

THE GENE FOR HUMAN CHROMOGRANIN A (CgA) IS LOCATED ON CHROMOSOME 14

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Chromogranin A (CgA) is a protein that is present in most neuroendocrine tissues and is co-secreted with their resident hormones. We have assigned the CgA gene to human chromosome 14 by hybridization of a CgA cDNA probe cloned from a cDNA library of human medullary thyroid carcinoma cells to spots of individual human chromosomes flow-sorted onto nitrocellulose filters. Southern analysis of human genomic DNA with the same probe revealed only 1-3 restriction bands. These studies indicate that the CgA gene is probably single copy and not a member of a dispersed, multigene family. The CgA gene is not co-localized with the genes of any of the CgA-associated hormones. © 1987 Academic Press, Inc.

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Chromogranin A (CgA) is a protein that is found in most hormone-producing tissues, including the pancreas, pituitary gland, parathyroid glands, C-cells of the thyroid gland, and the neuroendocrine cells of the lung and gut (1). It was originally described as the major soluble protein of adrenal chromaffin granules and in retrospect is indistinguishable from parathyroid secretory protein (2). In hormone-producing tissues it is co-localized and co-secreted with resident hormones of the tissue (3,4). It is found in elevated concentrations in the serum of patients with tumors of neuroendocrine tissues, including medullary thyroid carcinoma and small cell lung cancers (4,5). cDNA clones have been used to demonstrate the neuroendocrine specificity of CgA mRNA (6,7) and to predict the amino acid sequence of the bovine form (7,8).

#### Materials and Methods

The CgA-specific cDNA probe used in these studies is 339 base-pairs in length and encodes the first 113 amino acids of human CgA; its cloning from a lambda gt11

cDNA library of human medullary thyroid carcinoma cells has been described previously (6,9). For the hybridization studies the probe was labeled with  $^{32}\text{P}$ -dCTP using the method of Feinberg and Vogelstein (10).

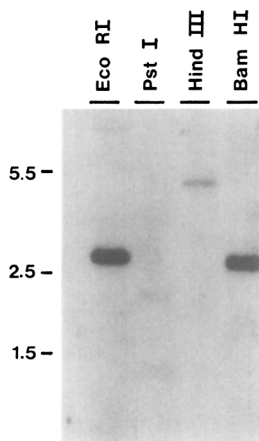
For Southern blot analysis, total cellular DNA was extracted from peripheral mononuclear cells using standard methods (11). The mononuclear cells were separated from the blood of normal human subjects by centrifugation with Histopaque-1077 (Sigma Chemical Co., St. Louis, Missouri) according to the method of Boyan (12). DNA from cultured human lymphoblastoid cells (GM130-A) was extracted using the same methods. Aliquots of the DNA samples were digested with the indicated restriction endonucleases, fractionated by agarose gel electrophoresis, transferred to nitrocellulose, and hybridized with the CgA-specific probe (13).

Human chromosomes were isolated and sorted as previously described (14-16) from diploid human lymphoblastoid cells (GM 130A) grown in suspension culture in RPMI medium supplemented with 10% fetal calf serum. Sorting was accomplished using an EPIC 5 V flow cytometer (Coulter Electronics, Inc., Hialeah, FL). The chromosomes were sorted directly onto nitrocellulose discs. Each chromosomal type can be sorted with greater than 90% purity, with the exception of numbers 9-12 and 14 and 15.

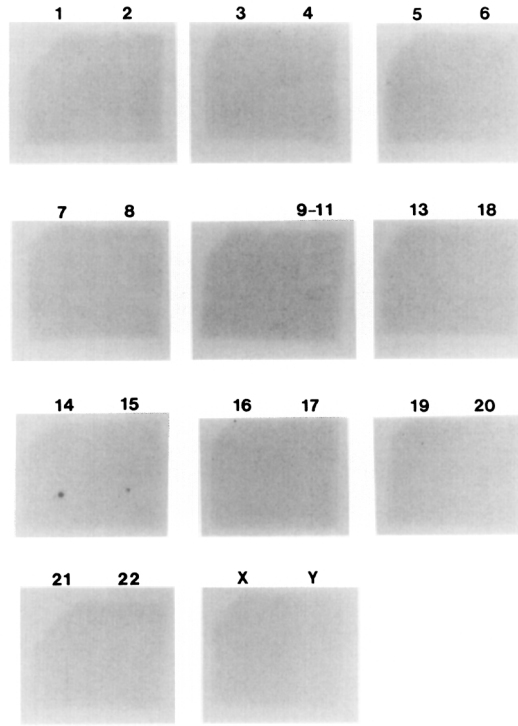
Chromosomes 9-12 overlap in one peak; 14 and 15 can be separated from each other, but with approximately 15 percent contamination of 15 with 14 (16). Thirty thousand chromosomes of each type were sorted onto the nitrocellulose discs, resulting in individual spots of  $3 \times 10^4$  chromosomes for 1-8, 13-22, x,y and 1 spot of  $1.0 \times 10^5$  chromosomes for 9-12. The DNA was denatured by washing the discs in 1.5 M NaOH, 1.5 M NaCl for 5 minutes, then neutralized in 0.5 M Tris, pH 7.5, 3.0 M NaCl, and baked at 80°C for 90 minutes under vacuum. The filter discs were hybridized with the CgA-specific probe for 24 hours (6,13).

## Results

The results of the Southern blot analysis of DNA from mononuclear cell DNA digested with various restriction endonucleases are shown in Figure 1. The appearance of a single dominant band of hybridization in some of the lanes suggests that the CgA gene occurs as a single copy; however, multiple bands are present in 2 of the 4



**Figure 1.** Southern blot analysis of DNA extracted from normal human mononuclear blood cells. Aliquots of 10  $\mu\text{g}$  of DNA were digested with the restriction endonucleases shown, fractionated by agarose gel electrophoresis, transferred to nitrocellulose, and hybridized with the CgA-specific cDNA probe. DNA size markers in kilobases are shown at the top.



**Figure 2.** Autoradiograph of flow-sorted human chromosomes hybridized with the CgA-specific cDNA probe. The intense signal in compartment 14 indicates that the CgA gene is located on chromosome 14. The faint signal in compartment 15 likely represents hybridization to chromosome 14 that commonly contaminates preparations of chromosome 15.

lanes. The cDNA did not contain restriction sites for any of the endonucleases used. Similar results were obtained from Southern blot analysis of DNA extracted from GM1289-A cells (data not shown).

Figure 2 shows the results of hybridization with the cDNA probe of human chromosomes sorted by flow cytometry and blotted on nitrocellulose (14-17). A strong hybridization signal is seen over the blot corresponding to chromosome 14. A much weaker signal is seen over the blot corresponding to chromosome 15. This is most likely due to the small amount of chromosome 14 DNA in the chromosome 15 spot because of the close proximity of chromosome 14 and 15 peaks in the flow histogram. It is also possible that a weakly cross-hybridizing species, such as a pseudogene for CgA, exists on Chromosome 15.

#### Discussion

CgA is a protein found in many neuroendocrine tissues (6-8). It is conserved with many peptide hormones and with catecholamines and its serum concentration is in-

creased in patients with eutopic and ectopic endocrine tumors (4). For example, serum CgA levels are increased in patients with medullary thyroid carcinoma and small cell lung cancer where its co-secreted hormone is calcitonin (4,5). Although its function is not known, CgA may play a role in the process of hormone secretion (18). There is some evidence to suggest that there may be multiple species of CgA and CgA-like proteins (17). However, our results suggest that the CgA gene is not a member of a dispersed, multi-gene family (Figures 1 and 2). Two of the peptides most clearly associated with CgA, calcitonin and parathyroid hormone, are encoded by genes of chromosome 11 (20,21). Furthermore, the genes for none of the other hormones or genetic abnormalities, such as Multiple Endocrine Neoplasia, that may be associated with CgA are assigned to chromosome 14 (20-30). Thus, it is not likely that cis-acting factors account for the co-regulation of CgA and its associated hormones (14). Further studies are in progress to define the function of CgA in neuroendocrine cells.

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