (Corbett, Qiagen, 276 Milan, Italy) using SYBRTM Green PCR Master Mix (Applied Biosystems, Milan, Italy). 277 Supplementary Table S1 shows the primers used for RT-qPCR. The melting curve was recorded 278 279 between 55°C and 99°C with a hold every 2s. The relative amount of gene expression in each sample was determined by the Comparative Quantification (CQ) method supplied as a part of the 280 Rotor Gene 1.7 software (Corbett Research) (48). The relative amount of each gene was 281 normalized with 18S and expressed as arbitrary units (AU), considering 1 AU obtained from 282 HepG2 cells used as a calibrator. 283

Measurement of C3a

The levels of C3a in PF from EM patients and C3 in cell culture supernatant were evaluated by ELISA kits purchased from Quidel.

287 **Co-culture and cell stimulation**

A confluent 24 well-plate of AN3CA cells was stimulated overnight (ON) with 100ng/mL of TNF- α , or 5ng/mL of IL-1 β (both from PeproTech, ListerFish, Milan, Italy), or 10% of pooled EM-PF. For the co-culture study, the cells were seeded at confluence, in the lower part of a 1 μ m pore TW system (Corning, Milan, Italy), whereas the HMC-1 (2x10⁵/TW) was present in the upper part, in the presence of 10% of pooled EM-PF, with or without blocking anti-human C3a (MyBiosource, Aurogene, Milan, Italy; 20 μ g/mL). Subsequently, the cells were lysed for RNA extraction, the supernatant recovered and stored at -80°C.

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435436 Figures and Tables





secretive normal endometrium (B), and endometriotic tissue sections (C). In the lower panel: C3
IHC on abdominal wall (D and F) and adnexal (E and F) endometriosis (EM). AEC (red)
chromogen was used to visualize the binding of anti-human C3 antibodies. Nuclei were
counterstained in blue with Harris Hematoxylin; scale bars, 100µm.



Fig. 2. C3 is locally produced in EM tissue. (A) The gene expression of C3 by endometriotic 445 cysts (n = 4) was investigated by RT-qPCR and compared to normal uterus (n = 3) using HepG2 446 hepatocyte cells as calibrator (AU = 1). 18S was used as the housekeeping gene. Data are 447 expressed as mean ± standard error. * p<0.05 (Mann-Whitney U Test). (B) Example of 448 morphological features of endometrial cells isolated from EM cyst biopsies. Images were 449 acquired by phase-contrast microscope, Leica original magnification: 200x. (C) The C3 mRNA 450 expression was measured by RT-qPCR in EM cells (n = 4) and compared to an endometrial cell 451 line, AN3CA. Data are expressed as mean ± standard error. * p<0.05 (Mann-Whitney U Test). 452 453 (D) Protein levels of C3 were assessed by ELISA in the supernatant of cells maintained in culture for 60 h. Data are expressed as mean \pm standard error. 454

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Fig. 3. C3 is up-regulated by pro-inflammatory cytokines. The stimulation of AN3CA cells 457 with pro-inflammatory cytokines induced the up-regulation of C3 expression. (A) AN3CA cells 458 459 were ON stimulated with TNF-a (100ng/mL) or IL-1β (5ng/mL) and the C3 expression was analyzed by RT-qPCR. Data are expressed as mean of three independent experiments conducted 460 in double \pm standard error. * p<0.05 (Wilcoxon matched pairs test). (B) Measurement of protein 461 level of C3 by ELISA in AN3CA cell culture supernatant stimulated for 36h with TNF-a 462 (100 ng/mL) or IL-1 β (5ng/mL). (C and D) The expression of C3 was evaluated in cell lysates by 463 western blot analysis and the intensity of the bands was measured with Odyssey-LICOR scanner. 464 Data are expressed as mean of three independent experiments conducted in double ± standard 465 error. * p<0.05. 466 467



Fig. 4. In vivo syngeneic mouse model of EM. (A) Treatment regimen of C3 WT and gene-469 deficient mice for generating EM in vivo model. Five C3-/- and WT mice each were injected (i.p.) 470 with minced uterus of a donor mouse; after 3 weeks, the peritoneal cyst formation was evaluated. 471 (B) Number of cysts counted in wild-type (WT, n = 5) or C3-/- (n = 5) mice. Mann-Whitney test 472 P = 0.03. (C and D) Representative images of cytocentrifuged peritoneal washing of untreated 473 WT, EM-induced WT and C3-/- mice (respectively), stained with Giemsa and counted with 474 ImageJ software (Particle Analysis Tool) to obtain relative percentage between total leukocytes 475 and mast cell (MC)/basophil number. MCs/basophils are identified as blue big dots indicated by 476 red arrows. Original magnification 100×. (E) Representative images of cytocentrifuged peritoneal 477 478 lavage of EM-induced WT mice stained with FITC conjugated anti-mouse CD117. Original magnification 100x. (F) Biochemical characterization of tryptase enzyme present in peritoneal 479 lavage of WT vs C3-/- mice by ELISA. * p<0.05. 480



Fig. 5. Peritoneal Fluid (PF) derived from EM patients presented elevated levels of C3a that likely acts on MCs of the EM tissue. (A) C3a ELISA evaluation of PF isolated from EM patients (n = 7) compared to control patient group (n = 6). (B to D) Toluidine blue staining of human EM tissue sections for the evaluation of MCs presence. Black arrows indicated MCs. (E and F) Immunohistochemical analysis of C3a in EM tissue sections. AEC (red) chromogen was used to visualize the binding of anti-human C3a antibodies.

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Fig. 6. The co-culture of MCs with endometrial cells, in the presence of EM-PF, induced the
 expression of C3, which was inhibited by C3a blocking antibody. (A) The C3 gene expression
 was evaluated by RT-qPCR on endometrial AN3CA cells alone (Resting), stimulated with TNF-α

- 496 $(+TNF-\alpha)$ or with a pool of EM peritoneal fluid (+EM-PF); or co-cultured with MCs alone 497 (+HMC-1), or stimulated with PFA as positive control (+HMC-1+PFA) or with EM-PF (+HMC-1)
- 498 1+ EM PF). Similar experiment was performed in the presence of anti-C3a blocking antibody (B),
- 499 and the explained by the graphical abstract in (C). (D) Graphical representation of the blocking
- 500 co-culture experiments. Data are expressed as mean of three independent experiments conducted
- 501 in double \pm standard error. * p<0.05.