

Biological and Therapeutic Implications of the Tumor Microenvironment in Pituitary Adenomas

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1	Biological and Therapeutic Implications of the Tumor Microenvironment in			
2	Pituitary Adenomas			
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43				
44	Abstract			
45	Pituitary adenomas are neoplasms derived from the endocrine cells of the anterior pituitary gland. Most			
46	frequently, they are benign tumors, but may sometimes display an aggressive course, and in some cases			
47	metastasize. Their biology, including their wide range of behavior, is only partly understood. In terms of			
48	therapeutic targeting, most pituitary adenomas are easily treated with the available medical treatments,			
49	surgery, and sometimes radiotherapy. Nevertheless, gonadotroph adenomas lack medical therapeutic			
50	options, and treatment of aggressive pituitary adenomas and pituitary carcinomas remains challenging.			

51 Here, we present an overview of the implications of tumor microenvironment in pituitary adenomas,

52 reviewing its composition and function, as well as published cases that have been treated thus far using 53 tumor microenvironment-targeting therapies. Additionally, we discuss emerging views, such as the concept 54 of non-angiogenic tumors, and present perspectives regarding treatments that may represent future potential 55 therapeutic options. Tumor-infiltrating lymphocytes, tumor-associated macrophages, folliculostellate cells, 56 tumor-associated fibroblasts, angiogenesis, as well as the extracellular matrix and its remodelling, all have complex roles in the biology of pituitary adenomas. They have been linked to hormone 57 58 production/secretion, size, invasion, proliferation, progression/recurrence, and treatment response in 59 pituitary adenomas. From a therapeutic perspective, immune-checkpoint inhibitors and bevacizumab have already shown a degree of efficacy in aggressive pituitary adenomas and pituitary carcinomas, and the use 60 61 of numerous other tumor microenvironment-targeting therapies can be foreseen. In conclusion, similar to 62 other cancers, understanding the tumor microenvironment improves our understanding of pituitary 63 adenoma biology beyond genetics and epigenetics, and constitutes an important tool for developing future therapies. 64

65

66 Introduction

The environment within which a tumor exists, the tumor microenvironment (TME), is composed of vascular structures, non-tumor cells such as immune cells, and non-cellular components such as extracellular matrix (ECM) macromolecules. It has long been accepted that the TME impacts tumor biology and has clinical implications, both as a prognostic tool and a therapeutic target in numerous cancers (1,2). In pituitary adenomas (PAs), the importance of the TME has emerged more recently, but there is now a constantly growing literature on the subject, and growing evidence of the complex and intricate roles that the TME plays in the biology of PAs (3,4).

Here, we review the biological and therapeutic implications of the TME in PAs. Briefly, we provide an
overview of the TME composition and function, and we review all of the published cases of aggressive
PAs and pituitary carcinomas (PCs) that have been treated with therapies that target the TME. Additionally,
we present perspectives regarding treatments that have not so far been used in PAs, but that may represent

potential therapeutic options in the future. Given that both the mechanisms of tumorigenesis and the composition of the TME, along with its biological and clinical implications, might differ depending on the histological subtype of the adenoma, where possible the available data is summarized separately for each major adenoma subtype. Figure 1 depicts T cells, macrophages, blood vessels, and folliculostellate cells in four different histological subtypes of PAs.

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84 1. Immune microenvironment

A. Biology

a) T cells

Tumor-infiltrating lymphocytes (TILs) have complex roles within the TME, that can translate into either 87 88 anti-tumorigenic or pro-tumorigenic effects (2). TILs that are considered to have an anti-tumorigenic effect 89 include CD56+ natural killer cells, CD8+ cytotoxic T cells, CD8+ memory T cells (CD8+CD45RO+), and 90 CD4+ T helper 1 T cells, while TILs generally considered to have an pro-tumorigenic effect include CD4+ 91 T helper 2 T cells and CD4+CD25+FoxP3+ regulatory T cells (1,2,5). However, these anti-tumorigenic and pro-tumorigenic effects are context-dependent, varying across tumor types and tumor stages, and are 92 93 dependent on the specific composition of the immune infiltrate of a tumor and composition of its TME in 94 general, as well as the location of the TIL, i.e. at the tumor invasive margin versus the tumor center (5). Indeed, although CD8+ T cells have been associated with a positive prognosis in most cancer types, they 95 96 have been associated with a negative prognosis in renal cell carcinoma. Conversely, regulatory T cells were 97 associated with a good prognosis in colorectal or gastric cancer (5). Therefore, TILs along with the other TME components need to be characterized in detail in PAs, in order to understand their respective roles. 98 99 To date, the majority of studies in PAs (6–16) have relied on only single/few immune markers, and the 100 quantification of the immune infiltrate varied across these studies. Nonetheless, these studies already 101 provide important information on the immune composition of the TME and its potential roles.

PAs have been shown to be infiltrated by CD3+, CD8+, CD4+, FoxP3+, CD45RO+, and CD56+ cells
(8,12–14,16,17). Compared to the normal pituitary gland (NPG), PAs had more CD4+ T cells and a lower

104 CD8+/CD4+ cell ratio (15). Functioning adenomas showed increased CD3+, CD8+, CD4+, and FoxP3+ 105 IHC infiltrates in comparison to non-functioning (NF) PAs (13,18). Regarding the quantity of various T 106 cell types relative to the different histological subtypes of adenomas, studies have thus far yielded 107 contradictory results (7,10,18). In terms of the genetic background, amongst GH-secreting adenomas, those 108 AIP-mutated had more FoxP3+ cells compared to sporadic adenomas (12). The potential impact of T cells 109 on aggressiveness-related characteristics (9,15,19) is summarized in **Table 1** (4,9–15,18–51).

110

111

b) Tumor-associated macrophages (TAMs)

Affected by cues coming from both tumor cells and TME cells, TAMs can acquire distinct phenotypes and have either anti-tumorigenic or pro-tumorigenic effects (52). TAMs with an M2 phenotype are considered to be pro-tumorigenic and have been consistently associated with a negative prognosis across various cancers, while TAMs with an M1 phenotype are considered anti-tumorigenic and have been associated with a positive prognosis (5). However, in some tumor types, the effect of TAMs on prognosis depends on their location, in the tumor stroma versus the tumor core (5).

PAs have been reported to contain more macrophages (CD68+), more M2-like macrophages (CD163+), 118 119 and a CD163/HLA-DR ratio three-times higher when compared to NPG (the HLA-DR marker being 120 indicative of M1-like TAMs) (15). This is similar to what is seen in other cancers, where M2-like TAMs also predominate (5). TAMs were also reported to be the most abundant immune cell in PAs (15,21,53), 121 122 most notably M2-like TAMs (15,53). The percentage of CD68+ TAMs showed a positive correlation with 123 adenoma size in somatotroph, lactotroph, NFPAs (21), and gonadotroph adenomas (14). In terms of TAM infiltration relative to the different histological subtypes of adenomas, studies have thus far yielded 124 125 contradictory results (10,14,15,18,21,53). Concerning the genetic background, amongst GH-secreting 126 adenomas, those AIP-mutated had more CD68+ cells compared to sporadic adenomas (12).

The potential impact of TAMs on aggressiveness-related characteristics (9–15,18,21,37) is summarized in
 Table 1. Taking into account both TILs and TAMs, it appears that aggressiveness is associated with an
 immunosuppressive TME. From a mechanistic point of view, bidirectional interactions between tumor cells

130 and TAMs take place, with tumor cells recruiting and polarizing TAMs preferentially towards an M2-like 131 phenotype, while M2-like TAMs promote PA aggressiveness, and most notably PA invasiveness 132 (12,14,15,21,37). These findings are consistent with the reports of M2-like TAMs being implicated in tumor 133 growth and development of an invasive phenotype in other types of cancer (5). Moreover, bidirectional 134 interactions take place between TAMs and other components of the TME, especially between M2-like TAMs and angiogenesis. For example, in several other cancers it has been shown that hypoxia leads to 135 136 macrophage recruitment, which in turn facilitates angiogenesis (54). Figure 2 summarizes the roles and 137 interactions of TAMs in PAs (12,14,15,18,19,21,37,55).

138

c) Other immune cells

There is limited published data concerning other immune cells in PAs. PAs have been reported to contain fewer neutrophils than NPG (15). Some studies have reported B cells to be only seen in exceptional cases (6,8,10), while other studies did identify B cells in PAs, but they were fewer compared to TAMs and T cells (14,15,53). Somatotroph adenomas have been reported to have more B cells compared to NFPAs (56), while NFPAs contained more neutrophils than somatotroph adenomas (15).

145

146 B. Therapy

a) Immune-checkpoint inhibitors

The immune-checkpoint inhibitors that have been used to date in aggressive PAs and PCs include: ipilimumab combined with nivolumab, used in six cases, and pembrolizumab, which was used in five cases (57–64). The mechanisms of action of these drugs are illustrated in **Figure 3A**. Their use in aggressive PAs and PCs has been extensively reviewed recently (65).

Six corticotroph carcinomas and one aggressive corticotroph adenoma have been treated with immunecheckpoint inhibitors. In terms of the radiological response, four of the carcinomas showed a partial response that was maintained for 6 to 42 months (duration not available in one case), two of the carcinomas showed stable disease that was maintained for at least 12 months in one of the cases (data not available in the second), while the aggressive adenoma showed progressive disease (57–61,64,65) (**Figure 4**). In two of the cases that initially showed a partial response, a dissociated radiological response was later noted, and in both cases, continuation of immune-checkpoint inhibitors in combination with local treatment (radiotherapy, surgery, and radiofrequency ablation, respectively) proved to be a good therapeutic option (65).

Three lactotroph carcinomas and one aggressive lactotroph adenoma have been treated with immunecheckpoint inhibitors. Regarding the radiological response, one carcinoma showed a complete response maintained for at least 13 months, one showed a partial response maintained for 8 months, the third showed progressive disease, while the aggressive adenoma also showed progressive disease (59,62–64) (**Figure 4**). Of note, the carcinoma that progressed was initially treated with pembrolizumab, and was subsequently treated with programmed cell death protein 1 (PD-1)/programmed death-ligand 1 (PD-L1) pathway inhibitors, and showed continued progression (64).

168 Regarding potential predictive factors of response, data is too scarce to allow firm conclusions. Currently, 169 the most promising predictive factor appears to be a high tumor mutational burden. Negative PD-L1 170 immunohistochemical staining should not preclude the administration of immune-checkpoint inhibitors, 171 since even negative cases responded well. Conversely, the presence of a mismatch repair deficiency did not guarantee a good response (65). Moreover, for corticotroph tumors, it appears important to achieve 172 eucortisolism, since glucocorticoids have an immunosuppressive effect that could lessen the efficacy of 173 174 treatment (65). Currently, too little data is available to reach a conclusion on the eventual superiority of 175 combined immunotherapy over monotherapy (65).

Concerning the tolerance of immune-checkpoint inhibitors, patients cumulatively experienced fatigue, diarrhea, nausea, vomiting, anorexia, fever, transaminitis, autoimmune hepatitis, autoimmune nephritis, rash, myalgia, and possible hypophysitis and progressive weight loss (57–64). As expected, monotherapy was better tolerated than combined treatment. Nevertheless, these adverse events were mostly mild, and led to interruption of dual immunotherapy in only two cases: in one in which progressive disease was simultaneously noted (59), and in a second case, where treatment was continued with nivolumab alone (62). 182

183

b) Perspectives

Potential leads for increasing the efficacy of immune-checkpoint inhibitors include a better patient selection, as well as combining immune-checkpoint inhibitors with angiogenesis-targeting drugs (9,66,67), with radiotherapy (68,69), or with peptide receptor radionuclide therapy (58).

187 At the same time, given both the presence and the important role TAMs appear to play in PAs, targeting 188 TAMs may prove to be a good option for PAs and PCs. This might be even more true for gonadotroph 189 tumors, which are, at least in theory, less amenable to immune-checkpoint inhibitors than other types of PA (70,71). There are numerous TAM-targeting agents and TAM-targeting strategies, including preventing 190 191 TAM recruitment by using, for example, colony stimulating factor 1 (CSF1)–CSF1 receptor inhibitors, C-192 C chemokine receptor (CCR)5–C-C motif chemokine ligand (CCL)5 inhibitors etc.; depleting TAMs by 193 using CSF1–CSF1 receptor inhibitors, triggering receptor expressed on myeloid cells (TREM) inhibitors 194 etc.; reprogramming TAMs by using CD40 agonists, TREM inhibitors etc.; exploiting the antitumor 195 functions of TAMs by using tyrosine protein phosphatase non-receptor type substrate 1 (SIRPa)-CD47 196 inhibitors or sialic acid-binding immunoglobulin-like lectin (SIGLEC)10-CD24 inhibitors; and infusing 197 macrophages that are engineered to eliminate tumor cells and/or reprogram the TME (72).

198

199 2. Folliculostellate cells

200 Apart from immune cells, the TME of PAs includes a number of additional cells, including folliculostellate 201 cells and tumor-associated fibroblasts. Folliculostellate cells are non-endocrine cells that are found in the 202 NPG, where they account for \sim 5% of the cells of the anterior lobe (73,74). These cells are recognized by 203 their morphology and by immunopositivity for S100B protein (75,76). Through both paracrine (76,77) and 204 physical actions (78,79), folliculostellate cells exert numerous roles in the NPG, including a supportive 205 role, phagocytic properties, regulation of hormone secretion, and the production of growth factors and cytokines (vascular endothelial growth factor (VEGF), interleukin 6, basic fibroblast growth factor, 206 207 follistatin, transforming growth factor β etc.) (74,77,80,81). A role as a stem/progenitor cell has also been

often discussed, however it is still debated whether the S100B/SOX2 positive cells, considered to be the
stem/progenitor cells, are a subset of folliculostellate cells or an entirely different population that happens
to also be positive for S100B protein (74,82).

211 PAs have been consistently shown to contain fewer folliculostellate cells than the NPG (35,73,83). 212 However, heterogeneity between tumors exists, and although the vast majority of tumors had very low percentages of S100B+ cells, other tumors were rich in S100B+ cells (35). Concerning their potential roles 213 214 in PAs, consistent with their role in the NPG, folliculostellate cells appear to potentially modulate hormone production/secretion: in acromegaly patients, relationships were shown between S100B+ cells and both 215 growth hormone and prolactin levels (84), while in gonadotroph tumors a positive correlation between 216 217 S100B staining and FSH staining was identified (35). A role of folliculostellate cells in maintaining a more 218 functional, differentiated state, thus closer to the normal state of gonadotroph tumor cells has also been 219 proposed (35). Another proposed role for folliculostellate cells in PAs, consistent with a similar role 220 proposed for a subset of folliculostellate cells in the NPG (74,85), is an antigen presentation function, this 221 proposition being based on the spatial proximity of immune cells and folliculostellate cells co-expressing 222 the S100 protein and MHC class II antigens that was observed in three PAs (86). The potential impact of 223 folliculostellate cells on aggressiveness-related characteristics (35) is summarized in Table 1. Therefore, it appears that folliculostellate cells might play important roles in tumorigenesis-related processes in PAs, but 224 these roles need to be further studied. Hopefully, ways in which these cells could be therapeutically 225 226 targeted/modulated will be developed in the future.

227

228 **3.** Tumor-associated fibroblasts

Tumor-associated fibroblasts are phenotypically and functionally heterogeneous cells, that are seen as a key TME component across various cancers (87,88). Tumor-associated fibroblasts have been mostly associated with tumor progression, but they can also exhibit cancer-restraining functions (88,89). Through the secretion of cytokines, chemokines and growth factors, as well as extracellular matrix components and remodeling enzymes, tumor-associated fibroblasts have been shown to be implicated in tumor growth, ECM remodeling, angiogenesis, tumor invasion, metastasis, immune and metabolic reprogramming of the TME,
as well as in chemoresistance (1,87–89).

236 Tumor-associated fibroblasts have also been identified in PAs (24), and although the literature is, so far, 237 scarce, promising biological and therapeutic implications can be foreseen. Cultured tumor-associated 238 fibroblasts, derived from PAs with cavernous sinus invasion, expressed higher levels of VEGF and 239 interleukin 6 compared to those derived from PAs without cavernous sinus invasion (23,24). Moreover, co-240 injecting GH3 cells with tumor-associated fibroblasts derived from PAs with cavernous sinus invasion resulted in xenografts with higher VEGF expression (expressed mainly by tumor cells) compared to co-241 injecting GH3 cells with tumor-associated fibroblasts derived from PAs without cavernous sinus invasion 242 (23). In addition, cultured tumor-associated fibroblasts derived from PAs secreted CCL2 - a molecule with 243 244 an angiogenic function, whose levels were strongly positively correlated with the microvessel area of PAs 245 (24). The potential impact of tumor-associated fibroblasts on aggressiveness-related characteristics (23,24) 246 is summarized in **Table 1**. These findings point towards a potential role of tumor-associated fibroblasts in 247 angiogenesis, tumor growth and tumor invasion, consistent with the roles observed for these cells in other 248 cancers.

249 From a therapeutic perspective, the somatostatin (SST) type1 receptor was shown to be the predominant 250 SST receptor expressed in tumor-associated fibroblasts from PAs, shown at the mRNA level, and pasireotide was able to decrease the release of interleukin 6 and CCL2 from these cells (24). This suggests 251 252 that pasireotide may exert its role by acting on both pituitary tumor cells and tumor-associated fibroblasts. 253 In pancreatic cancers, pasireotide was also shown to inhibit protein synthesis in tumor-associated fibroblasts, and in combination with gemcitabine it reduced the chemoresistance triggered by the tumor-254 255 associated fibroblast-derived secretome, as well as reducing tumor growth (90). Lastly, numerous other 256 therapeutic strategies are emerging for targeting of tumor-associated fibroblasts, ranging from cell depletion 257 (via targeting of cell surface markers), to altering their activation and/or their function, to tumor-associated 258 fibroblasts normalization (91).

259

260 4. Angiogenesis

A. Biology

Angiogenesis, i.e. the outgrowth of new blood vessels from pre-existing vessels, is a complex process that 262 263 involves numerous signaling pathways and molecules (92,93). Of these, the VEGF/VEGF-receptor 264 (VEGFR) pathway has remained, thus far, the most studied and most targeted pathway (67,94), including in PAs. These newly formed blood vessels are immature, disorganized, and with impaired functionality, 265 266 leading to poor tumor perfusion, hypoxia, and decreased immune cell infiltration and activity. These 267 characteristics further facilitate the acquisition of an aggressive phenotype by tumor cells, reduce the antitumoral immune response, but also decrease the beneficial effects of both chemotherapy and radiotherapy 268 269 (52,93). Therefore, a strategy of vascular normalization, which aims to restore the normal perfusion and 270 oxygenation of the tumor, has emerged in order to mitigate the aforementioned negative effects of 271 angiogenesis (93). It is noteworthy that while angiogenesis plays pivotal roles in tumorigenesis-related processes (52,95), a growing body of literature demonstrates that numerous human tumors in fact lack 272 273 angiogenesis, suggesting that their blood supply is acquired either by vascular co-option or by vasculogenic mimicry (92). These either completely non-angiogenic tumors or partly angiogenic/partly non-angiogenic 274 275 tumors, display biological differences compared to angiogenic tumors, which has important therapeutic 276 implications (92). A better characterization of the blood supply of PAs is therefore needed.

277 The vascularization of PAs (the microvessel density and area, but also the number and percentage of endothelial cells) was found to be lower compared to that of the NPG (49,96-99), and higher in PCs 278 279 compared to PAs (49,50), suggesting a potential role of angiogenesis during tumor progression and 280 malignant transformation of PAs. The exception to this is invasive macroprolactinomas, which have been 281 reported to have vascular densities similar to those of the NPG (28). VEGF expression, on the other hand, 282 was reported to be either lower in PAs compared to the NPG (100,101), or at similar levels (98). Regarding 283 vascular densities and the expression of angiogenesis-related molecules relative to the different histological 284 subtypes of adenomas, studies have so far yielded contradictory results (49,50,96–98,100,102,103). The

potential impact of angiogenesis and of angiogenesis-related molecules on aggressiveness-related
characteristics (9,20,22,28,30,34,36,38–41,48–51) is summarized in Table 1.

287

- B. Therapy
- 289

a) Anti-VEGF-A and anti-VEGFRs therapies

The anti-angiogenic therapies used to date in aggressive PAs and PCs include: bevacizumab, an anti-VEGF-A monoclonal antibody, which has been used in 21 cases; sunitinib, a tyrosine kinase inhibitor that targets multiple tyrosine kinases, including VEGFRs, and which was used in two cases; and apatinib, a tyrosine kinase inhibitor that is primarily a VEGFR2 inhibitor, and which was used in one case (3,64,104–107). The mechanisms of action of these drugs are illustrated in **Figure 3B**.

The 15 patients who received monotherapy with one of these molecules had all previously received at least one course of temozolomide, while for the nine patients receiving concomitant temozolomide, this included both first-course and second-course temozolomide (3,64,104–107). Given the already proven response to temozolomide, the response to combination treatments does not give a clear illustration of the potential efficacy of bevacizumab, therefore in **Figure 4** we present only the radiological response to monotherapy.

300 <u>Bevacizumab</u>

301 Five corticotroph carcinomas and three aggressive corticotroph adenomas received bevacizumab alone (six cases) or combined with temozolomide (two cases) (3,104,106,108). Of the six cases treated with 302 303 bevacizumab alone, the radiological response was reported in 4 cases, with two showing stable disease 304 (3,104,106) maintained for at least 26 months in one case (109), and for an unknown duration in the second (110), one showed progressive disease (108) (Figure 4), while the fourth patient died shortly after an MRI 305 306 performed following 2 cycles of bevacizumab showed radiological stable disease (106). Regarding the 307 cases receiving concomitant temozolomide, one corticotroph carcinoma received bevacizumab, 308 temozolomide, and radiotherapy, and demonstrated radiological complete response, while one corticotroph carcinoma received bevacizumab and temozolomide, and demonstrated radiological stable disease (104). 309

Following ICI failure, one lactotroph carcinoma and one aggressive lactotroph adenoma received bevacizumab alone and showed stable disease. Stable disease was maintained for at least 2 months in one case (59) (**Figure 4**), while in the second case stable disease was demonstrated 6 months after the administration of three cycles of bevacizumab (62) (**Figure 4**). A third case of a lactotroph carcinoma, progressed after ICI failure on combined therapy with bevacizumab, temsirolimus, and valproic acid (64). One aggressive somatotroph adenoma received concomitant bevacizumab and temozolomide, and showed partial response (111).

One aggressive non-immunoreactive adenoma was treated with bevacizumab monotherapy, and showed
stable disease, which was maintained for 18 months (106) (Figure 4).

Four tumors of unknown subtype were treated with bevacizumab alone and showed partial response in one case, stable disease in one case, and progressive disease in one case (3) (**Figure 4**). The duration of maintained partial response, and stable disease, respectively, were not reported. The radiological outcome is unknown in the fourth case (112). An additional four tumors of unknown subtype were treated with combined bevacizumab and temozolomide. The radiological responses consisted of a partial response in two cases, progressive disease in one case, and unknown radiological response in the fourth case (3).

The potential benefit of adding bevacizumab to first-course temozolomide is hard to evaluate at this time because of the generally good efficacy of first-course temozolomide, combined with the small number of cases that have been published with combination therapy. However, regarding the combination of bevacizumab with second-course temozolomide, in the European Society for Endocrinology survey, out of 18 patients who received second-course temozolomide, only two patients showed partial response, of whom one received concomitant bevacizumab (113). Therefore, if second-course temozolomide is envisioned,

331 given the generally poor response that has been reported, addition of bevacizumab might be considered.

Reported adverse events consisted of epistaxis and hypertension in one case (59), nephritis in another case

333 (62), and petechia and fatigue in a third case (106).

334

335 <u>Tyrosine kinase inhibitors</u>

13

Two tumors, one corticotroph carcinoma and a tumor of unknown type, were treated with sunitinib
monotherapy and showed progression (3,108) (Figure 4). Any eventual adverse events were not reported.
An aggressive somatotroph adenoma treated concomitantly with apatinib and temozolomide showed
radiological partial response. Adverse events included hypertension, and hand and foot syndrome (107).

- 340
- 341

b) Perspectives

342 While the VEGF/VEGFR pathway has been thus far the most targeted pathway in cancer, tumors can be inherently resistant or can become resistant to inhibition of the VEGF/VEGFR axis (67), and thus additional 343 therapeutic strategies are needed, for example vascular normalization (93). Strategies to achieve vascular 344 345 normalization include the use of agonists of dopamine receptor type 2, which have been found to have an 346 effect on both pericytes and endothelial cells (93,114,115). Interestingly, in a mouse prolactinoma model, 347 bromocriptine not only blocked tumor growth, but also restored the balance between anti-angiogenic and pro-angiogenic factors, and normalized blood vessels (116). Of note, in the same prolactinoma model, 348 349 axitinib (a TKI inhibiting VEGFRs) restrained tumor growth and improved vascular remodeling, while when bromocriptine was combined with axitinib, intratumoral hemorrhage was suppressed and blood vessel 350 351 perfusion was restored (116). This is consistent with the complementary effect that VEGF/VEGFR pathway 352 inhibitors and drugs targeting alternative angiogenesis pathways were shown to have, and highlight the role that combination therapies might have when targeting angiogenesis (116–118). 353

Interestingly, the angiopoietin-2/angiopoietin-1 receptor axis, classically regarded as an endothelial cell axis, was recently also shown to be active in pituitary tumor cells. Both *in vitro* and *in vivo* evidence indicated that this axis was an exploitable therapeutic target in PAs, most notably in NFPAs, and therefore the use of anti-angiopoietin biologicals or angiopoietin-1 receptor inhibitors might be warranted in these tumors (22).

359

360 5. Extracellular matrix and its remodeling

The ECM consists of a dynamic 3D network of macromolecules (proteins, proteoglycans, glycosaminoglycans etc.), in which soluble factors (cytokines, growth factors etc.) are embedded (1,119– 121). The ECM is implicated in the differentiation, adhesion, migration, proliferation, and survival of cells in both physiological and pathological conditions (121–123). In cancer, the ECM might also influence treatment efficacy, for example, ECM attachment to cancer cells might lead to resistance to apoptosis, a desmoplastic stroma might represent a physical barrier for chemotherapy, while soluble factors might activate survival signaling pathways resulting in resistance to cell death (52).

Regarding the presence and distribution of ECM macromolecules and of integrins (cell membrane receptors 368 369 that transduce ECM signals to cells), differences have been shown to exist between PAs and the NPG 370 (25,124,125). For example, vitronectin was found in PA TME, but not in the NPG, while tumor cells 371 showed a greater range of expression of integrin subunits, simultaneously with the downregulation of $\alpha_3\beta_1$ 372 expression and the loss of $\alpha_6\beta_4$ expression compared with the NPG (125). From a functional viewpoint, 373 laminin expression was found to gradually decrease from the NPG stage to the lactotroph hyperplasia stage 374 and then to the prolactinoma stage, and, in addition, laminin was found to inhibit GH3 and AtT-20 cell proliferation (25,32). Several other ECM macromolecules were also found to impact tumor cell 375 376 proliferation, invasion and migration (26,32) – see **Table 1**, suggesting ECM macromolecules might indeed play a role in PA tumorigenesis. Moreover, ECM macromolecules (laminin, collagen I, and fibronectin) 377 were found to inhibit hormonal production/secretion (25,32). Interestingly, the same ECM macromolecules 378 379 did not impact the hormonal production/secretion of normal rat pituitary cells, suggesting a potential change 380 in integrin expression during tumorigenesis (25,32).

Regarding ECM remodeling, numerous studies have looked at matrix metalloproteinases (MMPs), enzymes that degrade and remodel the ECM, while in the process releasing the molecules that are embedded in it, and also at tissue inhibitors of metalloproteinases (TIMPs), molecules that regulate MMPs (1). Lower TIMP levels and higher levels of MMP activity were found in PA-derived conditioned medium compared to normal pituitary explant-derived conditioned medium (27). *TIMP-3* expression was associated with the deposition of fibrous matrix in PAs (126). In both PAs and PCs, MMPs and TIMPs have been linked to 387 multiple characteristics related to aggressiveness (27,29,31,33,42,43) – see **Table 1**. In addition, by 388 releasing growth factors embedded in a synthetic matrix (matrigel), MMPs were shown, in vitro, to have a 389 role in hormone secretion by pituitary tumor cell lines (27). MMPs have also been linked to angiogenesis 390 in PAs, with MMP-9 positive prolactinomas showing higher vascular densities than MMP-9 negative 391 prolactinomas (29). These data are consistent with the roles the MMP-system plays in other cancers, notably in progression and angiogenesis (127). Other than MMPs, the urokinase plasminogen activator (uPA) 392 393 system is another key player in ECM remodeling and has been shown to be involved in cancer progression, 394 including in cell growth, cell migration, epithelial-mesenchymal transition, and angiogenesis (128). uPA, 395 uPA receptor, tissue-type plasminogen activator, and plasminogen activator inhibitor-1, were also found to 396 be expressed in PAs, and a higher uPA expression was noted in PAs when compared to the NPG (31). The 397 potential impact of the uPA-system, as well as of other ECM components, on aggressiveness-related 398 characteristics (4,25–27,29,31–33,42–47) is summarized in **Table 1**. Therefore, both the ECM and ECM 399 remodeling appear to have important roles in the biology of PAs, and may constitute potential treatment 400 targets. Appealing candidates include the targeting of the MMP-system (129,130) and targeting of integrins 401 (131, 132).

402

403 Conclusions

Similar to the case in other cancers, the TME represents a promising lead for understanding of PA biology 404 405 beyond genetics and epigenetics, and promises to be an important tool for therapeutic purposes. TILs, 406 TAMs, folliculostellate cells, tumor-associated fibroblasts, angiogenesis, as well as the ECM and ECM 407 remodelling, all have complex roles in the biology of PAs, being linked to their functionality, size, invasion, 408 proliferation, progression/recurrence, and to their response to treatment. From a therapeutic viewpoint, 409 immune-checkpoint inhibitors and bevacizumab have already shown some efficacy in treating aggressive 410 PAs and PCs, and the use of numerous other TME-targeting therapies can be foreseen. Further work is still 411 required for a better understanding of TME composition, its roles, and of the extensive crosstalk that takes 412 place between adenoma cells and the TME, as well as between the different components of the TME.

413 Ideally, future studies will integrate a more intricate and complete view of the network of interactions that

414 may exist between tumor cells and the components of the PA TME. Additionally, given the potential

differences in TME composition and involvement in different subtypes of PAs, it will be important to study

- 416 each of the histological subtypes of PA separately.
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792 Figure legends

Figure 1. Immunohistochemical stainings depicting T cells (CD3+), macrophages (CD68+), blood vessels
 (CD34+), and folliculostellate cells (S100B+) in four different histological subtypes of pituitary adenomas.
 Original magnification ×200.

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797 Figure 2. Roles and interactions of tumor-associated macrophages (TAMs) in pituitary adenomas. Derived 798 from data from (12,14,15,18,19,21,29,37,55). Dashed arrows indicate that only associations or correlations 799 (i.e., no causality, nor any mechanism demonstrated) are available so far. Question marks indicate that 800 additional, currently unknown, mechanisms might be also responsible. *In rat estrogen-induced 801 prolactinoma. Figure created using illustrations from Servier Medical Art (https://smart.servier.com/). Abbreviations: associated with (as.), correlated with (cor.), matrix metalloproteinase (MMP), S100 802 803 calcium-binding protein A9 (S100A9), epithelial-to-mesenchymal transition (EMT), enhancer of zeste 804 homolog 2 (EZH2), growth factors (GFs), C-C motif chemokine ligand (CCL), colony stimulating factor 1 805 (CSF1), granulocyte-macrophage colony-stimulating factor (GM-CSF), cluster of differentiation (CD), human leukocyte antigen – DR isotype (HLA-DR), vascular endothelial growth factor (VEGF), VEGF-806 807 receptor (VEGFR).

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Figure 3. Schematic representation of the mechanism of action of immune-checkpoint inhibitors (A), and 809 810 of anti-VEGF-A and anti-VEGFRs therapies (B) used in pituitary carcinomas and aggressive pituitary 811 adenomas. Immune-checkpoint inhibitors are monoclonal antibodies that block either CTLA-4 (ipilimumab) or PD-1 (nivolumab and pembrolizumab), that are found on T cells, or PD-L1, the ligand of 812 813 PD-1, which is found on antigen-presenting cells and on tumor cells. Bevacizumab is a monoclonal 814 antibody that blocks VEGF-A. Sunitinib and apatinib are tyrosine kinase inhibitors that target VEGFR1 815 and VEGFR2 (amongst other targets) for sunitinib, and VEGFR2 for apatinib. Figure produced using illustrations from Servier Medical Art (https://smart.servier.com/). *Not used so far in pituitary carcinomas 816 817 and aggressive pituitary adenomas. Abbreviations: programmed cell death protein 1 (PD-1), programmed

818	death-ligand 1 (PD-L1), cytotoxic T-lymphocyte-associated protein 4 (CTLA-4), vascular endothelial
819	growth factor A (VEGF-A), VEGF-receptor (VEGFR).
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821	Figure 4. Radiological response of pituitary carcinomas and aggressive pituitary adenomas treated with
822	therapies targeting the tumor microenvironment. Derived from data from (3,65,104,105). Abbreviation:
823	non-immunoreactive (NIR), immune-checkpoint inhibitors (ICIs).
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	Invasion (Patients)	Proliferation (Patients)	Progression/Recurrence/Resistance to treatment (Patients)	In vivo/in vitro
	F: CS inv. as. with less CD8+ cells (19) & more CD68+ TAMs (14)	F: Higher Ki-67 index in PAs with high VEGF expression vs. PAs with no VEGF expression (20)	F: Resistance to first-generation SRLs as. with less CD8+ cells (19)	Macrophage-conditioned media induced morphology changes, EMT activation & increased inv. of GH3 cells (12,15)
				CCL17 promoted increased EMT changes, migration, inv. & pf. of GH3 cells (21)
Somatotroph				Angiopoientin-2 promoted the pf. of GH3 cells (22)
20111100 0 PT				TAFs derived from PAs with CS inv. promoted pf. of GH3 cells <i>in vitro</i> , as well as the growth of GH3-derived xenografts in mice (23)
				TAF-conditioned medium induced morphology changes, EMT activation, & increased inv. & migration of GH3 cells (24)
	F: Inv. macroprolactinomas as. with higher microvessel density than non-inv. macroprolactinomas (28)F: Inv. macroprolactinomas expressed more MMP-9 than non-inv. macroprolactinomas (29)			Laminin inhibited GH3 cell pf. (25)
				Different collagen subtypes impacted differently GH3 cell inv. & migration strategy (26)
Lactotroph			F: Resistance to bromocriptine as. with higher endothelial expression of	MMP inhibition decreased GH3 cell pf. (27)
p			endocan (30)	IL-2, IL-6, BMP-4 & CXCL12 stimulated, while TGF- β inhibited GH3 cell pf. (4)
				CXCL12 stimulated, while TGF- β inhibited GH4 cell pf. (4)
	F: CS inv. as. with lower tPA expression (31)			Collagen IV, fibronectin & activation of β1 integrin stimulated, while collagen I & laminin inhibited AtT-20 cell pf. (32)
				TIMP-1 inhibited AtT-20 cell inv. & migration (33)
Corticotroph				MMP inhibition decreased AtT-20 cell pf. (27)
				RSUME silencing inhibited inv. of AtT-20 cells (34)
				IL-1 & IL-2 stimulated, while IFN-γ & LIF inhibited AtT-20 cell pf. (4)
Gonadotroph	CS inv. as. with more CD68+ & CD163+ TAMs, & less thick & thin collagen fibers (14)	Ki67 index \geq 3%, mitosis $>$ 2/10 HPFs & proliferative status as. with fewer S100B+ cells (35)	Progression after surgery as. with higher VEGF expression (36)	
			Less progression-free time as. with higher <i>VEGFR2</i> expression (36)	

844 Table 1. Potential impact of tumor microenvironment on aggressiveness-related characteristics of pituitary adenomas and carcinomas. Derived from data from (4,9–15,18–51).

Unspecified or combined NFPAs	CS inv. as. with more CD163+ TAMs (9), M2- like/M1-like TAMs ratio >1 (37) & higher FoxP3/CD8 ratio (9) CS inv. as. with more VEGF & VEGFR1 (9) Inv. of hormone-negative PAs as. with high endothelial expression of endocan (38)		Higher MMP-9 expression in recurrent NFPAs (for the second surgery compared to the first) (29)	Cultured M2 versus M1 macrophages promoted increased inv., motility & pf. of primary NFPA cultures (37)
	Knosp grades cor. with CD68+ TAMs number (10), endocan expression relative to CD34 expression (39) & tumor cell expression of endocan (40)	Ki67 index ≥3% as. with more FoxP3+ cells, lower CD8/FoxP3, CD68/FoxP3 & CD8/CD4 ratios, & a "deleterious immune phenotype" (CD68 ^{hi} CD4 ^{hi} FOXP3 ^{hi} CD20 ^{hi}) (15), as well as as. with fewer S100B+ cells (35) Ki67 index >3% as. with increased expression in CD45+ cells (13) Higher mitotic count & p53	Persistence/recurrence after surgery as. with the presence of CD45+ cells (11) Recurrent PAs had higher CD45+ $\&$	
	CS inv. as. with high infiltration of M2-like TAMs, higher expression of lactate-related genes, more		lower CD163+ infiltrates than primary PAs (18)	
Combined	 CCL17 & high CCL17 expression (21), lower VEGI levels (41), higher MMP-2, -9 (42) & MMP-14 expression (33), & lower TIMP-2 expression (31) 		Recurrence/progression after surgery as. with high CCL17 expression (21) & endothelial expression of endocan (48)	
subtypes	Inv. as. with more VEGF, HIF-1 α , RSUME (34), higher MMP-9 expression (43,46) & serum levels (43), lower TIMP-1 expression & serum levels (43), higher CVCI 12 & CVCP4 comparison more		Carcinomas as. with increased vascularization compared to PAs (49,50)	
	Inight CACL12 & CACR4 expression, morecells (13)CXCL12-positive & CXCR4-positive cells (44),cells (13)higher IL-6 & TNF- α expression (45), higher IL-17Higher mitotic count & p53expression & serum levels (46), higher IL-17expression as. with endothelialreceptor expression (46), & positively cor. with β 1integrin & FAK expression (47)		No recurrence/regrowth positively cor. with VEGF expression (51)	
		Lower post-operative survival rate as. with higher MMP-9 & lower TIMP-1 expression (43)		
845 Abbreviations: non-functioning nituitary adenoma (NEPA) functioning (F) cavernous sinus (CS) invasion (inv) was associated (as) tumor-associated macrophages				

Abbreviations: non-functioning pituitary adenoma (NFPA), functioning (F), cavernous sinus (CS), invasion (inv.), was associated (as.), tumor-associated macrophages (TAMs), matrix metalloproteinase (MMP), tissue-type plasminogen activator (tPA), vascular endothelial growth factor (VEGF), VEGF-receptor (VEGFR), pituitary

adenoma (PA), correlated (cor.), C-C motif chemokine ligand (CCL), endothelial growth inhibitor (VEGI), hypoxia-inducible factor-1α (HIF-1α), RWD-containing
sumoylation enhancer (RSUME), C-X-C motif chemokine (CXCL), C-X-C chemokine receptor (CXCR), interleukin (IL), tumor necrosis factor-α (TNF-α), vascular

849 somatostatin receptor ligands (SRLs), tissue inhibitors of metalloproteinases (TIMP), focal adhesion kinase (FAK), proliferation (pf.), epithelial-to-mesenchymal transition

850 (EMT), tumor-associated fibroblast (TAF), bone morphogenetic protein (BMP), transforming growth factor- β (TGF- β), interferon- γ (IFN- γ), leukaemia inhibitory factor

851 (LIF).

Graphical Abstract

Tumor Microenvironment Implications In Pituitary Adenomas



- tumor cells
- tumor-associated macrophages (TAMs)
- tumor-infiltrating lymphocytes (TILs)
- folliculostellate cells (FSCs)
- tumor-associated fibroblasts (TAFs)
- blood vessel
- enzymes
- cytokines/chemokines
- < extracellular matrix (ECM)

Biological → linked to pituitary adenoma:

- Functionality
- > Size
- Invasion
- Proliferation
- Progression/recurrence
- Treatment response

<u>Therapeutic \rightarrow targeting of:</u>

- The immune infiltrate: immune-checkpoint inhibitors, TAM-targeting
- Angiogenesis: bevacizumab, tyrosine kinase inhibitors, vascular normalization
- TAFs & FSCs
- The ECM & its remodeling

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Figure 2 Increase TC invasion (CCL17, upregulation of MMP9 & S100A9 in TCs, ?) Increase TC motility/migration (CCL17, upregulation of S100A9 in TCs, ?) Induce TC EMT activation (CCL17, ?) Increase TC proliferation (CCL17, upregulation of EZH2 in TCs, ?) Increase secretion of cytokines/chemokines/GFs by TCs

TAMs Tumor cells TCs) TAM recruitment (CCL5, CSF1, ?) M2-polarization (lactic acid/lactate, ?) Collagen degradation (MMP9, ?) M1-polarization (GM-CSF, ?)

CD163:HLA-DR ratio + cor. microvessel density & area Higher CD163 expression as. higher VEGF expression Higher CD68 expression as. higher VEGFR2 expression Endothelial endocan expression as. more CD68+ cells M2-like TAM infiltrate increased parallely with neovascularization, & then, hemorrhage*





Figure 3

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A. Immune-checkpoint inhibitors

B. Anti-VEGF-A and anti-VEGFRs therapies





Figure 4 Click here to access/download;Figure;Ilie_fig4.pdf * Radiological response of pituitary carcinomas and aggresive pituitary adenomas treated with tumor microenvironment-directed therapies



Essential points

- Tumor-associated macrophages are the most abundant immune cell in pituitary adenomas, and have been linked to the size, invasion, and proliferation of pituitary adenomas most notably to their invasion.
- Tumor-infiltrating lymphocytes have been linked to invasion and proliferation of pituitary adenomas, and to the response to treatment of somatotroph adenomas.
- Folliculostellate cells have been linked to hormone production/secretion and to the proliferation of pituitary adenomas.
- Angiogenesis and angiogenesis-related molecules have been linked to invasion, proliferation, and the progression/recurrence of pituitary adenomas, as well as to response to treatment in lactotroph adenomas.
- The extracellular matrix and its remodeling have been linked to hormone production/secretion, to the invasion and proliferation of pituitary adenomas most notably to their invasion.
- Substantial crosstalk takes place between adenoma cells and the tumor microenvironment, as well as between the different components of the tumor microenvironment.
- Immune-checkpoint inhibitors and bevacizumab have already shown some efficacy in treating aggressive pituitary adenomas and pituitary carcinomas.