# **INTERACTION OF** *H.2* **GENOTYPE AND BASAL** SERUM IMMUNOGLOBULIN **A**  LEVEL INFLUENCES LONGEVITY

#### DIANA M. POPP, J.A. OTTEN and R.A. POPP

*Biology Division, Oak Ridge National Laboratory, Oak Ridge, TN 3 7831 (U.S.A.)* 

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#### SUMMARY

The congenic pair of mice, C57BL/10 (BIO) and C57BL/10.F (B10.F), differ at the *H-2* locus and have mean ages at death of 706 and 456 days, respectively. B10.F also has reduced basal serum IgA levels compared with B10, 63 and 256 mg/dl, respectively. Controlled matings between the two strains of mice were used to identify genetic factors that govern longevity.  $F_2$  and backcross progeny from reciprocal  $F_1$  hybrids were classified for  $H-2$  genotype and serum IgA levels and allowed to live out their lifespan.  $F_2$  and backcross progeny homozygous for the *H-2* allele of BIO.F had a mean age at death (602 days) significantly reduced from that of progeny homozygous for the *H-2* allele of BIO (689 days). However, the greatest reduction of lifespan occurred among progeny of the  $(B10.F \times B10)F_1$  mothers, 693 compared with 540 days. The strain of the maternal parent also has been shown to affect the segregation of IgA phenotypes. An increased incidence of low IgA phenotype associated with *1t-2* genotype was observed among progeny of  $(B10.F \times B10)F_1$  mothers. Survival curves demonstrated a relationship between low serum IgA levels and shortened lifespan and no maternal effect was observed. The basis of the shortened lifespan among progeny of  $F_1$  hybrids in which the maternal parent was BIO.F was the increased incidence of offspring with low IgA phenotypes. The apparent association of *1-1-2* and shortened lifespan also was because the low IgA phenotype was more frequent among progeny that carried the *H-2* allele of the B10.F strain. The B10.F mice spontaneously shed an endogenous ecotropic retrovirus which may be responsible for the maternal effect on immunoglobulin levels and lifespan.

*Key words:* Lifespan; Genetics; Immunoglobulin; Retrovirus; Maternal Effect

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#### INTRODUCTION

The members of a congenic pair of mice, C57BL/10 (B10) and *C57BL/lO.F-H-2*  (BIO.F), have been reported to have significantly different life spans [1]. This pair has also been shown to differ in basal immunoglobulin A (IgA) levels  $[2]$ . Using  $F_2$  and backcross progeny from reciprocal  $F_1$  hybrids of B10 and B10.F, both life span [3] and serum IgA levels [4] have been shown to be influenced by an unidentified maternal effect. Genetic tests also showed that lifespan could be correlated with *H-2* genotype, i.e. progeny homozygous for the  $H-2^n$  allele of the short-lived strain (B10.F) had a mean age at death significantly reduced from that of progeny homozygous for the  $H-2^b$  allele of B10 [3]. It was also noted that *H-2* genotype and maternal parentage were synergistic in life span reduction; the  $H-2^{n/n}$  progeny from the (B10.F  $\times$  B10)F<sub>1</sub> survived less well than the *H*-2<sup>n/n</sup> progeny from the  $(B10 \times B10)$ F<sub>1</sub> mother. The basis for the observed maternal effect on life span is examined in this report and evidence is presented showing that low IgA levels in association with the  $H-2^n$  allele shortens longevity. The results of cell transfer and foster nursing experiments that were designed to examine the mechanism of the maternal effect are also reported. Data are presented that suggest that virus reintegration in BIO.F somatic cell DNA may account for the influence of the maternal parent on the inheritance of IgA phenotypes observed in progeny of B IO.F mice.

### MATERIALS AND METHODS

### *Mice*

The origin of the BIO and BIO.F congenic pair used in this study is described in a previous report  $[3]$ . Backcross and  $F_2$  progeny were produced from reciprocal hybrid females derived from strains BIO and B10.F to study the influence of *H-2* genotype, IgA phenotype and maternal parent on longevity. Each offspring was serotyped for *H-2*  genotype, classified according to IgA level, and the date of natural death was recorded. Parental strains and all progeny were maintained in the same room and under equivalent environmental conditions.

## *Serology*

The hemagglutination method of Gorer and Mikulska [5 ] was used to determine the *H-2* serotypes of the offspring. The antiserum used to detect the *H-2<sup>n</sup>* haplotype of the B10.F parent was made in  $(B10.A \times A.CA)F_1$  against B10.Y tissue (designated C-16) and was obtained from the Transplantation Immunology Branch of the National Institutes of Health. The antiserum used to detect the  $H-2<sup>b</sup>$  haplotype of the B10 parent was made in C3H mice against RFM tissues.

### *Immunoglobulin ( IgA ) quantitation*

The procedures for serum IgA quantitation are described in a previous publication  $[4]$ . In brief, the Mancini radial immunodiffusion method [6] was used, employing immunodiffusion plates obtained from Meloy Laboratories. All offspring were 4 months or older when basal serum IgA levels were determined.

## *Cellular studies*

Spleen cells were suspended in Eagle's MEM without bicarbonate [7]. Spleens were teased apart and single cell suspensions were prepared by aspirating through a 21-gauge needle. The cell concentrations were adjusted to  $75 \times 10^6$ /ml and 0.2 ml were injected intraperitoneally into suckling offspring. A B10.F spleen suspension was centrifuged at 30 000 g for 1 h to yield a cell-free preparation. The supernatant was injected into 15.day-old B10 neonates. These recipients received the same spleen-equivalent dose as the recipients of the whole cell inoculum.

Bone marrow was flushed from the femurs of donor mice and suspended in Eagle's MEM (without bicarbonate). Lethally irradiated (800 R) recipients received injections of  $15 \times 10^6$  cells via the tail vein. Litters selected to be foster nursed were removed from their mothers at birth and transferred to post-partum females whose own offspring had been removed prior to suckling.

The females that provided the blastocysts were mated and examined daily for copulation plugs. At 3.5 days post-plug the uterus was removed and flushed with Brinster's solution (Gibco Laboratories). The blastocysts collected in this manner were introduced into surrogate mothers by injection into the surgically exposed uterus. Six to ten blastocysts were introduced at a time. The appropriate hormonal state for blastocyst implantation in the recipient female was assured by pairing these females with vasectomized males and examining for copulation plugs.

### *Virus assays*

Spleens from young (4-month-old) B10 and B10.F mice were aseptically removed and frozen  $(-70^{\circ}\text{C})$  in 1.0 ml of phosphate buffered saline. The spleens were thawed, homogenized in a glass homogenizer, and the supernatant filtered through a 0.45  $\mu$ m filter. The filtered supernatant was assayed for virus on SC1 cells [8] by fluorescent antibody and XC assays [9,10]. At 48 h post-infection, the medium was removed from the SC1 cells and saved as small virus pools. The virus pools were used to infect A31 [11], NIH [12] and SC1 cells, and 5 days later the XC assay was performed as previously described [9].

### *Southern blot analysis*

DNA was prepared from the liver, spleen, and pooled peripheral and mesenteric lymph nodes of BIO and BIO.F mice. The DNA was digested by *PvuH* and electrophoresed in 1% agarose, transferred to nitrocellulose paper, and hybridized with a 0.4 kb probe specific for the *env* gene of ecotropic AKR MuLV [13].

## RESULTS

An analysis of  $F_2$  and backcross progeny of reciprocal  $F_1$  hybrids from B10 and B10.F

#### TABLE I

THE MEAN AGE AT DEATH  $\pm$  S.E.M. OF  $H-2$  SEGREGANTS IN  $F<sub>2</sub>$  AND BC POPULATIONS FROM RECIPROCAL  $F_1$  HYBRID FEMALES [3]

No. of animals in parentheses.

$H2$ genotype	Maternal parent					
	$\langle B10 \times B10.F \rangle F$	$(B10.F \times B10)F$ ,				
b/b	$687 \pm 18(58)$	$693 \pm 23(30)$				
$n/n^a$	$627 \pm 15(89)$	$540 \pm 19(35)$				

aHistorically, the *H-2* haplotype of B10.F was defined as *H.2 n.* Subsequent studies showed serological similarity with  $H_2P$  although skin grafts between  $H_2P$  and  $H_2P$  congenic lines were rejected in one direction. Thus, the *H-2* haplotype of B10.F has been designated a minor variant *of H-2P* [19]. We will use *H*-2<sup>n</sup> in this report to be consistent with the series of papers that have been published in the course of these studies.

mice showed that the maternal parent influences the segregation of serum IgA phenotypes [14]. Survival curves of these progeny sorted on the basis of their *H-2* types showed that the class with the *H-2n/n* genotype had a shortened life span and a maternal effect also influenced the severity of life shortening [3]. These data are summarized in Table I. For the purposes of this report, these data have been re-examined in Table II to show the distribution of IgA phenotypes among the progeny of each mating and the distribution of IgA phenotypes according to *H-2* genotype. Data in Table II show that a larger proportion of the F<sub>2</sub> and B10.F backcross offspring from (B10.F  $\times$  B10)F<sub>1</sub> females

#### TABLE II

THE DISTRIBUTION OF SERUM IgA PHENOTYPES  $%$  AMONG  $F<sub>2</sub>$  AND BACKCROSS (BC) PROGENY FROM RECIPROCAL F<sub>1</sub> HYBRID FEMALES

H-2 Genotype:	All		b/b		n/b		n/n	
IgA Phenotype <sup>a</sup> : High		Low	High	Low	High	Low	High	Low
$(B10 \times B10.F)F$								
$F2$ (193)	72	28	82	18	66	34	75	25
<b>B10 BC (41)</b>	73	27	77	23	68	32	$\overline{\phantom{a}}$	
B <sub>10</sub> .F BC(41)	84	16			92	8	77	23
$(B10.F \times B10)F$ ,								
F, (73)	49	51	93	7	49	51	16	84
<b>B10 BC (31)</b>	74	26	94	6	53	47	$\overline{\phantom{0}}$	
<b>B10.F BC (32)</b>	25	75			50	50	0	100

No. of animals in parentheses.

 $a_{\text{IgA high}} > 100 \text{ mg/dl}$ ; IgA low < 100 mg/dl (1).



Fig. 1. Survival curves of  $F_2$  progeny of reciprocal  $F_1$  hybrids classified according to IgA phenotype. Lo = Low level; Hi = high level;  $n/n = H-2^n$  homozygote.

have the low IgA phenotype than progeny from similar matings using the reciprocal  $F_1$ hybrid female. The distribution of IgA-low- and IgA-high-phenotypes is similar for each of the  $H-2$  segregant groups born by  $(B10 \times B10)$ . F)F<sub>1</sub> females. However, among the progeny of  $(B10.F \times B10)F_1$  females the distribution of IgA phenotype is different for each of the *H*-2 segregant groups and a greater proportion of  $H$ -2<sup>n/n</sup> and  $H$ -2<sup>n/b</sup> segregants have low IgA levels compared with similar progeny from the reciprocal  $F_1$  parent.

Since the populations that show reduced lifespans have an increased incidence of low-IgA-phenotypes, survival curves of  $(B10 \times B10)F_2$  and  $(B10.F \times B10)F_2$  selected by IgA phenotype were constructed (Fig. 1). The progeny that have high serum IgA levels survive better than those that have low IgA levels in both of the reciprocal  $F_2$  populations. The 50% lifespan of the high or low IgA groups is not influenced by the maternal parent; thus, there is no maternal effect imposed upon the survival data when expressed as a function of IgA levels. When the 50% lifespan of IgA-high- and IgA-low-progeny are compared among the reciprocal BC progeny, there is no maternal effect evident (Table III). The reduced survival associates with the IgA low phenotype, and the relative survivals of high and low groups are similar whether descended from (B10  $\times$  B10.F)F<sub>1</sub> or (B10.F  $\times$ B10)F<sub>1</sub> female parents (Table III), except for the  $(B10 \times B10)$ F<sub>1</sub> backcross to B10 where the survivals of the IgA low and high groups are equivalent. (See Discussion Section.) In addition, the survival of the  $H-2^{n/n}$  segregants of the IgA low populations from the reciprocal  $F_2$  matings is reduced equally (Fig. 1) and there is no evidence of a maternal

## TABLE III

THE 50% LIFESPAN (DAYS) OF  $F_2$  AND BACKROSS MICE FROM RECIPROCAL  $F_1$  HYBRIDS DERIVED FROM STRAINS BI0 AND B10.F CLASSIFIED ACCORDING TO IgA PHENOTYPE; 50% SURVIVAL WAS DETERMINED FROM SURVIVAL CURVES

$(B10 \times B10. F)F$ ,			$(B10.F \times B10)F$ ,			
IgA phenotype	No. mice	50% lifespan	IgA phenotype	No. mice	50% lifespan	
High	133	680	High	34	670	
Low	53	590	Low	37	580	
	$(B10 \times B10. F)F_1 \times B10$		$(B10.F \times B10)F \times B10$			
High	27	695	High	21	710	
Low	11	695	Low	8	600	
	$(B10 \times B10.F)F_1 \times B10.F$		$(B10.F \times B10)F_1 \times B10.F$			
High	44	650	High	8	710	
Low	9	480	Low	24	560	

effect. Thus, data on the lifespan of reciprocal progeny were pooled to analyze the survival of groups selected according to their IgA level and *H-2* genotype.

The mean age at death of *H:2* segregants sorted according to their IgA phenotype are given in Table IV. Among the progeny that have high serum IgA levels, the survival rate

#### TABLE IV

THE MEAN AGE AT DEATH (DAYS)  $\pm$  S.E.M. OF  $F_2$  AND BACKCROSS PROGENY OF RECI-PROCAL  $F_1$  HYBRIDS OF STRAINS B10 AND B10.F CLASSIFIED ACCORDING TO THEIR lgA PHENOTYPE AND *H-2* GENOTYPE

$IgA$ : $H-2$ types	No. mice	Mean age at death	P	
High IgA				
b/b	70	$685 \pm 17$		
n/b	130	$688 \pm 11$	<b>NS</b>	
n/n	67	$651 \pm 17$	<b>NS</b>	
Low IgA				
b/b	13	$688 \pm 27$	<b>NS</b>	
n/b	74	$623 \pm 16$	< 0.01	
n/n	54	$539 \pm 16$	< 0.001	

Statistical comparisons have been made with the high IgA,  $H-2^{b/b}$  group taken as representative of **the normal Ufespan of strain B10. Student's t-test was used to determine significant differences (P > 0.05 = NS, not significantly different). Hewlet-Packard 65 Star Pac 1-30A.** 



Fig. 2. Survival curves of homozygous  $H-2$  populations pooled from  $F<sub>2</sub>$  and backcross progeny of reciprocal F, hybrids, classified according to IgA phenotype. No. of animals in (). Lo = low level; Hi = high level;  $n/n = H-2^n$  homozygote;  $b/b = H-2^b$  homozygote.

is not different with respect to  $H-2$  genotype; the  $H-2^{n/n}$  progeny have a slightly reduced mean age at death but the difference is not significant  $(P > 0.05)$ . Among the progeny that have low serum IgA levels, the survival of the *H-2* segregants is different. The survival of *H-2<sup>b/b</sup>* progeny is comparable with that of progeny with high serum IgA levels, but progeny with low serum IgA levels and homozygous for the  $H-2^n$  allele are severely compromised with respect to lifespan. Individuals that are heterozygous at *H*-2 also have significantly reduced survival. Fig. 2 shows that  $H-2^{n/n}$ , high IgA and  $H-2^{b/b}$ , low IgA progeny have survival curves indistinguishable from that of  $H-2^{b/b}$ , high IgA progeny. Only the combination of  $H$ -2<sup>n/n</sup> and low serum IgA levels severely shortens lifespan. Survival of *H-2* heterozygotes that have low serum IgA levels is reduced but not as much as that of the  $H-2^n$  homozygotes.

A foster-nursing study was done in attempts to understand the maternal effect on serum IgA levels. Newborn B10.F,  $(B10 \times B10)$ .F)F<sub>1</sub> and B10 mice were placed with post-partum females in the combinations shown in Table V. When BIO.F neonates were foster nursed on B10 females, their adult IgA phenotypes were not altered. However, all B10 and  $(B10 \times B10)F_1$  offspring that had been foster nursed on B10.F mothers had reduced serum IgA levels that were similar to those of B10.F mice at 4 months of age. At 1 year, the IgA levels were still suppressed in the foster-nursed B10 mice and in 4 of the 6 foster-nursed (B10  $\times$  B10.F)F<sub>1</sub> mice.

To assess the possibility that maternally derived immunocompetent cells modulated the maturation of the humoral IgA response of immature mice, spleen cell transfers

	ABL		

BASAL SERUM IgA LEVELS OF 4-MONTH-OLD MICE THAT WERE FOSTER NURSED AT BIRTH



were made into 12-day-old neonates before IgA synthesis begins. Because B10 and B10.F differ at  $H-2$ , these cell transfers had to be done using  $F_1$  hybrids. The data are presented in Table VI. In three experiments in which spleen cells from adult  $(B10.F \times B10)F_1$ donors (low serum levels) were injected into  $(B10 \times B10)$ . F)F<sub>1</sub> neonates (potentially high serum IgA levels), 12 of 12 surviving recipients had low serum IgA levels at 4 months. The serum IgA levels were still low at 1 year; 6 survivors at 12 months had suppressed serum IgA levels and the last survivor was still suppressed at 19 months (later data not given).

### TABLE VI

Donor	$IgA$ (mg/dl) 4 months	$IgA$ (mg/dl) 12 months	
<i>Experimentals</i>			
$(B10.F \times B10)F_1$ Adult spleen	59, 55, 47, 47, 47	$D^b$ , 64, 44, 68, D	
$(B10.F \times B10)F_1$ Adult spleen	55, 55, 50, 55, 50	D, 77, 71, D, 55	
$(B10.F \times B10)F_1$ Adult apleen	65,65	56,56	
Controls			
$(B10 \times B10.F)F$ Adult spleen	75,75	97, 157	
$(B10 \times B10.F)F$ , Adult spleen	84, 71, 51, 56	200, 360, 77, 285	
Experimental $(B10.F \times B10)F,$ 17-day-old spleen	56, 56, 48, 41, 48	97, 68, 38, 77, D	

BASAL SERUM IgA LEVELS OF 4 MONTH OLD (B10 × B10.F)F, MICE THAT RECEIVED SPLEEN CELL INJECTIONS AS NEONATES<sup>8</sup>.

<sup>8</sup>All neonate recipients were 12 days old except for the 17-day-old recipients of 17-day-old spleen. bD, dead.

The controls,  $(B10 \times B10)F_1$  recipients of  $(B10 \times B10)F_1$  adult spleen, demonstrated the unexpected effect of adult isologous spleen to generally suppress (or delay) IgA expression. This effect was temporary; 4 of the 6 controls had achieved a high IgA serum level by 1 year (Table VI). To determine if the effect of injecting adult (B10.F  $\times$  $B10$ )F<sub>1</sub> spleen was due to the introduction of mature immunoregulatory cells (T suppressor cells) into a developing immune system or inherent in the genome of the B 10.F maternal strain, spleen cells from 17-day-old  $(B10.F \times B10)F_1$  mice were injected into 17-day-old  $(B10 \times B10)$ . Fig. offspring. All 5 recipients had low serum IgA levels when tested at 4 months of age. These recipients were still suppressed at 1 year and one survivor was still suppressed at 18 months. Adult bone marrow injected into neonatal mice was also capable of modulating the IgA levels (data not given).

A line of mice was developed from two  $(B10 \times B10)$ . F)F<sub>2</sub> that had the *H*-2<sup>b/b</sup> genotype of B 10 mice and the low serum IgA level of the B 10.F strain. Spleen cells from these mice were used to show that similar modulation of serum IgA levels could be induced in B10 mice (Table VII). Spleen transfers into 8-day-old neonates modulated serum IgA levels in the B10 recipients and they were still suppressed at 1 year. Spleen cell transfers into 30-day-old weanlings modulated serum IgA levels but the modulation was temporary. Irradiation of the spleen cells did not abrogate the effect on IgA levels, and 4 of 6 recipients of the irradiated spleen cells were still modulated at 1 year.

The development of the *H-2b/b,* IgA low line also permitted an experiment to assess the effect of bone marrow transplantation on serum IgA levels. The transfer of B10 bone marrow into lethally irradiated B10 adult recipients did not alter the normal serum IgA levels for 4 of the 5 B10 recipients (Table VIII). Reciprocal bone marrow transfers between B 10 and B IO.F mice resulted in death of the recipients due to graft versus host

### TABLE VII

Donor	Recipient	$IgA$ (mg/dl)			
		4 months	12 months		
$b/b$ Low	<b>B10</b>	72	ŋа		
Adult spleen	8 day neonate	56, 56, 46, 51, 46	54, 47, 54, D, D		
$b/b$ Low	<b>B10</b>				
Adult spleen	30 day weanling	56, 48, 56, 52, 56	400, 138, 250, 123, 110		
b/b Low	<b>B10</b>	77.56	52,84		
Adult spleen $2163r$ X-ray <sup>b</sup>	12 day neonate	61, 90, 56, 41	32, 123, 137, 84		

THE BASAL SERUM IgA LEVELS OF 4-MONTH-OLD B10 MICE THAT RECEIVED SPLEEN CELLS FROM DONORS THAT WERE  $H-2^{b/b}$  AND HAD LOW SERUM IgA LEVELS (b/b LOW)

**aD, dead.** 

**blrradiated** *in vitro. •* 



#### TABLE VIII

BASAL SERUM IgA LEVELS OF RADIATION-INDUCED CHIMERIC MICE

aLethal graft-versus-host disease.

disease. When the low IgA  $H-2^{b/b}$  mice were used as bone marrow donors, the B<sub>10</sub> recipients survived and expressed reduced levels of IgA in their serum.

Some of the IgA suppressed (B10  $\times$  B10.F)F<sub>1</sub> and B10 females were mated to produce progeny. The IgA suppressed  $(B10 \times B10)$ F)F<sub>1</sub> females (Table VI) produced two F<sub>2</sub> litters, all  $F_2$  progeny had low IgA levels; when these  $F_2$  produced  $F_3$  progeny, all  $F_3$ progeny had serum IgA levels normal for BIO mice. The 2 B10 females that were suppressed by the injection of irradiated spleen (Table VII) produced 19 progeny, 17 of these were suppressed but their progeny had normal serum lgA levels.

Blastocyst transfer was done to examine the influence of the uterine environment on IgA levels. Three survivors were raised following the transfer of  $(B10.F \times B10)F$ <sub>1</sub> blastocysts to a BIO surrogate mother. They were H-2 serotyped to establish that they developed from the transferred blastocysts. All 3 offspring had elevated lgA levels (180, 170 and 318 mg/dl), even though they would have had low IgA levels had they developed naturally in B10.F mothers.

The B10 neonate recipients of cell-free spleen supernatants were tested for IgA level at 4 and 7 months of age. At 4 months, 7 of the 9 recipients were suppressed and at 7 months 6 of the 9 were suppressed (data to be published). In addition, 1 female and 3 males showed signs of early greying.

Spleens from young B10 and BIO.F mice were tested for virus expression. Table IX shows that the proteins detected by the immunofluorescence assay and the infectious virus subsequently detected by the XC plaque assay were present in the spleens from both strains of mice. The immunofluorescence assay was performed by placing supernatant from a spleen homogenate on SC1 cells followed by examination of the cells 6 days later. No enhancement of virus in the spleen cells was done prior to the immunofluorescent assay. The data shown in Table IX indicate that more virus was present in the BIO.F spleen cells (94% positive cells) than in the B10 cells (20% positive cells). Comparable numbers of SC1 cells were treated with the supernatants and, in addition, the spleens used to prepare the supernatants were equivalent in weight. The SC 1 cells multiplied at similar rates and, at the time of coverslip harvest, equivalent numbers of cells were counted. The additional virus load in the BIO.F, as indicated by the immunofluorescent assay, is consistent with the suggestion that this strain of mouse, as opposed

#### TABLE IX



### THE GROWTH OF VIRUS ISOLATES FROM B10 AND B10.F SPLENIC HOMOGENATES ON A31 (Fv-1<sup>b</sup>), NIH/3T3 (Fv-1<sup>n</sup>) AND SC1 (Fv-1<sup>-</sup>) CELLS

aDirect infection of SC1 cells. Coverslips harvested at 6 days.

blnfection of SC1, A31 and NIH/3T3 cells with small virus pools prepared in SC1 ceils. XC at 5 days. CNot done.

dNumber of plaques at indicated dilution of virus pools are given in parentheses.

to the B10, carries with it an additional virus burden from birth that may be responsible for the observed maternal effect.

The tropism of the virus isolates from B10 and B10.F spleens was determined by infecting A31 (Fv-1<sup>b</sup>) and NIH/3T3 (Fv-1<sup>n</sup>) cells with the virus pools. In addition, SC1  $(Fv-I^-)$  cells were infected to determine the titers of the virus contained in the individual virus pools. The data in Table IX show that both virus isolates demonstrated preferential growth on the A31  $(Fv-I^b)$  cells as opposed to the NIH/3T3  $(Fv-I^n)$  cells which indicates that both virus isolates are B-tropic. When compared with the titers in SCI *(Fv-1-)*  cells, virus from the B10 and BI0.F mice show an apparent restriction in both the Nand B-tropic mouse cells. These results show that the virus is present in both strains of mice and that the B10.F strain has a heavier virus burden from an early age than the B 10 strain of mouse.

To explore the basis for the difference in viral shedding between the congenic pair, Southern blot analysis was carried out on B10 and B10.F DNA prepared from liver, spleen, and pooled mesenteric and peripheral lymph nodes. The 0.4 kb *env* specific probe used identifies the endogenous ecotropic locus *(Emv-2)* present in B10 mice as a 5.2 kb PvulI fragment located on chromosome 8 [14]. This probe also identifies the *Emv-3*  locus of DBA/2 as a 5.4 kb Pvull fragment located on chromosome 9 so DBA/2 DNA was also used as a control. The results of the hybridization are shown in Fig. 3. Lanes 1 and 2 show the 5.4 and 5.2 kb PvulI fragments from liver genomic DNA of DBA/2 and B10.F, respectively. Lanes 3, 4 and 5 contain PvuII digested DNA of liver, pooled peripheral and mesenteric lymph nodes, and spleen, respectively, of a B10.F mouse that was grey and had a low serum IgA. Several DNA fragments that hybridize with the MuLV probe are present in DNA from the lymph nodes and spleen.



Fig. 3. Southern blot of PvuII digests of DNA from liver, spleen and mesenteric lymph node of a B10.F following hybridization with the 0.4 kb MuLV probe.

#### DISCUSSION

The data shown that among the offspring of the long-lived BIO and the short-lived B10.F mice the  $H_2^{n/n}$  genotype is necessary but not sufficient to compromise lifespan. Only in combination with low levels of serum IgA is reduced survival observed. Thus, the previously reported maternal effect on the survival of progeny of reciprocal  $F_1$ hybrid females [3] is due to the increased number of progeny with low IgA phenotypes among offspring of  $(B10.F \times B10)F_1$  females (Tables II-IV), particularly among the  $H-2^{n/n}$  segregants: of the  $H-2^{n/n}$  segregants from the  $F_2$  and B10.F backcross matings with (B10.F  $\times$  B10)F<sub>1</sub> females, 84% and 100%, respectively, have low serum IgA levels compared with 25% and 23%, respectively, of the  $H-2^{n/n}$  segregants from similar matings with the  $(B10 \times B10)$ F<sub>1</sub> females. Thus, the basis for *H-2* association with lifespan previously reported [3] is due to the combined effect of the appropriate *H-2* genotype  $(H-2^{n/n})$  and the low serum IgA phenotype (Table V).

The one group of IgA low offspring in which no reduction in survival was seen (Table III) is from a mating in which no  $H-2^{n/n}$  progeny are produced, the (B10  $\times$  B10.F)F<sub>1</sub>  $\times$ BIO backcross. Because *H-2* segregates independently of IgA phenotype in this mating  $[4]$ ,  $1/2$  (5/11) of the progeny are homozygous for the *H*-2<sup>b</sup> allele (all of which lived 638 days or longer) and the 50% survival of this group was 695 days. The remainder of the progeny (6/11) are heterozygous at *H-2.* Although the effect is not as severe, the presence of the  $H-2^n$  allele in the heterozygous state in association with low IgA levels also significantly reduces lifespan (Table IV). It would take a larger population for the lifespan reduction due to *H-2* heterozygosity to significantly reduce the survival of this backcross population. Because of the association of *H-2* and IgA phenotype found in progeny from the  $(B10.F \times B10)F_1$  female parent [4], the preponderance of the IgA low offspring from the B10 backcross with this female parent (7/8) are *H*-2<sup>n/b</sup> and half of the mice are dead by 600 days. Therefore, the greater proportion  $(1/2)$  of  $H<sub>-2</sub>b$  homozygotes among the IgA low group from the  $(B10 \times B10)$ . F<sub>1</sub> female parent contributes to longer group survival. Conversely, the preponderance of *H-2* heterozygotes among the IgA low group from  $(B10.F \times B10)F_1$  female parent results in reduced group survival which is apparent even in small populations.

Several genetic models to explain a maternal effect have been suggested [4] and maternal physiology and the transmission of self-replicating particles were discussed. Short-term allotype suppression in rabbits and mice has been shown after inducing antipaternal-type immunoglobulin in the female parent [15]. However, long-term allotype suppression in mice was found to be brought about and maintained by cellular mechanisms [15] demonstrated by spleen cell transfers into neonates [7]. These experiments led to the suggestion that chronic allotype-specific suppression was induced by the transfer of suppressor cells to the immunologically immature neonate [15]. To assess an immunoregulatory role of maternal cellular components on IgA phenotype, we carried out several cell transfer experiments. Data show that neonatal offspring can be modulated to have low IgA serum levels by spleen cell transfer, foster nursing or bone marrow transfer (Tables V-VIII). The modulation was permanent throughout the lifespan of the

animal and could be passed by females to the next generation offspring. However, when blastocysts of potentially low IgA mice were gestated in the uterus of a high IgA female the mature offspring developed high serum IgA levels. The results of these experiments suggested that the basis of the IgA specific suppression is the presence of allotype specific suppressor cells in the B10.F mice and that suppressor cells are induced in  $F_1$ , and some  $F<sub>2</sub>$  and backcross progeny of suppressed females. The alternative hypothesis, i.e. transmission of a viral particle, is also consistent with the results of the cellular studies. The discovery of active virus shedding in the B 10.F required additional experiments to distinguish between these alternatives. Modulation of IgA levels following transfer of a cellfree splenic filtrate established that the mechanism for the long-term suppression in neonatally injected or fostered-nursed offspring was not cell mediated.

The experiments described in this report are consistent with the suggestion that a vertically transmitted virus either inherited or acquired from B10.F and (B10.F  $\times$  B10)F<sub>1</sub> mothers is responsible for the deficient serum IgA levels among their offspring. Yetter *et al.* [16] have reported that the BIO.F mouse, in contrast to its B10 congenic partner, sheds an endogenous, ecotropic virus. Tests on our B10.F and BIO colonies confirmed this observation. In addition, we have found that although the genomic DNA of B10.F does not differ from BIO with respect to the chromosomal location of the integrated virus, multiple restriction fragments were found in the DNA of spleen and lymph nodes, suggesting that shed virus had reintegrated at additional sites in the DNA of some cells of these tissues (Fig. 3). These data indicate that MuLV reintegration occurs in a minority component of the heterogenous lymphoid population and that a differentiation-specific restriction system may govern the expression of the MuLV. The data also suggest an etiological role for the ecotropic virus in regulation of IgA levels and life shortening. However, other physiological differences may also be related to viral burden [17,18] and the target cell for viral reintegration may be random leading to a variety of phenotypes. Studies are under way to determine if  $H-2^{n/n}$  is uniquely susceptible to the virus and if viral reintegration is associated with IgA phenotype and low survival of  $H-2^{n/n}$ progeny. If so, the viruses shed by the B10.F mothers and passed on to their offspring cause a deficient IgA humoral response that compromises survival.

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