BLOOD ACID-BASE BALANCE IN BROOK TROUT (*SALVELINUS FONTINALIS*)*

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SUMMARY

A nomogram is presented which allows the rapid assessment of acid-base status in brook trout (*Salvelinus fontinalis* Mitchill) at 18 °C.

Due to a P_{CO_2} of 3·5–4·3 mmHg in the respirometer water, the fish from which data were obtained for determination of zero base excess values were suffering from slight respiratory acidosis. Trauma of surgery and anaesthetization as well as confinement in the respirometer might also have generated a slight metabolic acidosis.

Base excess values for trout with a chronically implanted dorsal aortic cannula ranged from -3 to $+3$ m-equiv/l. Blood buffer values of $d[HCO_3^-]/d\text{pH} = 7·50 \pm 4·42$ slykes and $-d\log_{10} P_{CO_2}/d\,\text{pH} = 1·31 \pm 0·14$ were found. Mean plasma protein concentration was $3·6 \pm 0·8$ g/100 ml plasma and the buffering power of plasma was approximately 40% that of whole blood. Plasma proteins are more important blood buffers in fish than in humans.

INTRODUCTION

Various nomograms designed for assessment of acid-base balance in humans have been in use for a number of years. These nomograms, in their various forms, allow the derivation of a number of parameters pertaining to acid-base status from a few relatively easily measured factors such as blood pH, P_{CO_2}, and haemoglobin concentration ([Hb]) (Singer & Hastings, 1948; Siggaard-Andersen, 1962).

The aim of this research was the construction of a nomogram consisting of base excess (BE) and bicarbonate scales on a pH-P_{CO_2} co-ordinate system which could be used for the rapid assessment of acid-base status in brook trout (*Salvelinus fontinalis*). While producing the nomogram, we hoped to establish 'normal' ranges for some indicators of acid-base status such as BE, actual bicarbonate and standard bicarbonate in cannulated fish. Standard bicarbonate is a measure of what the plasma bicarbonate concentration would be at any given pH if P_{CO_2} was 'normal', and is used to assess the non-respiratory (or metabolic) component of blood acid-base status, as is BE. A further goal was to investigate blood buffering.

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* A portion of this work was submitted as a thesis by A. L. Sunkin to George Washington University in partial fulfilment of the requirements for the M.Sc. degree.

A. *Source and maintenance of fish*

The Pennsylvania Fish Commission Hatchery at Huntsdale, Pa. provided 44 trout of both sexes ranging in weight from 185 to 380 g. These trout are considered to be the Huntsdale strain (strain H). We also used seven strain O (wt 285–465 g) and three strain 15 (wt 235–485 g) trout of both sexes obtained from the Benner Spring Fish Research Station, Pennsylvania Fish Commission, Bellefonte, Pa. Strains 15 and O were derived from nine and seven generations of inbreeding, respectively, from the Bellefonte Open strain (Robinson *et al.* 1976).

Trout were maintained in the laboratory at $18 \pm 1 \cdot 5$ °C in a 680 l epoxy coated steel tank for at least 2 weeks prior to use in experiments. Approximately $8 \, \mathrm{lh^{-1}}$ of charcoal-filtered tap water flowed through the tank. Tank water was also recirculated through a glass wool and charcoal filter at a rate of approximately $7 \, \mathrm{lh^{-1}}$. An ultraviolet light was positioned over the filter. The fish were fed Purina Trout Chow three times per week and were exposed to the natural photoperiod. Trout were not fed for 48 h prior to an experiment. All experiments were conducted in a controlled temperature room at a water temperature of $18 \pm 0 \cdot 5$ °C.

B. *Instrumentation and techniques*

All measurements of blood pH and P_{CO_2} were made using a Radiometer Blood Micro System model BMS 2 Mk II in conjunction with a Radiometer PHM 71 Mk II acid-base analyser at $18 \pm 0 \cdot 1$ °C. This instrument was calibrated with Radiometer buffers S 1500 and S 1510 at pH 6·888 and 7·434. Buffer pH values were corrected to 18 °C using data provided by Radiometer. Blood samples were collected anaerobically and the pH of a 60 μl aliquot was measured. The original P_{CO_2} of the blood sample was then determined using the Astrup technique. Two 80 μl aliquots of the blood sample were equilibrated with water-saturated gases of 4% (Radiometer CCS) and 8% (Radiometer CCS) or 4% (Radiometer CCS) and 1·45% (Air Products) CO_2, 21% O_2, balance N_2. Following equilibration, the pH of each aliquot was measured and a pH–P_{CO_2} titration line was thus determined. When the Astrup technique is used the P_{CO_2} of the original sample, prior to equilibration, is determined by the position of the original pH on the pH-P_{CO_2} titration line which is plotted on a pH-P_{CO_2} co-ordinate system.

Blood [Hb] in gram percentage was determined by measuring cyanmethemoglobin using Hycel reagent and standards (Hycel, Inc.) and a Coleman Jr Spectrophotometer model 6A, following the methods of Blaxhall & Daisley (1973).

Dorsal aortic cannulations were done following the procedures of Smith & Bell (1964; 1967). Cannulae consisted of a 23-gauge Huber point syringe needle tip inserted in a length of PE-50 Intramedic polyethylene tubing. The cannulae were filled with heparinized (1000 i.u./ml) Courtland saline (Wolf, 1963). Following cannulation the fish were transferred to a closed respirometer system having a 35 l total capacity. The respirometer as well as the Radiometer blood gas system and the operating table used in cannulation were all situated in the controlled-temperature room at $18 \pm 0 \cdot 5$ °C. The aortic cannula was connected to a 23-gauge syringe needle tip which passed through a rubber stopper in the lid of the respirometer. A 6–10 cm

length of saline-filled PE-50 tubing was attached to the other end of this needle on the outside of the respirometer lid. Blood was sampled by releasing a clamp on this tubing. After a blood sample was taken, the blood remaining in the cannula was flushed back into the fish with heparinized saline. The exterior of the respirometer was painted so that the fish could not see the investigator. Cannulated fish were allowed to recover for at least 24 h prior to any data collection in all experiments. Houston, Czerwinski & Woods (1973) and Soivio, Nyholm & Huhti (1977) report that most physiological parameters altered by anaesthestization and cannulation stabilize after 24 h.

Haematocrit and plasma protein concentrations were measured in blood samples from cannulated trout. Haematocrit was determined by centrifugation of blood in capillary tubes at 11 500 rev/min for 5 min in an IEC model MB haematocrit centrifuge. Total protein concentration of 20 μl plasma samples was measured using the methods of Palma (1971).

C. *Determination of standard values of blood pH and P_{CO_2}*

Prior to the construction of the nomogram, it was necessary to establish values of pH and P_{CO_2} in dorsal aortic blood of cannulated trout which could be used to define the condition of zero BE. Aortic blood pH and P_{CO_2} were determined in anaerobically drawn blood samples from 19 trout with the dorsal aorta cannulated. Of this group, 9 were strain H, 7 were strain O, and 3 were strain 15.

D. *Construction of the nomogram*

The nomogram constructed was based on the curve nomogram of Siggaard-Andersen (1962). This form was chosen, rather than an alignment nomogram such as those of Singer & Hastings (1948) or Siggaard-Andersen (1963), because the curve nomogram is well suited for use in conjunction with the Astrup technique for P_{CO_2} determination. The pH-P_{CO_2} titration lines generated by the Astrup technique may be plotted directly on the pH-P_{CO_2} co-ordinate system of the curve nomogram. The original P_{CO_2} of the blood sample and the blood acid-base parameters given on the various nomogram scales may then be determined directly from the pH-P_{CO_2} titration line. In addition, the slope and position of the pH-P_{CO_2} titration line may be used to calculate blood buffer values.

The nomogram, consisting of BE and standard bicarbonate scales on a pH-P_{CO_2} axis system, was constructed following the methods of Siggaard-Andersen & Engel (1960) and Siggaard-Andersen (1962). Modifications of their methods are given in the following brief description of construction of the nomogram.

In order to construct the BE scale, blood was collected into a heparinized glass syringe from the dorsal aorta of 35 strain H trout which were anaesthetized in a solution of 50 mg/l of MS 222 (Crescent Research Chemicals). The blood of two or three fish was pooled to obtain sufficient quantities (6–7 ml) for titration. Blood was stored on ice for no more than 1 h until titrations and measurements were completed.

Each pool of blood was used in the titrations as whole blood, as plasma obtained by centrifugation, or as whole blood to which red blood cells had been added to produce a high [Hb]. Each pool was divided into a series of 490 μl aliquots, to which were added 12·5 μl of acid (HCl) or base (Na_2CO_3) solution. The amounts of acid added were

1, 3, 5, 7, 9, or 11 m-equiv/l of blood and the amounts of base were 1, 3, 5 or 7 m-equiv/l. Although the proportion of blood to acid or base added was the same as that used by Siggaard-Andersen (1962), we used smaller total volumes to conserve blood. A complete titration series consisted of three pools of blood of differing [Hb] to which the quantities of acid or base noted above were added. NaCl was added to the HCl and Na_2CO_3 solutions to give a total [Na^+] of 150 mM. Volumes of blood and acid-base solutions were measured using 1000 and 25 μl Hamilton syringes fitted with Chaney adaptors.

Following the addition of acid or base to a given blood or plasma sample, the pH was measured. Then, two 80 μl aliquots of blood were placed in the tonometers of the Radiometer apparatus and were equilibrated with water-saturated gases of 4% CO_2, 21% O_2, balance N_2 and 8% CO_2, 21% O_2, balance N_2. Following equilibration, the pH of each aliquot was measured and a pH-P_{CO_2} titration line was thus determined. The slope of this titration line is directly related to the [Hb] of the blood sample. Because of differences in slope resulting from differences in [Hb], the pH-P_{CO_2} lines intersect at a single point for aliquots of plasma, whole blood, and whole blood plus red blood cells to which, for example, 3 m-equiv/l of acid had been added. Any blood sample that had the same initial acid-base characteristics and to which 3 m-equiv/l of acid had been added should yield a pH-P_{CO_2} titration line which intersects the same point. Such intersection points were determined for each quantity of acid or base added. Four series of titrations were done, and yielded a minimum of three intersection points for each quantity of acid or base added.

All intersection points were then plotted and after standard values of arterial blood pH and P_{CO_2} were determined in cannulated trout, the BE curve was drawn on the nomogram. The BE value of a blood sample is read at the point the pH-P_{CO_2} titration line intersects the BE scale.

In addition to the BE curve, a standard bicarbonate scale was placed on the nomogram (see Results section).

From pH-P_{CO_2} titration lines plotted on the completed nomogram, we calculated two different blood buffer values. An assessment of non-bicarbonate buffering power was made using the relationship $d[HCO_3^-]/d$pH, which corresponds closely to $-\beta a(-\beta a = dB/d$pH$ + d[HCO_3^-]/d$ pH$ + 2d[CO_3^-]/d$ pH$)$ in titrations with carbon dioxide rather than with strong acids and bases. Due to this similarity we designated $d[HCO_3^-]/d$ pH as $-\beta_a^*$. We also calculated $-d \log_{10} P_{CO_2}/d$ pH which has been symbolized as $-\beta_{CO_2}$ (Burton, 1973). $-\beta_{CO_2}$ is a measure of the combined buffering power of bicarbonate and non-bicarbonate buffers.

RESULTS

Construction of the BE curve necessitated the establishment of 'normal' values for pH and P_{CO_2} of trout blood from the dorsal aorta at 18 °C in cannulated trout. Data from cannulated trout of all three strains yielded the following values: pH $= 7.797 \pm 0.065$ ($\bar{X} \pm$ S.D., $n = 19$); $P_{CO_2} = 5.2 \pm 1.1$ mmHg ($\bar{X} \pm$ S.D., $n = 18$). A one-way analysis of variance (Steel & Torrie, 1960) showed no significant differences in pH ($F = 1.50$; $P > 0.1$) and P_{CO_2} ($F = 1.818$; $P > 0.1$) among the three strains, so mean values were calculated by lumping data from fish of all three

Table 1. *Co-ordinates of points comprising the base excess scale*

Base excess (m-equiv/l)	Co-ordinates	
	pH	P_{CO_2}
− 11	7·662	1·94
− 9	7·694	2·70
− 7	7·717	3·28
− 5	7·742	3·85
− 3	7·766	4·43
− 1	7·789	4·98
0	7·797	5·20
+ 1	7·812	5·46
+ 3	7·842	5·98
+ 5	7·876	6·43
+ 7	7·917	6·70

strains. These values were then used to define the condition of zero BE. It must be pointed out that these trout were suffering from slight respiratory acidosis and perhaps metabolic acidosis as well. The P_{CO_2} of water in the respirometer ranged from 3·5 to 4·3 mmHg. This led to the relatively high blood P_{CO_2}. A slight metabolic acidosis may have been generated by the stresses of anaesthetization and surgery as well as by the stress of confinement in the respirometer. Determinations of true zero BE values could be obtained only from completely unstressed fish. After adjustment of all data from the acid or base titrations to these normal values, the intersection points of pH-P_{CO_2} titration lines were plotted for blood of different [Hb] to which a given amount of acid or base was added. A smooth curve, representing the BE curve, was fitted by eye through the point clusters (Fig. 1) and was placed on the nomogram (Fig. 2). The co-ordinates of points comprising the BE curve are given in Table 1. BE of any blood sample is then determined by the point of intersection of a pH-P_{CO_2} titration curve of the sample with the BE scale.

A standard bicarbonate scale was also constructed and placed on the nomogram. On this nomogram, standard bicarbonate is the plasma [HCO_3^-] in blood having a P_{CO_2} of 5·2 mmHg with the haemoglobin fully O_2 saturated. Using the Henderson–Hasselbalch equation, it is possible to calculate plasma [HCO_3^-] for any pH as follows:

$$[HCO_3^-] = \alpha_{CO_2}(P_{CO_2})\,(10^{pH-pK}).$$

From a plot of the data given by Severinghaus (1965), we determined that the solubility of CO_2 in human plasma at 18 °C is 0·0492 mmol/l per mmHg P_{CO_2}. Since trout plasma is similar in ionic strength to human, we felt safe using this value. Values for pK, which change with pH, were read from the nomogram of Severinghaus, Stupfel & Bradley (1956). Bicarbonate concentrations were calculated for pH ranging from 6·8 to 8·2 at a P_{CO_2} of 5·2 mmHg, and the scale was placed on the nomogram (Fig. 1). Standard bicarbonate values are read from the point of intersection of a pH-P_{CO_2} titration line of a blood sample with the standard bicarbonate scale.

The 'actual bicarbonate' value, i.e. the bicarbonate concentration of anaerobically drawn plasma, is found by connecting the original blood pH-P_{CO_2} point to the standard bicarbonate scale with a line having a slope equal to the slope of a pure bicarbonate buffer. The slope of an equilibrium line for a bicarbonate solution can be determined from calculation pH or P_{CO_2} values for a given constant bicarbonate
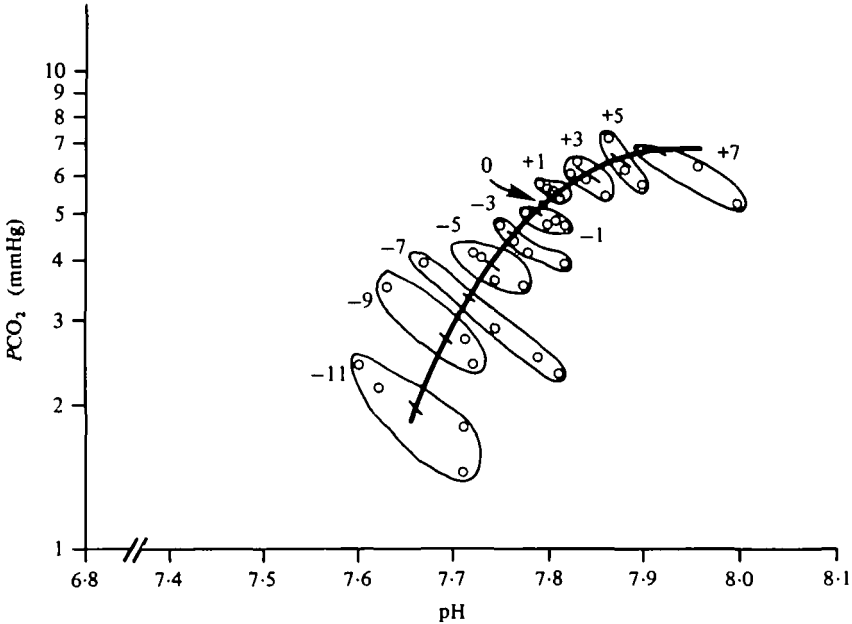
Fig. 1. Base excess curve. Individual points (open circles) represent intersection points of pH-P_{CO_2} titration lines of blood samples having varying [Hb], to which a known amount of acid or base had been added. All intersection points resulting from experiments in which a particular amount of acid or base was added are encircled and the amount of acid or base added (m-equiv/l) is shown.
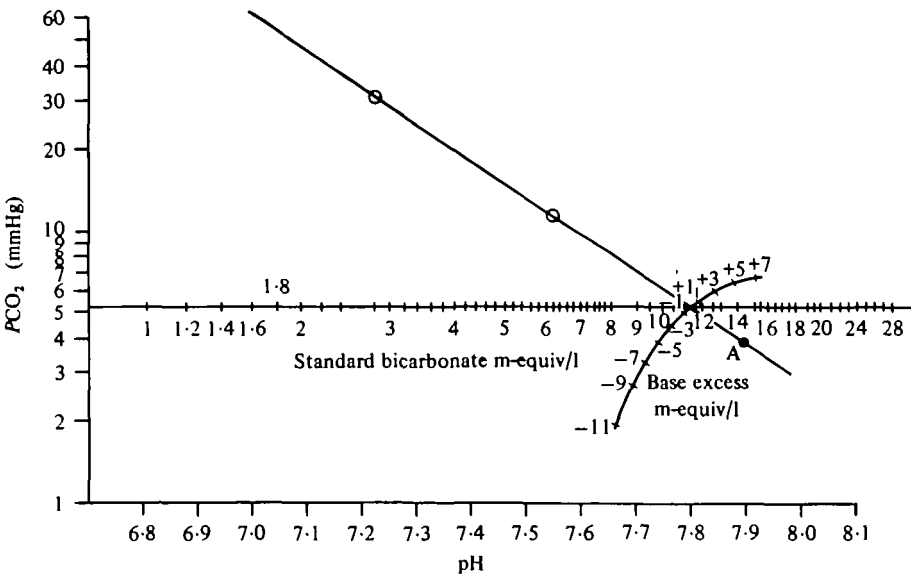

Fig. 2. The acid-base nomogram, with scales for standard bicarbonate and base excess; on which is shown a pH-P_{CO_2} titration line of a blood sample from a cannulated trout. Point A designates the original pH of the sample. Open circles indicate blood pH values after equilibration with 4 and 1·45 % CO_2.

concentration according to the modified Henderson–Hasselbalch equation (Siggaard-Andersen, 1964; Davenport, 1975).

$$\text{pH} = \text{p}K + \frac{\log[\text{HCO}_3^-]}{\alpha(P_{\text{CO}_2})},$$

where α is the solubility of CO_2. Rearranging,

$$\log P_{\text{CO}_2} = -\text{pH} + (\log[\text{HCO}_3^-] + \text{p}K - \log\alpha).$$

Plotting values of pH and P_{CO_2} obtained from this equation on the nomogram yielded a line having a slope of $-28°$. Therefore, to determine the actual bicarbonate value of any blood sample, a line with a slope of $-28°$ is drawn from the actual blood pH value to the standard bicarbonate scale. The intersection point is the actual bicarbonate value.

To illustrate the use of the nomogram, an example of a pH-P_{CO_2} titration line for a blood sample taken from a cannulated trout is shown on Fig. 2. Point A on the titration line designates the original pH of the sample, 7·881. From the nomogram, one can determine that original P_{CO_2} = 3·8 mmHg; standard bicarbonate = 11·4 m-equiv/l; base excess = 0; and actual bicarbonate = 10·7 m-equiv/l.

Table 2 shows mean values for various blood parameters for cannulated trout. These data were derived from not only the 19 cannulated trout which were used to determine the 'normal' pH and P_{CO_2} values needed to define zero BE, but also include data from an additional 21 cannulated trout used in experiments done after construction of the nomogram (see Packer, 1978).

The data were analysed for differences between strains using a one-way analysis of variance as described above. Values of F indicating no significant differences between strains were found for haematocrit, total plasma protein, base excess, and the two measures of blood buffering, $-\beta_a^*$ and $-\beta_{\text{CO}_2}$. F values indicating significant differences ($P < 0.05$) among strains were found for haemoglobin concentration, actual [HCO$_3^-$] and standard [HCO$_3^-$].

The application of Duncan's New Multiple Range Test (Steel & Torrie, 1960) indicated that [Hb] in strain H was lower ($P < 0.05$) than that of strain 15, but not strain O. There was no significant difference in [Hb] between strains O and 15. The standard and actual bicarbonate values for strain H were lower ($P < 0.05$) than both strains O and 15, but there were no differences between strains O and 15.

DISCUSSION

The nomogram we have constructed is valid only for brook trout at 18 °C having P_{CO_2} within the range of the fish used in this study. A further limitation is that, as may be seen in Fig. 1, the accuracy of the points on the BE curve decreases at numerically high BE values. Additionally we were unable to extend the BE curve beyond -11 m-equiv/l because the pH of acidified and CO_2 equilibrated blood was too low to be read on the expanded scale of the pH meter. Thus the BE scale of the nomogram is not useful for the assessment of severe acid-base disturbances. The standard bicarbonate scale may be used to assess the degree of metabolic disturbances in extreme conditions.

Table 2. *Values of various blood parameters for cannulated trout*

	Strain	Hematocrit (% cells)	[Hb] (g/100 ml blood)	Total protein (g/100 ml plasma)	$-\beta_a^*$ slykes	$-\beta_{CO_2}$	Actual [HCO$_3^-$] (m-equiv/l)	Standard [HCO$_3^-$] (m-equiv/l)	Base excess (m-equiv/l)
H	$\bar{X}$	33	6·8	3·7	6·76	1·305	10·3	10·4	−1·5
	S.D. (n)	±10 (18)	±2·0 (18)	±0·7 (7)	±5·03 (9)	±0·184 (9)	±1·8 (9)	±1·6 (9)	±2·3 (9)
	Range	20–60	4·2–9·9	2·6–5·0	1·23–15·40	1·109–1·730	7·2–13·2	7·6–13·0	−6 to +2
O	$\bar{X}$	37	7·4	3·7	9·12	1·349	12·3	12·1	+0·7
	S.D.	±5 (10)	±1·1 (10)	±1·0 (10)	±3·60 (6)	±0·107 (6)	±2·1 (6)	±1·8 (6)	±2·6 (6)
	Range	31–44	6·1–9·4	2·0–5·1	2·35–12·83	1·150–1·428	9·7–14·7	9·8–13·8	−3 to +3
15	$\bar{X}$	38	8·4	3·5	6·46	1·269	12·2	12·3	+1·3
	S.D.	±10 (12)	±1·9 (12)	±0·8 (7)	±1·92 (3)	±0·065	±1·2 (3)	±0·7 (3)	±1·2 (3)
	Range	20–54	6·0–11·9	2·1–4·3	4·63–8·45	1·212–1·340	10·9–13·3	11·5–12·7	0 to +2·5
	Pooled $\bar{X}$ ±S.D.	35±10	—	3·6±0·8	7·50±4·42	1·134±0·144	—	—	0·0±2·5

Table 3. *A comparison of values of arterial blood P_{CO_2} found in this study with those reported by other investigators*

Blood pH	Blood P_{CO_2} (mmHg)	Temp. (°C)	Species	Reference
7·83	1·01	15	Salmo gairdneri	Eddy, 1976
7·93	2	8·9	S. gairdneri	Janssen & Randall, 1975
7·5–7·75	1–1·5	15±1	S. gairdneri	Holeton & Randall, 1967
8·02	1·91	13±1	S. gairdneri	Cameron & Randall, 1972
7·82–7·85	—	20	S. gairdneri	Randall & Cameron, 1973
7·74	4·3	10±0·5	S. gairdneri (group I, Aug.)	Soivio et al. 1977
7·86	3·1	10±0·5	S. gairdneri (group II, Nov.)	Soivio et al. 1977
7·81	3·6	10±0·5	S. gairdneri (group III, Nov.)	Soivio et al. 1977
7·785	3·2	?	Salvelinus fontinalis	Milne & Randall, 1976
7·797	5·2	18	S. fontinalis	This study

The average pH of dorsal aortic blood from trout in neutral environments was 7.797 ± 0.065. Taking temperature differences into consideration, it can be seen in Table 3 that this value is within the expected range. Aortic blood P_{CO_2} (5·2 mmHg), on the other hand, is higher than values for rainbow trout reported by other workers (Table 3). The respirometer system used in this study was closed, and despite vigorous air bubbling in the reservoir, the P_{CO_2} of water entering the respirometer ranged from approximately 3·5 to 4·3 mmHg. Therefore the high P_{CO_2} of inspired water resulted in a blood P_{CO_2} that was high, but within a range easily tolerated by trout (Eddy & Morgan, 1969). Due to the elevated blood P_{CO_2}, actual plasma [HCO$_3^-$] (11·3 m-equiv/l) was higher than that reported by other workers for trout (Table 3).

BE values ranged from -6 to $+3$ m-equiv/l in fish in neutral environments, with most values falling between -3 and $+3$ m-equiv/l. The normal range of BE values in humans is -3 to $+3$ m-equiv/l (Tietz, 1976), indicating a similarity in control of the range of non-volatile acid concentrations in the blood of the two species.

We found blood buffer values of $-\beta_a^* = 7.50 \pm 4.42$ slykes and $-\beta_{CO_2}^* = 1.31 \pm 0.14$. In order to make comparisons with the data of other workers we are assuming that $-dC_{CO_2}/d$ pH (C_{CO_2} = total CO$_2$ content) would be only slightly greater than $-\beta_a^*$ for the same blood sample (Burton, 1973). As may be seen in Table 4, our buffer values are somewhat lower than those reported by other workers for rainbow trout. No obvious reasons for the differences, such as large differences in [Hb], are evident.

Average plasma protein concentration was 3.6 ± 0.8 g/100 ml plasma, and apparently plasma protein contributes significantly to blood buffering. $-\beta_a^*$ and $-\beta_{CO_2}$ were calculated from CO$_2$ titrations of seven separated plasma samples. $-\beta_a^* = 3.25 \pm 0.95$ slykes/l of plasma ($\bar{X} \pm$ S.D.), showing that the average non-bicarbonate buffering capacity of plasma was about 40% that of whole blood. In humans, the non-bicarbonate buffering power of plasma is about 24% that of whole blood (Tietz, 1976). The average [Hb] of whole blood samples for which $-\beta_a^*$ was determined was 73 g/l of blood, and the $-\beta_a^*$ was 7.5 ± 4.42 slykes/l ($\bar{X} \pm$ S.D.). Assuming an average *Hct* of 35%, and a plasma protein concentration of 36 g/l of plasma, there were 23 g plasma protein/l of blood. If the factor $-\beta_a^*$ for whole blood is divided by the g/l of plasma protein plus [Hb] in whole blood, a value of 0·08 slykes/g protein l^{-1} is obtained. The non-bicarbonate buffering power $(-\beta_a^*)$ of plasma, expressed as slykes/g protein l^{-1} = 0·09. A comparison of these two values indicates that plasma proteins and Hb have similar buffering power per unit weight.

$-\beta_{CO_2}$, which is a measure of both bicarbonate and non-bicarbonate buffering power, was 1.18 ± 0.04 in plasma, whereas in whole blood $-\beta_{CO_2}$ was 1.31 ± 0.14. Therefore, even though blood [Hb] is quite variable and leads to great variability in $-\beta_a^*$ in trout blood, changes in Hb content have a smaller effect on total blood buffering power (as measured by $-\beta_{CO_2}$) than might be expected.

No physiological significance could be ascribed to the differences in [Hb] and standard and actual bicarbonate values which were found between strain H and strains O and 15. A number of workers (Robinson *et al.* 1976; Falk & Dunson, 1977; Swarts, 1978) found strain 15 trout to be much more acid tolerant than strain O, and we had originally hoped to find a physiological basis for the difference in tolerance. We

Table 4. *A comparison of blood buffer values found by us and those reported by other investigators*

Species	Temp. (°C)	[Hb] (g/100 ml)	$-dC_{CO_2}/d\,pH$	$-\beta^*$	$-\beta_{CO_2}$	Reference
Salmo gairdneri	6	8 (estimate)	11·9	—	1·36	Eddy, 1974
Salmo gairdneri	15	8 (estimate)	10·0	—	2·14	Eddy, 1974
Salmo gairdneri	20	8 (estimate)	10·3	—	2·72	Eddy, 1974
Salmo gairdneri	15	—	8·47	—	2·48	Eddy, 1976
Salmo gairdneri	13	—	—	—	1·68	Cameron & Randall, 1972
Salvelinus fontinalis	18	7·4	—	7·50	1·31	This study

'were not successful. The fact that [Hb] and bicarbonate values were lower in strain H could not be interpreted in terms of resistance to acid environments since this strain had not been tested for acid tolerance as strains O and 15 had been.

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