

# Appendix A

## AP BIOLOGY EQUATIONS AND FORMULAS

| STATISTICAL ANALYSIS AND PROBABILITY   |      |   |       |       |                   |       |       |       |  |
|--|------|---|-------|-------|-------------------|-------|-------|-------|--|
| Standard Error   |      | Mean  |       |       |                   |       |       |       |  |
| $SE_{\bar{x}} = \frac{s}{\sqrt{n}}$  |      | $\bar{x} = \frac{1}{n} \sum_{i=1}^n x_i$                |       |       |                   |       |       |       |  |
| Standard Deviation   |      | Chi-Square  |       |       |                   |       |       |       |  |
| $s = \sqrt{\frac{\sum (x_i - \bar{x})^2}{n-1}}$  |      | $\chi^2 = \sum \frac{(o-e)^2}{e}$                       |       |       |                   |       |       |       |  |
| CHI-SQUARE TABLE   |      |   |       |       |                   |       |       |       |  |
| Degrees of Freedom   |      |   |       |       |                   |       |       |       |  |
| p  | 1    | 2   | 3     | 4     | 5                 | 6     | 7     | 8     |  |
| 0.05   | 3.84 | 5.99  | 7.82  | 9.49  | 11.07             | 12.59 | 14.07 | 15.51 |  |
| 0.01   | 6.64 | 9.32  | 11.34 | 13.28 | 15.09             | 16.81 | 18.48 | 20.09 |  |
| LAWS OF PROBABILITY  |      |   |       |       | METRIC PREFIXES   |       |       |       |  |
| If A and B are mutually exclusive, then $P(A \text{ or } B) = P(A) + P(B)$   |      |   |       |       | Factor            |       |       |       |  |
| If A and B are independent, then $P(A \text{ and } B) = P(A) \times P(B)$  |      |   |       |       | Prefix            |       |       |       |  |
|  |      |   |       |       | Symbol            |       |       |       |  |
| <b>HARDY-WEINBERG EQUATIONS</b>  |      |   |       |       | 10 <sup>9</sup>   |       |       |       |  |
| $p^2 + 2pq + q^2 = 1$  |      | $p$ = frequency of the dominant allele in a population  |       |       | giga              |       |       |       |  |
| $p + q = 1$  |      | $q$ = frequency of the recessive allele in a population |       |       | 10 <sup>6</sup>   |       |       |       |  |
|  |      |   |       |       | mega              |       |       |       |  |
|  |      |   |       |       | 10 <sup>3</sup>   |       |       |       |  |
|  |      |   |       |       | kilo              |       |       |       |  |
|  |      |   |       |       | 10 <sup>-2</sup>  |       |       |       |  |
|  |      |   |       |       | centi             |       |       |       |  |
|  |      |   |       |       | 10 <sup>-3</sup>  |       |       |       |  |
|  |      |   |       |       | milli             |       |       |       |  |
|  |      |   |       |       | 10 <sup>-6</sup>  |       |       |       |  |
|  |      |   |       |       | micro             |       |       |       |  |
|  |      |   |       |       | 10 <sup>-9</sup>  |       |       |       |  |
|  |      |   |       |       | nano              |       |       |       |  |
|  |      |   |       |       | 10 <sup>-12</sup> |       |       |       |  |
|  |      |   |       |       | pico              |       |       |       |  |
| Mode = value that occurs most frequently in a data set   |      |   |       |       |                   |       |       |       |  |
| Median = middle value that separates the greater and lesser halves of a data set                                   |      |   |       |       |                   |       |       |       |  |
| Mean = sum of all data points divided by number of data points   |      |   |       |       |                   |       |       |       |  |
| Range = value obtained by subtracting the smallest observation (sample minimum) from the greatest (sample maximum) |      |   |       |       |                   |       |       |       |  |

$s$  = sample standard deviation (i.e., the sample based estimate of the standard deviation of the population)  
 $\bar{x}$  = mean  
 $n$  = size of the sample  
 $o$  = observed individuals with observed genotype  
 $e$  = expected individuals with observed genotype

Degrees of freedom equals the number of distinct possible outcomes minus one.

| RATE AND GROWTH   |  | Water Potential ( $\Psi$ )   |
|---|--|--|
| <b>Rate</b><br>$dY/dt$<br><b>Population Growth</b><br>$dN/dt=B-D$<br><b>Exponential Growth</b><br>$\frac{dN}{dt} = r_{\max}N$<br><b>Logistic Growth</b><br>$\frac{dN}{dt} = r_{\max}N\left(\frac{K-N}{K}\right)$  | $dY$ = amount of change<br>$t$ = time<br>$B$ = birth rate<br>$D$ = death rate<br>$N$ = population size<br>$K$ = carrying capacity<br>$r_{\max}$ = maximum per capita growth rate of population   | $\Psi = \Psi_p + \Psi_s$<br>$\Psi_p$ = pressure potential<br>$\Psi_s$ = solute potential<br>The water potential will be equal to the solute potential of a solution in an open container, since the pressure potential of the solution in an open container is zero.<br><b>The Solute Potential of the Solution</b><br>$\Psi_s = -iCRT$<br>$i$ = ionization constant (For sucrose this is 1.0 because sucrose does not ionize in water.)<br>$C$ = molar concentration<br>$R$ = pressure constant ( $R = 0.0831$ liter bars/mole K)<br>$T$ = temperature in Kelvin ( $273 + ^\circ\text{C}$ ) |
| <b>Temperature Coefficient <math>Q_{10}</math></b><br>$Q_{10} = \left(\frac{k_2}{k_1}\right)^{\frac{10}{t_2-t_1}}$<br><b>Primary Productivity Calculation</b><br>$\text{mg O}_2/\text{L} \times 0.698 = \text{mL O}_2/\text{L}$<br>$\text{mL O}_2/\text{L} \times 0.536 = \text{mg carbon fixed/L}$   | $t_2$ = higher temperature<br>$t_1$ = lower temperature<br>$k_2$ = metabolic rate at $t_2$<br>$k_1$ = metabolic rate at $t_1$<br>$Q_{10}$ = the <i>factor</i> by which the reaction rate increases when the temperature is raised by ten degrees |  |
| SURFACE AREA AND VOLUME   |  | Dilution – used to create a dilute solution from a concentrated stock solution   |
| <b>Volume of a Sphere</b><br>$V = 4/3 \pi r^3$<br><b>Volume of a Cube (or Square Column)</b><br>$V = l w h$<br><b>Volume of a Column</b><br>$V = \pi r^2 h$<br><b>Surface Area of a Sphere</b><br>$A = 4 \pi r^2$<br><b>Surface Area of a Cube</b><br>$A = 6 a$<br><b>Surface Area of a Rectangular Solid</b><br>$A = \Sigma$ (surface area of each side) | $r$ = radius<br>$l$ = length<br>$h$ = height<br>$w$ = width<br>$A$ = surface area<br>$V$ = volume<br>$\Sigma$ = Sum of all<br>$a$ = surface area of one side of the cube   | $C_i V_i = C_f V_f$<br>$i$ = initial (starting)<br>$C$ = concentration of solute<br>$f$ = final (desired)<br>$V$ = volume of solution<br><b>Gibbs Free Energy</b><br>$\Delta G = \Delta H - T\Delta S$<br>$\Delta G$ = change in Gibbs free energy<br>$\Delta S$ = change in entropy<br>$\Delta H$ = change in enthalpy<br>$T$ = absolute temperature (in Kelvin)<br><b>pH = <math>-\log [H^+]</math></b>  |