Antibodies of all species are present in several classes of immunoglobulin. The serum level of each immunoglobulin class is determined by its rate of synthesis and catabolism. Thus catabolic properties are important biologic features of antibody because of the importance of catabolism in regulating serum level and in determining the survival of an antibody molecule.

Studies in man (1, 2), rabbit (3), and guinea pig (4) have shown that the 18S γ1-macroglobulins (IgM) are catabolized much more rapidly than 7S γ2-globulins (IgG). Thus, immunoglobulin classes may differ in their rates of turnover. Differences in the catabolic control of immunoglobulins are also discernible. The serum level of 7S γ2-globulin (IgG) helps to determine the catabolic rate; the higher the serum level, the greater the catabolic rate (5, 6). 18S γ1-macroglobulin (IgM) catabolism, however, appears to be unrelated to the serum γ1-macroglobulin level (2). Thus the catabolic properties of all classes of immunoglobulins need to be characterized and the catabolic interrelationships between immunoglobulin classes further defined.

Four major classes of immunoglobulins, the 7S γ2-globulins, the 7S γ1-globulins, the γ1A (β2A, IgA)-globulins and γ1M (IgM)-macroglobulins have been identified in recent studies in mice (7). These components are seen on immunoelectrophoresis of hyperimmune serum in Fig. 1. In addition, two subclasses of 7S γ2-globulin have been identified and tentatively designated as γ2a- and γ2b-globulins (8). These findings provided an opportunity to investigate further the catabolic relationship between different classes of immunoglobulin. In no species has the metabolism of five classes of immunoglobulin been compared. The present investigation was undertaken to compare the rate of catabolism of γ2a-, γ2b-, and 7S γ1-globulins, γ1A (β2A)-globulins, and γ1-macroglobulins in mice. Investigations were carried out in mice with low, normal, or high levels of all immunoglobulins, as well as in mice with selective immunoglobulin increases produced by plasma cell tumors.
Materials and Methods

Mice.—White Swiss-Webster (NIH-WS) mice were obtained from the general purpose supply colony of the National Institutes of Health, Bethesda. Germfree and conventional low pathogen white Swiss-Webster mice were classified on the basis of potential environmental exposure to bacteria (9), and have subnormal serum immunoglobulin levels (Fig. 1). A group of six NIH-WS mice were immunized with a total of 0.6 mg of alum-precipitated hemocyanin given in six doses.

C3H/HeN or BALB/c mice harboring nine individual plasma cell tumors were used in these studies. The tumors produced $\gamma_{2a}$-myeloma proteins (5563, Adj. PC-5), $\gamma_{2b}$-myeloma proteins (MPC-11, MPC-31, MPC-37), or $\gamma_{1A}$-myeloma proteins (SPC-1, MPC-1, MPC-36, MPC-40). Tumor weights were 3 to 7 gm at the time turnover studies were performed.

$^{131}I$-Labeled Proteins.—Nine individual myeloma proteins were isolated from the sera of tumor-bearing mice by block electrophoresis and column chromatography. Each preparation...
was tested by double diffusion in agar against potent rabbit antisera, both specific and polya-
ven, to determine the type of protein present and its purity. The nine myeloma proteins
included the following: \(\gamma_2\alpha\)- (5563 and Adj. PC-5); \(\gamma_2\mu\)- (MPC-11, MPC-31, and MPC-37); \(\gamma_7\alpha\)- (MPC-25); and \(\gamma_1\alpha\)- (MPC-1, MPC-40, and SPC-1). The preparations were separately trace labeled (< 1 mole I per mole of protein) with \(^{131}\text{I}\) by the iodine monochloride method of McFarlane (10). In addition a preparation of human \(\gamma_1\)-macroglobulin from a patient with Waldenström’s macroglobulinemia was also trace labeled with \(^{131}\text{I}\).

Experimental Protocols.—All animals were housed in plastic cages containing wood shavings
that were frequently changed or in cages with wire floors which allowed urine and fecal
droppings to pass through to a tray below. All animals were given drinking water containing
0.45 per cent NaCl and 0.01 per cent KI. Mice were injected intraperitoneally with from
0.2 to 1.0 \(\mu\text{g}\) of \(^{131}\text{I}\)-labeled protein in 0.1 ml 0.85 per cent NaCl. The whole body radioactivity
of the injected mice was measured in a gamma ray bulk spectrometer (Sharpe Laboratories,
La Jolla, California) for 1 minute. The radioactivity in each mouse was determined within 30
minutes of injection, then once every 1 to 2 days. A radioactive standard for each labeled
protein was prepared by injecting 1 dose of the given protein into 20 ml of 0.85 per cent NaCl
in a 30 ml plastic bottle. This standard was counted daily with the experimental animals and
the results used for correction of physical decay. The fractional rate of catabolism (\(T_\lambda\)) of
each of the \(^{131}\text{I}\) proteins in each animal was determined from the graphic plots of the decay
curves corrected for physical decay of the isotope. The percent \(^{131}\text{I}\) protein in the body de-
graded per day was obtained by dividing the \(T_\lambda\) into 0.593.

\(\gamma_1\)-Macroglobulin Catabolism.—\(\gamma_1\)-Macroglobulin catabolism was followed by determining
the serum hemolysin titers of mice that were injected with serum or a serum macroglobulin
fraction containing sheep cell hemolysin activity. \(\gamma_1\)-Macroglobulin hemolysin was obtained
from NIH-WS and C57B1/6JN mice immunized by intraperitoneal injections of 0.2 cc of a
50 per cent suspension of washed sheep cells every 2 to 3 days for 5 weeks. The mice were
bled from the retroorbital venous sinus at 3, 4, and 5 weeks after the start of immunization.
Each bleeding was done 3 days following the preceding sheep cell injection. The serum ob-
tained from each animal was tested for hemolysin titer by the microtiter method described below,
before and after incubation with an equal volume of 0.1 \text{M} 2-mercaptoethanol. The hemolysin
activity of the 3rd week bleedings of the NIH-WS mice and the 3, 4, and 5 week bleedings of
the C57B1/6J mice was completely destroyed by 2-mercaptoethanol indicating that all of the
antibody present was most likely 18S macroglobulin. The hemolysin titers of the sera from
the 4th and 5th week bleedings of the NIH-WS mice were only partially reduced follow-
ing mercaptoethanol treatment, indicating that both 7S and 18S antibodies were present. Ser-
ums containing high titers (1:1280 or greater) of mercaptoethanol sensitive antibody were
pooled and aliquots injected intravenously into NIH-WS mice. In addition, 2.0 ml of high titered
mouse hemolysin serum pooled from the 4th and 5th week bleedings of the NIH-WS mice
were passed through a 2.2 X 78 cm sephadex G-200 column. The eluate in the ascending por-
tion of the first protein peak was pooled to obtain the macroglobulin fraction. The pool was
concentrated by ultrafiltration until a hemolysin titer of 1:2500 was obtained. The he-
molysin activity of this preparation was completely removed by mercaptoethanol treatment.
Aliquots (0.2 ml) were injected intravenously into NIH-WS mice and the titers in each mouse
followed by the microhemolysin method.

Microhemolysin Technique.—The hemolytic activity of complement (C') or hemolysin is
related to the concentration of sensitized erythrocytes used in the test system (11). In order
to follow passively transferred 18S hemolysin activity in mice for more than 1 day, it was
necessary to use a system with very small numbers of erythrocytes. The hemolysin method
described previously (12) was modified by the use of the microtiter technique (13) for making
serial dilutions and the erythrocyte counting method of Sterzl and Kostka (14) for estimation
of the serum dilution giving 50 per cent hemolysis. 0.025 ml of a 1:100 dilution of normal mouse serum in veronal buffered saline (12) was added to each well of a plastic plate. The sera to be tested were then diluted with wire loops, 1 drop of a 0.01 per cent sheep cell suspension added to each well, followed by 1 drop of a 1:30 dilution of normal guinea pig serum (C'). The plates were incubated for 1 hour at 37°C and the reaction then stopped by the addition of 1 drop of isotonic citrate saline solution (1 part 0.075 M aqueous sodium citrate, 4 parts 0.15 M NaCl) (15).

The number of cells remaining in wells spanning 50 per cent lysis were counted microscopically by adding a sample from the appropriate wells to an erythrocyte counting chamber. The dilution of a given serum giving 50 per cent lysis was determined from the intersection of the 50 per cent lysis point by the linear graph of the cell counts of the given wells plotted against the appropriate serum dilutions.

The 50 per cent lysis dilutions from the serial bleedings of a given animal were then plotted against the time following antisera transfer. The rate of catabolism (T½) of the passively transferred γ1-macroglobulin antibody was estimated from the graphic plot of the 50 per cent lysis values.

### TABLE I

#### Turnover of Several Classes of Immunoglobulins

<table>
<thead>
<tr>
<th>Mice</th>
<th>Mean serum immunoglobulin levels</th>
<th>Half-lives of individual 125I immunoglobulins*</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>γ2 (a+b)</td>
<td>γ2</td>
</tr>
<tr>
<td>Normal (NH-WS)</td>
<td>4.0</td>
<td>2.5</td>
</tr>
<tr>
<td>Germfree (GF)</td>
<td>2.0</td>
<td>0.2</td>
</tr>
<tr>
<td>Low-pathogen (CVN-LP)</td>
<td>0.5</td>
<td>0.3</td>
</tr>
<tr>
<td>Hyperimmune (NH-H1)</td>
<td>15.5</td>
<td>19.5</td>
</tr>
</tbody>
</table>

* Mean value of observation in 4 to 6 mice.

### RESULTS

**Normal Mice.**—The catabolism of nine 125I-labeled mouse myeloma proteins, representing the 7S γ2a, 7S γ2b, 7S γ1A, and γ2A (IgA)-globulins, was measured in NIH-WS mice. The mean half-life of each myeloma protein is given in Table I. Representative decay curves for each immunoglobulin subgroup are shown in Fig. 2.

**7S γ-globulins:** The 7S γ2a, 7S γ2b, and 7S γ1-globulins differed in their catabolic rates. The two 7S γ2a-globulins had the longest mean half-lives (4.8 and
5.4 days) of the 5 classes of mouse immunoglobulins (Table I). The three \(\gamma_m\)-globulins had more rapid rates with half-lives of 2.5, 2.6, and 3.0 days. The half-life for the single 7S \(\gamma_1\)-myeloma protein available for study was 4.0 days.

These observations indicate that 7S \(\gamma\)-globulins are not uniform in their catabolic characteristics. The 7S \(\gamma_{2m}\)-globulins were catabolized at the rate of about 14 per cent per day. The 7S \(\gamma_2\)-globulins are catabolized at rates of 23 to 28 per cent each day; i.e., almost twice as fast as \(\gamma_{2m}\)-globulins.

\(\gamma_{1A}\) (\(\beta_{2A}, IgA\))-globulins: Three \(\gamma_{1A}\)-myeloma proteins had half-lives of 1.0, 1.0, and 1.3 days in normal mice. It is possible that the catabolic rate of the \(\gamma_{1A}\)-globulin is actually more rapid if the observed half-time reflects accumulation in the body of \(T^{IM}\), freed from catabolized protein, as well as non-catabolized labeled protein.
\(\gamma_1\)-Macroglobulin (IgM): Metabolism was measured by following the serum hemolysin titer after injection of mouse anti-sheep erythrocyte hemolysin antibodies in whole serum or in macroglobulin fractions prepared by sephadex G-200 filtration of immune serum. Data for a number of individual mice are shown in Fig. 3. Two patterns of decay in serum antibody activity were observed in both normal and low pathogen mice. In some mice a rapid rate of antibody removal was detected with a half-time of 0.2 days. In other mice the initial portion of the decay curve had a half-time of 0.6 days, but on the 2nd day the rate of antibody removal was more rapid and half-time of 0.2 days was observed.

Studies with \(^{131}\)I-labeled human 18S \(\gamma_1\)-macroglobulin revealed a half-time of 1.3 days for this protein. The half-time is based on whole body radioactivity measurements and calculation of the half-time from daily measurements of the per cent of the injected dose remaining in the whole body. The difference between this half-time (1.3 days) and that observed for the hemolysin antibody (0.2 to 0.6 days) may be due to delayed excretion of \(^{131}\)I and retention in the body fluids after catabolism of the protein or may be due to species differences in macroglobulin catabolism.

**Germfree and Low Pathogen Mice.**—Mice raised in a germfree or low patho-
gen environment have low levels of all mouse immunoglobulin components (Fig. 1, Table I).

The catabolism of each of the 7S globulins was prolonged in the low pathogen and germfree mice as shown in Table I. The comparative data for low pathogen mice are illustrated in Fig. 4 where the half-time of 7S $\gamma_2$-globulin was found to be 10 days; 7S $\gamma_6$-globulin was 5.5 days; and 7S $\gamma_1$-globulin was 14 days. In these mice, as in normal mice, the $\gamma_6$-globulins were catabolized more rapidly than $\gamma_2$-globulins.

![Graph showing catabolism of immunoglobulins](image)

**Fig. 4.** Catabolism of immunoglobulins, $\gamma_6$, $\gamma_2$, 7S$\gamma_1$, and $\gamma_1$ ($\beta_2A$) in low pathogen (CVN-LP) mice. Catabolic decay curves of the radioiodine-labeled immunoglobulins in mice with subnormal numbers of intestinal microorganisms (CVN-LP) are shown. The T $\frac{1}{2}$ values for the catabolism of $\gamma_2$, $\gamma_2$, 7S$\gamma_1$, and $\beta_2A$-globulins in normal mice (N) are given in brackets below the data obtained in this experiment with low pathogen mice.

The rates of catabolism of the $\gamma_1$-myeloma proteins were the same in the mice with low immunoglobulin levels as in normal mice. The half-times for $\gamma_1$-macroglobulin antibody (0.2 to 0.6 days) and for $\Gamma^H$ human macroglobulin (1.3 days) also were the same in low pathogen (CVN) as in normal mice.

**Hyperimmune Mice.**—In hyperimmunized mice the serum levels of all the immunoglobulin components were increased (Table I). The half-times for the 7S $\gamma_2$-globulins (2.2 and 2.6 days), the 7S $\gamma_6$-globulins (1.1, 1.4, and 2.1 days), and the 7S $\gamma_1$-globulin (1.9 days) were shorter when injected into hyperimmunized mice than in normal mice. The rate of catabolism of $\gamma_1$ (IgA)-globulin was not altered in the hyperimmunized mice.

**Effect of Specific Immunoglobulin Increases.**—Plasma cell tumors synthesize only one class of immunoglobulin, and mice bearing plasma cell tumors have
<table>
<thead>
<tr>
<th>Recipient of ¹³¹I-labeled proteins</th>
<th>Mean serum immunoglobulin levels</th>
<th>Half-life of individual IgM immunoglobulins (days)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>131I γ₇₅⁺⁺⁺</td>
<td>131I γ₇₁</td>
</tr>
<tr>
<td>Normal mice:</td>
<td>(mg/ml)</td>
<td>(mg/ml)</td>
</tr>
<tr>
<td>NIH-WS</td>
<td>4.0</td>
<td>2.5</td>
</tr>
<tr>
<td>Tumor bearing mice:</td>
<td></td>
<td></td>
</tr>
<tr>
<td>7Sγ₇₅⁺⁺⁺ S563</td>
<td>44.0</td>
<td>0.6</td>
</tr>
<tr>
<td>Adl. PC-5</td>
<td>17.7</td>
<td>1.1</td>
</tr>
<tr>
<td>7Sγ₇₁</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MPC-11</td>
<td>64.7</td>
<td>1.5</td>
</tr>
<tr>
<td>MPC-31</td>
<td>15.3</td>
<td>1.2</td>
</tr>
<tr>
<td>MPC-37</td>
<td>23.2</td>
<td>1.4</td>
</tr>
<tr>
<td>γ₇₁ (IgA)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MPC-1</td>
<td>1.0</td>
<td>1.3</td>
</tr>
<tr>
<td>MPC-40</td>
<td>1.2</td>
<td>2.3</td>
</tr>
<tr>
<td>MPC-36</td>
<td>0.5</td>
<td>1.1</td>
</tr>
<tr>
<td>SPC-1</td>
<td>1.6</td>
<td>1.6</td>
</tr>
</tbody>
</table>
a selective increase in serum concentration of the myeloma protein formed by the tumor (Table II). Because of this selective increase, the effects of large amounts of one protein on the catabolism of this protein and other immunoglobulins may be studied (Table II). Mice bearing a \( \gamma_{2a} \)-globulin-producing plasma cell tumor (Adj.PC-5) had a high level of 7S \( \gamma_{2a} \)-globulin (Table II) and showed very rapid catabolism of the \( \gamma_{2a} \)-myeloma protein (T \( \frac{1}{2} \) = 2.5 days, in contrast to the normal value of 5.4 days) (Fig. 5). Not only was the catabolism of \( \gamma_{2a} \)-globulin accelerated but also the catabolism of \( \gamma_{2b} \)-globulin and 7S \( \gamma_{1} \)-globulin was increased by the high serum \( \gamma_{2a} \)-globulin level (Fig. 5). The \( \gamma_{1}A \)-globulin catabolism, however, was unaltered by large quantities of \( \gamma_{2a} \)-myeloma protein (Fig. 5) (Table II).

Large quantities of \( \gamma_{2a} \)-myeloma protein accelerated the catabolism of all three classes of 7S \( \gamma \)-globulin (Table II). The turnover of \( \gamma_{2a} \)-globulin, \( \gamma_{2b} \)-globulin, and 7S \( \gamma_{1} \)-globulin in mice bearing \( \gamma_{2b} \)-type plasma cell tumor MPC-37 is shown in Fig. 6, where they are seen to be catabolized at a much greater

Fig. 5. Catabolism of immunoglobulins, \( \gamma_{2a} \), \( \gamma_{2b} \), 7S\( \gamma_{1} \) and \( \gamma_{1}A(\beta_{2a}) \) in mice bearing \( \gamma_{2a} \) myeloma protein. The catabolic curves of \( ^{125}I \)-labeled \( \gamma_{2a} \), \( \gamma_{2b} \), 7S \( \gamma_{1} \), and \( \beta_{2a} \)-globulins in a group of mice bearing plasma cell tumor Adj.PC-5-producing \( \gamma_{2a} \)-myeloma protein are indicated. The T \( \frac{1}{2} \) values for the catabolism of each immunoglobulin in normal mice (N) is given in brackets.
Fig. 6. Catabolism of immunoglobulins, $\gamma_2a$, $\gamma_2b$, $\gamma_1\lambda$, and $\gamma_1\lambda(\beta_{2A})$ in mice bearing $\gamma_2b$-myeloma protein. The catabolic curves for each $^{131}I$-labeled immunoglobulin in a group of three mice with plasma cell tumor MPC-37-producing $\gamma_2b$-myeloma protein is shown.

Fig. 7. Catabolism of immunoglobulins, $\gamma_2a$, $\gamma_2b$, $\gamma_1\lambda$, and $\gamma_1\lambda(\beta_{2A})$, in mice bearing $\beta_{2A}$-myeloma protein. Catabolic curves for $^{131}I$-labeled $\gamma_2a\gamma$, $\gamma_2b\gamma$, $\gamma_1\lambda\gamma$, and $\beta_{2A}\lambda$-globulins in a group of mice bearing plasma cell tumor MPC-1 which produces a $\gamma_1\lambda(\beta_{2A})$-myeloma protein are indicated. The catabolic rate for each immunoglobulin in normal mice (N) is given in brackets.
rate than in normal mice. Similar observations were made with \( \gamma_{2b} \)-type tumors MPC-11 and MPC-31 (Table II).

The catabolism of \( \gamma_1 \) (IgA)-globulins on the other hand appeared to be unaffected by a large quantity of \( \gamma_{1b} \)-globulin. Mice bearing \( 7S \) \( \gamma_1 \)-globulin-producing tumors were not available for study so that a direct comparison of

![Graph showing the effect of exogenous \( \gamma \)-globulin on the catabolism of individual mouse \( 7S \) immunoglobulins.](image)

**Fig. 8.** Effect of exogenous (human) \( \gamma \)-globulin on the catabolism of individual mouse \( 7S \) immunoglobulins. Low-pathogen mice were used. Six mice were injected with each protein. Three served as controls and the remaining three in each group were injected intraperitoneally with human \( \gamma \)-globulin (50 mg on day 6, and 10 mg on days 7 and 8).

the effect of these tumors on each of the immunoglobulin components could not be made.

Large amounts of \( \gamma_1 \) (IgA)-myeloma protein did not have any appreciable effect on the catabolism of \( 7S \) \( \gamma \)-globulin components (Fig. 7). Each of the \( 7S \) \( \gamma \)-globulin components was catabolized at a rate close to normal in mice bearing \( \gamma_2 \) (IgA)-type plasma cell tumors (Table II). \( \gamma_1 \)-globulin catabolism was also unaltered by the presence of large amounts of \( \gamma_1 \)-myeloma protein (Table II).

**Effect of Exogenous \( 7S \) \( \gamma \)-Globulin.**—Prior studies (5, 16, 17) have shown that
the administration of human 7S γ-globulin to mice accelerates the catabolism of normal 7S γ-globulin of mouse origin. The effect of human 7S γ-globulin on each of the mouse 7S γ-globulin components was investigated in the present work. Administration of human γ-globulin in large amounts caused a marked acceleration of the catabolism of mouse 7S γ1-globulin, γ2s-globulin, and γ2b-globulin (Fig. 8). These findings in mice with low immunoglobulin levels were confirmed in normal mice (Table III). The effects of human γ-globulin on the catabolism of individual mouse myeloma proteins are similar to the effect on normal mouse 7S γ-globulin catabolism. Exogenous human 7S γ-globulin had no effect on mouse γ1A-globulin metabolism (Table III), in accord with previous observations (5).

### Table III

<table>
<thead>
<tr>
<th></th>
<th>γ1a</th>
<th>γ2b</th>
<th>7S γ1</th>
<th>7S γ1</th>
</tr>
</thead>
<tbody>
<tr>
<td>NIH-WS (Control)</td>
<td>5.4</td>
<td>4.8</td>
<td>3.0</td>
<td>2.0</td>
</tr>
<tr>
<td>Injected† NIH-WS</td>
<td>2.2</td>
<td>2.0</td>
<td>1.7</td>
<td>1.0</td>
</tr>
<tr>
<td>CVN or GF (Control)</td>
<td>10.2</td>
<td>11.8</td>
<td>1.8</td>
<td>2.0</td>
</tr>
<tr>
<td>Injected† CVN or GF</td>
<td>2.1</td>
<td>1.9</td>
<td>0.7</td>
<td>0.6</td>
</tr>
</tbody>
</table>

* Half-life in days.
† 10 mg human IgG (7S γ2-globulin)/day/animal is maintenance dose.

### DISCUSSION

**Control of Catabolism.**—Five immunoglobulin components of mice were found to have different catabolic properties.

The γ2s-globulins had a fractional rate of catabolism of approximately 13 per cent/day in normal mice while the γ2b-globulins were catabolized at almost twice this rate; i.e., about 25 per cent per day. These observations were confirmed with several myeloma proteins, and the catabolic rates of the γ2s- and γ2b-protein groups did not overlap. The γ2s- and γ2b-myeloma proteins differ in other respects. The genetically determined Iga-1 isoantigens are present on γ2a-globulins but not on γ2b-globulins (8). The γ2a-globulins sensitize guinea pig skin for reverse passive cutaneous anaphylaxis, which γ2b-globulins will not do (18). The γ2s- and γ2b-globulins also differ in antigenic determinants which can be detected with rabbit antiserum (8). The catabolic differences observed in the present study are another parameter of difference between these molecules.
The 7S $\gamma_1$-myeloma protein was catabolized at a rate of about 17 per cent per day. The 7S $\gamma_1$-protein clearly differs in catabolic properties from the $\gamma_2\beta$-myeloma proteins. Before an exact comparison of 7S $\gamma_1$- and $\gamma_2\beta$-proteins can be made, however, additional myeloma proteins of each type would have to be studied. The 7S $\gamma_1$-globulins differ from both the $\gamma_2\beta$- and $\gamma_2\alpha$-globulins in antigenic determinants (7) and in capacity to sensitize homologous (mouse) skin (18–20).

Prior studies of the turnover of I$\alpha$-labeled normal 7S immunoglobulins have shown a progressive change (flattening) of the decay curve; i.e., a progressive reduction in the fractional rate of catabolism of the labeled protein (5, 16). This finding could be accounted for by the existence of molecules with different catabolic properties in the normal population. Those molecules with the more rapid rates of catabolism (shorter half-lives) would be removed most rapidly. On each succeeding day, there would be a progressively greater proportion of molecules with slower rates of catabolism among the remaining population of labeled 7S immunoglobulins. This would produce a flattening of whole body decay curves. The catabolic heterogeneity might have been artifactual due to molecular differences introduced during purification or radioactive labeling of the globulins. On the other hand, the changing curve of normal 7S immunoglobulin catabolism could have resulted because the normal population was heterogeneous in terms of catabolic rates. This last interpretation is supported by the present observations of catabolic heterogeneity of 7S immunoglobulins in the mouse.

The catabolic curves of normal IgG (7S $\gamma$-globulin) preparations in man also show a notable change (flattening) of the decay curve (1, 6) indicating that in man, as in the mouse, the normal IgG population is catabolically heterogeneous. Investigation of the four subclasses of human IgG ($\gamma_2\alpha$, $\gamma_2\beta$, $\gamma_2\delta$, and $\gamma_2\alpha$), which differ on the basis of specific heavy polypeptide chain features (21, 22) may be fruitful in this respect.

The evidence for metabolic heterogeneity in the normal 7S globulin population indicates that data obtained from studies of I$\alpha$-labeled normal 7S immunoglobulins represent mean values for mixtures of proteins. The finding of normal 7S immunoglobulin half-lives of about 4 days in normal mice and 10 days in low pathogen mice (5, 16) indicate that the short-lived 7S $\gamma_2\beta$-molecules may represent only a small part and $\gamma_2\alpha$- and 7S $\gamma_1$-globulin molecules a larger part of the normal IgG (7S $\gamma$-globulin) population.

Differences in catabolism of several classes of 7S immunoglobulin have been emphasized in the preceding discussion. Similarities in control of catabolism also need to be emphasized.

The catabolic rates for $\gamma_2\alpha$, $\gamma_2\beta$, and 7S $\gamma_1$-globulins showed a similar dependence on the serum level of 7S immunoglobulin. When the serum immunoglobulin levels were low, about 10 per cent of normal as in mice with little
exposure to pathogens, the half-time was prolonged and the fractional rate of catabolism was reduced as low as 5 per cent per day for \( \gamma_2 \)-globulins and 7S \( \gamma_1 \)-globulins (one-third of the normal rate). When all serum immunoglobulin levels were increased by hyperimmunization, the half-time for each 7S immunoglobulin was shortened. The fractional rate of catabolism was increased up to 30 per cent per day for \( \gamma_2 \)- and 7S \( \gamma_1 \)-globulin (twice the normal rate). The \( \gamma_2 \)-globulin showed parallel changes at low and high immunoglobulin levels. Similar changes occurred when the serum 7S \( \gamma \)-globulin level was raised by the injection of human \( \gamma \)-globulin (IgG). These findings show that all three classes of 7S immunoglobulin had a similar catabolic relationship to the total serum level of 7S immunoglobulin.

The studies of catabolic control in low pathogen and hyperimmunized mice did not indicate whether each class of 7S immunoglobulin was responsive to the level of all 7S immunoglobulins or was sensitive only to the level of a single class of 7S immunoglobulin. This question was investigated by taking advantage of the selective immunoglobulin increases due to serum myeloma protein in mice with plasma cell tumors. Observations conducted in mice with high myeloma protein levels indicated that selective increase of one class of 7S immunoglobulin increased the catabolism of all three classes. This similarity indicated that catabolism was regulated by a property common to these classes of immunoglobulin. Other studies have shown that this is a property of the heavy polypeptide chain of the molecule (5).

The mechanism controlling the fractional rate of 7S \( \gamma \)-globulin catabolism remains to be identified. Brambell, Hemmings, and Morris (23) have proposed that some of the 7S \( \gamma \)-globulins become isolated from the general pool each day, and that at the time of this isolation some molecules become attached to specific receptors which protect these molecules (from catabolism) and return them undamaged to the general pool, whereas the unprotected molecules are degraded. At low serum \( \gamma \)-globulin levels, many protector sites would be available in relation to the total number of 7S immunoglobulin molecules and, thus, account for a low rate of catabolism (6, 16). At high serum \( \gamma \)-globulin levels, the protective sites would be relatively fewer and a greater fraction of 7S immunoglobulin would be catabolized (5, 6, 16). The present data are in accord with this hypothesis and indicate that the \( \gamma_2 \)-, \( \gamma_3 \)-, and 7S \( \gamma_1 \)-globulin molecules may be protected from catabolism in a similar way.

The \( \gamma_1 \) (IgA, \( \beta_1 \))- and \( \gamma_1 \)M (IgM, \( \beta_2 \))-globulins were notable for their rapid rates of catabolism and for their independence from the factors controlling 7S immunoglobulin catabolism. The fractional rate of catabolism was estimated at 55 per cent per day for IgA \( (\gamma_1 \), \( \beta_1 \)) and 140 per cent per day for IgM \( (\gamma_1 \)M). Catabolic measurements of these proteins presented major problems and the present data can only be regarded as approximations. The rapid rate of IgA catabolism (55 per cent per day) could have led to accumulation
of free IgM in the body and produced apparent half-times of survival that are greater than the true value (as well as falsely low estimates of the fractional rate of catabolism). There was, however, no evidence of an artifact from the shape of the whole body radioactivity decay curves. The basis for the two different macroglobulin antibody half-times observed (0.2 and 0.6 days) remains to be explained. Both half-times were found in at least two separate experiments. In any event, rapid rate of catabolism was characteristic of the IgM (\(\gamma_1\)M) molecules.

**Rates of Synthesis.**—The rates of synthesis for each of the immunoglobulin components were found to be of the same order of magnitude (Table IV). In

<p>| TABLE IV |
|-------------------------|-------------------------|-------------------------|
| Approximate Synthetic Rates for Immunoglobulins (mg/day/25 gm mouse) |</p>
<table>
<thead>
<tr>
<th></th>
<th>Normal mice</th>
<th>Hyperimmune mice</th>
<th>Low pathogen mice</th>
</tr>
</thead>
<tbody>
<tr>
<td>7S (\gamma_2)-: if 50 per cent</td>
<td>0.65</td>
<td>5.6</td>
<td>0.025</td>
</tr>
<tr>
<td>if 10 per cent</td>
<td>0.13</td>
<td>1.1</td>
<td>0.005</td>
</tr>
<tr>
<td>7S (\gamma_6)-: if 50 per cent</td>
<td>1.15</td>
<td>9.2</td>
<td>0.078</td>
</tr>
<tr>
<td>if 10 per cent</td>
<td>0.23</td>
<td>1.9</td>
<td>0.015</td>
</tr>
<tr>
<td>7S (\gamma_1)</td>
<td>1.06</td>
<td>17.5</td>
<td>0.025</td>
</tr>
<tr>
<td>(\gamma_M) ((\gamma_1)A, (\beta_M))</td>
<td>0.64</td>
<td>4.5</td>
<td>(\dagger)</td>
</tr>
<tr>
<td>(\gamma_M) (IgM, (\beta_M))</td>
<td>1.4</td>
<td>7.7</td>
<td>(\dagger)</td>
</tr>
</tbody>
</table>

* Two calculations are given for 7S\(\gamma_2\)- and 7S\(\gamma_6\)-globulins based on the possibility that these represented 50 or 10 per cent of the total 7S \(\gamma_1\)-globulin population. The exact serum level of these two components was not determined.

† These components could not be detected in the serum of low pathogen mice. If the serum level was one-half of the minimal level that could be detected, the rates of synthesis would be about 0.005 mg/day for these proteins.

normal mice the IgM (\(\gamma_1\)M-globulin) is synthesized at about the same rate as the 7S \(\gamma_2\)- and 7S \(\gamma_1\)-globulins. The serum level of IgM (\(\gamma_1\)M), however, is normally only 10 or 20 per cent of the level of IgG (7S \(\gamma\)-globulins) because of the rapid rate of IgM catabolism.

The amount of IgA (\(\gamma_1\)A, \(\beta_2\)A-globulins) synthesized daily is not fully reflected in the relative serum concentration (Table IV). The rapid rate of catabolism causes the serum level of IgA to be lower than the serum level of 7S \(\gamma_2\)- and 7S \(\gamma_1\)-globulins.

The rates of immunoglobulin synthesis in hyperimmunized mice are markedly increased, the data in Table IV showing 5- to 10-fold increases in rate of synthesis for all immunoglobulins. The data for IgA (\(\gamma_1\)A) synthesis in hyperimmunized mice (Table IV) is based on observations in C57BL and BALB/c mice which typically show higher IgA (\(\gamma_1\)A) levels (24) and probably have a correspondingly greater rate of IgA (\(\gamma_1\)A) synthesis than the white Swiss
(NIH-WS) mice used for the present studies. With this exception, the remaining values in Table IV are average values from observations in this study.

The observations in low pathogen (germfree or CVN) mice indicate the importance of antigen exposure in determining the production rate of immunoglobulins. Mice with a minimum of antigenic experience had low serum levels and very low rates of synthesis (Table IV). Many of the germfree mice available at the time of these studies evidently had been exposed to considerable amounts of antigen because they had relatively abundant serum 7S 7S-globulin levels (Table I). The CVN-low pathogen mice, however, had low 7S 7S-globulin levels, as did germfree mice investigated previously (16). The rates of 7S- and

TABLE V

| Estimates of Molecular Synthetic Rates (molecules of Immunoglobulin/day/25 gm mouse) |
|---------------------------------|---------------------------------|
|                                 | Normal mice                     | Low pathogen mice               |
|                                 | (Whole molecules*) | (6.6S units) | (intact molecules) |
| 7S γ (γH + γm)  | 3.6 × 10^{15} | 3.6 × 10^{12} | 2 × 10^{14} |
| 7S γ     | 4.2 × 10^{15} | 4.2 × 10^{12} | 1 × 10^{14} |
| γM(IgA, βA) | 1.5 × 10^{15} | 3.0 × 10^{12} | 2 × 10^{13} |
| γM(IgM, βM) | 8 × 10^{14} | 4.0 × 10^{12} | 5 × 10^{13} |

* Based on amounts calculated in Table IV, and molecular weights of 150,000 for 7Sγ, and 7Sγ, globulins, 300,000 for γM(IgA) and 900,000 for γM(IgM), the latter two made up of 2 and 5 units respectively.

† Based on the assumption that serum concentration is 1/2 of the minimal amount of immunoglobulin detectable by the quantitative techniques used. The actual rate of synthesis for these two proteins might be much less than these amounts but could not be more than twice the levels given above.

7S-globulin synthesis calculated in Table IV for low pathogen mice reflect the experience with this larger group of animals.

In the low pathogen mouse, the rates of immunoglobulin synthesis are much reduced (Table V). Even in the mouse with the lowest 7S γ-globulin levels, however, the calculated rate of 7S γ-globulin synthesis was 10^{14} molecules per day (16). This rate of synthesis in germfree mice is low in comparison to normal but is not low in absolute terms.

The rate of synthesis of IgA (γA, βA) and IgM (γM, βM) cannot be calculated in low pathogen mice because no serum level can be detected. This does not mean, however, that such proteins are not formed. The lower limit of detectability of mouse IgA and IgM was estimated to be 0.01 mg/ml. If the serum level in low pathogen mice was one-half of this, i.e. 0.005 mg/ml, then it was calculated that the rate of synthesis of these proteins would be about 10^{15} molecules each day (Table V). These calculations are not meant to imply that these are the rates of IgA and IgM synthesis in low pathogen mice, but the calculation emphasizes that as many as 10^{13} molecules of these im-
munoglobulins can be synthesized per day without being detected in mouse serum.

SUMMARY

The metabolic properties of immunoglobulin were investigated by comparing five classes of mouse immunoglobulin. Three forms of 7S immunoglobulin had different rates of catabolism. The fractional rates of catabolism were found to be about 13 per cent per day for 7S γ2b-globulin; 25 per cent for 7S γ2a-globulin; and 17 per cent for 7S γ1-globulin. Catabolism of the three classes of 7S γ-globulin (γ1m, γ2b, and γ1) were prolonged at low serum 7S γ-globulin levels and accelerated at high serum 7S γ-globulin levels. Each of the 7S γ-globulin components was influenced by the serum level of the other mouse 7S γ-globulin components and by exogenously administered human 7S γ-globulin. They were not appreciably altered, however, by the serum level of IgA (γ1A-, β2A-globulin).

The progressively changing (longer) half-times observed in turnover studies of normal IgG (7S γ-globulin) may be caused by catabolic heterogeneity of normal 7S immunoglobulins which are immunochemically and catabolically related to γ1m, γ2b, and 7S γ1-myeloma proteins.

These studies indicate that the 7S γ2b-, 7S γ2a-, and 7S γ1-globulins share a common catabolic control mechanism. This mechanism is influenced by the serum level of each of these components, but is independent of the serum level of IgA (γ1A-globulin) and probably is independent of IgM (γ1M-globulin).

Catabolism of IgA (γ1A-, β2A-globulin) and IgM (γ1M-globulin) was much more rapid than the catabolism of the 7S γ-globulins. The half-times of the IgA and IgM were approximately 1.2 and 0.5 days respectively. The fractional rate of catabolism of IgA and IgM seemed to be independent of their serum concentration.

The rate of catabolism, as well as the rate of synthesis, was shown to play a major role in determining the serum level of each class of immunoglobulin.

BIBLIOGRAPHY


