# Biochemical Properties of Tracheobronchial Mucins from Cystic Fibrosis and Non-Cystic Fibrosis Individuals

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ABSTRACT. Tracheobronchial mucins from healthy individuals and from patients with bronchial asthma or cystic fibrosis (CF) were isolated from lung mucus, purified, and their chemical and physical properties compared. Normal and asthmatic mucins required both a dissociating and a reducing agent for solubilization and exhibited identical chromatographic behavior on Sepharose 4B, Sepharose 2B, and hydroxylapatite and similar amino acid and carbohydrate compositions. In contrast, 1) CF lung mucins were solubilized in the absence of dissociating and/or reducing agents and 2) the majority of the CF mucins analyzed was eluted in the included volume of Sepharose 4B with  $K_d$  values of 0.3 ± 0.1 rather than in the void volume and thus appeared smaller than normal and asthmatic mucins. The lower molecular weight mucins in CF sputum apparently are produced by bacterial or inflammatory cell proteinases since radiolabelled asthmatic mucin was digested to smaller fragments when incubated with crude CF lung mucosal samples. Furthermore, mucins secreted by tracheal explants from CF and from non-CF individuals eluted in the void volume on Sepharose 4B, suggesting that CF tracheobronchial mucins were not inherently smaller than non-CF mucins. (Pediatr Res 22: 545-551, 1987)

### Abbreviations

BGA, blood group activity BA, bronchial asthma CF, cystic fibrosis DTT, dithiothreitol ME, 2-mercaptoethanol N, normal SDS, sodium dodecyl sulfate TBM, tracheobronchial mucin V<sub>o</sub>, void volume MEM, minimal essential medium PMSF, phenylmethylsulfonyl fluoride

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mucus of exocrine glands or epithelial tubular systems of the respiratory, gastrointestinal, and reproductive tracts (2) which is the basis for the generally held assumption that CF mucus is "abnormal." Since mucus glycoproteins (mucins) are the macromolecules responsible for the viscoelastic properties of mucus (3–5), alterations in mucin structure could effect the physiological behavior of mucus.

To date, the most comprehensive studies on CF mucins (6-8) have utilized TBM isolated from CF sputum as copious amounts of such material are readily available. There is limited information available on TBM from healthy airways primarily because of the difficulties in obtaining sufficient amounts of adequate material for analysis. Thus, insufficient information currently exists to determine whether CF and non-CF mucins are different (see Ref. 9 for review) although available studies suggest differences in sulfation of CF respiratory tract mucins (10). However, recent studies have suggested that TBM from healthy airways exhibit physical properties (size, chromatographic behavior on molecular sieving, solubility) different than those of mucins from pathological airways (11-13). While isolating and purifying human mucins from normal and pathological mucus, we obtained results that indicated that CF TBM were considerably more heterogeneous and smaller in size than non-CF TBM. The smaller size of CF mucins in the samples analyzed could result from in vivo proteolysis, as CF airways are chronically infected with pathogens that produce proteinases (14, 15) and both CF sputum and bronchial washings contain high levels of extracellular proteinase activity (16-18). Alternatively, the smaller mucins could reflect inherent differences in CF mucins. These possibilities were examined; the former interpretation was supported by studies reported herein.

#### MATERIALS AND METHODS

*Materials*. All chemicals were obtained from commercial sources and were of the highest available purity. Mucosal samples were obtained from the airways of adult volunteers (N-1, 21-yr-old man; N-2, 34-yr-old woman) via the hypertonic saline stimulation procedure of Barton *et al.* (11), whereas BA and CF samples (BA, 29-yr-old woman; CF-1, 21-yr-old woman; CF-2, 22-yr-old man; CF-3, 18-yr-old man) were obtained on expectoration. All samples were collected on ice and were frozen at  $-20^{\circ}$  C. Tracheas for explant cultures were obtained within 3 to 6 h after death of a 16-year-old girl with CF and from a 69-yr-old man with no chronic obstructive pulmonary disease.

Isolation and purification of mucins. The procedures used for the solubilization and chromatographic resolution of the components of LM-gel and for the purification of tracheobronchial mucins have been described (5). Briefly, the samples were thawed, diluted with saline, homogenized, heated in a boiling

CF is the most prevalent recessive genetic disease of Caucasians (1). A major pathological manifestation of CF is the blockage by

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water bath for 10 min., dialyzed overnight *versus* water, separated by centrifugation into soluble and insoluble phases, lyophilized, and weighed. Samples were either 1) reduced and carboxymethylated and dialyzed as previously described (5) and chromatographed on Sepharose 4B with buffer A (0.15 M ammonium acetate, pH 7.0, 0.1% SDS) or 2) solubilized in buffer A that had been made 1% in SDS and 0.02 M ME and chromatographed on Sepharose gels with eluent buffer A which had been made 0.02 M in ME. The mucin fractions were purified as described in the text.

Radiolabeling of TBM. Radiolabeled mucin was prepared by reductive carboxyamidomethylation with iodo[1-<sup>14</sup>C]acetamide. Five mg of BA mucin that had been purified by procedure B was solubilized in 1 ml of reducing buffer (1% SDS, 0.1 M tris, 0.2% EDTA, 0.01 M DTT, pH 8.5) at room temperature for 4 h. Thirty  $\mu$ mol of sodium arsenite was added to the sample to destroy excess DTT. After 5 min, 100  $\mu$ l containing 5  $\mu$ mol of iodoacetamide, and 5  $\mu$ mol iodo[1-<sup>14</sup>C]acetamide (19.1  $\mu$ Ci/ $\mu$ mol; New England Nuclear) was added; alkylation was carried out in the dark for 1 h. The pH was adjusted to 4.5; the sample was dialyzed extensively (5) and lyophilized.

Analytical procedures. Assays were performed as previously described (5) except as noted below. Amino acid analyses were performed with a Glenco custom modular analyzer using a dual temperature, single column system with a 9  $\mu$ m, 12% crosslinked resin (Glenco CX-9-12). The components of the mixtures were resolved with Pierce Picobuffer system II, sodium citrate form. Duplicate samples for amino acid analyses were hydrolyzed at a concentration of 50-500 µg/ml in 6 N HCl for 20 h under vacuum at 100° C (optimal conditions selected from hydrolysis curves of BA mucin). Samples for hexosamine analyses were hydrolyzed for 6 h at 100° C and resolved at 56° C on the Glenco analyzer by elution with sodium citrate-borate buffer (19). Monosaccharides were analyzed by gas liquid chromatography of the trimethylsilyl derivatives of the methyl glycosides using a slight modification of the procedure of Bhatti et al. (20). Sialic acid was determined by the thiobarbituric acid method of Aminoff (21). Sulfate was determined by the barium chloranilate procedure of Spencer (22) after hydrolysis for 6 h in 1 N HCl at 100° C and by the rhodizonate method of Silvestri et al. (23) after alkaline hydrolysis. Both procedures yielded values that agreed within 5%; the rhodizonate procedure was more sensitive, faster, easier to perform, and required considerably less material. Sulfate values were corrected, where necessary, for trace amounts of SDS as determined by the methylene blue reaction of Takagi et al. (24). Blood group titers were measured with 2 hemaglutinating units of anti-A or anti-B serum or Ulex Europas agglutinin-1 (Miles-Yeda Ltd.) and a 2% suspension of A, B, or O red blood cells (25). Electrophoresis was carried out on 5% polyacrylamide or 3% polyacrylamide/0.5% agarose gels according to the procedure of Holden et al. (26) and stained with Coomassie blue or periodic acid-Schiff reagents. The mucin samples were incubated overnight in electrophoresis sample buffer that contained ME before loading.

Enzymatic assays. Proteinase activity was assayed by a colorimetric procedure. The reaction mixture contained 100 µl of homogenized sample and 100 µl of casein yellow (0.3 mg) suspended in 0.01 M NaCl. After incubation at 37° C for 0-3 h, the reaction was stopped by adding 0.5 ml of ice cold 5% phosphotungstic acid. The insoluble protein was sedimented by centrifugation at  $3000 \times g$  for 5 min. The supernatant was removed, made alkaline by adding 150 µl of 4 M NaOH, and the absorbance measured at 428 nm. Hyaluronidase assays were carried out with bovine testicular hyaluronidase (300 U/ml) (Calbiochem) which had no detectable proteinase activity when tested with albumin and which was active when tested with hyaluronic acid and monitored by orcinol assay. To 0.5 ml of sample in buffer (0.15 M NaCl/0.1 M sodium acetate, pH 5.2), 25  $\mu$ l of enzyme was added at 0, 4, 8 h. Incubation was carried out for 24 h at 37° C. The enzyme was inactivated by increasing

the pH to 7.0. Chromatography was carried out on Sepharose 4B with 0.15 M ammonium acetate, pH 7.0, as eluent buffer.

Tracheal explant cultures and isolation of radiolabeled mucin in vitro. Tracheas were rinsed several times in ice-cold transport media which contained 100  $\mu$ g of tobramycin sulfate and 1.1  $\mu$ g of amphotericin per ml of media. The submucosa-mucosa layers were microdissected from the underlying cartilage and serosa. The tissues were cultured as 2–4 mm<sup>2</sup> explants in 35-mm Petri dishes as described by Boat and coworkers (27, 28) except for the following modifications. MEM (Eagle's base) supplemented with nonessential amino acids (at the concentrations in Media 199) was used and the sulfate concentration was reduced to 0.1% of that normally used in MEM. The media contained tobramycin base (gift of Lilly Research Laboratories), 50  $\mu$ g/ml; amphotericin B, 0.25  $\mu$ g/ml; and carbenicillin 250  $\mu$ g/ml. Explants were cultured in a 5% CO<sub>2</sub>/95% air water-saturated environment at 37° C.

Radiolabels were added to the culture media at the time of incubation. Every 24 h the culture media were removed, the tissue was washed with unlabeled media, and fresh media and radiolabel were added. The culture media and washes were combined and centrifuged briefly. Two volumes of a proteinase inhibitor solution (0.01 M EDTA, 0.1 M  $\epsilon$ -amino caproic acid, 5 mM benzamidine HCl, 1 mM phenylmethylsulfonyl fluoride, 10 mM N-ethylmaleimide) were added and the media was dialyzed *versus* 100 ml of the proteinase inhibitor solution in the cold overnight and then *versus* 2 l of water for 24 h with two changes of water and lyophilized.

Labeled mucins were isolated from other labeled macromolecules by chromatography on Sepharose 4B-CL. The void volume fractions were dialyzed *versus* urea/NaCl and then water to remove SDS (5) and then lyophilized. The samples were dissolved in a 10 mM sodium phosphate pH 6.5/4 M GuHCl buffer containing proteinase inhibitors (5 mM NaEDTA; 5 mM Nethylmaleimide; 1 mM PMSF; 2.5 mM benzamidine HCl; 0.05 M  $\epsilon$ -aminocaproic acid). The density was adjusted to 1.42 g/ml with cesium chloride. Isopycnic density centrifugation was carried out at 15° C for 72–80 h at 32,000 rpm in a Beckman L565 ultracentrifuge (8 × 12 ml rotor). Fractions were collected from the bottom of the tubes and 100-µl aliquots were weighed for density calculations. Aliquots of the eluates were counted in Aquasol-2 on a Packard Scintillation Counter.

#### RESULTS

Extraction of TBM. Lung mucosal samples were separated by centrifugation into soluble and insoluble/gel phases and analyzed for protein, hexose, and units of BGA. The latter is a more reliable estimate of mucin than neutral hexose as neutral hexose is present in both mucins and in serum-type glycoproteins, whereas serum-type glycoproteins do not contain blood group determinants (29). Blood group activity was specific to mucin in these samples; no glycolipids with BGA were detected in lung mucosal samples. In two (N-1 and BA-1-xv) of the five samples reported in Table 1 the distributions of protein and hexose between the soluble and the gel phases and the specific activities of the blood group hapten (measured either as BGA/mg hexose or BGA/mg of lyophilized powder) were in close agreement with values previously reported for several mucosal samples from a BA patient (5), *i.e.* about two-thirds of the total TBM remained in the gel phase. In contrast, the three CF lung mucosal samples differed markedly from both normal and asthmatic samples: 1) the insoluble phases of the CF samples were precipitates rather than gels; and 2) the ratio of BGA units in the two phases indicated that the CF insoluble phases contained only traces of mucin; essentially all of the TBM in the CF samples was in the soluble phase.

The distribution of mucin between the two phases of a single sample could be estimated by measuring BGA units; however, this procedure could not be used for comparing the mucin content of different samples since the number of completed carbohydrate chains (chains carrying the blood group determinants) vary (32). Thus, the mucin content of the dialyzed, lyophilized mucosal samples was determined on a weight basis after chromatographic resolution of the mucins on Sepharose 4B (Figs. 1 and 2). The mucin content was 40-50% for normal (fraction I; Fig. 1A), 20-30% for asthmatic (fraction 1; Fig. 1B), and 12-14% for CF mucosal samples (fractions Ib and II; Fig. 2).

*Isolation of TBM by molecular sieving.* Typical chromatographic profiles obtained with the gel phase of normal and of asthmatic samples are shown in Figure 1. The  $V_0$  fractions

 Table 1. Analyses of soluble and insoluble fractions of human
 lung mucus

Sample	Wt* (mg)	Protein† (mg)	Hexose‡ (mg)	$BGA \times 10^{-5}$ §
Normal				
N-1 sol	46	9.1	6.6	4.7 (A)
insol	54	6.5	8.9	11
Bronchial asthma				
BA-1-xv sol	33	5.4	3.9	1.7 (AB)
insol	67	17	6.6	3.4
CF				
CF-1 sol	53	17	3.8	22 (A)
insol	47	11	1.7	0.08
CF-2 sol	64	20	8.1	13 (B)
insol	36	15	2.4	0.06
CF-3 sol	55	16	5.3	0 (NS)
insol	45	15	2.3	0

\* Results are expressed per 100 mg of dialyzed, lyophilized lung mucus.

† Assayed by procedure of Lowry et al. (30).

‡ Neutral hexose, assayed by procedure of Dubois et al. (31).

§ Units and types of BGA: A, B, AB, NS (nonsecretor). Assayed by hemagluttination assay of Kabat (32).

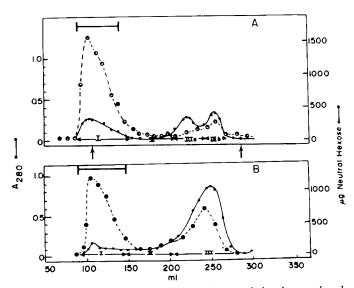


Fig. 1. Chromatography of reduced carboxymethylated normal and asthmatic tracheobronchial mucus on Sepharose 4B (90  $\times$  2 cm). The carboxymethylated gel phase material was dissolved in 7.5 ml of a solution of 1% SDS, 0.15 M ammonium acetate. The solution was layered onto a Sepharose 4B bed and the components were eluted with eluent buffer (0.1% SDS, 0.15 M ammonium acetate, pH 7.0). The eluate was monitored for protein by A<sub>280</sub> ( $\bigcirc$ ) and for neutral hexose ( $\bigcirc$ ) by the phenol sulfuric acid assay (31). The fractions were pooled as indicated, dialyzed to remove the buffer, and assayed for blood group activity ( $\_$ ). The V<sub>o</sub> and salt peak are indicated by *arrows*. *A*, 80 mg of N-1; *B*, 114 mg of BA-1-xv.

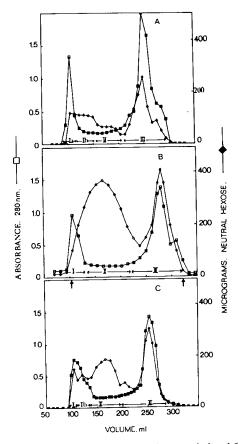


Fig. 2. Chromatography of reduced carboxymethylated CF lung mucosal samples on Sepharose 4B (93  $\times$  2 cm). Chromatography was carried out with 100-mg samples as described in the legend to Figure 1. Absorbance, 280 nm (–––); ug of neutral hexose (– $\bullet$ –). *A*, CF-1, *B*, CF-2; *C*, CF-3.

contained the mucus glycoprotein components: they were high in neutral hexose, low in protein, and contained all of the BGA activity in the starting sample. The V<sub>o</sub> fractions (90–150 ml, Fig. 1) eluted as a broad, skewed peak. When the tail (120–150 ml, Fig. 1A) of the peak was rechromatographed on Sepharose 4B, its hexose and A<sub>280</sub> profiles were similar to that of the V<sub>o</sub> peak observed for fraction I (Fig. 1A). Mucin isolated by method B as described in "Materials and methods" exhibited profiles on Sepharose 4B identical to those of reduced carboxymethylated samples shown in Figure 1 A and B. The profiles of the soluble phases of the samples (not shown) were similar to those of the samples eluted in the V<sub>o</sub> fraction. However the concentration (w/w) of fraction I decreased and that of fraction III increased by 20 to 30%.

In contrast to the normal and asthmatic samples, CF mucins were isolated from the soluble rather than the insoluble phases of the CF samples as the soluble phases were elevated in BGA and hexose (Table 1) and the insoluble pellets of the CF samples contained essentially no mucin. The chromatographic profiles of three CF mucosal samples on Sepharose 4B (Fig. 2) clearly differed from those of normal and asthmatic samples (Fig. 1) and from one another. Hexose was present in all fractions; the sharp  $A_{280}$  peaks in the V<sub>o</sub> of samples CF-1 and CF-2 did not cochromatograph with hexose and disappeared on treatment with DNAase. Since 80% of the BGA in the sample was recovered in fraction Ib of CF-1 (Fig. 2A) and 95% in fraction II of CF-2 (Fig. 2B), fractions Ib and II of CF-3 (Fig. 2C) were considered to be mucin-rich.

Purification of human tracheobronchial mucins. The mucinrich fractions of normal and asthmatic mucus (the  $V_o$  fractions

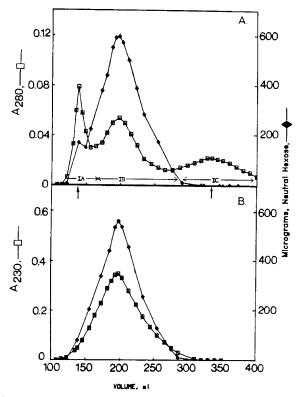


Fig. 3. *A*, chromatography of carboxymethylated normal (N-1) lung mucin on Sepharose 2B. Fraction I (A) was pooled, dialyzed, lyophilized, and dissolved in 10 ml of eluent buffer (0.1% SDS and 0.15 M ammonium acetate, pH 7.0). The solution was layered onto a Sepharose 2B bed (108 × 2 cm) and the components eluted with eluent buffer. The fractions (4.7 ml) were monitored for protein by  $A_{280}$  (-D-) and for neutral hexose (31) (- $\bullet$ -). The V<sub>o</sub> and salt peaks are indicated by *arrows*. B, rechromatography of fraction IB on Sepharose 2B. The IB fraction from *A* was pooled, lyophilized, taken up in 7 ml of water, dialyzed *versus* the eluent buffer, and chromatographed as described above, except that the absorbance was monitored at 230 nm (-D-).

of Sepharose 4B, Fig. 1) were further purified by chromatography on Sepharose 2B and hydroxylapatite. Most of the hexose eluted with a  $K_d$  of 0.3 (fraction Ib, Fig. 3A) on Sepharose 2B. Rechromatography of IB yielded a single, somewhat broad, included peak; no  $V_o$  or smaller included components were observed (Fig. 3B).

The purity of the IB mucin fraction (Fig. 3B) was investigated by electrophoresis. The material was free of smaller protein or glycoprotein components; no bands that stained with Coomassie blue or Schiffs reagent were observed to enter 3% polyacrylamide/0.5% agarose gels even when 500  $\mu$ g of mucin was loaded in a sample buffer that contained 2% ME. The rechromatographed mucin fraction (Fig. 3B) was further purified by chromatography on hydroxylapatite; hexose and A<sub>280</sub> coeluted as sharp peaks with 0.3 M phosphate buffer; a minor component (10% by weight of the applied sample) eluted with 0.5 M phosphate (data not shown).

The LM-gel sample from N-2 (a nonsecretor) was processed without centrifugation into two phases; otherwise TBM was isolated by method B as described in "Materials and methods." Its profiles on Sepharose 4B and 2B were almost identical to those shown for N-1 (Fig. 1A). Increasing the mercaptoethanol concentration 10-fold (to 0.2 M in both the sample and eluent buffer) did not alter the  $K_d$  of TBM (fraction IB) on Sepharose 2B nor did it release detectable lower molecular weight protein fractions.

The CF lung mucins in fractions Ib and II (Fig. 2) were also purified by chromatography on Sepharose 2B and on hydroxylapatite. They eluted as a broad peak on Sepharose 2B with a  $K_d \sim 0.5$ , and often appeared to elute as two or more closely overlapping fractions. The mucin samples did not enter 3–5% polyacrylamide gels; no smaller protein or glycoprotein components were observed. Chromatography of the CF mucins on hydroxylapatite yielded fractions that eluted with 0.15, 0.3, and 0.5 M phosphate buffer. Fifty percent of the mucin in the three CF samples examined was eluted with 0.3 M phosphate and varying amounts were eluted with 0.15 and 0.5 M phosphate buffer.

Characterization of TBM. Analyses of all mucin samples were carried out; only the compositions of purified mucins isolated from blood group A secretors (N-1, BA-1, CF-1) are reported (Table 2). All mucins exhibited a similar spectrum of amino acids: threonine, serine, and proline accounted for 40-51% of the amino acid residues; only low concentrations of hydrophobic and basic amino acids were detected.

The mucins contained N-acetylgalactosamine, N-acetylglucosamine, galactose, fucose, and N-acetylneuraminic acid. Neither mannose nor glucose were detected, with the limit of detection being 1% of the galactose content. Sulfate was variable and present at moderate concentrations (210–780 nmol/mg); these values were within the ranges previously reported for mucins isolated from nonpathological, bronchitic, and CF airways (33).

TBM that had been isolated by method B from healthy (N-2) or asthmatic airways was capable of forming viscoelastic gels and exhibited measurable viscous and elastic moduli at mucin concentrations  $\geq 12.5$  mg/ml after removal of dissociating and reducing agents (34). TBM from CF airways was not viscoelastic even at concentrations of 50 mg/ml. The CF mucosal samples analyzed did, however, exhibit viscoelastic behavior before fractionation (34) presumably because of the high concentrations of DNA found in such samples (35).

Proteinase activity in lung mucus. Proteinase activity was not detected in the asthmatic and normal samples used in this study. Appreciable proteinase activity was detected in the three CF

Table 2. Compositional analyses of human TBM\*

	N-1	BA-1-xv	CF-1	
Amino acid	Residues/1000 residues			
Asp	50.0	43.8	42.8	
Thr	196	222	200	
Ser	129	163	178	
Glu	74.6	68.2	90.0	
Pro	79.4	129	85.7	
Gly	78.4	64.0	109	
Ala	94.1	68.2	86.2	
Cys†	21.5	29.8	8.29	
Val	49.7	37.7	36.2	
Ileu	20.6	22.4	16.8	
Leu	56.0	33.5	39.8	
Tyr	21.7	16.4	23.6	
Phe	28.8	17.4	19.7	
Lys	25.5	25.7	23.2	
His	27.8	26.4	23.4	
Arg	41.1	32.8	17.9	
Carbohydrate	nmol/mg			
GlcNH <sub>2</sub>	535	312	428	
GalNH <sub>2</sub>	443	443	465	
Galactose	884	653	874	
Fucose	715	519	751	
NeuAc	63.2	280	92.5	
Sulfate	550	295	250	
BGA	A	AB	A	

\* TBM fraction eluted from hydroxylapatite column with 0.3 M phosphate buffer.

† Determined as S-carboxymethylcysteine.

mucosal samples examined and it decreased to baseline values on heating. Proteinase levels were generally four times higher in the CF-2 sample than in the CF-1 and CF-3 samples. However, samples collected at different times from the same CF individuals exhibited varying levels of proteinase activity.

In order to determine whether the proteinases detected in CF lung mucosal samples were capable of digesting larger tracheobronchial mucins, purified BA mucin (whose thiol groups had been radiolabeled by carboxyamidomethylation with iodo [1-<sup>14</sup>C] acetamide) was incubated at 37° C for 16 h in freshly thawed sputum samples from CF-1 and CF-2 individuals. After incubation the samples were processed as usual, solubilized in a buffer containing SDS and ME, and chromatographed on Sepharose 4B. No fragmentation of the radiolabeled mucin incubated in saline was observed; it eluted in the  $V_{\circ}$  on Sepharose 4B (Fig. 4A). Approximately 87% of the radioactivity of the BA mucin incubated in the CF lung mucosal samples was lost during dialysis, suggesting that most of the BA mucin was fragmented during the 16-h incubation. The nondialyzable radioactive material contained several fragments (Fig. 4B and C); the major nondialyzable radioactive fraction generated by both CF samples eluted with a K<sub>d</sub> of 0.37 on Sepharose 4B, a value similar to that of CF mucins (Fig. 2).

Biosynthetically labeled CF and non-CF TBM. Tracheal explants from CF and non-CF tissue were radiolabeled with [ $^{35}$ S] sulfate which is a marker for human TBM (10, 36). The secretions from both CF and non-CF tracheal explants yielded two radiolabeled fractions when analyzed by chromatography on Sepharose 4B (Fig. 5). The V<sub>o</sub> fractions, the presumptive mucin-containing fraction, eluted as single peaks on analytical density centrifugation in CsCl/4 M Gu HCl (data not shown) with

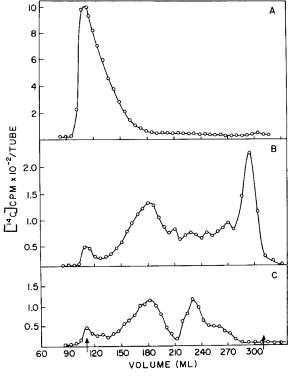


Fig. 4. Chromatography on Sepharose 4B of <sup>14</sup>C-mucin. Radiolabeled BA mucin (7.5 × 10<sup>5</sup> cpm) was incubated in 2 ml of CF mucosal gel samples or saline at 37° C for 16 h. The mixture was then processed (homogenized in 3 vol of isotonic saline, heated in a boiling water bath for 10 min, dialyzed, and lyophilized), solubilized in eluent buffer and chromatographed on Sepharose 4B. Total cpm per tube is plotted on the ordinate. A, <sup>14</sup>C-mucin incubated in saline; B, <sup>14</sup>C-mucin incubated in a CF-1 mucosal sample; C, <sup>14</sup>C-mucin (8.2 × 10<sup>3</sup> cpm) incubated in a CF-2 mucosal sample. The V<sub>o</sub> and salt peaks are indicated by *arrows*.

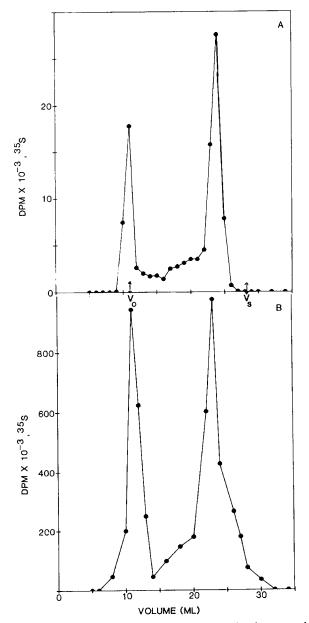


Fig. 5. Chromatography of radiolabeled macromolecules secreted by tracheal explants. Four and 100  $\mu$ Ci of carrier-free H<sub>2</sub><sup>35</sup>SO<sub>4</sub> (New England Nuclear) were added per ml of media, respectively, to the non-CF (*A*) and CF (*B*) explants every 24 h. The secretions were collected as described in "Materials and methods." After dialysis and lyophilization, the radiolabeled secretions were solubilized in 0.5 ml of column buffer (0.1% SDS, 0.15 M ammonium acetate pH 7.0, 0.02 M ME) and chromatographed on Sepharose 4B (30 × 1 cm). Appropriate aliquots were counted and dpm per tube plotted on the ordinate, A, non-CF tracheal explant; *B*, CF tracheal explant.

buoyant densities characteristic of mucins (1.42-1.45 g/ml). The purified radiolabeled mucins were further examined to determine whether the radiolabel had been incorporated into glycosaminoglycans by digesting the sample with bovine testicular hyaluronidase. Hyaluronidase will hydrolyze  $\beta$ 1-4 linkages between N-acetylhexosamine and D-glucuronate residues in hyaluronic acid, chondroitin, and chondroitin 4- and 6-sulfated glycosaminoglycans. When hyaluronidase-digested mucin samples were analyzed by gel chromatography on Sepharose 4B, the radiolabeled mucin eluted in the void volume and less than 5% of the radioactivity eluted in the salt peak, indicating the absence of secreted radiolabeled glycosaminoglycans that cofractionated with the biosynthetically labeled mucin fraction.

The radiolabeled secretions of several nasal polyp epithelial explants from CF and non-CF individuals have also been examined in detail as such material is easier to obtain than human tracheal explants and has characteristics of upper respiratory tract mucosa (10). In 10 CF and non-CF nasal polyp secretions examined to date, a mucin-rich fraction has eluted in the  $V_o$  on Sepharose 4B.

#### DISCUSSION

This study has demonstrated that TBM obtained from the airways of healthy individuals had similar solubility and chromatographic properties to TBM from a patient with bronchial asthma whereas TBM isolated from CF sputum had markedly different properties. In comparison to normal and asthmatic TBM, CF TBMs were 1) present in the soluble rather than the insoluble phase after centrifugation of lung mucus, 2) smaller, as determined by their elution positions on Sepharose 4B; and 3) did not form viscoelastic gels. The profiles that the CF lung mucins exhibited on Sepharose 4B (Fig. 2) suggested the presence in CF lung mucus samples of a spectrum of glycoproteins of different sizes. While some CF TBM eluted in the void volume on Sepharose 4B, most of the mucin in the samples eluted with apparent K<sub>d</sub> values of 0.2-0.4 which are within the range observed for TBM glycopeptide fragments obtained by proteolysis of mucins from patients with asthma, bronchitis and/or bronchorrhea (37-39).

Possible explanations for this variable chromatographic behavior were 1) CF TBM is synthesized as considerably smaller macromolecules than normal or asthmatic TBM, or 2) CF TBM isolated from sputum has already been digested in vivo by proteinases present in CF lung samples as a result of chronic bacterial infection and inflammation (14, 15). The former possibility appears unlikely as TBM synthesized in vitro by tracheal explants from both a non-CF and a CF individual eluted in the  $V_0$  on Sepharose 4B (Fig. 5 A and B) as does mucin secreted by nasal epithelial explants (data not shown). The latter possibility appears more likely as 1) proteinase activity was detected in the three CF LM-gel samples analyzed and was higher in samples enriched in mucin II (CF-2, Fig. 2B) relative to mucin Ib (CF-1, Fig. 2A). 2) CF lung mucus contained proteinases capable of digesting 88% of the purified radiolabeled BA lung mucin to dialyzable components in an 18 h incubation. 3) The K<sub>d</sub> of the major fraction of the radiolabeled nondialyzable mucin fractions produced by incubation in CF lung mucus samples (Fig. 4 B and C) were similar to K<sub>d</sub> values on Sepharose 4B of TBM glycopeptide generated by proteolytic enzymes (37-39). 4) The cysteine contents of the CF lung mucins analyzed were considerably lower than those of normal and asthmatic mucins, and similar to values obtained on analysis of the proteolytically-digested glycopeptide fragments of asthmatic mucin (39, 40). However, considerable variability in the cysteine concentrations of tracheobronchial mucins has been reported (5, 6, 33, 38-43) and cysteine is often not measured quantitively as it requires almost a mg of purified mucin for accurate measurements in duplicate.

This present study has shown that fragmentation can occur *in vitro* when TBM is incubated with CF lung mucus samples, presumably because of the presence of extracellular proteinases in such material (16–18). Although CF sputum contains high levels of bacteria (1, 2, 14) which are known to produce proteinases (15), recent studies (16–18) have indicated that the major proteinase activity in CF sputum is from host defense cells (neutrophils and alveolar macrophages) and that bacterial proteinases are likely to be effectively inhibited by the large amount of bacterial specific immunoglobulins (44). The consequences of proteinase activity in infected lungs is still under intense investigation by several laboratories. In addition to fragmenting TBM, bacterial and cellular proteinases may also increase mucin secretion *in vivo* in humans. Proteinases from *Pseudomonas aerugi* 

nosa increase the secretory index of mucin secretion in rabbit tracheal explants (45). However, any resultant increase in mucin secretion may be offset by proteolysis of TBM as the mucin concentration (weight % of nondialyzable sputum) of CF samples is lower than that of normal and asthmatic samples. Since proteinases in the CF LM-gel samples fragmented radiolabeled tracheobronchial mucin *in vitro* (Fig. 4), proteinases in LM-gel samples from pathological airways may account for the variations in molecular weight, solubility, and chromatographic behavior reported for tracheobronchial mucins (6, 37–43, 46).

This study has also demonstrated that TBM from healthy individuals exhibited physical and chemical properties similar to those of TBM from a patient with bronchial asthma (5). However, our results on TBM from healthy airways are at variance with results on normal TBM reported by some laboratories (11-13). Although the normal TBM isolated in this study was obtained from healthy airways by the procedure of Barton et al. (11) our observations on the chromatographic behavior of TBM are somewhat different than theirs although results cannot eaily be compared as the procedures used for sample preparation and chromatography by each laboratory were quite different. Barton et al. (11) pooled mucosal samples from an unspecified number of volunteers and solubilized their sample by incubation at room temperature for 4 h in 0.2% ME and then overnight at 4° C. After dialysis and lyophilization, 6 M urea and 0.2% ME were required to dissolve the sample in loading buffer. However, these reagents were not included in the eluent buffer. (In our experience both dissociating and reducing agents must be included in both the loading and eluent buffers to prevent aggregation of the components of LM-gel.) Two hexose-rich components eluted on Sepharose 4B (11): a Vo component and an included component that somewhat overlapped the tail of the Vo fraction. While these fractions presumably contain TBM, they were not further purified and no compositional data were reported. In contrast to the study of Barton et al. (11), we were able to purify normal mucins without pooling samples; 10-15 mg of mucin were isolated from LM-gel samples obtained from healthy individuals by hypertonic saline aerosol stimulation.

The TBM samples we have isolated from lung mucus of healthy airways appeared to be both smaller (13) and large (12) than TBM obtained by fiberoptic bronchoscopy. Williams et al. (13) reported that alcian-blue reactive peaks from human tracheobronchial mucus eluted in the Vo and included volumes on Sepharose 2B when chromatography was carried out in the presence of urea and a reducing agent. While the Vo peak was referred to as mucin, no characterization or analyses were reported to substantiate this claim. On the other hand, the mucin sample isolated from bronchial washings of healthy airways by Sachdev et al. (12) has a molecular weight of 69,400 daltons which is markedly smaller than the molecular weight values of 2 to  $7 \times 10^6$  obtained with asthmatic, bronchitic and/or bronchorrhea TBM (38, 42, 46) as well as with TBM isolated from tracheal aspirates of nonpulmonary patients (33, 43). The normal mucin isolated by Sachdev et al. (12) is also smaller than the tracheobronchial glycopeptide fragment (MW = 150,000) obtained on pronase digestion of BA mucin (38). Unless normal tracheobronchial mucin isolated from lavage samples obtained by fiberoptic bronchoscopy is inherently smaller than mucin isolated from expectorated LM-gel samples and from tracheal aspirates, some step in the processing procedure of the lavage fluid likely results in fragmentation of the mucin. It has recently been shown by electron microscopy and molecular sieving studies that sonication of human TBM yields smaller molecules (47). Presumably, the mucin molecules in the lavaged samples (11) could be fragmented either by sonication (which procedure was used by Sachdev et al. (11) to dissolve the lyophilized sample prior to chromatography) or by proteinases released by freezing and thawing of lung cells present in lavaged secretions (48).

In summary, this study demonstrates that TBM isolated from CF lung mucus is not an ideal material for investigating or

comparing the structure of the polypeptide backbone of non-CF and CF mucins. TBM isolated from pooled CF lung mucus samples has, however, provided information on the structure of the oligosaccharide chains (6-8). Insufficient information is, as yet, available to allow a comparison of the non-CF and CF oligosaccharide TBM structures. If marked differences, e.g. a shorter size and decreased length of the CF oligosaccharides, are found, then the possibility of partial degradation of CF TBM oligosaccharide chains by bacterial glycosidases may have to be examined.

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