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COMMENTARY

THE CHROMOGRANINS A AND B: THE FIRST 25 YEARS AND FUTURE PERSPECTIVES

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CONTENTS

1. INTRODUCTION	497
2. ISOLATION AND QUANTITATIVE DETERMINATION	499
3. PHYSICO-CHEMICAL PROPERTIES	499
4. ORGANIZATION OF THE GENES	501
5. DISTRIBUTION IN NEUROENDOCRINE AND IMMUNE SYSTEM	501
6. PRESENCE IN THE NERVOUS SYSTEM	505
7. SUBCELLULAR LOCALIZATION	506
8. PHYLOGENY	506
9. ONTOGENY	507
10. SECRETION, PRESENCE IN SERUM AND CEREBROSPINAL FLUID AND FINAL FATE	507
11. CELL-FREE SYNTHESIS	508
12. CELLULAR SYNTHESIS	508
12.1. Rate of synthesis	508
12.2. Post-translational modifications	509
12.2.1. Glycosylation	509
12.2.2. Phosphorylation	509
12.2.3. Sulfation	509
12.2.4. Pyroglutamylation	510
12.2.5. Carboxymethylation	510
12.2.6. α -Amidation	510
12.2.7. Formation of disulfide bonds	510
13. PROTEOLYTIC PROCESSING	510
13.1. Rate and degree of proteolysis	510
13.2. Pathways of proteolysis	511
13.3. Proteases involved in processing	511
14. REGULATION OF BIOSYNTHESIS	512
14.1. Regulation of synthesis by hormones	512
14.1.1. Chromogranin A—a peptide regulated by steroids	512
14.1.2. Regulation of chromogranin B	512
14.2. Regulation of synthesis by stimulation of cells	512
14.3. Regulation by second messengers	513
15. FUNCTION	513
15.1. Involvement in catecholamine storage	513
15.2. Enzyme inhibition	513
15.3. Binding of calcium and possible consequences for granule formation	514
15.4. Precursors of functional peptides	514
16. CONCLUSIONS	515
ACKNOWLEDGEMENTS	516
REFERENCES	516

1. INTRODUCTION

Twenty-five years ago, in 1967, Blaschko, Comline, Schneider, Silver and Smith²⁷ named the soluble proteins of bovine chromaffin granules chromo-

granins, and in the same year Schneider, Smith and Winkler^{28,5} coined the term chromogranin A (CgA) for their major component (see Fig. 1). When this Journal, *Neuroscience*, was started in 1976, the first commentary^{36,2} in volume 1 dealt with the early investigations on the chromogranins. It now appears timely to have a commentary derived from two questions. (i) What is finally established after 25 years? (ii) What has still to be done? Early studies have already revealed that the chromogranins were a

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Abbreviations: CgA, chromogranin A; CgB, chromogranin B; PSP, parathyroid secretory protein; RFLP, restriction fragment length polymorphism; SDS, sodium dodecyl sulfate.

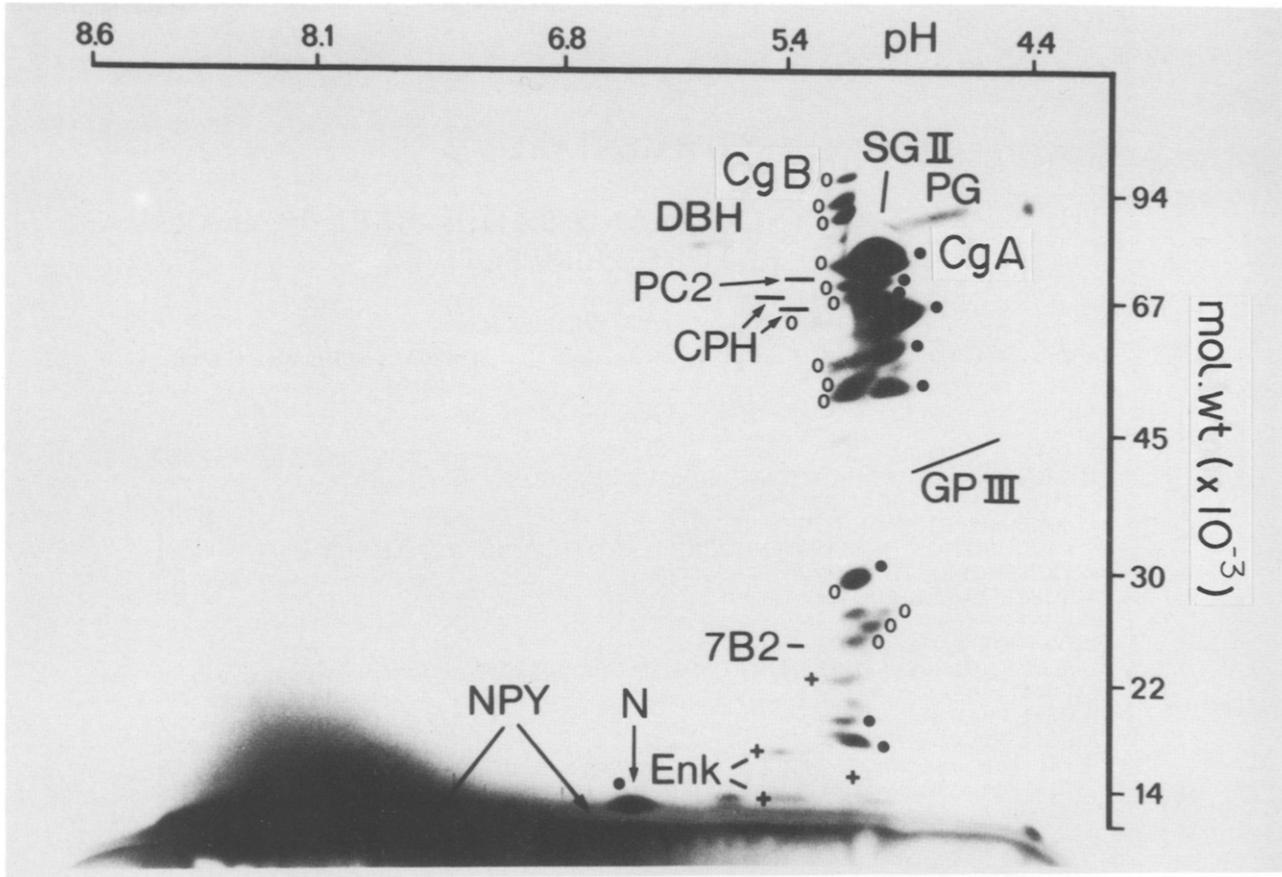


Fig. 1. Chromogranins in bovine chromaffin granules. The soluble proteins of bovine chromaffin granules were separated by two-dimensional electrophoresis.⁸ For CgA (●), CgB (○) and enkephalins (Enk: +) the immunologically cross-reacting spots are marked. For CgA and CgB the largest molecules represent the respective proproteins. The spot (marked N) represents the N-terminal peptide, derived from CgA.^{14,373} Further spots are: NPY, neuropeptide Y; PG, proteoglycan form of CgA; DBH, dopamine β -hydroxylase;³⁶⁴ SGII, secretogranin II; 7B2: see Ref. 223. For the following minor components of the soluble lysate their position is marked; however, spots cannot be visualized with Coomassie Blue, but only after enrichment with lectin affinity chromatography (CPH, carboxypeptidase;¹⁹³ GPIII, glycoprotein III;⁸⁹ PC2, endoprotease^{183a}).

complex mixture of proteins³⁶² (see Fig. 1). This is due to the presence of a large number of distinct proteins, but is also caused by the fact that these proteins are processed within the chromaffin granules by endogenous proteases (see below). Thus in addition to CgA several breakdown products of this proprotein are present¹⁴⁹ (see Fig. 1).

In 1984^{79,366} we described a second group of acidic proteins, i.e. the chromogranins B (CgB) which could be immunologically differentiated from CgA. These proteins were subsequently also called secretogranin I;²⁷⁴ however, since they are related to CgA (see below) in some respects this name is less appropriate (see nomenclature agreement).⁶⁹ In addition (see Fig. 1) chromaffin granules contain secretogranin II^{81,274} (originally discovered in the pituitary gland).²⁷⁶ This protein has also been called chromogranin C⁸¹ but a nomenclature proposal agreed on the name secretogranin II.⁶⁹ Furthermore,

another protein isolated from pituitary, 7B2,²²² is also present in chromaffin granules²²³ and so are numerous neuropeptides and their precursors. Higher concentrations are found for the enkephalins and neuropeptide Y.^{364,371} Finally the secretory proteins include dopamine β -hydroxylase, glycoprotein III,³⁶⁴ carboxypeptidase H^{93,193} and endoproteases of the kex2 type.^{44,290,183a} The recent proposal¹⁵⁴ to call the acidic proteins of secretory organelles "granins" would at present add a further, quite unnecessary name, since the original name chromogranins already covered the same proteins. Further nomenclature proposals should await a better understanding of the function of all these proteins.

In the present commentary we will concentrate on the two closely related proteins, CgA and CgB, and will not deal with the other secretory proteins. Several reviews on this topic have been published.^{57,82,132,143,261,306,315,355,362,364,365,371}

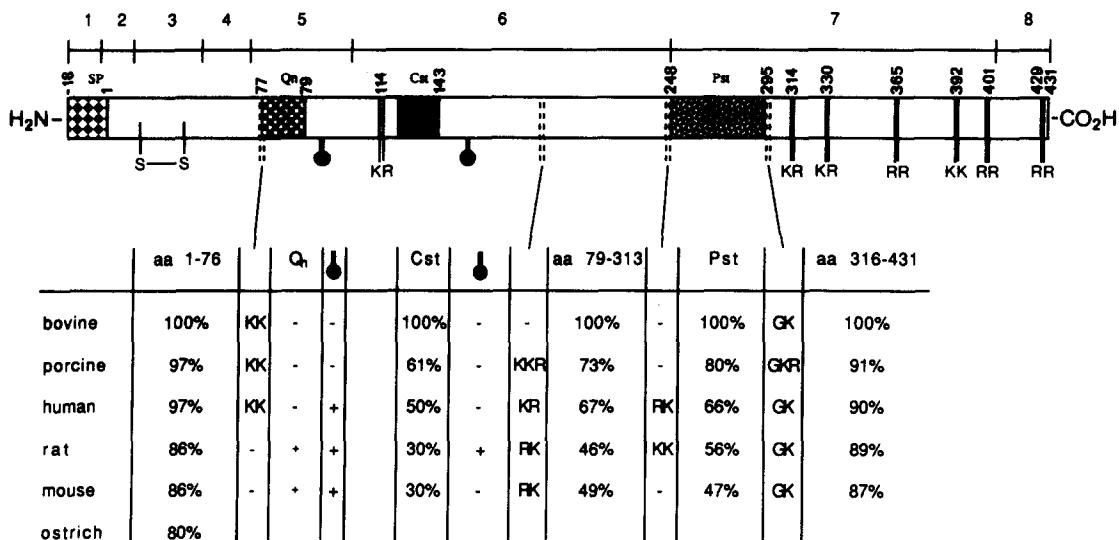


Fig. 2. CgA: amino acid sequences and gene structure. The top part of this figure presents the gene structure^{162,374} with the exons of the CgA gene numbered from 1 to 8. Below, a schematic consensus sequence is shown. The numbers on top indicate amino acid positions of bovine CgA.^{1,20,160} In addition, the sequences for rat,^{158,163,256} human,^{138,186} pig¹⁶¹ and mouse³⁷⁴ CgA and for a peptide from ostrich¹⁹⁹ are given. Certain peptides are indicated: SP, signal peptides; Q_n, polyglutamyl region; Cst, chromostatin; Pst, pancreastatin. Glycosylation sites are marked by black dots. Dibasic cleavage sites are marked by two lines for those conserved in the five species (the single-letter amino acid code is given for these sites). Cleavage sites not conserved are marked by double broken lines. The species comparison demonstrates which features of the sequence are conserved. A high degree of conservation is found in the N- (aa, amino acid 1-76) and C-terminals (aa, amino acid 316-431) whereas in the middle part of the molecule, where the defined peptides are located, the conservation is rather limited.

2. ISOLATION AND QUANTITATIVE DETERMINATION

For the isolation of chromogranins several methods have employed chromatography on molecular sieve, ionic exchange or butyl Sepharose columns.^{49,134,180,317,318,323} More recent purification schemes are based on high pressure liquid chromatography with molecular sieve, ionic exchange and reverse phase columns,^{86,123,296} chromatography on calmodulin columns,³⁷⁵ immunoaffinity chromatography³⁵⁹ and preparative one- or two-dimensional electrophoresis.^{251,252,274} The heat stability²⁷⁴ of these proteins is very useful for preventing proteolysis during isolation and also provides a very efficient step of purification since many other proteins can simply be removed by boiling tissue.²⁸⁶ Since clones for both CgA and CgB are available, pure chromogranins can now also be produced by recombinant techniques.¹⁴⁴

Radioimmunoassays^{70,232,247,253} and enzyme assays^{64,180} have been developed with antisera for CgA, but also for the CgA-derived polypeptide pancreastatin^{31,53,99,178,255,284,345} and the GAWK peptides derived from CgB.^{279,294} These assays are very sensitive (100 pg for radioimmunoassay, 1 pg for enzyme assay); however, it is not clear which peptide moieties are actually measured, since the antisera react both with intact CgA and its breakdown products irrespective of whether they were originally raised against CgA or peptides derived from it. With quantitative immunoblotting^{79,350} one can measure the relative concentrations of CgA or its larger breakdown

products but one misses smaller peptides (<10,000 mol. wt) formed from them. An ideal assay would depend on an antiserum against a synthetic peptide (flanked by basic pairs in the CgA sequence) used in a radioimmunoassay before and after digestion of all larger CgA molecules to smaller peptides (cf. enkephalin assays).³³⁷ For the chromogranins this approach has apparently only been used once.³⁴⁵

3. PHYSICO-CHEMICAL PROPERTIES

In the following we will only deal with some general features related to the primary amino acid sequence of these proteins. Further properties acquired by post-translational modification (e.g. glycosylation) and the calcium binding ability of CgA will be discussed below.

The primary amino acid sequences of these proteins have been determined for several species (see Figs 2 and 3). For bovine CgA the mRNA has a length of 2100 bases, for CgB 2600. This message codes for mature proteins of 431 amino acids (48,000 mol. wt) for bovine CgA and 626 amino acids (71,500 mol. wt) for CgB. As already recognized previously from amino acid analyses (see Ref. 362) common features of these proteins are the high glutamic acid (25%) and proline (about 10%) content yielding an acidic pI of about 4.5-5.0.^{8,362} The homology between CgA and CgB is rather limited, in agreement with the obvious lack of immunological cross-reaction;^{79,274}

4. ORGANIZATION OF THE GENES

A Southern blot analysis of genomic DNA digested with various restriction enzymes revealed that both CgA^{1,158,186,230,240,256} and CgB²⁶³ are present as single genes in the haploid genome. However, it has also been claimed that there are genomic differences in the amino acid sequences of CgA which may have arisen from a duplicated CgA gene.³⁴⁵ The gene for human CgA was assigned to chromosome 14q32^{230,240} and that of mouse CgB to chromosome 2.¹⁷¹ In the human genome a two-allele restriction fragment length polymorphism (RFLP) was reported for CgA. Nine out of 10 restriction enzymes did not detect a RFLP; however, Bgl II digests of genomic DNA from 56 Caucasians did identify two alleles of 24.0 and 8.0 kb with frequencies of 0.34 and 0.66, respectively, and a constant band of 6 kb.²³⁰

The bovine and mouse CgA gene comprise eight exons separated by seven introns (Fig. 2) which span approximately 13.6 kb¹⁶² and 11 kb.³⁷⁴ The mouse CgB gene consists of five exons and four introns²⁶³ and the coding region spans 12.1 kb (Fig. 3). A comparison of the mouse CgA gene with the mouse CgB gene reveals homologies in the structural organization of three exons. These three exons code for the N-terminal and C-terminal domains which share significant sequence homologies between both proteins. Exon 2 of mouse CgA and CgB encodes the last three amino acids of the signal peptide plus the first 13 amino acids of the N-terminus of mature CgA and CgB; exon 3 encodes for the next 33 amino acids containing the disulfide loop motif present in both CgA and CgB. Exon 8 of CgA and exon 5 of CgB code for the 27 and 25 C-terminal amino acids of both proteins. The variable middle domain of CgA and CgB that differs substantially between species and shares no significant homology between CgA and CgB is reflected also by a different genomic organization.

The region upstream of the Cg promoter contains one cyclic AMP response element in both CgA and CgB. With the exception of two Sph I sites for CgA and one Sph I site for CgB no further consensus matches for other known transcription factors were found.^{162,263,374} The elements responsible for tissue-specific expression in multiple neuroendocrine cells await identification; for CgA the region containing this information was shown to reside within 1.2 kb upstream of the coding region.³⁷⁴

5. DISTRIBUTION IN NEUROENDOCRINE AND IMMUNE SYSTEM

Originally CgA was isolated from the chromaffin granules^{134,317,318} of adrenal medulla. It was also found in the large dense-core vesicles of sympathetic nerve^{12,15,56} and in the brainstem rich in adrenergic neurons.¹⁴⁷ Based on these studies it was assumed that CgA was an "adrenergic" protein confined to

sympathetic neurons and to the adrenal medulla. In 1982 we discovered⁵⁰ that CgA and secretory protein I (also called parathyroid secretory protein, PSP, Ref. 323) independently characterized (see Ref. 47) in the parathyroid gland were apparently identical proteins. Lloyd and Wilson²¹⁶ had raised a monoclonal antibody against a protein present in pheochromocytoma tissue which reacted with a large number of endocrine tissues and their tumors and the responsible antigen was then identified as CgA.³⁵⁸ Within a short time several groups firmly established that CgA had a widespread distribution in endocrine tissues^{46,84,248,358} (for further references see Table 1). When CgB was characterized in adrenal medulla in 1984^{79,366} it was clear from the beginning that CgB shared this property with CgA^{84,274} (for further references see Table 1).

The occurrence of immunoreactive CgA and CgB, of peptides derived from them and of their mRNAs in most endocrine tissues is well established and we will only comment on a few points (for references for the presence of CgA and CgB in tumor tissue see Refs 210, 217, 348, and an extensive review, 355).

Chromogranin immunoreactivity as defined by immunohistochemistry exhibits great variations depending on the fixation method^{116,203,237,244,273,277} and dilution of antisera.¹¹⁸ This capricious behavior makes it rather difficult to draw definite conclusions on the presence or absence of chromogranins in various cell types and therefore explains discrepancies found in the literature. For Table 1 we have not included negative results when positive immunostaining was reported by other groups, since a positive result would appear more reliable. In any case, for many tissues independent methods (immunohistochemistry, immunoblotting, *in situ* hybridization and radioimmunoassay) provide us with concordant results.

Immunohistology does not reveal the molecular properties of the detected antigens, but for several tissues immunoblotting (see Table 1) has established that these endocrine tissues contain CgA and CgB of the same size as in the adrenal medulla but also a variable amount (see below) of breakdown products.^{84,122,159,210,244,277} Furthermore, for two tissues (pituitary and endocrine pancreas) the cDNA clone for CgA was found to be practically identical to the adrenal one.^{1,158}

It is interesting to consider the relative size of the CgA stores in neuroendocrine tissues.³²⁷ Thus in human beings the largest store of CgA is found in the adrenal medulla; the pituitary contains 25% of this amount (which appears surprisingly high), the pancreas and stomach plus intestine 5% each, whereas all other endocrine glands represent less than 1%. In addition to the classical endocrine tissues, CgA has also been found in scattered neuroendocrine cells in several organs like the breast, the lung or the prostate (see Table 1). Its presence in the storage granules of the myoendocrine cells of the heart storing the atrial

Table 1. Distribution of chromogranins and related peptides in neuroendocrine tissues

	Human	Bovine	Rat	Other
Anterior pituitary				
Unspecified cells	IB: A, ¹²² B ¹²² RIA: A, ³²⁷ GAWK ²⁵ IH: A, ^{314,216,249,282,358} B, ^{122,282} GAWK ^{25a} EM: A ¹³¹ NB: A ^{138,139} IS: A, ^{211,212} B ^{211,212} IH: A, ^{33,215} B, ³³ GAWK ²² EM: GAWK ²⁵ IH: A, ^{33,215} B ³³ EM: GAWK ²⁵ IH: A, ³³ B, ³³ GAWK ²² EM: GAWK ²²	IB: A, ^{14,84,180,202,210,274,319} B, ^{94,106} RIA: A, ^{14,180,245,249,251} IH: A, ^{195,248} B ¹⁹⁵ NB: A, ^{1,160} IS: A, ³⁰² EM: A, ¹⁶ B ¹⁶ IH: A, ²⁴⁹ EM: A, ¹⁶ B ¹⁶ IH: A, ²⁴⁹ EM: A, ¹⁶ B ¹⁶ RIA: A, ^{245,249} IS: A, ³⁰² RIA: A, ^{245,249} IB: A, ^{14,68,378} B ³⁷⁸ RIA: A, ^{245,249} IH: A, ^{68,249} Pst, ³⁴⁵ B ³⁷⁸ IH: A, ^{67,116,117,248,378} EM: A, ⁶⁷ IH: A, ^{67,116,378} B ²⁷³ EM: A, ⁶⁷ IH: A, ^{67,116,378} EM: A, ⁶⁷ IH: A, ^{67,116}	IB: A, ^{5,88,190,350} β G ¹⁵⁹ B ⁸⁸ RIA: Pst, ⁵³ GAWK ²⁷⁹ IH: A, ⁸⁴ Pst, ⁵³ β G, ¹⁵⁹ B ⁷⁴ NB: A, ^{87,88,163} B ^{87,92} IS: A, ⁸⁷ B ⁹² IH: A, ⁴⁶ IH: A, ^{46,88,344} EM: A, ³⁴⁴ IH: β G, ¹⁵⁹ B ⁸⁸ IH: β G ¹⁵⁹ IB: A, ³⁵⁰ RIA: Pst ⁵³ IH: Pst, ⁵³ B ⁸⁴ IB: β G ¹⁵⁹ IH: β G ¹⁵⁹ IH: A, ¹¹⁶ β G ¹⁵⁹ IH: A, ⁴⁶ IH: A, ⁴⁶ RIA: Pst ⁵³ IH: A, ²⁷⁰ β G, ¹⁵⁹ Pst ^{53,270}	IB: A, ^{319,277} B ²⁷⁷ RIA: Pst, ^{31,178,284} GAWK ²⁷⁹ IH: Pst ^{191,267,284} EM: Pst ¹⁹¹ IH: A, ^{273,277} Pst, ^{31,178,191} B ²⁷³ IH: A, ^{273,277} Pst, ^{31,178,191} B ^{273,277} IH: B ²⁷⁷ RIA: Pst ¹⁷⁸ IH: Pst ¹⁷⁸ IB: Pst ^{31,284} IH: A, ²⁷⁰ Pst ^{191,270,284} IH: A, ^{39,34,116,117,118} Pst, ^{31,267} B ³⁹ EM: A, ⁴¹ Pst ^{191,267} IH: A, ^{34,116,273} Pst, ³¹ B ^{34,273} EM: A, ³⁹ Pst ¹⁹¹ IH: A, ^{39,116,117,118} Pst, ^{31,267} B ¹¹⁸ EM: A, ³⁹ Pst ^{191,267} IH: A, ^{34,39,116,117,118,273} Pst, ³¹ B ¹¹⁸ EM: A, ^{39,41} IH: A, ¹⁴⁰ RIA: Pst ³¹ IH: A, ²⁷⁰ Pst ^{169,270}
TSH cells				
LH/FSH cells				
GH cells				
Multihormonal cells				
Intermediate pituitary				
Posterior pituitary				
Pancreas				
Unspecified cells	IB: A, ¹²² RIA: A, ^{259,327} Pst, ²⁸⁴ GAWK ^{25,284} IH: A, ^{314,216,270,358} Pst, ^{249,270} B ¹²² IS: A, ²¹² B ²¹² IH: A, ^{33,116,215,282} B ³³ EM: A, ¹³¹			
Insulin cells				
Glucagon cells				
Somatostatin cells				
PP cells				
Esophagus				
Stomach				
Unspecified cells	RIA: A, ³²⁷ Pst, ²⁸⁴ GAWK ²⁵ IH: A, ^{270,356,358} Pst, ²⁷⁰ B ³⁵⁶ GAWK ²⁵ EM: A, ¹³¹			

Enterochromaffin	IH: A, 33,42,72,214,282 B ^{25,282} GAWK ²⁵	IH: A, 42 B ⁸⁴	IH: A, 40,42 Pst ^{31,191} EM: A, 41 Pst ¹⁹¹
Somatostatin	IH: A ^{42,72,356}	IH: A ⁴⁶	IH: A, 40,42 Pst ¹⁹¹ EM: Pst ¹⁹¹
Substance P EC-like cell	IH: A ^{33,42,72,273}	IH: A ⁴⁶ IH: A ²⁷³	IH: A, 40,42,273 EM: A ⁴¹
Gastrin cell	IH: A, 33,72,273,282,356 B, 2533 GAWK ²⁵	IH: A, 179,273 B ²⁷³	IH: A, 34,40,117,273 Pst, 31,191 B ^{34,40,117,273}
Glucagon cell PP cell	EM: A ³³⁹ EM: A ³³⁹	IH: A ¹⁷⁹	EM: A, 41 Pst, 191 B ⁴¹
Small intestine Unspecified cells	RIA: A, 326 GAWK ²⁵ IH: A, 42,249,262,282,356,358 B, 282,356	IB: B ²⁷⁴ RIA: Pst ⁵³ IH: B, 274 β G ¹⁵⁹ NB: B ⁹² IH: A, 273 B ²⁷³	RIA: Pst ^{31,284} IH: A, 42,300 Pst ^{267,284,300}
Enterochromaffin	EM: A ¹³¹ IH: A, 33,72 B ³³	IH: A, 195 B ^{195,273}	IH: A, 34,40,273 Pst, 31,191 B ^{34,273}
Gastrin cell	IH: A, 33,72,356 B ³³	IH: A ¹⁷⁹	IH: Pst ^{31,191}
Peptide YY cells	IH: A ^{33,42}	IH: A ⁴²	IH: A ^{40,42}
Motilin cells	IH: A ⁷²	IH: A ¹⁷⁹	IH: A, 42 Pst ³¹
Neurotensin cells	IH: A ^{33,42,72}	IH: A ⁴²	IH: A, 40,42 Pst ³¹
Secretin cell	EM: A ³³⁹ IH: A, 33,42,72,338 B, 25 GAWK ²⁵	IH: A ⁴²	IH: A, 34,40,42 Pst ^{31,191}
Substance P	EM: A ³³⁸	IH: A ¹⁷⁹	IH: A, 34,40,42 Pst ^{31,191}
Somatostatin cells	IH: A ^{72,356}	IH: A ¹⁷⁹	IH: Pst ³¹
CCK cell	IH: A ⁷²	IH: A ⁴²	IH: A, 40,42 Pst ³¹
Enteroglucagon	IH: A ^{39,72}	IH: A ⁴²	IH: Pst ^{31,191}
GIP cell	IH: A ^{42,72}	IH: A ⁴²	IH: A, 40,42 Pst ³¹
PP cell	IH: A ²⁶⁵	IH: A ²⁶⁵	IH: B ³⁴
Cells in lamina propria			
Large bowel			
Unspecified cells	RIA: GAWK ²⁵ IH: A ^{399,270}	RIA: Pst ⁵³ IH: β G ¹⁵⁹	RIA: Pst ^{31,284} IH: A, 270 Pst ²⁷⁰
Enterochromaffin	EM: A, 72 B, 25 GAWK ²⁵ IH: A ⁷²	IH: A, 34,40 Pst, 31 B ³⁴	IH: A, 34,40 Pst, 31 B ³⁴
Enteroglucagon	IH: A ⁷²		IH: Pst ³¹
Peptide YY	IH: A ⁷²		IH: Pst ³¹
Somatostatin	IH: A ¹²²		IB: A ¹⁰⁷
Parathyroid	RIA: A ³²⁶ IH: A, 33,216,249,273,282,358 B ³³	IB: A ^{195,202} RIA: A ¹⁸⁰ IH: A, 195,244,248,249,273 B ⁸⁴	IH: A ²⁷³
	EM: A ¹³¹ IS: A, 211 B ²¹¹	EM: A ⁹ NB: A ^{1,160} IS: A ³⁰²	

Table 1—continued

	Human	Bovine	Rat	Other
Thyroid C-cells	RIA: A ³²⁶ IH: A ^{197,216,249,273,382,358} B ^{33,282}	IB: A ²⁴⁴ RIA: A ²⁴⁹ IH: A ^{195,244,248,249} B ¹⁹⁵	RIA: Pst ⁵³ IH: Pst ⁵³	IB: A ³⁴⁷ B ³⁴⁷ RIA: Pst ³¹ IH: A ²⁷³ Pst ^{31,169} IH: A ^{104,127} EM: A ^{128,127,125} Pst ^{126,125,169}
Merkel cells	IH: A ¹²⁷ EM: Pst ¹²⁶			
Neuroendocrine cells of Breast	IH: A ³⁷		IB: A ³²¹ B ³²¹ IH: A ³²¹ B ³²¹	
Heart (ANF cells)				
Lung, trachea	IH: A ^{197,358}			IH: A ¹⁹⁷ IH: A ^{124,340} EM: A ⁵⁵ IH: A ³⁴¹
Urether, bladder				
Prostate				
Uterus				
Lymphoreticular tissue		IB: A ²⁴⁴ IH: A ²⁴⁴	IH: A ¹⁴⁵	
Central nervous system				
Neurons	RIA: A ³²⁶ GAWK ²⁷⁹ IH: A ^{236,235,237} B ²⁸²	IB: A ^{14,180,319} RIA: A ^{180,249,251} IH: A ²⁴⁴ NB: A ¹⁶⁰ IS: A ³⁰²	IB: A ^{190,350} B ^{274,350} RIA: Pst ⁵³ GAWK ²⁷⁹ IH: A ¹⁹⁰ B ²⁷⁴ NB: A ^{92,219} B ^{92,219} IS: A ²¹⁹ B ^{92,219}	IB: A ³¹⁹ RIA: Pst ^{178,284} GAWK ²⁷⁹ IH: A ³¹⁹ Pst ^{169,178}
Glia	IH: A ²²⁵ IH: B ²⁸²	IH: A ²⁴⁴	IH: A ²⁹	IH: A ³¹⁹ Pst ¹⁷⁸ RIA: Pst ¹⁷⁸
Spinal cord				IH: A ^{141,176} Pst ¹⁶⁹
Ganglia and peripheral nerves	RIA: A ³²⁶ IH: A ¹⁴¹	IB: A ^{84,121,242,289} B ^{84,121} RIA: A ^{251,249} IH: A ¹⁹⁵ B ¹⁹⁵ IB: A ²⁴⁴ IH: A ²⁴⁴	IH: A ^{29,46,141,342} B ⁸⁴	
Retina	IH: A ²⁴⁴			IH: Pst ¹⁶⁹
Liquor	RIA: A ³⁰¹ GAWK ²⁷⁹	RIA: A ^{180,249,251}		
Serum	RIA: A ^{52,64,151,172,325,326,327,246,247} GAWK ²⁹⁴			RIA: Pst ³
Urine	RIA: A ¹⁵¹ Pst ^{100,329,330}			

The table gives the distribution for the three species as defined, whereas all other species are included in the "other" column. A, chromogranin A; B, chromogranin B; Pst, pancreaticatin; β G, beta-granin; IB, immunoblot; RIA, radioimmunoassay; IH, immunohistochemistry; EM, immunoelectron microscopy; NB, Northern blot; IS, *in situ* hybridization histochemistry; TSH, thyroid stimulating hormone; LH, luteinizing hormone; FSH, follicle stimulating hormone; GH, growth hormone; PP, pancreatic polypeptide; EC, enteropancreatic chromaffin; CCK, cholecystokinin; GIP, gastric inhibitory peptide; ANF, atrial natriuretic factor.

natriuretic factor³²¹ raises the question whether it is also found in the renin granules of the kidney. One might also ask whether the erythropoietin cells of the same organ contain CgA. Thus CgA presents a general neuroendocrine marker and that has made it useful for a histopathological classification of neuroendocrine tumors.^{244,355} However, recent publications for the first time also reported an immunostaining for CgA in exocrine cells, i.e. alveolar type II cells of rat lung¹⁷³ and for CgA and CgB in duct cells of the submandibular gland whose nature was, however, not well defined.^{204,205,244}

As far as the immune system is concerned, one study reported the presence of scarcely distributed CgA-positive cells in spleen, lymph node, thymus and fetal liver.¹⁴⁵ By Western blotting small amounts of immunoreactive CgA were also found in these organs. Unfortunately this interesting observation has not been followed up. Are these special cells natural killer cells which are known to store and secrete enzymes and peptides by exocytosis?³²⁴

The relative concentration of CgA and CgB varies from tissue to tissue; however, in general, both proteins are found to be co-stored, but there is also evidence for a dissociation of their storage. Examples are the high concentration of CgA in the parathyroid gland (see Table 1) but the presence of only small amounts of CgB^{210,212} if any^{33,122,195,273} or the presence of CgA in ovine and bovine intermediate pituitary but the absence of CgB^{195,277} or the differential staining of some enterochromaffin cells for A, but not for B in the stomach,^{33,282} however, a staining of at least some of these cells for the GAWK protein derived from CgB has also been reported.²⁵ Analogous results, i.e. a differential localization of CgA and CgB, have been obtained for brain (see below).

6. PRESENCE IN THE NERVOUS SYSTEM

The presence of CgA in the central nervous system was already established in early studies.^{147,244,251,319} A detailed immunohistochemical mapping of ovine brain and spinal cord³¹⁹ revealed a widespread distribution of the antigen. The immunoreactivity partly overlapped with those of several neuropeptides and other markers like tyrosine hydroxylase and dopamine β -hydroxylase. However, the conclusion reached was that CgA is present in neurons using many different transmitters. This protein, originally thought to be "adrenergic" is also found in cholinergic neurons like the motoneurons of the spinal cord.³¹⁹ In agreement, CgA was also found in cholinergic nerve terminals in the muscles of the diaphragm.³⁴² For human brain, Munoz and collaborators demonstrated that if methodical difficulties are overcome, a widespread distribution of CgA can be found.²³⁵⁻²³⁷ Interestingly, a high proportion of senile plaques in Alzheimer's disease,^{32,196,233,234,349} and also of Pick's bodies in another brain disease leading

to dementia, consistently immunostain for CgA.³⁴⁹ Furthermore, the CgA/synaptophysin ratio in such diseased brains is high.^{196,349}

In porcine brain, CgA and a peptide derived from it, i.e. pancreastatin, have an identical, widespread distribution.¹⁷⁸ CgA immunoreactivity has also been found in bovine²⁴⁴ and porcine¹⁶⁹ retina. Quantitative analyses of CgA by radioimmunoassay in bovine brain^{180,251} and by immunoblotting in rat brain³⁵⁰ revealed that the concentration of CgA was only about 0.01–0.2% of that in the respective adrenal medulla. The claim¹⁹⁰ that the rat olfactory bulb as measured by immunoblots contains CgA in amounts comparable to that of adrenal medulla and the pituitary is likely to be wrong since at least in ox the pituitary gland has only 3% of the CgA concentration found in adrenal medulla.^{180,251}

Bovine and rat brain contain CgA mRNA which has been measured by Northern blotting.^{160,190,219,350} With tissue hybridization CgA mRNA was found in the caudate nucleus of bovine brain.³⁰² A detailed mapping analysis of rat brain²¹⁹ revealed the highest concentration in the pyramidal cell layer of the hippocampus, in the subiculum and the septum, with lower concentrations in many other brain areas, in agreement with the immunohistochemical data. For CgB a first comprehensive study measured the distribution of GAWK, a peptide derived from CgB, in the brain of several species, by radioimmunoassay.²⁷⁹ In human brain the highest concentrations were found in the hypothalamus, hippocampus, amygdala, nucleus accumbens, the basal ganglia and the cerebellum; much lower concentrations were present in rat and guinea-pig brain. Immunoblotting measuring the proprotein CgB in rat brain revealed the highest concentration in the rat cerebellum, followed by the hypothalamus and amygdala.³⁵⁰ Only very limited immunohistochemical data are available for human brain with a positive staining found in the Purkinje cells of the cerebellum²⁸² and for rat brain.²⁷⁴ Two studies^{92,219} used *in situ* hybridization to study the distribution of CgB mRNA in rat brain. In agreement with the immunochemical data, a wide distribution was reported with high concentrations in parts of the cortex, pyramidal cell layer of the hippocampus, in the dorsal raphe, pontine nucleus and several other regions. A comparison of the distribution of the mRNA of CgA and CgB²¹⁹ revealed that some nuclei apparently synthesized only CgA but not CgB, e.g. the subiculum, whereas the dorsal raphe and the granule cell layer of the cerebellum were only positive for the CgB-message. Immunoblotting indicated that CgA and CgB found in brain are identical to the adrenal proteins in size and migration behavior (ovine CgA;³¹⁹ rat CgA and CgB).^{274,350} However, a relatively higher concentration of the proteoglycan form of CgA was present when compared with adrenal medulla and especially with endocrine tissue like the anterior pituitary.³⁵⁰

7. SUBCELLULAR LOCALIZATION

Two methods are particularly useful to establish the subcellular localization of a component, i.e. subcellular fractionation and immunohistochemistry at the ultrastructural level. With the first method the specific localization of CgA in bovine^{79,251,278} and human²⁵² and of CgB in bovine⁷⁹ chromaffin granules has been established. The same approach was used for localizing CgA in the storage vesicles of anterior pituitary²⁴⁵ and of betagranin, a peptide derived from CgA, in the insulin granules of pancreas¹⁵⁷ and of CgA^{56,242,289} and CgB in the large dense-core vesicles of splenic nerve.^{121,289} One of these studies²⁴² also revealed that small dense-core vesicles of adrenergic nerves were devoid of CgA. What is the concentration of CgA in these vesicles relative to other peptide constituents? In bovine chromaffin granules the CgA family represents the major component, although the values obtained rather differ (70% as calculated from densitometric scans and the degree of degradation⁸² vs 46% as determined by radioimmunoassay).¹⁴ We have calculated that in a single bovine chromaffin granule there are about 5000 molecules of the proprotein CgA present.³⁷¹ A quite similar value (2763 molecules) was found by quantitative immunoelectron microscopy.²²⁴ In the anterior pituitary granules, CgA is only a minor (4.5%) component¹⁴ and this is also true for the endocrine pancreas where the insulin/CgA ratio is about 2000.⁶⁸ Even considering that in this tissue, most of the CgA is processed (see below) this would make CgA a minor component in insulin granules (taking 90% breakdown into account: only 3 mg total CgA/100 mg insulin). The observation²⁵¹ that after subcellular fractionation of the bovine brain the cytosol contained the majority (69%) of CgA apparently suggested that in this tissue CgA is not localized within the secretory compartment. However, immunoelectron-microscopy³¹⁹ revealed that CgA reactivity is confined to Golgi stacks and large dense-core vesicles, but absent from translucent synaptic vesicles and from the cytosol. Most likely during homogenization of brain a considerable part of the Golgi stacks and large dense-core vesicles release their content including CgA.

In 1974 we published the first data on the immunoelectron microscopic localization of CgA in adrenal medulla.³⁷⁰ Immunoreactivity was confined to the content of the vesicles; there was no staining of their membranes. Several studies^{10,41,191,224,320} have confirmed this finding and extended it to CgB^{41,320} and to other endocrine tissues (Table 1). By double immunolabeling or labeling of serial sections not only the presence of CgA and CgB in vesicles but also their co-localization with transmitters or neuropeptides within the same vesicles was established. Combinations of co-localization were: CgA/CgB/secretogranin II/neuropeptide Y,³²⁰ CgA/parathyroid hormone,⁹ CgA/insulin,⁶⁷ CgA/glucagon,⁶⁷ CgA/

atrial natriuretic factor,³²¹ CgA/calcitonin,³⁰⁴ thyroid stimulating hormone/CgA/CgB/secretogranin II,¹⁶ and luteinizing hormone/CgA/CgB/secretogranin II.^{16,344}

Within the same cell the respective hormones and the chromogranins are not always stored together in the same vesicles. This was first demonstrated for secretogranin II, prolactin and growth hormone for bovine somatomammotroph cells¹²⁹ and has now been extended to CgA and CgB.¹⁶ In the somatomammotroph cells, two types of granules are present, i.e. small (150–300 nm) and larger (400–550 nm) ones. Only the small ones apparently contained CgA, CgB (and also secretogranin II, thyroid stimulating hormone and luteinizing hormone), whereas only the large ones contained growth hormone and prolactin. Apparently, in this special type of cell with multi-hormone storage, quite distinct sorting takes place. An analogous result was obtained for gonadotrophs of the anterior pituitary, in which large-sized granules reacted for follicle stimulating hormone, luteinizing hormone and CgA, whereas smaller ones were only positive for luteinizing hormone and secretogranin II.³⁴⁴ It will be interesting to see whether detailed studies on other cells will reveal further subpopulations of granules exhibiting a differential storage of the various hormones and chromogranins.

Immunoelectron microscopy has also revealed some uneven labeling for CgA within the vesicles. This was particularly marked for the glucagon-storing vesicles of endocrine pancreas where CgA or pancreastatin labeling was preferentially localized on a less electron-dense halo surrounding a dense core^{67,131,191,339} whereas for insulin granules a peripheral immunostaining was reported only by one group⁶⁷ but not by others.^{131,191} A labeling of the peripheral part of the vesicles has also been reported for Merkel cells¹²⁷ and for secretin-storing cells of the intestine,³³⁸ and for CgB in small granules of the somatomammotroph cells,¹⁶ and in the granules of gastrin cells⁴¹ but apparently not for all the other secretory granules where CgA and CgB have been localized by immunostaining (see Table 1).

8. PHYLOGENY

The initial studies on CgA were performed with bovine tissues.^{139,317,318} However, it was soon recognized that analogous components to CgA as shown by electrophoresis or by immunoreaction were present in horse, pig and sheep adrenals^{133,264,372} and in human adrenals and pheochromocytoma.³²² CgA was then isolated and characterized from human^{252,359} and also rat material.^{86,274} In immunohistochemical studies a cross-reaction with antigens in several species including pig, cat, guinea-pig, dog and rabbit was observed.^{42,118,273} For CgB in the first study⁷⁹ the occurrence of similar (electrophoresis, immunoreaction) components in several species (pig, horse,

cat, guinea-pig and rat) was recognized and this was soon extended to human material.^{106,122}

The cloning of CgA and CgB from several species provides now the molecular basis for these immunological results. In some parts of the molecules (especially the N- and C-terminals) there is a high degree of homology between the individual mammalian species in agreement with the immunological cross-reaction (see Fig. 2). Unfortunately, only clones from mammalian species have been analysed. However, it is already clear that analogous proteins are present much earlier in phylogeny. Rieker *et al.*²⁷² found immunoreactive components comparable in size to CgA and CgB in chicken, frog, fish and lobster. In chicken ultimobranchial gland,¹⁷⁵ in toad urinary bladder⁵⁵ and in several fish species, CgA immunoreactivity (with antimammalian CgA) could also be demonstrated by immunohistology and radio-immunoassay.^{58,333} In the ostrich pituitary a large peptide (see Fig. 2) with an 80% homology to the N-terminal of bovine CgA has been described¹⁹⁹ and in *Drosophila* heads an immunoreactive band for CgA could be detected by immunoblotting (Kapelari and Winkler, unpublished observation). A detailed immunohistochemical study on CgA and pancreastatin in the gastro-entero-pancreatic system revealed positive reaction apart from mammalian species in birds, reptiles and fish but not in deuterostomian invertebrates.²⁷⁰

A molecular cloning of chromogranins from non-mammalian species would seem extremely useful for defining regions with a high degree of conservation. Peptides in these regions would be good candidates for a relevant function which has evolved during phylogeny. Finally, even in the protozoan, *Paramecium tetraurelia*, peptides immunologically cross-reacting with antisera against CgA and synthetic peptides derived from it have been found.²⁶⁰ However, in contrast to all other species only small peptides of 20,000 mol. wt were present. On the other hand, the mRNA hybridizing with a CgA probe suggests a much larger molecular weight at least for an unprocessed proprotein. Since the CgA antisera immunostained components which represented the major soluble proteins of the trichocysts of *Paramecium* an artefact is very difficult to exclude. Therefore these data should be confirmed by an independent approach, e.g. sequencing of some of these peptides.

As compiled by Fischer-Colbrie *et al.*,⁸² the relative amounts of CgA vs CgB in the different species vary. In ox adrenals, CgA is by far the dominating component, whereas in rat and human material CgB is prominent.^{86,122} Thus the CgA/CgB ratio in ox is 9, whereas in human material it is about 0.8.¹⁰⁶

9. ONTOGENY

Several immunohistochemical studies investigated the ontogenic development of CgA. In rat fetuses,

CgA immunoreactivity appeared at day 18 of the gestation period and at 15 days of postnatal age was similar to the adult staining pattern.¹⁸¹ In human fetuses a positive reaction for CgA²³¹ and CgA-mRNA¹³⁸ was found in adrenal primordia at six to eight weeks of the gestational period, slightly preceded by immunostaining for tyrosine hydroxylase. At nine weeks, positive staining was confined to "large cells" considered to be progenitors of chromaffin cells.²³¹ In the endocrine cells of the human intestinal tract, CgA immunoreactivity appeared together with peptides typical to these cells, e.g. somatostatin, at eight weeks of gestation⁷¹ and also in the ultimobranchial gland of the chicken¹⁷⁵ and in the carotid body¹⁷⁶ there was a concomitant appearance with other peptides and with tyrosine hydroxylase. For CgB the appearance of the mRNA has been studied in developing rat brain.⁹² It was first detected at embryonic day 14 and then increases till three weeks postnatally, when it starts to drop again to a level close to that found at birth.

A detailed study by immunoblotting of the development of CgA and CgB in rat adrenal medulla revealed two additional interesting features.²⁸⁷ Both chromogranins were detected on the 17th prenatal day. Their levels increased together with that of the catecholamines. However, the relative concentration of CgA vs CgB showed a significant change. At the end of the gestational period the ratio of CgA to CgB increases strongly at exactly the same time when the adrenaline/noradrenaline ratio becomes elevated. This latter effect is known to be due to the induction of methylation caused by the increase in cortisone production at this gestational time point.⁴⁵ For CgA we have shown^{83,303} that its biosynthesis is increased by corticosteroids. The observation of an increase of CgA biosynthesis together with that of adrenaline establishes the ontogenic significance of this concept. A second finding, described by Schober *et al.*,²⁸⁷ concerns the proteoglycan form of CgA. Whereas fetal adrenal contains a relatively high concentration of this CgA form, in adult animals it is much less prominent. Since rat brain also contains a prominent proteoglycan form of CgA³⁵⁰ this result was interpreted as a sign of the developmental relationship between early chromaffin cells and neurons.

10. SECRETION, PRESENCE IN SERUM AND CEREBROSPINAL FLUID AND FINAL FATE

Secretion of catecholamines from the adrenal medulla and chromaffin cell culture is accompanied by the release of all soluble constituents of chromaffin granules^{13,363} including CgA^{14,70,185,285,308,346} and CgB.⁷⁹ This is also true for noradrenaline release from the sympathetic nerve.³¹⁶ From the parathyroid gland, CgA is released with the parathormone⁴⁸ and beta-granin, derived from CgA, is co-released with insulin from endocrine pancreas¹⁵⁷ and so is another CgA-derived peptide, pancreastatin.^{3,255} This latter peptide

is also secreted from a human pancreastatin-producing cell line.⁹⁹ GAWK, a peptide derived from CgB, is released when rat anterior pituitary or bovine chromaffin cells are depolarized.^{164,279} CgB is also released from PC12 cells²⁷⁴ and a medullary thyroid carcinoma cell.¹⁶⁴

Since secretion from all endocrine tissues and neurons occurs by exocytosis, it is obvious that in all these tissues chromogranins must be co-secreted with the respective hormone. After secretion from endocrine organs, CgA reaches the bloodstream via the capillary or the lymph. For the adrenal medulla it has been shown that, in contrast to the catecholamines, most of CgA is first directed into the lymph vessels.³⁸ In human serum, O'Connor and collaborators found by radioimmunoassay about 30–50 ng/ml CgA in normal controls,^{250,253} whereas in patients with endocrine tumors these levels were significantly elevated.²⁵⁰ Differential stimulation of various endocrine organs revealed that only stimulation of the adrenal gland led to measurable increases in plasma CgA levels.³²⁷ On the other hand, in adrenalectomized patients, CgA levels in serum are in the normal range.³²⁵ Furthermore, blockade of the sympathetic system by a ganglion blocker and non-selective blockade of neuroendocrine secretion by somatostatin led to a decline of the basal levels by 25 and 48%, respectively.³²⁵ Thus basal CgA levels in plasma are mainly derived from these sources and not from the adrenal medulla (see also Ref. 52). The half-life of CgA in plasma is about 18 min.²⁴⁷ Plasma levels of pancreastatin-like immunoreactive material, mainly representing high molecular weight peptides with only small amounts of free pancreastatin have also been determined.^{3,31}

For GAWK (20–38), a peptide derived from CgB, a level of 150 pmol/l was found for human plasma.²⁹⁴ Also, these levels are elevated in patients with neuroendocrine tumors. This concentration corresponds to 10 ng/ml of the proprotein CgB which is in the same range as the values for CgA. This seems reasonable considering that equal amounts of CgA and CgB are present at least in human adrenal medulla.

Human cerebrospinal fluid contains 170 ng/ml CgA³⁰¹ which is higher than the serum values (40 ng). This finding indicates either significant release of CgA from brain or a longer half-life of CgA in the cerebrospinal fluid. For GAWK (20–38) a value of 1.1 pmol/ml has been reported (85 ng CgB/ml).²⁷⁹

The studies discussed above were based on radioimmunoassays which do not define the molecular size of the measured molecules. However, in a recent study with immunoblotting the presence of the unprocessed proprotein CgA could be demonstrated in bovine serum.³⁵² Apparently, neuroendocrine tissues release unprocessed CgA which is not quickly degraded in serum. O'Connor and collaborators^{151,253,379} reported that patients with renal failure had high CgA levels in serum. This indicated that the kidney might be essential for the removal of plasma CgA.

Kidney tissue contains CgA, but no CgA-mRNA.³⁵² Immunoelectron microscopy revealed that CgA immunoreactivity is found in proximal tubule cells. Within the cells it was present in small vesicles with or in close proximity to the brush border and closer to the nucleus in typical lysosomal structures. These results make it likely that CgA reaches kidney tubule cells by glomerular filtration and is taken up into the endocytotic lysosomal pathway followed by degradation. On the other hand, smaller peptides derived from CgA may reach the urine. At least, pancreastatin-like immunoreactivity has been detected in it.³³⁰

11. CELL-FREE SYNTHESIS

After *in vitro* translation of adrenal medullary mRNA, two translation products of very similar molecular weight (apparent molecular weight in SDS electrophoresis of about 70,000) can be immunoprecipitated with antisera against bovine CgA.^{74,183} In analogous findings, four products of very similar size were reported for parathyroid mRNA.^{220,221} When the *in vitro* translation for adrenal medulla was performed in the presence of dog pancreas microsomes only one immunoprecipitable translation product is found.⁷⁴ The two or more spots seen after *in vitro* translation could be due to different lengths of the signal peptides, but they may represent *in vitro* artefacts, since there is no evidence that signal peptides of different length are present in the bovine CgA sequence.^{20,160} In any case, in agreement with the fact that there is apparently only one gene for CgA, only one *in vitro* translated product is found after the removal of the signal peptide. The claim²⁹⁵ that several translation products are produced in such a system is therefore disproven. For CgB, cell-free systems yielded one translation product in bovine adrenal medulla.⁷⁴ For rat adrenal (PC12 cells) two products of very similar size are apparently produced.²⁷⁴

12. CELLULAR SYNTHESIS

12.1. Rate of synthesis

Early studies on bovine adrenal medulla had established that after labeling with tritiated amino acids the chromogranins were the major labeled products. Only 30 min after a radioactive pulse could the secretion of labeled CgA be induced by stimulation of the adrenal gland.^{200,367,369} Apparently, the synthesis rate of the secretory protein CgA was much higher than that of the membrane proteins which led to the suggestion that the membranes of chromaffin granules are re-used for several secretory cycles from the Golgi region to the plasma membrane.^{361,369} These data were recently extended in a careful study³⁶⁰ comparing the synthesis rates of CgA, enkephalins and dopamine β -hydroxylase in bovine chromaffin cell cultures. Whereas CgA and enkephalin had about

the same rate of synthesis, that of dopamine β -hydroxylase was lower by a factor of 10. The most likely explanation is that this enzyme which is partly membrane bound³⁶⁸ can be replenished at a lower rate since the membranes are re-used.

Further studies investigated the rate of amino acid incorporation in the presence of various secretagogues. In one series of experiments a direct correlation of this synthesis to the rate of secretion, as induced by several secretagogues, was found.³⁰⁷ In contrast, Wilson *et al.*³⁶⁰ did not discover any changes in CgA synthesis under very similar conditions. There was also no increase in total (cells and medium) CgA levels after stimulation of chromaffin cells in culture by nicotine treatment.⁷⁰ Apparently, there are some discrepancies and we will discuss this topic further in the section on the regulation of biosynthesis.

12.2. Post-translational modifications

Pulse-chase labeling of chromaffin cells of bovine adrenal medulla followed by two-dimensional electrophoresis revealed that the early labeled CgA and CgB spots behaved identically to the *in vitro* translation products formed in the presence of microsomes.⁷⁴ However, at later intervals significant changes in size and pI occurred which were especially marked for CgB. These modifications were obviously due to the post-translational processes which we will now discuss.

12.2.1. *Glycosylation.* Bovine CgA of adrenal medulla is a glycoprotein (5.4% carbohydrate) with galactose, *N*-acetylgalactosamine and sialic acid as major components.^{85,105,317} The sugars are mainly present as O-glycosidically linked tri- and tetrasaccharides composed of *N*-acetylgalactosamine, galactose and sialic acid.¹⁸² The primary amino acid sequence does not contain potential sites for N-glycosylation in this species; such sites are present in human, rat and mouse sequences (see Fig. 2), but it is not known whether N-glycosidically linked sugars are actually added in these species. For CgA of the parathyroid gland a very similar sugar composition has been reported;⁵⁰ however, with one exception, i.e. a much lower content of galactosamine. It seems unlikely that this low value is correct since the tri- and tetrasaccharides require about equal amounts of galactose and galactosamine. It would therefore seem premature to suggest that CgA outside the adrenal medulla is differently glycosylated, but further studies should settle this question.

The sugar composition of rat chromogranins has been investigated but a separation of CgA and CgB was not performed at that time.²⁸⁰ Bovine CgB has a sugar composition similar to that of CgA but with more fucose and mannose (see Ref. 85; CgB in this early paper was called A1). In contrast to CgA it reacts with *Pisum sativum* lectin, but like CgA it binds peanut agglutinin after neuraminidase treatment.⁷ Sites for N-glycosylation are present in the amino acid sequences of ovine, mouse and human CgB. The

presence of fucose and mannose in bovine CgB may indicate that this molecule is actually N-glycosylated. In other tissues the glycosylation of CgB has apparently not been investigated.

12.2.2. *Phosphorylation.* Settleman *et al.*²⁹⁶ have established that adrenal CgA is a phosphoprotein containing five phosphoserine residues per molecule. In agreement, labeling of chromaffin granules with [³²P]phosphate leads to incorporation of this isotope into CgA^{51,274} mainly into serine and to a small extent into threonine.²⁷⁴ An analogous finding has been reported for CgB.^{51,274} For other tissues it is only established that CgA of parathyroid gland is also phosphorylated.²⁴

12.2.3. *Sulfation.* Early studies have shown that [³⁵S]sulfate labeling of the adrenal medulla leads to the incorporation of this isotope into proteoglycans (see Ref. 182 for earlier references) and into chromogranins.¹⁸ Detailed recent studies have elucidated these phenomena in more detail. Lee and Huttner²⁰¹ demonstrated in PC12 cells the sulfation of secretory proteins which were later identified as CgB.^{73,274} Most of the sulfate was found to be linked to tyrosine, but some was also present in O-linked oligosaccharides.²⁷⁴ Since such sulfation mechanisms are widespread post-translational modifications¹⁵² there is no reason to assume that CgB in other tissues is not tyrosine sulfated. Adrenal CgA also incorporates [³⁵S]sulfate during biosynthesis.^{18,73,274} Most if not all of this sulfate appears to be bound to carbohydrates.²⁷⁴ For CgA of porcine parathyroid gland it was originally reported that it contained sulfate linked to tyrosine;¹⁸⁹ however, in a more recent study both bovine and porcine parathyroid CgA were found to be sulfated on oligosaccharides.¹⁰⁹

Chromaffin granules also contain proteoglycans in which the glycosaminoglycan component consists of dermatan sulfate, chondroitin 4- and 6-sulfate and heparan sulfate.^{11,26,105,182} When bovine chromaffin cells are labeled with [³⁵S]sulfate and analysed by two-dimensional electrophoresis a spot moving slower and to a more acidic pH than CgA (see Fig. 1) becomes labeled.^{73,274} This spot was identified as a proteoglycan form of CgA by its sensibility to chondroitinase treatment⁷³ and by its immunostaining for CgA.^{183,274} This proteoglycan form of CgA represents only a minor component of the total proteoglycans in bovine chromaffin granules and only about 1–2% of CgA is present in this form.¹¹² Proteoglycan CgA contains probably only a single chondroitin sulfate or dermatan sulfate chain of approximately 15,000 mol. wt.¹¹² In rat PC12 chromaffin cells CgA represents a significant portion of the larger proteoglycans. There is no evidence that bovine CgB is present in a proteoglycan form.¹¹² In the parathyroid gland a proteoglycan form of CgA was only found in bovine, but not in porcine tissue.¹⁰⁹ On the other hand, in rat tissues the relative amounts of proteoglycan CgA varied from tissue to tissue, being low or absent in anterior pituitary, with

intermediate amounts in the adrenal and a higher concentration in brain.³⁵⁰ During ontogeny of the rat adrenal medulla the relative amount of the proteoglycan form of CgA starts from a high level and then declines.²⁸⁷

12.2.4. *Pyroglutamylation*. This post-translational modification of peptides leads to the formation of pyroglutamyl residues from N-terminal glutamic acid³⁶ and the responsible enzyme, glutamic cyclase, is also present in adrenal medulla. A first analysis of bovine chromaffin granules for the presence of pyroglutamyl peptides revealed a 14-amino acid peptide derived from CgB.⁹⁰ It will be interesting to see whether this post-translational modification is an essential prerequisite for a possible function of this peptide.

12.2.5. *Carboxymethylation*. CgA is a carboxymethylated protein.²⁴³ Since carboxymethylase activity in adrenal medulla is cytosolic,⁶³ the modification of the CgA molecules as suggested^{102,243} would have to occur co-translationally which, however, is difficult to imagine.

12.2.6. *α -Amidation*. Pancreastatin, a peptide derived from CgA has been shown to possess a C-terminal glycine amide structure in pig,³³¹ bovine²⁴¹ and human^{100,284,293} tissues.

12.2.7. *Formation of disulfide bonds*. Based on reduction studies it has been proposed²¹ that the two N-terminal cysteines are present *in vivo* in a disulfide-bonded loop structure. This probably requires the action of a disulfide isomerase.

13. PROTEOLYTIC PROCESSING

In 1974 we observed¹⁴⁹ that in bovine chromaffin granules there were several smaller proteins immunologically cross-reacting with CgA.^{183,251,296} Based on these findings we suggested³⁶² "that these smaller proteins are formed in chromaffin granules *in vivo* due to the presence of proteases. One is of course reminded of the fact that in insulin-containing secretory granules, proteases are present which are essential for forming insulin from proinsulin". By now this concept is well established not only for CgA but also for additional components of chromaffin granules, e.g. the enkephalins³³⁷ and CgB.⁷⁹ Several defined small peptides proteolytically derived from CgA and CgB have been shown to occur in neuroendocrine secretory granules: pancreastatin³³¹ and betagranin¹⁵⁶ from CgA; and GAWK and CCB,^{22a} BAM 1745⁹⁰ and OA-8²⁰⁷ from CgB.

13.1. *Rate and degree of proteolysis*

Apparently, the processing of the chromogranins in different neuroendocrine vesicles does not occur at the same rate. Thus in adrenal medulla a relatively slow processing is obvious. In pulse-label experiments newly synthesized CgA appeared unprocessed after two⁷⁴ and even after 18 h,^{14,308} whereas for CgB a limited proteolysis occurred earlier⁷⁴ with some large

processed peptides being present after 2 h. On the other hand, in endocrine pancreas (insulinoma cells) a significant breakdown of CgA was already seen within much shorter times. Thus the half-time of conversion of CgA to betagranin (representing the N-terminal end of CgA) was only about 30 min.¹⁵⁵

To what degree the chromogranins are processed in various tissues depends both on the rate of processing and on the average life-span of the vesicles before they release their content, since even a low rate of processing may lead to considerable breakdown if vesicles do not secrete their content. In adrenal medulla the half-life of the secretory content (catecholamines) of chromaffin granules has been estimated to be close to seven days.³³⁶ Despite this long half-life, CgA in bovine adrenal medulla is only processed on average (in total adrenal medulla) to about 50% whereas for CgB a value of 85% has been reported.^{14,82} Obviously the rate of processing might be very low (in agreement with the pulse-labeling data discussed above), but one cannot exclude that processing does not occur throughout the whole life-span of chromaffin granules. For bovine neuroendocrine tissues it is only in the anterior pituitary that the degree of CgA processing is apparently less than in adrenal medulla,^{14,82} whereas in the intestinal mucosa (enterochromaffin cells) and especially in the endocrine pancreas, proteolytical degradation of CgA is extensive.³⁴⁵ This conclusion was established by a radioimmunoassay with antisera against synthetic peptides and measurements before and after trypsin digestion. The ratio of free peptide to total (after trypsin) peptide was 0.11 in adrenal medulla, 0.52–0.76 for intestinal mucosa and 1.09 for pancreas. Similar results were obtained for other species: in the porcine adrenal, CgA is little processed, whereas in the pancreas considerable breakdown has occurred as measured by pancreastatin radioimmunoassays and gel permeation chromatography,²⁸⁴ while in hypothalamus and especially in anterior pituitary, processing is limited.¹⁷⁸

For rat tissues the degree of breakdown increases from the adrenal medulla to the pituitary and finally to the endocrine pancreas.¹⁵⁹ In these studies, antibodies against betagranin and the intact CgA molecule were used to probe the degree of proteolysis. In another study⁵³ only an antibody against pancreastatin was employed for this purpose. In this case most immunoreactive material eluted close to the void volume when extracts of adrenal, pituitary, brain but also pancreas were subjected to gel permeation chromatography. Apparently, depending on what antisera are used the results may differ; however, the general conclusion seems valid that the degree of proteolytic processing of CgA in adrenal medulla and anterior pituitary is low; in other tissues especially the endocrine pancreas it is high. For CgB a study using antibodies against the GAWK peptides indicated a more extensive proteolysis in pituitary when compared with adrenals.¹⁶⁵

Proteolytic processing of CgA may also occur after their release from the neuroendocrine cells.³⁴⁶ In primary cultures of bovine chromaffin cells such a processing was observed in two studies;^{308,346} but not in another one.¹⁴ In any case it is not known whether such processing can also occur *in vivo*. The first evidence that the rate of proteolytic processing of CgA can be upregulated was recently obtained for chromaffin cells treated with reserpine.^{346a}

13.2. Pathways of proteolysis

One of the best-defined cleavage points in the processing of propeptides is at dibasic sites and in fact as shown in Figs 2 and 3, several of such cleavage sites are found in the CgA and CgB sequences and several of them are phylogenetically conserved at least in mammalian species. A detailed study using microsequencing and region-specific antibodies established actual processing at such sites for bovine CgA at residue 79 (preceded by KK) and 116 (by KR) and for human CgA at the second dibasic (KR) site counting from the N-terminus¹⁴ (for the latter cleavage see also Ref. 328). For CgA of the latter species (isolated from carcinoid tumors) there was also processing at position 210 (preceded by KR) and at position 340 (by KR).²⁸³ For the endocrine pancreas cleavage at dibasic site 131 (KR) in rat CgA has been established leading to the release of the N-terminal peptide called betagranin.¹⁵⁵ An analogous peptide (22,000 mol. wt) seems to be present in chromaffin granules.¹³⁶

For the well-characterized peptide, pancreastatin, which has been originally isolated from porcine pancreas³³¹ but is also found in human^{293,328} and bovine tissue,²⁴¹ processing at the C-terminal end requires cleavage (see Fig. 2) at a Gly-Lys site and on the N-terminal site cleavage at a monobasic amino acid at least for the porcine³³¹ and the bovine peptide.²⁴¹ A peptide found in bovine adrenal medulla³⁰ which is now known to be derived from CgA, also requires processing at monobasic site 16 (lysine) for its formation. In contrast, it is noteworthy that human CgA-derived peptides require for their formation cleavage at Asp-Pro and Trp.²⁸³ For CgB, processing at dibasic sites is well established, since several peptides detected in various tissues depend on such a cleavage, i.e. human GAWK (420–493) and CCB (597–657),²² bovine BAM-1745 (547–560),⁹⁰ OA 21,²⁰⁷ ovine OA 8²⁰⁷ and OA 60.²²⁶

Proteolytic processing of large proteins like the chromogranins could occur in successive steps either from the C- or N-terminal end or also independently from both sites. For CgA this latter process is apparently operating in bovine chromaffin granules.^{14,373} In fact, a preferential cleavage site is the dibasic site at the N-terminal end which leads to the formation of a smaller peptide (1–77) representing a significant spot in 2D-electrophoresis (see Fig. 1) and a larger size residue of the CgA molecule.^{14,373} It seems likely, although it has not been definitely demon-

strated, that in rat pancreas the formation of beta-granin can also occur by first removing the N-terminal end of CgA from the intact molecule by cleavage at a dibasic site which represents in this species the first cleavage site from the N-terminal end. For CgB of several species, processing of the intact molecule starting either from the N- or C-terminal site has also been established for several neuroendocrine tissues.¹⁰⁶

We have already pointed out above that in the various endocrine tissues the degree of proteolytic processing of chromogranins varies. This may lead to an accumulation of peptides of different sizes within the various tissues as, for example, shown in a detailed study on bovine tissues¹⁴ or in endocrine tumor cells lines.⁵⁹ These authors concluded that the "pattern of processing was qualitatively similar across tissues". However, we cannot yet exclude that in one of these tissues, CgA is specifically processed to a particular peptide (see Ref. 53a). For CgB significant differences between proteolysis in adrenal medulla vs pituitary have been found^{106,165} with the appearance of a unique 40,000 mol. wt C-terminal fragment in the anterior pituitary.¹⁰⁶

13.3. Proteases involved in processing

The first study on secretory granule enzymes responsible for the processing of CgA were performed by Hutton *et al.*¹⁵⁶ They found in insulin granules of the endocrine pancreas a calcium-activated enzyme which together with a carboxypeptidase H produced betagranin from CgA. In bovine chromaffin granules we described an analogous enzyme activity which produced a small peptide (22,000 mol. wt) from CgA probably also presenting betagranin.¹⁹² In the mean time two kex2-related proteases have been cloned for which an involvement in prohormone processing has been established.^{23,290,314,332} The mRNA of both PC1 (high concentration) and PC2 (low concentration) have been found in rat adrenal medulla²⁹⁰ and both PC1 and PC2 sequences have been shown to be present in glycoprotein H of bovine chromaffin granule membrane.⁴⁴ By using antisera against synthetic peptides derived from the primary amino acid sequence we could immunostain glycoprotein H for PC2, but the PC1-reactive spot corresponded to a larger (85,000 mol. wt) glycoprotein.^{183a} Thus it seems quite likely that the enzymes processing CgA in insulin and chromaffin granules as discussed above are in fact PC1 and PC2. These enzymes are capable of processing prohormones at pairs of basic residues and we have already pointed out that such cleavage occurs for both CgA and CgB. The involvement of those enzymes in CgA processing does not exclude the action of other enzymes which have been described previously in chromaffin granules (see Ref. 364 for references). Thus CgA can apparently also be processed by a trypsin-like enzyme,¹⁹² first described by Lindberg *et al.*²⁹⁹ This enzyme was reported to have

a molecular weight of 31,000.²⁹⁹ Furthermore, a thiol protease of 33,000 mol. wt splitting at dibasic sites has also been found in chromaffin granules.¹⁸⁸ A possible action of this enzyme on chromogranins has not been established. As already pointed out, CgA processing does not only occur at dibasic sites, thus proteases acting at monobasic sites or in other positions might also be involved. An endopeptidase with such a specificity (monobasic sites) has recently been purified from pituitary.⁶² When all these proteases are defined in molecular terms it will be interesting to see whether a concerted action of them is necessary to produce all the peptides formed from chromogranins and whether different activities of these enzymes in various tissues will explain the varying degrees of processing and possibly various ways of peptide formation.

Finally, the unusual possibility has to be considered, that the enzyme acetylcholinesterase may be involved in CgA breakdown. It has been claimed that highly purified acetylcholinesterase has proteolytic activity³¹¹ and can also process CgA.³¹³ However, this concept is still controversial³¹² and evidence against such proteolytic properties has also been presented.⁴³ Whether acetylcholinesterase is present in chromaffin granules and therefore in a position to attack CgA has also been controversial (see Ref. 364); however, at least this topic now seems settled, since a subspecies of this enzyme appears to be specifically localized in chromaffin granules.²⁸

14. REGULATION OF BIOSYNTHESIS

The chromogranins are constituents of the secretory content of neuroendocrine vesicles. In principle the biosynthesis of the chromogranins could be regulated *en bloc* together with other secretory components like the neuropeptides or in differential patterns allowing modifications of the secretory cocktail. We here present evidence that the latter mechanism is operating.

14.1. Regulation of synthesis by hormones

14.1.1. *Chromogranin A—a peptide regulated by steroids.* In 1987 we provided the first evidence that the biosynthesis of CgA in adrenal medulla was specifically regulated by corticosteroids.³⁰³ After hypophysectomy in rats the CgA but not the CgB content of adrenal medulla declined markedly, apparently due to the atrophy of the adrenal cortex leading to a lack of corticosteroids. This concept was confirmed when it was demonstrated that after hypophysectomy the lowered mRNA levels of CgA (but not that of CgB) in adrenal medulla could be restored to control levels by cortisone treatment.⁸³ The specificity of the action of cortisone could be seen by its lack of effect on the CgB message. Similarly, in PC12 cells and in bovine chromaffin cells in culture, cortisone treatment specifically increases the levels of the CgA message.^{162,266} For the anterior pituitary

an analogous influence of cortisone could also be demonstrated,^{88,115} on the other hand, in brain dexamethasone treatment or adrenalectomy did not change CgA levels or that of its mRNA.³⁵⁰ In the corticotrophic cell line, AtT-20, cortisone treatment increases the CgA mRNA levels to 250% whereas those for POMC peptides declines (to 40%)³⁴³ and an analogous regulation was observed for the CgA message in pituitary tumor cell cultures^{319a} and in an insulinoma cell line.¹⁶² It has not been established by what mechanism cortisone induces the biosynthesis of CgA. The induction by cortisone of the CgA message in the pituitary occurred relatively late, i.e. with a time-lag of 24 h after the injection, interpreted as the action of cortisone being indirect via protein synthesis.⁸⁸ However, a direct effect of the cortisone receptor on the gene cannot be excluded, although gene analysis did not reveal a perfect match to a glucocorticoid responsive element in the promoter region.^{162,374}

Another example of steroids regulating CgA synthesis has been found in the parathyroid cell cultures. 1,25-Dihydroxycalciferol-treatment induced a several-fold increase of the CgA mRNA^{232,277a} via an effect on CgA gene transcription, whereas CgA mRNA stability was not affected. Finally oestrogen steroids are also involved in the regulation of CgA synthesis in the pituitary gland. In female rat pituitaries, CgA drops with maturation of the animals.⁶ After ovariectomy CgA and its mRNA rises and is suppressed again by oestrogen treatment.^{4,5} In male rats oestrogen treatment markedly lowered the pituitary mRNA for CgA.⁸⁷ Furthermore, male rats had higher CgA mRNA levels in the pituitary than female rats which are of course under the constant influence of oestrogens, whereas CgB message was equal for both male and female. Thus the biosynthesis of CgA in the pituitary is suppressed by endogenous or exogenous oestrogen. These changes take apparent place in the gonadotrophic cells as indicated by immunohistochemistry.⁸⁷ However, for these oestrogen effects it has not yet been demonstrated that they are due to a direct action on these cells. They may also be indirect via hypothalamic regulation. In adrenal chromaffin cells, oestrogen is without influence on CgA mRNA levels.^{4,87}

14.1.2. *Regulation of chromogranin B.* When GH₄C₁ cells (a rat pituitary tumor cell line) are treated with humoral factors (E₂, insulin and epidermal growth factor, EGF) the number of secretory granules increases several-fold. This leads to increased levels of prolactin, but also of CgB and secretogranin II.²⁸¹ In GH₃B₆ pituitary cell lines, CgB-mRNA is reduced by dexamethasone, whereas oestrogen increased it.¹⁹⁸

14.2. Regulation of synthesis by stimulation of cells

When chromaffin cells of the rat adrenal medulla are stimulated reflexly *in vivo* by insulin-induced hypoglycemia, the levels of enkephalins,^{177,303} but also

of several other neuropeptides, i.e. calcitonin gene-related peptide, neuropeptide Y, neurotensin and substance P,^{78,194} increase several-fold. On the other hand, those of CgA and CgB remain unchanged³⁰³ clearly demonstrating an apparent dissociation of the biosynthesis of chromogranins and neuropeptides. This conclusion was confirmed at the mRNA level.⁸³ Whereas in these experiments with insulin-induced hypoglycemia the biosynthesis of chromogranins was apparently unchanged, treatment of rats with reserpine induced such effects.¹⁴² A high dose of reserpine leads to a prolonged stimulation of the adrenal medulla. This treatment induced a significant increase of the levels of CgB and its mRNA (600%), whereas for CgA mRNA a smaller increase (230%) was observed. No definite explanation can be offered as to why indirect stimulation of the adrenal by insulin or by reserpine affects chromogranin biosynthesis differently; however, it has been suggested¹⁴² that the time of stimulation may be important with only a prolonged stimulation as provided by reserpine leading to such biosynthesis changes. For the chromogranin biosynthesis in brain the first data are already available. Thus in rats in which seizures have been induced by the injection of kainic acid, changes in the mRNA levels for CgA³⁵⁰ and for CgB^{219a} have been observed. Apparently, in this model intensive stimulation of certain neurons in brain does not only change the levels of neuropeptide Y but also those of the chromogranins.

For bovine chromaffin cells in culture, results on the biosynthesis of CgA are controversial. Two groups of authors measured the incorporation of radioactively labeled amino acids into CgA. Whereas one group³⁰⁷ reported an increased incorporation after depolarization (nicotine, K⁺), another one found no change³⁶⁰ for CgA, but the expected change for the enkephalins. Possibly, *in vitro* factors like the length or intensity of stimulation (see above) may lead to different results. In any case, in pancreatic endocrine islets, amino acid incorporation into both insulin and CgA is increased several-fold in the presence of glucose which was interpreted as an effect at the translational level;¹¹⁹ however, in all these studies, *in vitro* mRNA levels of CgA were not yet determined.

14.3. Regulation by second messengers

In cultured chromaffin cells, amino acid incorporation into CgA is increased in the presence of phorbol esters (short incubation), but not by forskolin.³⁰⁷ At the mRNA level, forskolin has been shown to induce an insignificant rise after 36–48 h,⁸⁰ but after more than two days of treatment a doubling of the mRNA levels was apparent.¹⁶² This result is consistent with the finding of a cAMP-responsive element in the CgA-gene.^{162,374} Varying effects of phorbol ester on mRNA levels have been reported in different cells. In a calcitonin-producing cell line, phorbol ester but not forskolin treatment for 48 h led to a doubling of

the CgA mRNA level.^{238,239} On the other hand, in bovine chromaffin cells, treatment for two days with the same agent significantly decreased the CgA message^{80,162} and an analogous finding has been obtained for human neuroblastoma cells.³⁵¹ Since phorbol esters, depending on the time of incubation, can activate or down-regulate protein kinase C it is not clear by what mechanisms the CgA gene is regulated. Gene analysis has not revealed an established AP-I site for CgA.^{162,374}

In conclusion, the chromogranin genes can apparently be activated in a specific way. Thus the biosynthesis of the secretory content of neuroendocrine vesicles is not regulated *en bloc* (see also Ref. 103a). Much further work is required before we will understand the detailed mechanisms at the cellular and genetic level which regulate the biosynthesis of this secretory cocktail. Certainly, such sophisticated mechanisms argue for specific functions of these regulated peptides, but are we yet in a position to understand them?

15. FUNCTION

15.1. Involvement in catecholamine storage

When CgA was still considered to be a specific component of chromaffin granules which also contain a high concentration of catecholamines, their possible involvement in the storage of these molecules was a much discussed topic. As outlined above, the chromogranins are now known to be present in many neuroendocrine granules which store only small amounts if any of biogenic amines. This finding indicates that chromogranins cannot be proteins with an exclusive function for amine storage, but it does not exclude such a function in some vesicles. As discussed in detail previously³⁷¹ (see also Ref. 132) the chromogranins cannot be involved in a stoichiometric binding of catecholamines or ATP and there is also, as shown by nuclear magnetic resonance studies, no stable storage complex made up of chromogranins, nucleotides and catecholamines. However, a certain interaction between all these molecules apparently takes place (see Ref. 371) and could help to reduce the high osmolar pressure inside chromaffin granules,¹³⁷ although catecholamine/ATP complexes fulfil this purpose.¹⁸⁷ Direct evidence for an interaction of nucleotides with purified CgA has recently been obtained with a fluorescence probe.³⁷⁶ It was concluded that adenosine bases play an important role in this interaction and if this interaction also occurs *in vivo* it might provide a basis for some involvement in catecholamine storage. From solutions containing CgA, catecholamines and ATP in stoichiometric concentrations comparable to the granule content, an opaque precipitate can be sedimented which also indicates some interaction.⁹⁴ Finally, calcium, which interacts with the chromogranins (see below), increases the binding of catecholamines to the chromogranins.^{339a,354}

15.2. Enzyme inhibition

CgA was shown to inhibit the proteolytic cleavage of prohormones by trypsin or serine proteases.²⁹¹ It would be interesting to see whether such inhibition can also be observed with the newly defined endoproteases PC1 and PC2. However, to prove whether chromogranins have a role as regulators of post-translational processing of prohormones²⁹¹ would require more direct experiments, e.g. by introducing the chromogranin genes into secretory cells originally lacking it and subsequently defining their effects on proteolytic cleavage rates.

15.3. Binding of calcium and possible consequences for granule formation

In 1977 we reported³⁶¹ the following unpublished observation and some speculations based on it: "In this context it is interesting to note that *in vitro* calcium can precipitate the acidic proteins of the granule content, i.e. the chromogranins (Winkler, unpublished observations), which is reminiscent of the fact that calcium precipitates casein, an acidic protein of the milk secretion granules.⁷⁵ Furthermore, the condensation process, at least in exocrine pancreas, is apparently not energy dependent.¹⁷⁰ This would be consistent with the involvement of a calcium uptake which does not require energy as found for chromaffin granules. It therefore seems justified to speculate that during "the condensation process" calcium enters newly formed chromaffin granules before the other small molecules, i.e. ATP and catecholamines are accumulated. However, there is no direct evidence to support such a sequence of events." In the meantime, detailed studies on the relationship of calcium and chromogranins have appeared. Gratzl *et al.*^{268,269} have established that CgA can bind considerable amounts of calcium (152 nmol/mg protein with a dissociation constant of 54 μ M). This property explains why in intact chromaffin granules despite the high concentration of total calcium (40 mM) the free concentration is low being about 20 μ M.³⁵ Not surprisingly, parathyroid secretory protein I which is in fact CgA was also found to bind calcium.^{108,202} In the presence of this cation, CgA binding to membranes was also enhanced.^{108,202} CgB is apparently also able to bind calcium.¹¹¹ The binding of calcium to CgA induces pH-dependent conformational changes.^{375,377} In these studies, CgA was found to bind up to 1150 nmol of Ca²⁺/mg protein (dissociation constant 2.7–4 mM).

What are the functional implications of this calcium-binding by chromogranins? Several groups suggested^{108,110,367,371} that the aggregation of CgA by calcium may be involved in the condensation process in the early phase of granule formation in agreement with the speculation quoted above. However, more complicated schemes for the function of chromogranins (and also secretogranin II) have been proposed: thus it has been suggested that these acidic

proteins "function as helper proteins in the packaging of various hormones".²⁷⁴ Although it has been demonstrated²⁷⁵ that antibodies against CgB when co-expressed in secretory cells became co-packed with CgB in the regulated pathway this does not provide direct evidence for co-packing of hormones and chromogranins which do not interact specifically like antigen/antibodies. Furthermore, the fact that the concentration of chromogranins/secretogranins in secretory granules other than chromaffin granules is low makes it difficult to envisage a stoichiometric "helper process". The suggestion that the tyrosine sulfation of these proteins²⁷⁴ may act as a sorting mechanism has not been supported by subsequent studies.¹⁴⁸ In a more detailed modification of these proposals it has been suggested that chromogranins are partly membrane bound in the trans-Golgi network and that these molecules together with calcium, aggregate with further chromogranins forming together with other secretory peptides the "granule core" for vesicles which subsequently pinch off.^{153,262a} This scheme assumes that sorting occurs concomitant with condensation as proposed previously;³³⁴ however, evidence to the contrary, i.e. sorting precedes condensation has also been obtained.³³⁵ One can only re-emphasize that many neurosecretory granules contain only small amounts of chromogranins, thus a general hypothesis which concentrates on these minor components apparently neglects the role of the major granule constituents, i.e. the respective hormones, for the granule formation process.

15.4. Precursors of functional peptides

Two features of chromogranins discussed above are consistent with a function of chromogranins as precursors of active peptides: chromogranins, like prohormones or proneuropeptides, are proteolytically processed *in vivo* and their biosynthesis can be regulated by humoral factors or by stimulation of the cells storing them. Direct support for this concept was provided when pancreastatin, a 49-amino acid peptide derived from CgA, was isolated from porcine pancreas and found to have a defined function in inhibiting the stimulated secretion of insulin from endocrine pancreas.^{66,331} In the mean time pancreastatin peptides were isolated from other species including human^{101,283,293} and bovine²⁴¹ material and pancreastatin immunoreactivity was found to be widespread in the neuroendocrine system (see Table 1), although this does not prove the presence of the free peptides in all these organs since the antisera against pancreastatin can also react with CgA. However, in addition to the pancreas the free peptide has also been found in the pituitary gland^{241,284} and chromatographic analysis suggests its presence also in porcine thyroid gland and duodenum.³¹

Many subsequent studies, but not all of them (see below and Ref. 288), have confirmed the original finding, that the 49-amino acid porcine pancreastatin

and also its 16-amino acid C-terminal peptide can inhibit glucose-stimulated insulin release from perfused rat pancreas or isolated acini^{66,331} and from a pancreas tumor cell line.²¹⁸ Apparently, porcine pancreastatin is not only effective in the rat pancreas,^{66,113,167,258,305,331} but also in mouse,^{2,209} whereas in canine pancreas an effect²⁷¹ or no effects²⁵⁴ have been reported.

For porcine pancreas "no effect at all on endocrine secretion of insulin, glucagon and somatostatin" was found.¹⁴⁶ Inhibiting effects on rat pancreas were also exhibited by human pancreastatin^{95,292} and not surprisingly also by rat pancreastatin and its 26-residue C-terminal peptide.^{96,228,257} For glucagon-mediated insulin release, pancreastatin was reported either as inhibitory²⁵⁸ or as stimulatory.¹⁶⁸ For glucagon release itself, increases caused by pancreastatin were reported in two studies^{66,271} whereas another one, even with the homologous pancreastatin in rat, was unable to find any effect.²⁵⁷ Effects of pancreastatin on exocrine pancreatic (e.g. amylase release) secretion have also been reported;^{96,227} however, on isolated acini no effect could be seen^{65,98,229} and it has been suggested that pancreastatin acts on the secretion by inhibiting vagal nerve activity.²²⁹ On the other hand, for guinea-pig, isolated acini pancreastatin inhibited cholecystokinin-induced amylase release.¹⁶⁶ Several further tissues have been tested for effects of pancreastatin. In rabbit isolated parietal cells of the stomach, pancreastatin had a direct inhibitory effect on secretion through interference with multiple second messenger pathways;²⁰⁶ on the other hand, in dogs, gastric acid secretion was increased.¹³⁰ Secretion is also inhibited by this peptide in the parathyroid gland.^{76,77} Interestingly, antibodies against pancreastatin potentiate the secretion from these cells at different calcium concentrations, apparently by neutralizing CgA peptides released into the medium from these cells.⁷⁶ This appears to be the first evidence that CgA-related peptides when secreted have a "physiological" function.

Some actions on the central nervous system are also already indicated. Peripherally administered pancreastatin modulates memory processing in tests on mice⁹¹ and intracranial microinfusion¹²⁰ leads to changes in blood glucose and free fatty acids.

A second peptide derived from CgA, i.e. chromostatin, has been shown to be an effective inhibitor of catecholamine secretion from chromaffin cells.^{103,309} In a first study³⁰⁹ it was established that CgA inhibits catecholamine secretion induced by nicotine, but not that by high K^+ , only after pre-incubation with the cells or after trypsin treatment indicating an involvement of a breakdown product. In a second study CgA was processed by proteases and the peptides inhibiting catecholamine secretion were analysed. A shorter peptide present in these sequences was synthesized and found to be a potent inhibitor of catecholamine secretion induced both by carbamoylcholine or by depolarizing K^+ concentration. The peptide, i.e.

chromostatin, inhibited secretagogue-induced calcium influx.^{103,103b} It has not been demonstrated that chromostatin is actually formed *in vivo* or whether a larger molecule has this function on catecholamine release. Chromostatin is not flanked by basic pairs of amino acids (Fig. 1); thus the actual processing of this or larger peptides has still to be elucidated.

Finally, some actions of further peptides derived from CgA are likely. The N-terminal peptide (1-40) stimulated the secretion of CGRP and inhibited the secretion from calcitonin and of parathyroid hormone-related protein from a cell line.^{60,61,65a} CgA, or a peptide formed from it during incubation, inhibits the secretion of POMC from the AtT-20 mouse cell line. Furthermore, CgA antisera increased basal secretion indicating a physiological inhibitory feedback of CgA secreted from these cells as discussed above.³⁴³

Two recent findings raise interesting new possibilities for the function of chromogranins. CgA, or peptides derived from it, were shown to have neurotrophic activity for embryonic chick dorsal root ganglia.³⁵³ With the availability of synthetic peptides it should be possible to define this observation in more detail. In endocrine pancreas, pancreastatin inhibited beta-cell DNA synthesis and lowered the content of spermidine in these cells which has been implicated in beta-cell replication.³¹⁰ An interaction with other peptides has also been indicated, since the N-terminal segment (1-46) of CgA significantly suppressed contraction of vascular smooth muscle caused by endothelin.¹³⁵

We can conclude that there seems to be a good indication that peptides derived from CgA are functional. Thus the CgA molecule, which is proteolytically processed *in vivo* and whose biosynthesis is regulated (see above) seems to belong to the ever-growing group of pro-peptides like those giving rise to neuropeptides. In apparent contrast to many of the neuropeptides, the phylogenetic conservation of the functional peptides (pancreastatin and especially chromostatin) is rather limited (see Fig. 2). If these peptides have specific receptors, then it is difficult to see how the lack of conservation of the peptide can be correlated in phylogeny with such a specific receptor. Furthermore, we will still need more data on the actual formation of these peptides (especially chromostatin) in the various tissues *in vivo*.

The indications that these peptides have trophic functions, or influence cell replication or can interact with other peptides should stimulate more research. Finally, regulatory functions, e.g. on synapse formation during development and on the regulation of postsynaptic receptors, should be considered.

16. CONCLUSIONS

Twenty-five years of research on the chromogranins have led to many established facts on their physico-chemical and molecular properties and on their widespread distribution in the neuroendocrine

system. The function of these proteins is most likely to be as precursors for active peptides. In this context, future research to yield the most rewarding results should answer the following questions. Which functional peptides are actually formed *in vivo*? Which proteases are involved? Is the function of the peptides, at present only defined by their exogenous addition, also of physiological significance? Are functional peptides (and their yet undefined receptors) conserved during phylogeny going beyond the mam-

malian system? Do changes in the biosynthesis of these proteins control the functions of peptides formed from them? Is the proteolytic processing of the chromogranins regulated?

Acknowledgements—The original research of the authors quoted in this paper was supported by the Dr Legerlotz-Stiftung and by the Fonds zur Förderung der wissenschaftlichen Forschung (Austria). We are grateful to Mrs M. Viehweider for typing the manuscript and to Mr Ch. Trawöger for preparing the photographs.

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(Accepted 16 March 1992)