

Renal tubular absorption of β_2 microglobulin

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Renal tubular absorption of β_2 microglobulin. 125 Iodinated human β_2 microglobulin (β_2m , 5 to 30 mg) was administered to anesthetized rats. Clearance studies showed a low threshold of excretion of injected β_2m and a high T_m of 400 to 600 $\mu g \cdot min^{-1} \cdot kg^{-1}$. A glomerular sieving coefficient of 0.97 was calculated as the slope of the curve: β_2m excretion rate = F (plasma $\beta_2m \times$ glomerular filtration rate) for values above saturation. Electrophoresis analysis of proteinuria in agarose gel and sodium dodecyl sulfate polyacrylamide gel showed that injection of saturating doses of β_2m induced the excretion of proteins of similar size but different charge and that of other proteins of different size. Among the latter, some were excreted transiently in association with β_2m , whereas others had a delayed excretion suggesting existence of a complex mechanism of reabsorption whose steps remain to be elucidated.

Absorption tubulaire rénale de la β_2 microglobuline. De 5 à 30 mg de β_2 microglobuline (β_2m) humaine marquée à l'Iode 125 ont été injectés à des rats anesthésiés. Des études de clairance ont montré un seuil d'excrétion bas et un T_m élevé de 400 à 600 $\mu g \cdot min^{-1} \cdot kg^{-1}$. Un coefficient de tamisage de 0,97 a été mesuré à partir de la pente de la courbe: excretion de $\beta_2m = F$ (concentration plasmatique de $\beta_2m \times$ filtration glomérulaire) pour les points au-dessus de la saturation. L'analyse de la protéinurie par électrophorèse sur gel d'agarose et sur gel de polyacrylamide avec dodecyl-sulfate de sodium a montré que l'injection de doses saturantes de β_2m provoque l'excrétion de protéines de même taille mais de charge différente, ainsi que de protéines de taille différente. Parmi ces dernières, certaines sont excrétées de manière transitoire et en même temps que la β_2m , tandis que d'autres ont une excrétion retardée suggérant l'existence d'un mécanisme de réabsorption complexe dont les étapes restent à étudier.

β_2 microglobulin (β_2m) is a small protein with a molecular weight of 11,800 daltons [1], a Stokes-Einstein radius of 16 Å [2], and a negative charge [1]. It has been identified as a component of class I histocompatibility antigens [3]. When human β_2m is administered to rats by injection, part of the protein binds to a carrier-protein of molecular weight 40 to 50,000 daltons [4]. Due to its size, complexed β_2m does not cross the glomerular filter and it is catabolized in an extrarenal site. Conversely, free β_2m is filtered by the kidney with a glomerular sieving coefficient (GSC) of 0.94 as calculated in studies utilizing the isolated perfused kidney [5]. The unique site of catabolism of free β_2m is the kidney [4].

The kidney is one of the major sites of catabolism of plasma low-molecular weight proteins; hence, it plays a role in regulating the metabolism of substances such as peptide hormones, enzymes, and immunoproteins [6]. However, the precise mechanism of uptake and degradation by the tubules has not yet been investigated completely. Filtered proteins are taken up by micropinocytosis [7, 8], and then degraded in the lysosomes. Owing to their nearly complete reabsorption, filtered proteins are thought to be bound to the brush border prior to absorptive transport into the cell. The questions then arise as to whether fixation to the brush border is specific or not, and what are the parameters of its specificity. Sumpio and Maack [5], and Cojocel et al [9] have shown that charge plays a role in the specificity of reabsorption since cationic cytochrome inhibits transport of cationic lysozyme but not that of anionic β_2m and growth hormone.

We focused our present work on studying renal handling of β_2m as a model of protein tubular reabsorption because its lack of biological activity or toxicity allows one to raise the plasma β_2m concentration far above that necessary for tubular saturation. T_m and GSC were determined; the nature of proteinuria was also studied after saturating injections of β_2m in order to obtain preliminary estimates of the size and charge of endogenous rat proteins which compete with β_2m for tubular reabsorption.

Methods

β_2 microglobulin. Human β_2m was purified from the pooled urine of several renal transplant patients and isolated according to the procedure described by Vincent and Revillard [10]. After preparation, salt-free β_2m was lyophilized and kept at $-40^\circ C$ until the day before the experiment; it was then redissolved in distilled water and kept overnight at $+4^\circ C$. β_2m was labelled with 125 Iodine by the lactoperoxidase method of Thorrell and Johansson [11]. Specific activity was about 0.75 $\mu Ci/\mu g$ protein (9.9×10^5 cpm/ μg). More than 95% of the radioactivity was precipitable by a rabbit antiserum against human β_2m and at least 99% was precipitable in 10% trichloroacetic acid (TCA). After chromatography of a sample of labelled β_2m on Sephadex S200, all the radioactivity was recovered as a single peak in the 12,000-dalton position, thus indicating the homogeneity of the preparation.

Animal experiments. Seven nonfasting male Wistar rats, weighing 180 to 300 g, were given potassium iodide (2 g/liter)

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in their drinking water 24 hr before the experiment to prevent thyroid fixation of nonprecipitable iodine.

Early on the day of the experiment, each rat was weighed and anesthetized with the administration of sodium pentobarbital (Nembutal, Abbott Laboratories Hospital Products Division, North Chicago, Illinois, 5% w/v), 50 mg/kg i.p., then as often as necessary during the experiment. Each animal was placed on a heated board. The bladder was catheterized carefully with a funnel-shaped PE-50 polyethylene tubing for urine collection. After tracheotomy, the carotid artery was catheterized for blood sampling and 0.5 ml heparinized saline was infused via the carotid. Isotonic saline solution was infused into the jugular vein at a constant rate of 4.5 or 6 ml/hr according to body weight; surgical losses were replaced by a single intravenous injection of isotonic saline solution (2.0 ml).

A constant infusion of Polyfructosan (Inutest, Laevosan Gesellschaft, Linz, Austria) in 0.9% saline solution (0.03 g/ml) was administered and subsequent measurements of Polyfructosan clearance were used to measure glomerular filtration rate (GFR). Mean GFR was 10.6 ± 2.2 ml/min/kg ($N = 44$) in our experimental conditions and was not modified significantly during the course of the experiment, even after β_2 m injection. After a 60-min equilibration period, a control collection of urine was obtained for 20 min, and a blood sample was drawn at the mid-point of the urine collection period.

Labelled β_2 m (20 to 50 μ g, 10^6 cpm/ μ g) and cold β_2 m (in doses ranging from 5,000 to 30,000 μ g) were then injected through a lateral connection into the jugular vein at zero time, followed immediately by 0.1 ml saline. Blood samples were taken at 1, 3, and 10 min after β_2 m injection, then every 10 min, and liquid losses were replaced by saline. Urine was collected during consecutive 10-min periods, starting 5 min after β_2 m injection (urine collected 5 min after injection was not used for clearances determination). Urine pH was about 6, thus insuring no significant urinary degradation of β_2 m.

Analytical methods. Polyfructosan was determined by a fluorimetric method [12] using a dimedon reagent. Total and TCA precipitable radioactivity in blood and urine was determined on an autogamma scintillation spectrometer (Packard Instruments, Downers Grove, Illinois) as previously described [4].

Free versus total β_2 m ratio was determined by chromatography of 100 μ l samples as described previously [4].

Urine electrophoresis was performed on agarose gel following the procedure of Jeppsson, Laurell, and Franzen [13] and on polyacrylamide gel in the presence of sodium dodecyl sulfate (SDS) as described by Laemmli [14]. Autoradiography of desiccated agarose plates was performed at -70°C with Kodak X-O MATR films and an intensifying screen Cronx lighting plus (Du Pont) [15].

Calculation of β_2 m clearance. The lag time between plasma and urine, as well as the nonsymmetric distribution of β_2 m plasma content during each period of time had to be taken into account for β_2 m clearance determination. Since the lag time between filtration and the appearance of urine in the collection tubing depends on urine flow rate, six rats in the same experimental conditions as described above were injected with 0.1 ml lissamine green. The lag time between injection and dye appearance in the collection tubing was measured; lag time between injection and coloration of the kidney was negligible. The results are shown in Figure 1. This plot was used to de-

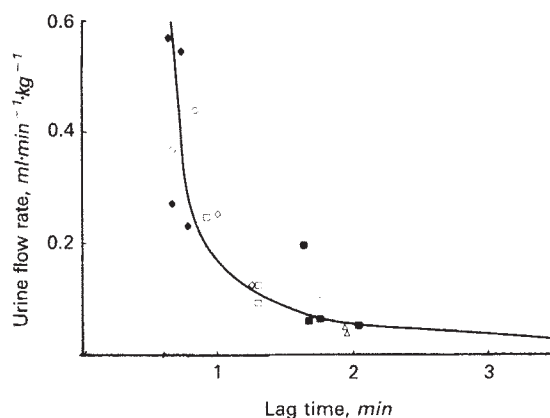


Fig. 1. Lag time between injection and urinary output of lissamine green. Each symbol represents experimental points from one animal.

termine lag time for each experimental period after β_2 m injection and thus to shift the limits of the urine period on the corresponding plasma kinetic curve. Mean plasma β_2 m concentration of each period was calculated from the area of the plasma kinetic curve divided by the time. Filtered load was calculated as plasma free β_2 m \times GFR \times GSC, with plasma free β_2 m calculated as described above.

Results

β_2 m clearance data: glomerular sieving coefficient T_m and kinetics of degradation. Injection of human β_2 m administered to rats in amounts up to 30 mg per animal allowed plasma content of total β_2 m to rise up to about 600 μ g/ml, that is, about 150 times the normal value of rat plasma β_2 m [16]. Chromatography of each plasma sample and some urine samples showed that part of the injected β_2 m bound gradually to the plasma carrier protein, whereas complexed β_2 m was not found in urine. Free β_2 m was rapidly eliminated from plasma with a mean metabolic half-life of $30 \text{ min} \pm 7$ ($N = 7$) which was very similar to that obtained when small doses of β_2 m were injected [4].

The relationship between plasma free β_2 m \times GFR and urinary free β_2 m excretion, is presented in Figure 2. β_2 m appeared in urine as soon as the filtered load was increased by twofold. Then β_2 m excretion increased linearly as filtered load increased; the slope of 0.97 represents the glomerular sieving coefficient for β_2 m.

The titration curve could be drawn from the above values, taking 0.97 as GSC value for β_2 m (Fig. 3). That curve shows that reabsorption was saturated above $1,200 \mu\text{g} \cdot \text{min}^{-1} \cdot \text{kg}^{-1}$ of filtered β_2 m. It gives a T_m value of about 400 to $600 \mu\text{g} \cdot \text{min}^{-1} \cdot \text{kg}^{-1}$.

Figure 4 depicts the kinetics of events after β_2 m injection in four rats with the highest injected doses: Filtered load of free β_2 m decreased sharply after injection whereas reabsorbed amounts were grossly constant in the range of time considered. Degradation was estimated by the appearance of nonproteic radioactivity in urine. Urinary excretion of nonTCA-precipitable activity paralleled plasma content and clearance values averaged GFR (not shown), showing that the nonproteic degradation products of β_2 m are freely filtered. Urinary output of nonTCA-precipitable activity increased during the

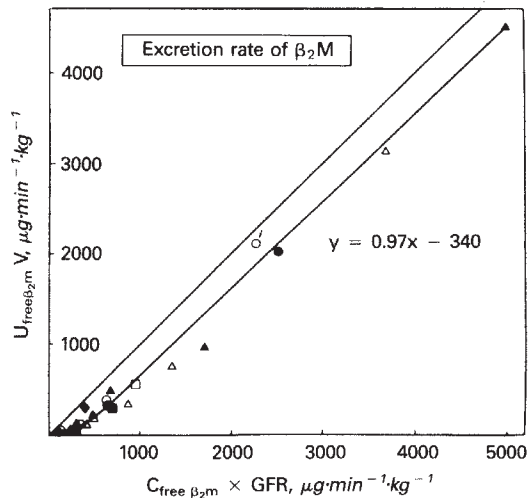


Fig. 2. Urinary excretion rate of β_2m . Each symbol represents experimental points from one animal. Experimental points below $300 \mu g \cdot min^{-1} \cdot kg^{-1}$ were excluded for calculation of the regression curve.

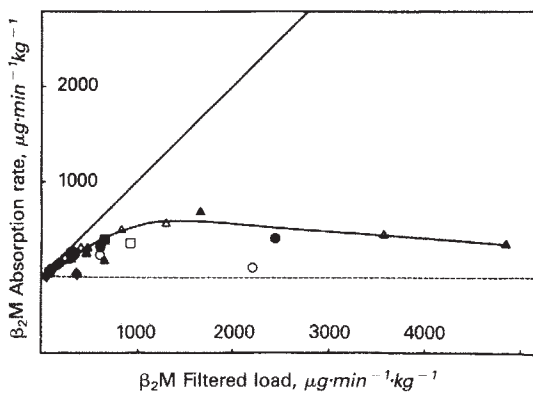


Fig. 3. Titration curve of free β_2m . Reabsorption rate: filtered load-urinary excretion rate. Filtered load: plasma $\beta_2m \times GFR \times GSC$. The symbols are the same as in Figure 2. The lower line was drawn from the data for one rat (\blacktriangle).

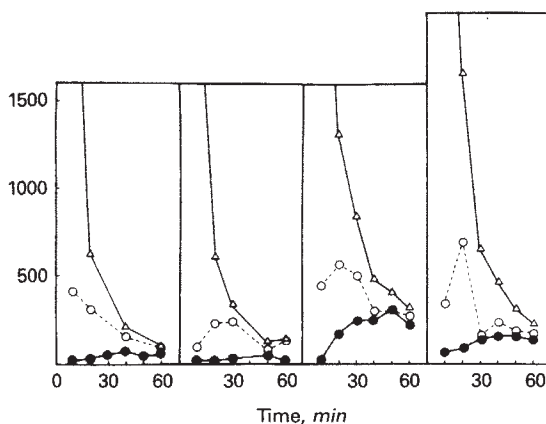


Fig. 4. Kinetics of renal handling of injected β_2m (highest doses): β_2m filtered load (Δ - Δ), plasma free $\beta_2m \times GFR \times GSC$; β_2m absorption rate (\circ - \circ), filtered load-excretion rate; degraded β_2m (\bullet - \bullet), equivalent amount of β_2m estimated from urinary excretion rate of nonproteinic radioactivity. All forms of β_2m are expressed as $\mu g \cdot min^{-1} \cdot kg^{-1}$.

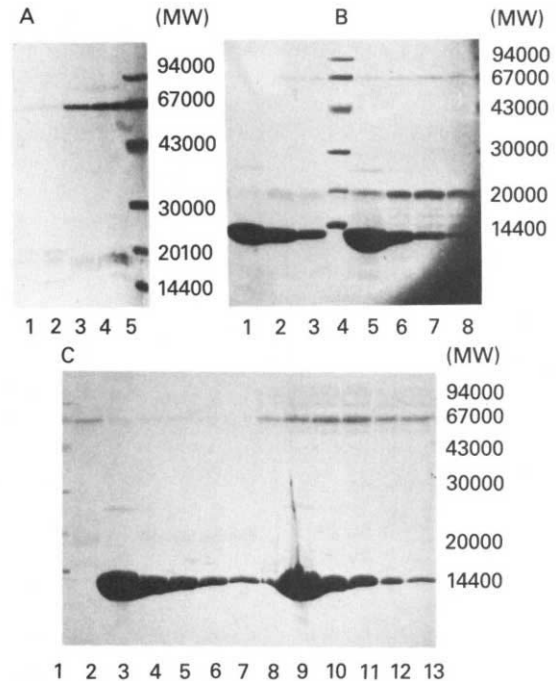


Fig. 5. Gel electrophoresis on polyacrylamide + SDS. Calibration mixture (A5, B4, C1) was obtained from a low molecular weight calibration kit (Pharmacia Diagnostics, Uppsala, Sweden). Molecular weights are indicated on the right side of each photograph. **A** Electrophoresis on 15% gel of control urine from two rats (1, 2, and 3, 4, respectively) with $15 \mu l$ (1,3) and $100 \mu l$ (2,4) inclusion. **B** Electrophoresis on 20% gel of urine from two rats (1-3 and 5-7, respectively) after β_2m injection. Inclusions of $15 \mu l$ were ordered from the left to the right in the chronologic order, always beginning by the first period following β_2m injection. **C** Electrophoresis on 20% gel of urine from two rats (2-7 and 8-13, respectively). Inclusion volume was $30 \mu l$ for every inclusion of the first rat, $15 \mu l$ for every inclusion of the second rat. Inclusions were ordered from the left to the right in chronologic order, always beginning by a control period (2 and 8), then the first period following injection (3 and 9), and so on. The lower band of 8 (control) is an artefact due to side diffusion of the very intense band 9.

course of the experiment and, after 1 hr, was close to the reabsorbed amounts. There was a delay between the bulk of reabsorption of filtered β_2m and the appearance of degradation products.

Analysis of proteinuria after β_2m injection. SDS polyacrylamide gel electrophoresis of urine before and after β_2m injection are shown in Figure 5. Control samples of urine from rats just before an injection was administered (Fig. 5 A and C) were found to contain mainly albumin (upper band) and a number of thin bands between the 12,000- and the 20,000-dalton positions. One rat (Fig. 5C, right) had a glomerular proteinuria during the control period, that was slightly increased by β_2m injection and remained after 1 hr. After β_2m injection (Fig. 5 B and C) urine contained so much β_2m that the electrophoresis plate was overloaded in the 12,000-dalton zone, thus preventing identification of additional proteins in that zone. β_2m band intensity decreased during the course of the experiment but never returned to the control pattern within the range of time chosen. The albumin band was not inflected by the β_2m injection. Besides, the excretion pattern of smaller proteins was markedly modified by the β_2m injection: One band appeared in

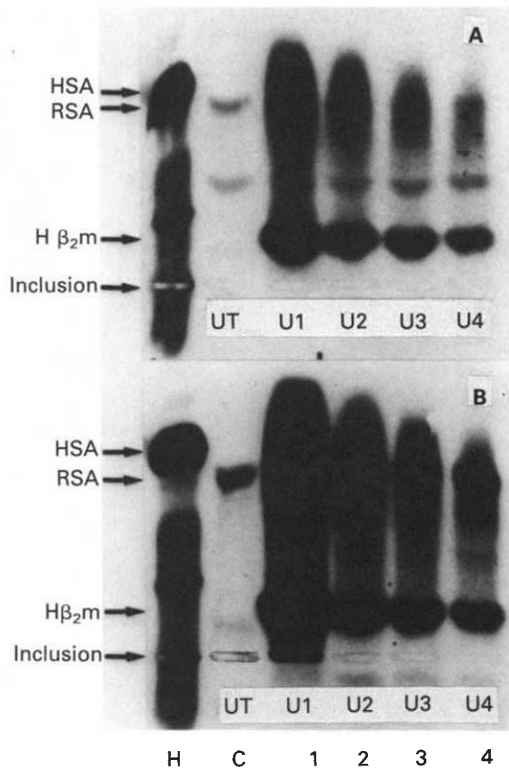


Fig. 6. Electrophoresis on agarose gel at pH 8.6 of urine from two rats (*A and B*, respectively). From the left to the right, abbreviations are: H, human serum; C, control rat urine; 1, 2, 3, 4, the first four urine samples from the same rat after β_2m injection; HSA, human serum albumin; RSA, rat serum albumin; H β_2m , human β_2 microglobulin. Each sample was collected during 10 min.

each rat in a 27,000-dalton zone and decreased in parallel with β_2m . In the 17,000 to 20,000-dalton zone, some bands appeared or were enforced after the β_2m injection, but their intensity did not decrease. Some of these bands seemed to appear after a delay of one period or to increase during the second or third period.

The observed changes in the proteinuria pattern after β_2m injection were quantitatively but not qualitatively different from one animal to another.

Figure 6 shows agarose gel electrophoresis of rat urine before and after the β_2m injection. The main band was β_2m which decreased during the course of time. The bands detectable in control urine, albumin (Fig. 6 *A and B*), and a component of lower mobility (Fig. 6*A*), were also detectable after β_2m injection and did not decrease during the course of time. Two additional bulks of proteins appeared in the albumin zone and the region of middle mobility: These bulks were very intense and represent several intricately bands. Their intensity decreased in the same way as did the β_2m band. A very pale band appeared after β_2m injection on the anodic migration side of the plate. Autoradiography of the agarose plates (not shown) showed that only the band in β_2m position was radioactive. In addition some precipitated material in the inclusion well was also radioactive in the case of Figure 6*B*. From agarose gel electrophoresis of serially diluted (1:2 to 1:640) urine samples, it was estimated that the amount of β_2m was at least 30 times

greater than that of the bulk of proteins migrating in the albumin position.

Discussion

Renal clearance of β_2m . Our results show a low threshold of β_2m urinary excretion, which is a common feature for small proteins [6]. From determinations of urine and serum β_2m in patients with normal or altered GFR, Wibell [17] suggested a β_2m plasma threshold of 4.5 $\mu\text{g/ml}$ (normal serum levels in humans are 1.7 $\mu\text{g/ml}$), above which urinary excretion of β_2m was increased. However, it is, to our knowledge, not possible to estimate the extent of tubular alterations in patients with decreased GFR.

Due to the large amount of purified material needed for each experiment, little is known about the T_m of proteins in the kidney tubules. The generally low threshold of excretion of filtered proteins, added to the increasing-type relationship between filtered load and excretion, led to suppose that their reabsorption was saturated for filtered loads close to normal values [18]. Further studies [6, 19] on lysozyme and albumin showed that when filtered load increases above the threshold of excretion, there occurs a "splay" region [5] where part of the filtered load is increasingly reabsorbed whereas the remaining is excreted. Plasma protein level may have to be very largely increased before the real T_m is reached, since T_m for lysozyme is 50 to 100 times larger than the normal filtered load. The absolute value of T_m by itself may be very different from a protein species to another. For instance Sumpio and Maack [5] showed a T_m of 800 $\mu\text{g/min}$ per rat kidney for lysozyme and in the same conditions, a T_m of 6.5 $\mu\text{g/min}$ per rat kidney for cytochrome C. The rate of reabsorption depends on the charge and configuration of the molecule [20]. These findings support the view that protein tubular absorption is a complex phenomenon of low affinity, high capacity, and low specificity.

Our results provide further evidence for this conception: There is a "splay" region beyond the threshold of excretion where β_2m reabsorption is not saturated. The T_m value is not very high (four times the normal filtered load) when compared to the lysozyme T_m value. However, one should keep in mind that elevated plasma β_2m levels are observed in many common pathologic situations such as inflammatory disorders [21, 22], viral infections [23], and hepatitis [24]; in such cases plasma β_2m level is generally included in the splay region. Therefore, increased urinary excretion which may be encountered in such situations does not provide an estimate of the increased biosynthesis of β_2m . The glomerular sieving coefficient was drawn from the curve $UV = f(P \times \text{GFR})$ considering only filtered loads far beyond T_m . The calculation of T_m is usually drawn from clearance measurements performed in steady-state conditions by using continuous intravenous infusion. Such a method could not be used because it would require much greater amounts of highly purified β_2m than those available for the present study. Therefore, the single injection technique was chosen to achieve very high plasma levels at least during a short period. Errors in GFR determination by this technique depend mainly on the nonequilibrium of the test substance among fluid compartments [25]. In the present study no assumption on the diffusion of β_2m was required because for each period the plasma concentrations of β_2m were graphically derived from the measured plasma levels. Kinetics of re-

absorption (Fig. 4) would suggest that reabsorption is not saturated by the initially high filtered β_2 m load, but that maximum of the transport capacity is achieved after a delay of sometimes more than 5 min. If β_2 m itself could trigger an increase in its reabsorption, then the slope in Figure 2, that is, the GSC, would be slightly overestimated. Calculation of GSC was therefore performed with one rat in which saturation was with certainty achieved 5 min after injection (Fig. 4, left). The calculated GSC value was of 0.91 in this case. Whatever are the potential pitfalls of the single injection technique, the values obtained (0.97 for the seven rats or 0.91 for the latter) are close to those (0.94) obtained by Sumpio and Maack [5] in the isolated rat kidney in presence of iodoacetate, an inhibitor of tubular reabsorption. The similarity of the values provides a certain confidence to them and indicates that free β_2 m is almost not hindered by the glomerules. This may be explained by the small size of the molecule (16 Å) since Brenner, Hostetter, and Humes [26] have shown that sieving of macromolecules appears beyond a 20-Å radius. The negative charge of β_2 m does not seem to interfere very much, although negatively charged small molecules are significantly hindered by the glomerulus.

The kinetics data agree with previous findings in humans [28] and rat [29] showing that there is a lag time between β_2 m reabsorption and appearance of the degradation products. Autoradiography of urine electrophoresis showed that after β_2 m injection, the migration properties of β_2 m excreted in urine remained identical to that of intact β_2 m. This suggests that brushborder proteases which were shown to degrade small peptidic hormones [30] do not affect the β_2 m molecules that are not reabsorbed.

Competition experiments. Electrophoresis of urine on SDS polyacrylamide gel shows that no competition occurs between β_2 m and albumin, although the small filtered fraction of albumin is extensively reabsorbed by the kidney [19]. Two groups of small proteins appeared in urine after β_2 m injection. Some appeared and disappeared at the same time as β_2 m whereas some appeared after a delay and were still excreted after 1 hr. This result could be explained by the hypothesis that the first group may share the same site of fixation or transport on the cell wall and the same degradation pathway whereas the second group quantitatively more important only shared the same degradation pathway.

Electrophoresis in agarose gel showed a very intense bulk of protein in addition to the β_2 m band. Because in SDS polyacrylamide gel electrophoresis no such important band of protein was seen outside the β_2 m zone, the bulk of protein in agarose gel must be proteins of the same molecular weight as β_2 m but of a different charge. Autoradiography demonstrated that these proteins are not degradation products of β_2 m. Thus, β_2 m must be reabsorbed in competition with several proteins of similar molecular weight, which represents a major part of the competition between proteins. Agarose gel electrophoresis pointed out a possible competition between proteins of a different charge at a very high dose of β_2 m, although at a more physiological concentration no competition could be seen between cationic and anionic proteins [5, 9, 31]. In conclusion, the survey of proteinuria after injection of a single high dose of β_2 m allowed the discrimination between several groups of proteins, suggesting the existence of a complex mechanism of reabsorption whose steps remain to be elucidated.

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