

Letter to the Editor: Classification of Pituitary Adenomas

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In their article, Sanno *et al.* [15] address an important issue concerning the classification of pituitary adenomas. Their proposed classification is based on the tumour's histological data. Yet, pituitary adenomas are treated by a number of specialists, not all of whom have histological data available. Therefore, I would like to present an alternative classification scheme which can be shared by endocrinologists, gynaecologists, ophthalmologists, neurosurgeons, pathologists, and researchers alike.

Advances in the understanding of an adenoma's hormone synthesis (genotype), hormone storage (immunophenotype), and hormone release into the blood stream (clinical phenotype) have led to several parallel and sometimes confusing classifications of pituitary adenomas. The present confusion on what exactly is, e.g., a prolactinoma or a corticotroph cell adenoma (does it produce, store, or release ACTH?) has led to the occasional use of such terms as "PRL-oma", "GH-oma", "ACTH-oma", and "TSH-oma" [2, 16] or "somatotropinoma" and "corticotropinoma" [3]. Unfortunately these terms do not clarify whether in the case of "PRL-omas" the adenomatous cells produce, store, or release PRL or if the excessive release of PRL is caused by the stalk effect or raised intrasellar pressure which would mean that the adenomatous cells do not even derive from the mammotroph cell line. Currently, due to these difficulties and to overlap created by the various terminologies, tumours of the same cell line may be classified differently depending on whether one refers to their genotype, immunophenotype, or clinical phenotype. Vice versa, tumours deriving from different cell lines may incorrectly be classified as identical tumours. In an attempt to overcome the difficulties in making a correct diagnosis

based on clinical data, some authors have developed regression equations in order to extract most information from the clinical data [5]. The introduction of mathematical tools to improve the accuracy of the clinical diagnosis demonstrates the great difficulties encountered when trying to determine an adenoma's true nature in the absence of histological data. On the other hand, classifications based on a tumour's histological examination can often not be shared with clinicians who treat their patients before or without surgery.

Therefore, I would like to emphasize an unequivocal classification of the various adenoma types by crossreferencing an adenoma's cell line of origin, its immunoreactivity, and its clinical phenotype. At the same time, existing anatomical classifications should be used to complement this classification. The aim is to come to a diagnosis that can be shared by researchers, pathologists, and clinicians.

Proposed Classification of Pituitary Adenomas

The system is based on the work by Kovacs [8] and Landolt [9] whose contributions replaced the classical histological designation of pituitary adenomas as acidophilic, basophilic, and chromophobic with the more meaningful designation as adenomas of a certain anterior lobe cell line. It takes into account new findings that link somatotroph, mixed mammotroph and somatotroph, and mammosomatotroph adenomas to one basic cell line committed to the production of growth hormone [12].

The classification presented in Table 1 has been published elsewhere [18]. It takes into account an adenoma's direct and indirect (stalk effect, raised

Table 1. Classification of Pituitary Adenomas According to ^a

Cell line(s) (hormone synthesis ^b)	Immunophenotype (hormone storage ^c)	Clinical phenotype (hormone release ^d)
Clinically endocrine-active pituitary adenomas ^e		
Mammotrophic cell	PRL	Prolactinoma
Corticotrophic cell	ACTH	Cushing's disease
Corticotrophic cell	ACTH	Nelson's syndrome
Somatotrophic cell	GH, or GH and PRL	Acromegaly, prolactinoma, or both
Acidophil stem cell	GH and PRL	Prolactinoma, acromegaly, or both
Thyrotrophic cell	TSH	TSH-secreting PA
Unclassified plurihormonal cell(s)	Any combination of PRL, GH, ACTH, LH, FSH, TSH, and α SU	Plurihormonal PA of one dominant clinical phenotype ^f
Clinically endocrine-inactive ("nonsecreting") pituitary adenomas ^g		
Null cell	Nonreactive	Null cell adenoma
Oncocytic cell	Nonreactive	Oncocytoma
Gonadotrophic cell ^e	LH, FSH, or LH and FSH	Gonadotrophic PA
Corticotrophic cell ^e	ACTH	Silent corticotrophic PA
Somatotrophic cell	GH	Silent somatotrophic PA
Acidophil stem cell	GH and PRL	Nonsecreting acidophil stem cell PA
Thyrotrophic cell	TSH	Silent thyrotrophic PA
Unclassified pure α SU cell	α SU	Pure α SU-secreting PA
Unclassified plurihormonal cell(s)	Any combination of PRL, GH, ACTH, LH, FSH, TSH, and α SU	Nonsecreting plurihormonal PA

^aPRL prolactin; ACTH adrenocorticotrophic hormone; GH growth hormone; TSH thyroid-stimulating hormone (thyrotropin); LH luteinizing hormone; FSH follicle-stimulating hormone; α SU α subunit.

^b As determined by in situ hybridization and electron microscopy.

^c As determined by electron microscopy and immunostaining.

^d As determined clinically by symptoms, signs, and serum hormone levels.

^e Cosynthesis of the clinically inactive α SU and its release into the bloodstream is possible.

^f LH and FSH are clinically inactive; their release into the bloodstream can be measured.

^g May present clinically as prolactinoma when stalk effect or raised intrasellar pressure leads to an increased release of prolactin by non-adenomatous cells or when PRL is cosynthesized by tumorous cells.

intracellular pressure) endocrine activities. The direct endocrine activities are: hormone synthesis (genotype), hormone storage (immunophenotype), and hormone release (clinical phenotype). The indirect endocrine activities are: compression of non-adenomatous anterior lobe tissue resulting in partial pituitary insufficiency and excessive release of PRL by non-adenomatous cells caused by the stalk effect or raised intracellular pressure.

The genotype is determined by molecular genetic techniques like in-situ hybridization [13] allowing one to identify both DNA and mRNA in adenomatous cells revealing their genetic fingerprint. It is the most accurate information obtainable concerning the cell line of origin and is mostly used in research.

The immunophenotype is determined by the tumour's immunostaining properties when adenomatous tissue is stained with antibodies to the various anterior lobe hormones or hormonal compounds such as the alpha-subunit. This technique reveals which hormone(s) or hormonal compound(s) are stored by tumorous cells. It is a less accurate way of determining an adenoma's cell line of origin since hormonal products may be released without the delay of tissue storage. Immunostaining for the various anterior lobe hormones should be part of today's routine diagnostic procedures.

The clinical phenotype is the least accurate information concerning the nature of a pituitary adenoma because: (a) hormone excess may be insufficient to cause any clinical symptoms or signs, (b) release of hormonal products may be clinically silent, (c) due to raised intracellular pressure or the stalk effect prolactin may be released in sufficient amounts to cause amenorrhea and galactorrhea even though the tumour might not produce prolactin [1, 10, 11], (d) the clinical phenotype may be determined by only one of multiple co-existing tumours of differing immunophenotypes [7], (e) tumours may change clinical and immunophenotype [14], and (f) in addition to hormone excess, tumours may cause partial pituitary insufficiency.

The proposed classification scheme keeps the doctor aware of the degree of certainty concerning the true nature of a pituitary adenoma. For obvious reasons, a classification of a patient's adenoma based on clinical data only leads to the least reliable diagnosis. One might grade the three groups represented in the three columns to which a pituitary adenoma may be designated as fair evidence (clinical phenotype), good

evidence (immunophenotype), or best evidence (cell line) for the correct diagnosis.

The anatomical classification schemes developed by Hardy [4] and modified by Wilson [17] grade the tumour's extent of sellar destruction and its extrasellar extension. A newly developed classification by Knosp *et al.* [6] based on magnetic resonance imaging (MRI) grades an adenoma's invasion into the cavernous sinus space. At surgery, tumour extending beyond the intercarotid line in coronal MRI proved very likely to be associated with invasion [6]. These purely anatomical classifications should be used in addition since the data contained in these schemes have prognostic value and are helpful in choosing the best treatment.

The proposed classification in combination with the various grading schemes should facilitate interdisciplinary and interinstitutional communication on what tumour exactly clinicians, pathologists, different institutions and articles in scientific publications are referring to. It is designed to heighten the awareness that there are several aspects characterizing a pituitary adenoma ranging from its cell line of origin to its immunophenotype and to its clinical presentation. A heightened awareness of such aspects may help in improving study designs and in the understanding of the tumour biology and the clinical course of the various adenoma types.

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