The Series of a Four-node Motif can Provide Sensitive Detection over Arbitrary Range of Signal, thereby Explain Weber's Law, and Compute Logarithm

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Abstract

Weber's law states that the ratio of the smallest perceptual change in a signal and the background signal is constant. The law is observed across the perception of weight, light intensity, and sound intensity and pitch. Two models of perception have been proposed to explain Weber's law, namely the logarithmic and the linear model. This paper argues in favour of the linear model, which requires the sensory perception system to generate a linear signal-response relationship over several orders of magnitude. To this end, a four-node motif is constructed from first principles which in series provides a linear relationship between signal and response over arbitrary range of signal. This series provides a general mechanism to sensitively detect signal over arbitrary range. Such a series also provides a general basis for a class of bow-tie architecture where the number of receptors is much lower than the range of signal and response. Two examples of this class of bow-tie architecture are numerosity processing and absorption spectra of cone opsins of humans. Further, the series can also compute logarithm over arbitrary range of signal.

20 **Keywords**: Sensitive detection, Weber's law, linear model, bow-tie architecture, numerosity
21 processing, human cone cell, logarithm

22 1. Introduction

In the papers [1,2], Weber's law is formulated as the outcome of a system generating equal

maximum amplitude as a response to an input if the ratio of the current input to the background

signal is constant. However, it is doubtful that the processes which show Weber's law involve

scale-invariance with respect to the maximum amplitude. For example, consider the case of

weight perception. Suppose the just noticeable difference (JND) for weight perception is 5% of

the background, so that a subject holding a weight of 100 gm can feel the change in the weight

if at least 5 gm of weight is added. The JND when the subject is holding 1kg of weight is 50

gm. Although the ratio of the JND to the background signal is constant, the subject is able to

sense the background weight and the additional weight in the absolute sense. The subject is

not, if at all, just calculating the ratio of the two signals.

It has also been claimed that fold-change detection (FCD) entails Weber's law [1]. A system

capable of providing FCD also shows exact adaptation, in which, as a response to a constant

signal, the system generates an output which returns to the resting value of the system. That

FCD implies exact adaptation can be seen in the following way. Consider a system equilibrated

at a stimulus X_0 produces a constant output of Z_0. It generates a certain output when a stimulus

X_1 is provided. The ratio X_1/X_0 being the fold change, f. If the system does not show exact

adaptation, the resting output of the system, equilibrated at stimulus X_1 (or fX_0), will not be

equal to Z_0. If the system is capable of providing FCD, it will show identical response when the

two stimuli are replaced by kX_0 and kX_1, respectively, where k is a scalar. Clearly, the output

of the system equilibrated at stimulus kX_0 should be Z_0. This, then, leads to a contradiction if k

is replaced with f. That is, in the latter scenario the resting output of the system when stimulus

is fX_0 is Z_0, whereas in the former scenario it is not equal to Z_0. Hence, a system which shows

FCD will also show exact adaptation.

However, it is doubtful that the processes which show Weber's law exhibit exact adaptation.

Again considering the case of weight perception, a subject holding a certain amount of weight

continues to perceive the same amount of signal; that is, the perception of the weight does not

come back to the previous equilibrium value. As an example, the intensity of perception of two

different weights (say, 100 gm and 1 kg) does not come back to a previous value after some

time. The same argument applies also to the perception of sound intensity and brightness.

Figure 1: (**A**) Two pairs of rectangular strips which differ by the same length. (**B**) Two pairs of collections of discs numbering (i) 4 and 5; (ii) 24 and 25. (**C**) Two pairs of pseudo-randomly placed dots numbering (i) 50 and 100; (ii) 500 and 550

Weber's law is also observed in size (length/area/volume) perception, for which three examples

are presented here. The first example demonstrates that the detection of difference between

the length of two rectangular strips becomes more difficult with the increase in their lengths

keeping the difference constant (Figure 1A). The second example shows that the human eye can

easily notice the difference of one disc between collections of 4 and 5 such identical discs on

a quick glance, whereas doing so is virtually impossible between collections of 24 ad 25 discs

(Figure 1B). The third example is that of dot counting. As the figures 1C-F show, the difference

between two pictures containing pseudo-randomly placed 50 and 100 dots, respectively, is

noticeable, but that between two pictures containing 500 and 550, respectively, is not. This

example may look like the previous one, but here since the ratio of the dot's size to the total

area is much smaller, the brain is probably estimating empty spaces between the dots. The

size reduction in the empty spaces between the figures 1C and 1D is much greater than that

between the figures 1E and 1F. Hence, the third example is a two-dimensional analogue of the

first example. Weber's law is also observed in numerosity processing [3]. Because it is not

clear how scale-invariance with respect to the maximum amplitude is present in size perception

and numerosity processing; arguably, a general mechanism which yields Weber's law is needed

which does not include scale-invariance with respect to the maximum amplitude or involves

exact adaptation. Generalizing the example of numerosity processing, such a mechanism

should be able to generate linear signal-response relationship over several orders of magnitude.

2.   Results

2.1.   Linear Signal-Response Relationship Over Arbitrary Range

In order for a system to respond sensitively throughout the whole range of signal, the ideal

relationship between a signal and the response should be a straight line with the slope of one.

Such a formulation is presented now. Because sensory systems operate over several orders of

magnitude, the formulation should be able to generate linear signal-response relationship over

several orders of magnitude. This is achieved in the following way. Consider a collection of five

receptors which bind a certain ligand, representing the signal, with the equilibrium dissociation

constant (K_{da}) equal to 10^1, 10^2, 10^3, 10^4, and 10^5 arb. units, respectively. The signal, X, causes

the *Z* producer (an imaginary generic form of gene) to produce Z. In order to have sensitive

detection, that receptor should contribute to the output, Z, whose K_{da} is an order of magnitude

larger than the strength of the signal (the concentration of X). Because the receptors meant to

respond to signal of higher strength have lower binding affinities (larger K_{da}), their contribution

to the output (or the production rate of Z, β_1) should be proportionally larger.

$$\frac{dZ}{dt} = \frac{\beta_1 \dfrac{X}{Kd_{XZ}}}{1 + \dfrac{X}{Kd_{XZ}}} - \alpha_1 Z \tag{1}$$

87 where, β_1 is the rate constant for production of Z by X. α_1 is the degradation rate of Z. Kd_{XZ} is

88 the equilibrium dissociation constant for the binding of X to the producer element (an imaginary

89 generic form of promoter) of Z, which causes Z production.

91 The equilibrium value of Z is

$$Z = \left[\frac{\dfrac{\beta_1}{\alpha_1} \dfrac{X}{Kd_{XZ}}}{1 + \dfrac{X}{Kd_{XZ}}} \right] \tag{2}$$

92 Choosing appropriate values of β_1/α_1 and Kd_{XZ}, the total output from the five receptors is

$$Z = \left[\frac{10\dfrac{X}{10^1}}{1 + \dfrac{X}{10^1}} \right] + \left[\frac{10^2\dfrac{X}{10^2}}{1 + \dfrac{X}{10^2}} \right] + \left[\frac{10^3\dfrac{X}{10^3}}{1 + \dfrac{X}{10^3}} \right] + \left[\frac{10^4\dfrac{X}{10^4}}{1 + \dfrac{X}{10^4}} \right] + \left[\frac{10^5\dfrac{X}{10^5}}{1 + \dfrac{X}{10^5}} \right] \tag{3}$$

Let's assume that the value of the signal is of the order of magnitude of two. The receptor

with Kd_{xz} equal to 10^3 is ought to generate output from the signal of this value. However, the

receptors with Kd_{xz} equal to 10^1 and 10^2 arb. units, respectively, saturated by the signal, also

contribute to the output. Thus, output from the receptors whose Kd_{xz} is either equal to or less

than the order of magnitude of the signal should be removed. This is achieved by adding to the

motif a "negative arm": the signal X engenders production of the molecule Y which suppresses

the output Z from such receptors. The signal X and the molecule Y bind to the producer element

of Z in mutually exclusive manner. The resulting motif is type-1 incoherent feed-forward loop.

Parameters of the negative arm are such that it suppresses the output mainly from the receptors

whose Kd_{xz} is either equal to or less than the order of magnitude of the signal.

$$\frac{dY}{dt} = \frac{\beta_2 \frac{X}{Kd_{XY}}}{1 + \frac{X}{Kd_{XY}}} - \alpha_2 Y \tag{4}$$

$$\frac{dZ}{dt} = \left[\frac{\beta_1 \frac{X}{Kd_{XZ}}}{1 + \frac{X}{Kd_{XZ}}} \right] \cdot \left[\frac{1}{1 + \left(\frac{Y}{Kd_{YZ}}\right)^n} \right] - \alpha_1 Z \tag{5}$$

103 where, β_2 is the rate constant for production of Y by X. α_2 is the degradation rate of Y. Kd_{XY} is

104 the equilibrium dissociation constant for the binding of X to the producer element of Y, which

105 causes Y production. Kd_{YZ} is the equilibrium dissociation constant for the binding of Y to the

106 producer element of Z, leading to suppression of Z production.

107 The equilibrium value of Y and Z are

$$Y = \frac{\frac{\beta_2}{\alpha_2} \frac{X}{Kd_{XY}}}{1 + \frac{X}{Kd_{XY}}} \tag{6}$$

$$Z = \left[\frac{\frac{\beta_1}{\alpha_1} \frac{X}{Kd_{XZ}}}{1 + \frac{X}{Kd_{XZ}}} \right] \cdot \left[\frac{1}{1 + \left(\frac{\left(\frac{1}{Kd_{YZ}} \frac{\beta_2}{\alpha_2}\right) \frac{X}{Kd_{XY}}}{1 + \frac{X}{Kd_{XY}}} \right)^n} \right] \tag{7}$$

The expression in the second pair of brackets on the right hand side of the equality in equation

7, representing suppression of the output by the signal, reaches an asymptotic value because the

denominator in the expression reaches an asymptotic value. On the other hand, the formulation

demands the expression to keep decreasing with increasing X. That is, the expression should

be of the form: $1/(1+X)^n$. This is achieved in the following way. The expression in the second

pair of brackets on the right hand side of the equality in equation 7 is approximately equal to

the expression 8, which is of the required form, at least in the parameter range of interest if the

order of magnitude of $\left(\dfrac{1}{Kd_{YZ}}\dfrac{\beta_2}{\alpha_2}\right)$ is at least two (Figure 2A). This is because X/Kd_{XY} remains

much less than 1 until the saturation has occurred.

$$\left[\frac{1}{1+\left(\left(\dfrac{1}{Kd_{YZ}}\dfrac{\beta_2}{\alpha_2}\right)\dfrac{X}{Kd_{XY}}\right)^n}\right] \tag{8}$$

$$\left[\frac{1}{1+\left(\dfrac{\left(\dfrac{1}{Kd_{YZ}}\dfrac{\beta_2}{\alpha_2}\right)\dfrac{X}{Kd_{XY}}}{1+\dfrac{X}{Kd_{XY}}}\right)^n}\right] \approx \left[\frac{1}{1+\left(\left(\dfrac{1}{Kd_{YZ}}\dfrac{\beta_2}{\alpha_2}\right)\dfrac{X}{Kd_{XY}}\right)^n}\right] \tag{9}$$

Figure 2: (**A**) The blue curve represents the expression $1/(1 + 10((x/100)/(1 + (x/100))))$. The green curve: $1/(1 + 100((x/1000)/(1 + (x/1000))))$. The red curve: $1/(1 + (x/10))$. (**B**) The blue curve represents the expression $10((x/100)/(1+(x/100)))/(1+10((x/100)/(1+(x/100))))$. The green curve: $100((x/1000)/(1 + (x/1000)))/(1 + 100((x/1000)/(1 + (x/1000))))$. The red curve: $(x/10)/(1 + (x/10))$.

117 To fulfill the aforementioned demand on the value of parameters of the negative arm, $Kd_{YZ}\left(\dfrac{\alpha_2}{\beta_2}\right)Kd_{XY}$,

118 the effective equilibrium dissociation constant of this arm, is set equal to Kd_{XZ}. The total output

119 from the five receptors is

120
$$Z = \left[\frac{10^1\,\dfrac{X}{10^1}}{1+\dfrac{X}{10^1}}\right]\cdot\left[\frac{1}{1+\dfrac{X}{10^1}}\right] + \left[\frac{10^2\,\dfrac{X}{10^2}}{1+\dfrac{X}{10^2}}\right]\cdot\left[\frac{1}{1+\dfrac{X}{10^2}}\right] + \left[\frac{10^3\,\dfrac{X}{10^3}}{1+\dfrac{X}{10^3}}\right]\cdot\left[\frac{1}{1+\dfrac{X}{10^3}}\right]$$

$$+ \left[\frac{10^4\,\dfrac{X}{10^4}}{1+\dfrac{X}{10^4}}\right]\cdot\left[\frac{1}{1+\dfrac{X}{10^4}}\right] + \left[\frac{10^5\,\dfrac{X}{10^5}}{1+\dfrac{X}{10^5}}\right]\cdot\left[\frac{1}{1+\dfrac{X}{10^5}}\right] \tag{10}$$

121 The following is non-approximated form of the above equation

$$Z = \left[\frac{10^1\,\dfrac{X}{10^1}}{1+\dfrac{X}{10^1}}\right]\cdot\left[\frac{1}{1+\dfrac{10^2\,\dfrac{X}{10^3}}{1+\dfrac{X}{10^3}}}\right] + \left[\frac{10^2\,\dfrac{X}{10^2}}{1+\dfrac{X}{10^2}}\right]\cdot\left[\frac{1}{1+\dfrac{10^2\,\dfrac{X}{10^4}}{1+\dfrac{X}{10^4}}}\right] + \left[\frac{10^3\,\dfrac{X}{10^3}}{1+\dfrac{X}{10^3}}\right]\cdot\left[\frac{1}{1+\dfrac{10^2\,\dfrac{X}{10^5}}{1+\dfrac{X}{10^5}}}\right] +$$

$$\left[\frac{10^4\,\dfrac{X}{10^4}}{1+\dfrac{X}{10^4}}\right]\cdot\left[\frac{1}{1+\dfrac{10^2\,\dfrac{X}{10^6}}{1+\dfrac{X}{10^6}}}\right] + \left[\frac{10^5\,\dfrac{X}{10^5}}{1+\dfrac{X}{10^5}}\right]\cdot\left[\frac{1}{1+\dfrac{10^2\,\dfrac{X}{10^7}}{1+\dfrac{X}{10^7}}}\right] \tag{11}$$

However, a signal can generate output from those receptors also whose Kd_{xz} is larger than the

signal by more than one order of magnitude. In the current case, this means that the signal of

10^2 arb. units being able to generate output from the receptors whose Kd_{xz} is equal to 10^4 and

10^5 arb. units, respectively. Output generated by each of the two receptors is approximately

equal to 10^2 arb. units, which is of the same order as output generated by the receptor whose

Kd_{xz} is equal to 10^3 arb. units. Output by the receptors whose Kd_{xz} is larger than the signal

by more than one order of magnitude is removed by adding to the motif a "positive arm": the

signal X engenders production of the molecule V, which must also be bound along with X to

the producer element of Z in order to engender Z production. Parameters of the positive arm

are such that it lets mainly those receptors generate the output whose Kd_{xz} is at most one order

of magnitude larger than the signal.

$$\frac{dV}{dt} = \frac{\beta_3 \frac{X}{Kd_{XV}}}{1 + \frac{X}{Kd_{XV}}} - \alpha_3 V \tag{12}$$

$$\frac{dY}{dt} = \frac{\beta_2 \frac{X}{Kd_{XY}}}{1 + \frac{X}{Kd_{XY}}} - \alpha_2 Y \tag{13}$$

$$\frac{dZ}{dt} = \left[\frac{\beta_1 \frac{X}{Kd_{XZ}}}{1 + \frac{X}{Kd_{XZ}}} \right] \cdot \left[\frac{\left(\frac{V}{Kd_{VZ}}\right)^m}{1 + \left(\frac{V}{Kd_{VZ}}\right)^m} \right] \cdot \left[\frac{1}{1 + \left(\frac{Y}{Kd_{YZ}}\right)^n} \right] - \alpha_1 Z \tag{14}$$

where, β_1, β_2, and β_3 are the rate constants for production of Z, Y, an V, respectively, by X.

α_1, α_2, and α_3 are the degradation rates of Z, Y, and V, respectively. Kd_{XZ}, Kd_{XY}, and Kd_{XV}

are the equilibrium dissociation constants for the binding of X to the element of Z, Y, and V,

respectively, which causes production of Z, Y, and V, respectively. Kd_{YZ} and Kd_{VZ} are the

equilibrium dissociation constants for the binding of Y and V, respectively, to the element of Z,

which causes Z production.

The equilibrium value of V, Y, and Z

$$V = \frac{\frac{\beta_3}{\alpha_3} \frac{X}{Kd_{XV}}}{1 + \frac{X}{Kd_{XV}}} \tag{15}$$

$$Y = \frac{\dfrac{\beta_2}{\alpha_2} \dfrac{X}{Kd_{XY}}}{1 + \dfrac{X}{Kd_{XY}}} \qquad (16)$$

$$Z = \left[\frac{\dfrac{\beta_1}{\alpha_1} \dfrac{X}{Kd_{XZ}}}{1 + \dfrac{X}{Kd_{XZ}}} \right] \cdot \left[\frac{\left(\dfrac{\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}}}{1 + \dfrac{X}{Kd_{XV}}} \right)^m}{1 + \left(\dfrac{\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}}}{1 + \dfrac{X}{Kd_{XV}}} \right)^m} \right] \cdot \left[\frac{1}{1 + \left(\dfrac{\left(\dfrac{1}{Kd_{YZ}} \dfrac{\beta_2}{\alpha_2} \right) \dfrac{X}{Kd_{XY}}}{1 + \dfrac{X}{Kd_{XY}}} \right)^n} \right] \qquad (17)$$

The formulation demands that the expression in the second pair of brackets on the right hand

side of the equality in equation 17, representing the generation of the output by the signal X

and the molecule V according to the AND logic, should be of the form $X^n/(1 + X)^n$. This is

achieved in the following way. As previously, the expression in the second pair of brackets on

the right hand side of the equality in equation 17 is approximately equal to the expression 18

at least in the parameter range of interest if the order of magnitude of $\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right)$ is at least

two (Figure 2B). This is because X/Kd_{XV} remains much less than 1 until the saturation has

occurred.

$$\left[\frac{\left(\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}} \right)^m}{1 + \left(\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}} \right)^m} \right] \qquad (18)$$

$$\left[\frac{\left(\dfrac{\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}}}{1 + \dfrac{X}{Kd_{XV}}} \right)^m}{1 + \left(\dfrac{\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}}}{1 + \dfrac{X}{Kd_{XV}}} \right)^m} \right] \approx \left[\frac{\left(\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}} \right)^m}{1 + \left(\left(\dfrac{1}{Kd_{VZ}} \dfrac{\beta_3}{\alpha_3} \right) \dfrac{X}{Kd_{XV}} \right)^m} \right] \qquad (19)$$

151 To fulfill the aforementioned demand on the value of parameters of the positive arm, $Kd_{VZ}\left(\dfrac{\alpha_3}{\beta_3}\right)Kd_{XV}$,

152 the effective equilibrium dissociation constant of this arm, is set equal to $Kd_{XZ}/10$. The total

153 output from the five receptors is

$$
Z = \left[\frac{10^1\frac{X}{10^1}}{1+\frac{X}{10^1}}\right]\cdot\left[\frac{\frac{X}{10^0}}{1+\frac{X}{10^0}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^1}}\right] + \left[\frac{10^2\frac{X}{10^2}}{1+\frac{X}{10^2}}\right]\cdot\left[\frac{\frac{X}{10^1}}{1+\frac{X}{10^1}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^2}}\right] + \left[\frac{10^3\frac{X}{10^3}}{1+\frac{X}{10^3}}\right]\cdot\left[\frac{\frac{X}{10^2}}{1+\frac{X}{10^2}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^3}}\right]
$$

$$
+ \left[\frac{10^4\frac{X}{10^4}}{1+\frac{X}{10^4}}\right]\cdot\left[\frac{\frac{X}{10^3}}{1+\frac{X}{10^3}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^4}}\right] + \left[\frac{10^5\frac{X}{10^5}}{1+\frac{X}{10^5}}\right]\cdot\left[\frac{\frac{X}{10^4}}{1+\frac{X}{10^4}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^5}}\right] \tag{20}
$$

155 The following is non-approximated form of the above equation

$$
Z = \left[\frac{10^1\frac{X}{10^1}}{1+\frac{X}{10^1}}\right]\cdot\left[\frac{10^2\frac{\frac{X}{10^2}}{1+\frac{X}{10^2}}}{1+\frac{10^2\frac{X}{10^2}}{1+\frac{X}{10^2}}}\right]\cdot\left[\frac{1}{1+\frac{10^2\frac{X}{10^3}}{1+\frac{X}{10^3}}}\right] + \left[\frac{10^2\frac{X}{10^2}}{1+\frac{X}{10^2}}\right]\cdot\left[\frac{10^2\frac{\frac{X}{10^3}}{1+\frac{X}{10^3}}}{1+\frac{10^2\frac{X}{10^3}}{1+\frac{X}{10^3}}}\right]\cdot\left[\frac{1}{1+\frac{10^2\frac{X}{10^4}}{1+\frac{X}{10^4}}}\right] +
$$

$$
\left[\frac{10^3\frac{X}{10^3}}{1+\frac{X}{10^3}}\right]\cdot\left[\frac{10^2\frac{\frac{X}{10^4}}{1+\frac{X}{10^4}}}{1+\frac{10^2\frac{X}{10^4}}{1+\frac{X}{10^4}}}\right]\cdot\left[\frac{1}{1+\frac{10^2\frac{X}{10^5}}{1+\frac{X}{10^5}}}\right] + \left[\frac{10^4\frac{X}{10^4}}{1+\frac{X}{10^4}}\right]\cdot\left[\frac{10^2\frac{\frac{X}{10^5}}{1+\frac{X}{10^5}}}{1+\frac{10^2\frac{X}{10^5}}{1+\frac{X}{10^5}}}\right]\cdot\left[\frac{1}{1+\frac{10^2\frac{X}{10^6}}{1+\frac{X}{10^6}}}\right] +
$$

$$\left[\frac{10^5\,\dfrac{X}{10^5}}{1+\dfrac{X}{10^5}}\right]\cdot\left[\frac{10^2\,\dfrac{\dfrac{X}{10^6}}{1+\dfrac{X}{10^6}}}{1+\dfrac{10^2\,\dfrac{X}{10^6}}{1+\dfrac{X}{10^6}}}\right]\cdot\left[\frac{1}{1+\dfrac{10^2\,\dfrac{X}{10^7}}{1+\dfrac{X}{10^7}}}\right] \qquad (21)$$

Figure 3: (**A**) The four-node motif which in series can generate linear signal-response relationship and compute logarithm. (**B**) The series for generating linear signal-response relationship. (**C**) The series for computing logarithm.

Hence, together the negative arm and the positive arm create a window which shows maximum

response to signal of a particular order of magnitude and whose response declines progressively

as the distance between the preferred order of magnitude and the order of magnitude of signal

increases in either direction. Further, the signal-response curve is right-skewed. Figure 4

shows signal-response relationship of a single four-node motif with four combinations of the

Hill-coefficients. Increasing the value of m introduces steepness at the leftmost end of the

output, whereas increasing the value of n causes the output to have higher and sharper peak

and decline quickly. Notably, because a) β_1/α_1 is equal to Kd_{xz}, b) the effective equilibrium

dissociation constant of the negative arm is equal to Kd_{xz}, c) the effective equilibrium dissociation

constant of the positive arm is equal to $Kd_{xz}/10$, the value of Kd_{xz} is sufficient to specify all other

parameters of the four-node motif whose series produces linear signal-response relationship, as

shown below. Since the first receptor does not need the positive arm (nor it should have the

positive arm because the arm causes the output of the series to further deviate from linearity in

the leftmost end of the output) and the last receptor should not have the negative arm (otherwise

a very large signal will generate a very small output), the expression for the output should be:-

$$
Z = \left[\frac{10^1 \frac{X}{10^1}}{1 + \frac{X}{10^1}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^1}} \right] + \left[\frac{10^2 \frac{X}{10^2}}{1 + \frac{X}{10^2}} \right] \cdot \left[\frac{\frac{X}{10^1}}{1 + \frac{X}{10^1}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^2}} \right] + \left[\frac{10^3 \frac{X}{10^3}}{1 + \frac{X}{10^3}} \right] \cdot \left[\frac{\frac{X}{10^2}}{1 + \frac{X}{10^2}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^3}} \right]
$$

$$
+ \left[\frac{10^4 \frac{X}{10^4}}{1 + \frac{X}{10^4}} \right] \cdot \left[\frac{\frac{X}{10^3}}{1 + \frac{X}{10^3}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^4}} \right] + \left[\frac{10^5 \frac{X}{10^5}}{1 + \frac{X}{10^5}} \right] \cdot \left[\frac{\frac{X}{10^4}}{1 + \frac{X}{10^4}} \right] \tag{22}
$$

172 The following is non-approximated form of the above equation

$$
Z = \left[\frac{10^1 \frac{X}{10^1}}{1 + \frac{X}{10^1}} \right] \cdot \left[\frac{1}{1 + \frac{10^2 \frac{X}{10^3}}{1 + \frac{X}{10^3}}} \right] + \left[\frac{10^2 \frac{X}{10^2}}{1 + \frac{X}{10^2}} \right] \cdot \left[\frac{10^2 \frac{X}{10^3}}{1 + \frac{X}{10^3}} \cdot \frac{1}{1 + \frac{10^2 \frac{X}{10^3}}{1 + \frac{X}{10^3}}} \right] \cdot \left[\frac{1}{1 + \frac{10^2 \frac{X}{10^4}}{1 + \frac{X}{10^4}}} \right] +
$$

$$
\left[\frac{10^3 \frac{X}{10^3}}{1 + \frac{X}{10^3}} \right] \cdot \left[\frac{10^2 \frac{X}{10^4}}{1 + \frac{X}{10^4}} \cdot \frac{1}{1 + \frac{10^2 \frac{X}{10^4}}{1 + \frac{X}{10^4}}} \right] \cdot \left[\frac{1}{1 + \frac{10^2 \frac{X}{10^5}}{1 + \frac{X}{10^5}}} \right] + \left[\frac{10^4 \frac{X}{10^4}}{1 + \frac{X}{10^4}} \right] \cdot \left[\frac{10^2 \frac{X}{10^5}}{1 + \frac{X}{10^5}} \cdot \frac{1}{1 + \frac{10^2 \frac{X}{10^5}}{1 + \frac{X}{10^5}}} \right] \cdot \left[\frac{1}{1 + \frac{10^2 \frac{X}{10^6}}{1 + \frac{X}{10^6}}} \right] +
$$

$$\left[\frac{10^5 \dfrac{X}{10^5}}{1 + \dfrac{X}{10^5}}\right] \cdot \left[\frac{10^2 \dfrac{\dfrac{X}{10^6}}{1 + \dfrac{X}{10^6}}}{1 + 10^2 \dfrac{\dfrac{X}{10^6}}{1 + \dfrac{X}{10^6}}}\right] \tag{23}$$

Figure 4: (**A-D**) Output of the four-node motif defined by the equation $[10 \cdot (X/10)/(1+X/10)] \cdot [(X/1)^m/(1 + (X/1)^m)] \cdot [1/(1 + (X/10)^n)]$ with different values of the Hill-coefficients m and n.

It could be observed that output of equation 22 with the Hill-coefficients m and n being equal

to one was a curve with a slight concavity for signal 's values up to one order of magnitude

less than Kd_{xz} of the last receptor. As the signal's value increases further, the output for all the

combinations of m and n starts to plateau.

In order to assess how well equation 23 approximates equation 22, output of equation 23

was fitted with equation 22 having each of its term multiplied with a coefficient (equation 24).

If equation 23 is a good approximation of equation 22, the value of the coefficients should

turn out to be around one, besides goodness-of-fit (r^2) being close to one. Because output of

Figure 5: Output from the first three receptors for the approximated form (**A**) and non-approximated form (**B**) when m=1, n=1 (blue); m=1, n=2 (green); m=1, n=4 (red); m=1, n=8 (cyan); m=1, n=16 (pink).

Figure 6: Output from the first three receptors for the approximated form (**A**) and non-approximated form (**B**) when m=1, n=1 (blue); m=2, n=1 (green); m=4, n=1 (red); m=8, n=1 (cyan); m=16, n=1 (pink).

Figure 7: Output from the first three receptors for the approximated form (**A**) and non-approximated form (**B**) when m=1, n=1 (blue); m=2, n=2 (green); m=4, n=4 (red); m=8, n=8 (cyan); m=16, n=16 (pink).

equation 22 (and hence also equation 23) begins to plateau as the signal's value goes beyond

Kd_{xz} of the penultimate receptor (whose value is 10^4), the signal's values at which output of

equation 23 was generated for the purpose of fitting were 1, 2, 5, 10, 20, 50, 10^2, $2 \cdot 10^2$, $5 \cdot 10^2$,

10^3, $2 \cdot 10^3$, $5 \cdot 10^3$, 10^4. Except for the value of the coefficient A for m=2, n=1; m=4, n=1;

m=8, n=1; and m=16, n=1, the values of all the coefficients for all the combinations m and

n were close to one and goodness-of-fit was equal to one (Supplementary Information, Table

1-3). Specifically, for m and n equal to one, the value of A, B, C, D, E were 1.0075, 0.9948,

0.9974, 0.9968, 0.9963, respectively. It should be noted that the output plateaus for very large

signal. This may explain why humans do not have a refined sense for very large numbers, and

thus they are represented by a common term, infinity.

$$
Z = A \cdot \left[\frac{10^1 \frac{X}{10^1}}{1 + \frac{X}{10^1}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^1}} \right] + B \cdot \left[\frac{10^2 \frac{X}{10^2}}{1 + \frac{X}{10^2}} \right] \cdot \left[\frac{\frac{X}{10^1}}{1 + \frac{X}{10^1}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^2}} \right] + C \cdot \left[\frac{10^3 \frac{X}{10^3}}{1 + \frac{X}{10^3}} \right] \cdot \left[\frac{\frac{X}{10^2}}{1 + \frac{X}{10^2}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^3}} \right]
$$

$$
+ D \cdot \left[\frac{10^4 \frac{X}{10^4}}{1 + \frac{X}{10^4}} \right] \cdot \left[\frac{\frac{X}{10^3}}{1 + \frac{X}{10^3}} \right] \cdot \left[\frac{1}{1 + \frac{X}{10^4}} \right] + E \cdot \left[\frac{10^5 \frac{X}{10^5}}{1 + \frac{X}{10^5}} \right] \cdot \left[\frac{\frac{X}{10^4}}{1 + \frac{X}{10^4}} \right] \tag{24}
$$

As stated above, the ideal relationship between a signal and the response should be a straight

line with the slope of one, but output of equation 22 with m=1, n=1 has a slight concavity up

until the signal's value is less than Kd_{xz} of the penultimate receptor. Thus, in order to assess

if, given a range of Kd_{xz}, there is a range of signal in which the series could generate linear

signal-response relationship, the output was fitted with straight line for various ranges of Kd_{xz}

and signal (Table 1). For a given range of signal, the values at which the output was generated

for the purpose of fitting were taken as powers of ten. For example, for the range $10^{-1} – 10^3$, the

output was generated at the following values: 10^{-1}, 10^0, 10^1, 10^2, and 10^3.

For the middle intervals, the slope converges to the value 0.7466. In other words, if the output

is multiplied by the reciprocal of this value, one gets a straight line with the slope of 1 for

Table 1: Slope and goodness of fit (R^2) for the fit of the output with straight line for various ranges of signal and Kd_{xz}.

Signal Range (across) Kd_{xz} Range (down)	10^0–10^2	10^{-1}–10^3	10^{-2}–10^4	10^{-3}–10^5	10^{-4}–10^6
10^{-2}–10^4	0.736 (1.0000)	0.6867 (1.0000)	0.4573 (0.9975)	NA	NA
10^{-3}–10^5	0.7455 (1.0000)	0.7363 (1.0000)	0.6872 (1.0000)	0.4585 (0.9975)	NA
10^{-4}–10^6	0.7465 (1.0000)	0.7455 (1.0000)	0.7364 (1.0000)	0.6875 (1.0000)	0.4591 (0.9975)
10^{-5}–10^7	0.7466 (1.0000)	0.7465 (1.0000)	0.7455 (1.000)	0.7365 (1.0000)	0.6876 (0.9999)
10^{-6}–10^8	0.7466 (1.0000)	0.7466 (1.0000)	0.7465 (1.0000)	0.7455 (1.0000)	0.7365 (1.0000)

the middle intervals. Because there is no limit on the number of the receptors, the series can

generate linear signal-response relationship over arbitrary range of signal. For the values of m

and/or n being more than one, the output becomes oscillatory around the output generated with

m=1 and n=1 (Figures 5, 6, and 7). Therefore, I posit that V and Y molecules should be present

as monomers in a real system.

The exact same result was got when the range of Kd_{xz} and concomitantly the range of signal

were shifted by the factor of 0.01, 0.1, 10, and 100. That is, for example, for the second case the

new ranges of Kd_{xz} were 10^{-3}–10^3, 10^{-4}–10^4, 10^{-5}–10^5, 10^{-6}–10^6, and 10^{-7}–10^7. And the new

ranges of signal were 10^{-1}–10^1, 10^{-2}–10^2, 10^{-3}–10^3, 10^{-4}–10^4, and 10^{-5}–10^5. Similarly, for the

third case the new ranges of Kd_{xz} were 10^{-1}–10^5, 10^{-2}–10^6, 10^{-3}–10^7, 10^{-4}–10^8, and 10^{-5}–10^9.

And the new ranges of signal were 10^1–10^3, 10^0–10^4, 10^{-1}–10^5, 10^{-2}–10^6, and 10^{-3}–10^7. As

the ranges of Kd_{xz} were multiplied by the factor of 0.01, 0.1, 10, and 100, the y-intercept of the

fitted line were 0.01, 0.1, 10, and 100 times the previous y-intercept, respectively. Explanation

of this observation is as follows. The first transformation, for example, is f(x) → 0.01·f(x/0.01).

Because the y-intercept should be zero when x is zero, it can be considered as the deviation of

the output from the straight line. The multiplicative factor of 0.01 implies that the error also

gets multiplied by 0.01, hence the y-intercept is 0.01 times it was before the transformation. In

other words, y = mx + c → y = 0.01·mx/0.01 + 0.01·c

223 2.2. Weber's Law

Now I present a formalism to get Weber's law from equation 22, which, when multiplied by

the factor of (1/0.7466), yields Z = X for intermediate range of values of X. Given a signal

(X=μ), the output (Z) is considered to be the result of sampling from a Gaussian distribution

with mean equal to μ and standard deviation linearly proportional to μ; that is, $\sigma = f\mu$, where

f is the Weber fraction, the ratio of the smallest perceptual change in a signal (ΔX_{min}) and the

background signal ($X_{background}$): $\Delta X_{min}/X_{background} = f$. It is posited here that the subject is able

to perceive a change in the signal when the ratio of non-overlapping region of the Gaussian

centered on μ_2 to the right of the Gaussian centered on μ_1 and the area under the Gaussian

centered on μ_1, is above a certain threshold (figure 8). Because the Gaussians in consideration

are normalized, the ratio is equal to the non-overlapping region.

Figure 8: The Gaussian on the left has mean μ_1=10, and the Gaussian on the right has mean μ_2=15. The shaded portion is the non-overlapping region, R

As is defined now, the non-overlapping region is just below the threshold when $\mu_2 - \mu_1 = \sigma_1$;

that is, when the difference between mean of the current signal and mean of the background

signal is equal to the standard deviation of the background signal. In other words, σ_1 is the

just noticeable difference (JND). The non-overlapping area (R) is $F_1(p_2)$ - $F_2(p_2)$, where $F(X)$

is the cumulative distribution function of Gaussian distribution and p_2 is the farther point of

intersection of the two Gaussians when $\mu_2 - \mu_1 = \sigma_1$. That is, this point is the bigger-valued

solution of the quadratic equation below. It should be noted that the total output, given by

equation 22, consists of the sum of outputs by individual receptors. But that still does not affect

the formalism above because the individual outputs are independent, and thus the sum of the

individual means is μ and the sum of the individual standard deviations is σ.

$$\frac{1}{\sqrt{2\pi\sigma_1^2}}e^{-\frac{(p-\mu_1)^2}{2\sigma_1^2}} = \frac{1}{\sqrt{2\pi\sigma_2^2}}e^{-\frac{(p-\mu_2)^2}{2\sigma_2^2}} \tag{25}$$

$$\ln\left|\frac{\sigma_2}{\sigma_1}\right| = \frac{(p-\mu_1)^2}{2\sigma_1^2} - \frac{(p-\mu_2)^2}{2\sigma_2^2} \tag{26}$$

$$\frac{\ln\left|\frac{\sigma_2}{\sigma_1}\right|}{\left(\frac{1}{\sigma_1} - \frac{1}{\sigma_2}\right)} = p\left[\left(\frac{1}{\sigma_1} + \frac{1}{\sigma_2}\right)p - \frac{2}{f}\right] \tag{27}$$

245 Setting $C = \dfrac{\ln\left|\frac{\sigma_2}{\sigma_1}\right|}{\left(\frac{1}{\sigma_1} - \frac{1}{\sigma_2}\right)}$, $A = \left(\frac{1}{\sigma_1} + \frac{1}{\sigma_2}\right)$, $B = \dfrac{2}{f}$

246 Equation 27 becomes

$$C = p[Ap - B] \quad \text{or} \quad Ap^2 - Bp - C = 0 \tag{28}$$

$$p_{1,2} = \frac{B \pm \sqrt{B^2 4AC}}{2A} \tag{29}$$

$$p_2 = \frac{\dfrac{2}{f} + \sqrt{\left(\dfrac{2}{f}\right)^2 + 4\left(\dfrac{1}{\sigma_1} + \dfrac{1}{\sigma_2}\right)\dfrac{\ln\left|\dfrac{\sigma_2}{\sigma_1}\right|}{\left(\dfrac{1}{\sigma_1} - \dfrac{1}{\sigma_2}\right)}}}{2\left(\dfrac{1}{\sigma_1} + \dfrac{1}{\sigma_2}\right)} \tag{30}$$

247 Setting $\dfrac{\sigma_2}{\sigma_1} = 1 + f$

$$p_2 = \frac{\mu_1\left(1 + \sqrt{1 + f(2 + f)\ln|1 + f|}\right)}{\left(1 + \dfrac{1}{1 + f}\right)} \tag{31}$$

248 The expression 31 can be written as $p_2 = \mu_1 g(f)$

249 The CDF of a Gaussian distribution with mean μ and variance σ^2 is

$$F(Y) = \frac{1}{2}\left[1 + \text{erf}\left(\frac{Y - \mu}{\sigma\sqrt{2}}\right)\right]$$

250 Hence

$$R = \frac{1}{2}\left[1 + \text{erf}\left(\frac{p_2 - \mu_1}{\sigma_1\sqrt{2}}\right)\right] - \frac{1}{2}\left[1 + \text{erf}\left(\frac{p_2 - \mu_2}{\sigma_2\sqrt{2}}\right)\right] \tag{32}$$

251 Inserting the value of p_2 in equation 32

$$R = \frac{1}{2}\left[1 + \text{erf}\left(\frac{\mu_1 g(f) - \mu_1}{\sigma_1\sqrt{2}}\right)\right] - \frac{1}{2}\left[1 + \text{erf}\left(\frac{\mu_1 g(f) - \mu_2}{\sigma_2\sqrt{2}}\right)\right] \tag{33}$$

$$R = \frac{1}{2}\left[1 + \text{erf}\left(\frac{g(f) - 1}{f\sqrt{2}}\right)\right] - \frac{1}{2}\left[1 + \text{erf}\left(\frac{g(f)/(1 + f) - 1}{f\sqrt{2}}\right)\right] \tag{34}$$

252 Being a function of f alone, R can be written as R = h(f). It also implies that the value of

253 R remains the same when the two means are multiplied by a scalar. This is because the two

254 Gaussians become flatter, whereas the point p_2 goes farther. Hence, the non-overlapping region

255 stays constant when the two signals are multiplied by a scalar, which is Weber' law.

$$\frac{\Delta X_{min}}{X_{background}} = \frac{\mu_2 - \mu_1}{\mu_1} = \frac{\sigma_1}{\mu_1} = f = h^{-1}(R) \tag{35}$$

In the formalism presented here, Weber's law is the result of internal noise whose standard deviation is proportional to the signal. On the other hand, Fechner's explanation is based on the assumption that the response is equal to the logarithm of the signal, and JND therefore is a constant increment in the response.

$$Z_0 = \log(X_0) \tag{36}$$

$$Z_1 = \log(X_0 + \Delta X_{min}) \tag{37}$$

$$\Delta Z = Z_1 - Z_0 = \log\left(1 + \frac{\Delta X_{min}}{X_0}\right) \tag{38}$$

$$e^{\Delta Z} - 1 = \frac{\Delta X_{min}}{X_0} = f \tag{39}$$

However, Fechner's formulation does not provide a biological basis of the relation between the Weber fraction and ΔZ. Also, the response being logarithm of the signal does not seem natural. Two Wildebeests butting their head against each other should not compute logarithm of the force that they receive from the other in the process. Otherwise, a force four times as large will appear larger by only log4. With such a sensory system the weaker animal will be much more injured at the end of the fight than if the sensory system were to convert the signal to an equally valued response.

In order to validate the above-stated formalism for Weber's law, which assumes output of equation 23 (approximated as equation 22) to be noisy and thus considers it to be equivalent to sampling from Gaussian distribution centered on the signal's value, output of equation 23 should be shown to be a Gaussian distribution with mean equal to the signal's value and standard deviation proportional to the mean when output generated by individual components of the four-node motif is considered to be the result of sampling from their corresponding Gaussian

278 distributions. That is, given the signal X, the value of V and Y are generated from Gaussian

279 distribution with mean equal to $\dfrac{\beta_3}{\alpha_3}\dfrac{X}{Kd_{XV}}\Big/\left(1+\dfrac{X}{Kd_{XV}}\right)$ and $\dfrac{\beta_2}{\alpha_2}\dfrac{X}{Kd_{XY}}\Big/\left(1+\dfrac{X}{Kd_{XY}}\right)$ respectively,

280 and the standard deviation equal to Weber fraction times the respective mean.

282 Then, with the value of X, V and Y, Z is generated from Gaussian with mean equal to

283
$$\left[\frac{\beta_1\dfrac{X}{Kd_{XZ}}}{1+\dfrac{X}{Kd_{XZ}}}\right]\cdot\left[\frac{\left(\dfrac{V}{Kd_{VZ}}\right)^m}{1+\left(\dfrac{V}{Kd_{VZ}}\right)^m}\right]\cdot\left[\frac{1}{1+\left(\dfrac{Y}{Kd_{YZ}}\right)^n}\right]$$

and the standard deviation equal to Weber fraction times the expression. Adding output of

all receptors and multiplying by the reciprocal of 0.7466 should turn out to be equivalent to

sampling from a Gaussian centered on the signal X and standard deviation equal to a constant

time mean. It should be noted that because the signal-response curve is linear only for the

middle values of signal, the order of magnitude of X should be the preferred order of magnitude

of the middle receptors.

The expanded form of the expression is

291
$$\left[\frac{\dfrac{\beta_1}{\alpha_1}\dfrac{X}{Kd_{XZ}}}{1+\dfrac{X}{Kd_{XZ}}}\right]\cdot\left[\frac{\left(\dfrac{\left(\dfrac{1}{Kd_{VZ}}\dfrac{\beta_3}{\alpha_3}\right)\dfrac{X}{Kd_{XV}}}{1+\dfrac{X}{Kd_{XV}}}\right)^m}{1+\left(\dfrac{\left(\dfrac{1}{Kd_{VZ}}\dfrac{\beta_3}{\alpha_3}\right)\dfrac{X}{Kd_{XV}}}{1+\dfrac{X}{Kd_{XV}}}\right)^m}\right]\cdot\left[\frac{1}{1+\left(\dfrac{\left(\dfrac{1}{Kd_{YZ}}\dfrac{\beta_2}{\alpha_2}\right)\dfrac{X}{Kd_{XY}}}{1+\dfrac{X}{Kd_{XY}}}\right)^n}\right]$$

The value of $\beta_2/\alpha_2 Kd_{YZ}$ and $\beta_3/\alpha_3 Kd_{VZ}$ were set equal to 100 in order to carry out the

approximations (equations 9 and 19), respectively. However, values of β_2/α_2 and β_3/α_3,

required to generate the value of V and Y from the signal X, are not known individually.

Therefore I considered three combinations of β_2/α_2 and Kd_{YZ} while keeping the value of their

product equal to 100. The combinations are $\beta_2/\alpha_2=100$ and $Kd_{YZ}=1$, $\beta_2/\alpha_2=10$ and $Kd_{YZ}=10$,

and $\beta_2/\alpha_2=1$, $Kd_{YZ}=100$. The same was done with β_3/α_3 and Kd_{VZ}.

Given a value of mean (X) and Weber fraction (f), ten thousand values of Z were generated from

this sampling procedure and fitted with a Gaussian distribution. This procedure was carried out

for thirty times. The average of mean and standard deviation of the fitted Gaussian for mean

equal to 0.1, 1, and 10 (and therefore, standard deviation equal to f·0.1, f·1, and f·10) were fitted

with straight line for a given value of f. The value of f taken were 0.01, 0.02, 0.05, 0.1, 0.2, 0.5,

and 1. The slope of the fit was very close of 1, the y-intercept was of the order of magnitude

-5, and R^2 was equal to 1 for all the nine combinations up to f equal to 0.2 (data not shown).

Curiously, the straight line fit of standard deviation of the Gaussian fit of ten thousand values

sampled from the Gaussian random variable generator for mean equal to 0.1, 1, and 10 and a

given value of f was consistently approximately 1.4 times the value of f.

In 2002, two studies [4,5], investigating parietal cortex and prefrontal cortex, respectively,

observed the presence of neurons whose firing rate was tuned to a specific numerosity. In

a study by Nieder and Miller [6], the activity of neurons from the lateral prefrontal cortex

displayed a maximum at a preferred number and declined progressively as the numerical distance

from that preferred number was increased, thereby forming numerosity filter function. The

tuning curves, which were skewed towards the right on a linear scale, were plotted on nonlinear

scales, namely a power function with an exponent of 0.5, a power function with an exponent of

0.33, and a logarithmic scale. The new distributions along with the original distribution (linear

scale) were fitted with a Gaussian distribution. The mean goodness-of-fit (R^2) values were in

ascending order when arranged from left to right for the linear scale, the power function with

an exponent of 0.5, the power function with an exponent of 0.33, and the logarithmic scale.

On the logarithmic scale, the variance of numerosity filters was approximately constant with

increasing preferred numerosity.

Receptors considered in the current study until now produce the maximum at values which

are powers of ten (for example, the four-node motifs which constitute the series represented

by equations 22 and 23 have maximum at the values 10^1, 10^2, 10^3, 10^4, and 10^5, respectively).

Thus, in order to have the receptors produce maximum at the values 1, 2, 3, etc, and also possess

a certain mathematical property, a few combinations of equilibrium dissociation constant and

Hill-coefficient were tried. The mathematical property that the expression of output of the

individual receptors having constant β_1/α_1 needed to possess is $f_n(X) = f_{n+1}(X/c)$, where f_n

and f_{n+1} are outputs of consecutive receptors, respectively. This property makes output of the

individual receptors have equal variance when plotted on a logarithm scale [Nieder and Miller

Neuron 37 149–157, 2003]. The following expression turned out to satisfy both requirements

$$\left[\frac{1\frac{X}{1}}{1+\frac{X}{1}}\right] \cdot \left[\frac{\frac{X}{1}}{1+\frac{X}{1}}\right] \cdot \left[\frac{1}{1+\left(\frac{X}{1}\right)^2}\right] \tag{40}$$

$$\left[\frac{2\frac{X}{2}}{1+\frac{X}{2}}\right] \cdot \left[\frac{\frac{X}{2}}{1+\frac{X}{2}}\right] \cdot \left[\frac{1}{1+\left(\frac{X}{2}\right)^2}\right] \tag{41}$$

$$\left[\frac{3\frac{X}{3}}{1+\frac{X}{3}}\right] \cdot \left[\frac{\frac{X}{3}}{1+\frac{X}{3}}\right] \cdot \left[\frac{1}{1+\left(\frac{X}{3}\right)^2}\right] \tag{42}$$

As in the study [5], R^2 values of the Gaussian fit of output of the expressions 40-42 plotted

on the linear scale, the power function with an exponent of 0.5, the power function with an

exponent of 0.33, and the logarithmic scale were in ascending order when arranged from left

to right (Supplementary Information, Table 4). The same result was obtained with respective

output of the first three terms of equation 20 (Supplementary Information, Table 5).

The following are non-approximated forms of the above three equations

$$\left[\frac{1\dfrac{X}{1}}{1+\dfrac{X}{1}}\right] \cdot \left[\frac{\dfrac{10^2\dfrac{X}{10^2}}{1+\dfrac{X}{10^2}}}{1+\dfrac{10^2\dfrac{X}{10^2}}{1+\dfrac{X}{10^2}}}\right] \cdot \left[\frac{1}{1+\left(\dfrac{10^2\dfrac{X}{10^2}}{1+\dfrac{X}{10^2}}\right)^2}\right] \tag{43}$$

$$\left[\frac{2\dfrac{X}{2}}{1+\dfrac{X}{2}}\right] \cdot \left[\frac{\dfrac{10^2\dfrac{X}{2\cdot 10^2}}{1+\dfrac{X}{2\cdot 10^2}}}{1+\dfrac{10^2\dfrac{X}{2\cdot 10^2}}{1+\dfrac{X}{2\cdot 10^2}}}\right] \cdot \left[\frac{1}{1+\left(\dfrac{10^2\dfrac{X}{2\cdot 10^2}}{1+\dfrac{X}{2\cdot 10^2}}\right)^2 \cdot 10^2}\right] \tag{44}$$

$$\left[\frac{3\dfrac{X}{3}}{1+\dfrac{X}{3}}\right] \cdot \left[\frac{\dfrac{10^2\dfrac{X}{3\cdot 10^2}}{1+\dfrac{X}{3\cdot 10^2}}}{1+\dfrac{10^2\dfrac{X}{3\cdot 10^2}}{1+\dfrac{X}{3\cdot 10^2}}}\right] \cdot \left[\frac{1}{1+\left(\dfrac{10^2\dfrac{X}{3\cdot 10^2}}{1+\dfrac{X}{3\cdot 10^2}}\right)^2}\right] \tag{45}$$

In their experiments, monkeys were shown a set of dots, and then, after a short delay, were
presented with another set of dots, which they had to distinguish from the first set. The
percentage of trials on which they judged the second number to be indistinguishable from
the first number, which varied between 2-6, was plotted as a function of the second number.
Monkeys' performance –percentage of trials on which they reported the two sets to be of the
same numerosity– declined progressively as the two sets became numerically more distant.
For larger numerosities, the two sets had to be proportionally larger in order to keep the level
of performance the same, thereby translating to Weber' law. The distributions of both monkey's
performance had positive skew, but assumed a more symmetric shape when plotted on nonlinear
scales, namely a power function with an exponent of 0.5, a power function with an exponent
of 0.33, and a logarithmic scale. The mean goodness-of-fit values for the linear scale, the

power function with an exponent of 0.5, the power function with an exponent of 0.33, and the
logarithmic scale were 0.93, 0.97, 0.98, and 0.98, respectively. Further, the variance of the
distributions for each numerosity was constant when the data was plotted on a power function
scale with 0.33 exponent and the logarithmic scale.

Because the above-stated formulation of Weber's law requires the signal-response relationship
to be linear, the series of the receptors which produce maximum at integral values must be able
to produce a linear signal-response relationship in order to explain behavioural data of the study
by Nieder and Miller [6]. The values of β_1/α_1, which were determined by fitting the output
with straight line, vary over 5 orders of magnitude, which does not seem biologically plausible.
Thus, the present work is unable to explain the behavioural data.

Figure 9: Output of the receptors which produce the maximum at integral values. The first three outputs are of the expressions 40, 41, and 42, respectively.

363 After logarithmic-compression of abscissa, the expression 40 and 41 become

$$\left[\frac{\frac{\exp(X)}{2}}{1+\frac{\exp(X)}{2}}\right]\cdot\left[\frac{\frac{\exp(X)}{2}}{1+\frac{\exp(X)}{2}}\right]\cdot\left[\frac{1}{1+\left(\frac{\exp(X)}{2}\right)^2}\right] \tag{46}$$

$$\left[\frac{\frac{\exp(X)}{3}}{1+\frac{\exp(X)}{3}}\right]\cdot\left[\frac{\frac{\exp(X)}{3}}{1+\frac{\exp(X)}{3}}\right]\cdot\left[\frac{1}{1+\left(\frac{\exp(X)}{3}\right)^2}\right] \tag{47}$$

365 The two expressions can be written as

$$\left[\frac{\exp(X - \log_e 2)}{1 + \exp(X - \log_e 2)}\right] \cdot \left[\frac{\exp(X - \log_e 2)}{1 + \exp(X - \log_e 2)}\right] \cdot \left[\frac{1}{1 + \exp(X - \log_e 2)}\right] \tag{48}$$

$$\left[\frac{\exp(X - \log_e 3)}{1 + \exp(X - \log_e 3)}\right] \cdot \left[\frac{\exp(X - \log_e 3)}{1 + \exp(X - \log_e 3)}\right] \cdot \left[\frac{1}{1 + \exp(X - \log_e 3)}\right] \tag{49}$$

The latter function can be got from the former by the transformation $X \to X + \log_e 2 - \log_e 3$,

hence the standard deviation remains the same. Notably, barring the first and last receptor, the

expression of output of a receptor (i.e., the four-node motif) which in series generates linear

signal-response relationship is related to the expression of output of the preceding receptor by

the relation $f_n(X) = c \cdot f_{n+1}(X/c)$. Therefore, the receptor follows the above-stated mathematical

property when β_1/α_1 is kept constant, and the output of these receptors have constant variance

when plotted on a logarithm scale.

On a logarithmic abscissa, the expressions

$$\left[\frac{1\frac{X}{10}}{1 + \frac{X}{10}}\right] \cdot \left[\frac{\frac{X}{1}}{1 + \frac{X}{1}}\right] \cdot \left[\frac{1}{1 + \frac{X}{10}}\right] \tag{50}$$

375 and

$$\left[\frac{1\frac{X}{100}}{1 + \frac{X}{100}}\right] \cdot \left[\frac{\frac{X}{10}}{1 + \frac{X}{10}}\right] \cdot \left[\frac{1}{1 + \frac{X}{100}}\right] \tag{51}$$

376 become

$$\left[\frac{\frac{\exp(X)}{10}}{1 + \frac{\exp(X)}{10}}\right] \cdot \left[\frac{\frac{\exp(X)}{1}}{1 + \frac{\exp(X)}{1}}\right] \cdot \left[\frac{1}{1 + \frac{\exp(X)}{10}}\right] \tag{52}$$

$$\left[\frac{\dfrac{\exp(X)}{100}}{1 + \dfrac{\exp(X)}{100}}\right] \cdot \left[\frac{\dfrac{\exp(X)}{10}}{1 + \dfrac{\exp(X)}{10}}\right] \cdot \left[\frac{1}{1 + \dfrac{\exp(X)}{100}}\right] \tag{53}$$

The two expressions can be written as

$$\left[\frac{\exp(X - \log_e 10)}{1 + \exp(X - \log_e 10)}\right] \cdot \left[\frac{\exp(X)}{1 + \exp(X)}\right] \cdot \left[\frac{1}{1 + \exp(X - \log_e 10)}\right] \tag{54}$$

$$\left[\frac{\exp(X - 2\log_e 10)}{1 + \exp(X - 2\log_e 10)}\right] \cdot \left[\frac{\exp(X - \log_e 10)}{1 + \exp(X - \log_e 10)}\right] \cdot \left[\frac{1}{1 + \exp(X - 2\log_e 10)}\right] \tag{55}$$

The latter function can be got from the former by the transformation $X \rightarrow X\text{-}\log_e 10$, which leaves the standard deviation unchanged.

2.3. The Four-node Motif and Bow-Tie Architecture

For the series generating linear signal-response relationship (figure 3B), assuming the signal's range is 10^n-10^m (n > m and both are integers), the number of receptors (i.e., the four-node motifs) is equal to n-m+10. When m $\geqslant$ 0, the generation of linear signal-response relationship becomes an example of bow-tie architecture. Therefore, a class of bow-tie architecture could be defined: a class in which the number of receptors is much smaller than the range of signal and consequently the range of response.

The range of output generated by the series in figure 3B is identical to signal's range. In contrast to that, now I consider a hypothetical requirement of sensitive detection over wide range of signal while keeping the range of output constant. It should be noted that, although the output's range is constant, the range of response is still proportional to signal's range as in the previous example. That's because, even though the output lies within a constant range, the receptors that respond to a higher valued signal are different from those receptors that respond to a lower

valued signal. Thus, a system able to provide such a detection belongs to the above-defined

class of bow-tie architecture. Black line in figure 10A depicts that, when the output's range is

kept constant, sensitivity of a mechanism generating single linear signal-response relationship

reduces as the signal's range increases. However, the sensitivity is maintained if a mechanism

generates multiple linear signal-response outputs, each of which is from its corresponding

receptor and covers corresponding limited range of signal. The blue, green, and red lines

are ideal hypothetical outputs which together provide sensitive detection over a wide range of

signal. However, the outputs are not biologically realistic, as they terminate abruptly at their

maximum. Figure 10B shows a biologically realistic form of the outputs.

Figure 10C shows output of a series of the four-node motif which qualitatively matches figure

10B and, thus, provides sensitive detection over arbitrary range of signal. The four-node motif

in this series has the following properties: β_1/α_1 is constant, the Hill coefficients m and n

are equal to 2 and 8, respectively, all equilibrium dissociation constants of a given receptor

are equal, and the value of equilibrium dissociation constants of a receptor are twice those of

the preceding receptor. The expressions 56-58 define the first three receptors. Output of such

a four-node motif fits Gaussian distribution very well. R^2 for the Gaussian fit of output of

the three expressions are 0.9967, 0.996, and 0.9959, respectively. Because eqs. 56-58, when

keeping β_1/α_1 constant, possess the mathematical property $f_n(X) = f_{n+1}(X/c)$, the coefficient

of variation of Gaussians fitted by them are equal. Further, output of such a series oscillates

with constant amplitude and decreasing frequency around a flat line. It should be noted that the

individual outputs in figure 10B have constant width at half height (WAHH), whereas, because

WAHH is equal to $4 \cdot \ln 2 \cdot \sigma$, WAHH of the outputs in figure 10C are proportional to mean of the

approximate Gaussians they form.

Assuming the equilibrium dissociation constants of the first and the last receptor are 2^m and

2^n, respectively; arguably, the range in which signal can be detected sensitively is $\mu_1 \cdot (2^n - 2^m) +$

$(1/2) \cdot \sigma_1 \cdot (2^m + 2^n)$, where $n > m$ and both are integers, and μ_1 and σ_1 are mean and standard

deviation of the Gaussian distribution which fits output of the defining expression of the receptor
whose equilibrium dissociation constants are equal to 1. When m $\geqslant$ 0, the signal-response
relationship becomes an example of bow-tie architecture, as for the signal's range being $\mu_1 \cdot (2^n - 2^m)$
$+ (1/2) \cdot \sigma_1 \cdot (2^m + 2^n)$, the number of receptors is equal to n-m+1. Figure 10D shows that, with
some modifications, a series of three such four-node motifs is able to produce a signal-response
relationship which qualitatively matches absorption spectra of three cone opsins of humans [6].
The defining expressions of the three receptors are in the figure's legend.
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$$\left[\frac{5\dfrac{X}{1}}{1 + \dfrac{X}{1}} \right] \cdot \left[\frac{\left(\dfrac{X}{1}\right)^2}{1 + \left(\dfrac{X}{1}\right)^2} \right] \cdot \left[\frac{1}{1 + \left(\dfrac{X}{1}\right)^8} \right] \tag{56}$$

$$\left[\frac{5\dfrac{X}{2}}{1 + \dfrac{X}{2}} \right] \cdot \left[\frac{\left(\dfrac{X}{2}\right)^2}{1 + \left(\dfrac{X}{2}\right)^2} \right] \cdot \left[\frac{1}{1 + \left(\dfrac{X}{2}\right)^8} \right] \tag{57}$$

$$\left[\frac{5\dfrac{X}{4}}{1 + \dfrac{X}{4}} \right] \cdot \left[\frac{\left(\dfrac{X}{4}\right)^2}{1 + \left(\dfrac{X}{4}\right)^2} \right] \cdot \left[\frac{1}{1 + \left(\dfrac{X}{4}\right)^8} \right] \tag{58}$$

432 **2.4. The Four-node Motif and Logarithm Computation**

A series of four-node motif in which the last receptor lacks the negative arm is also able to
compute logarithm of signal. To achieve this, the value of β_1/α_1 should increase linearly,
instead of as powers of ten, with the motif's number. However, a strict linear increase in β_1/α_1
gives a very poor fit with the logarithm function. Because a series was sought which could
compute logarithm of signal ranging between 1 and 10^4, series of five four-node motifs having
Kd_{xz} equal to 10^1, 10^2, 10^3, 10^4, and 10^5, respectively, was considered. The values of β_1/α_1 of
the series which compute logarithm with the base ten and the natural logarithm were estimated

Figure 10: A The black line shows that, when the range of response is constant, sensitivity of linear signal-response relationship reduces with increase in the range of signal. The blue, green, and red lines are minimal ideal hypothetical signal-response relationship, i.e., which maintain high sensitivity throughout signal's range. **B** Biologically realistic form of ideal hypothetical signal-response relationship. **C** Output of the first four receptors, first three of which are defined by eqs. 44-46. The total output of the series oscillates around a flat line. The outputs are approximately Gaussian and have constant coefficient of variation. **D** A series of three four-node motifs is able to qualitatively generate the absorption spectra of human cone cells. The equations of the three receptors are: $[3.6 \cdot (X/4)/(1 + X/4)] \cdot [(X/8)^2/(1 + (X/8)^2)] \cdot [1/(1 + (X/8)^8)]; [5 \cdot (X/20)/(1 + X/20)] \cdot [(X/20)^2/(1 + (X/20)^2)] \cdot [1/(1 + (X/20)^8)]; [5 \cdot (X/24)/(1 + X/24)] \cdot [(X/24)^2/(1 + (X/24)^2)] \cdot [1/(1 + (X/24)^8)]$

by linear regression. $\log_{10}$ and $\log_e$ of powers of 2 ranging from 0 to 14 (i.e., the values ranging

from 1 to 8192) were regressed against output of equation 59. It was observed that β_1/α_1 of

all the receptors were positive only when the signal range was from 1 to a value which was

slightly larger than Kd_{xz} of the antepenultimate. On the other hand, when $\log_{10}$ and $\log_e$ of

powers of 10 were regressed against output of equation 59, the signal range had to be from 1 to

slightly larger than Kd_{xz} of the penultimate receptor in order for β_1/α_1 of all the receptors to

be positive. β_1/α_1 used to generate figures 13 and 14 are from regressing logarithm of powers

of 2, because this way more data points are available for the procedure.

Because the logarithm of values that lie between 0 and 1 is negative, a biologically unrealistic

scenario, the output in this range was suppressed. This was done by adding a positive arm with

m=2 to the first receptor. This raises the possibility of a series of the four-node motif being

able to compute the exponential of signal. To achieve this, the value of β_1/α_1 of the successive

receptor should increase much faster than the corresponding value of Kd_{xz}. As before, linear

regression was used to find out the values of β_1/α_1. However, β_1/α_1 of the first receptor turned

out to be negative. This result alone rules out the possibility of a series of the four-node motif

to compute exponential. Moreover, the regression gave a poor R^2 value. The four-node motif

in series being able to compute logarithm but unable to compute exponential might be due to

positive skew in its output.

$$Z = \left[\frac{A\frac{X}{10^1}}{1+\frac{X}{10^1}}\right]\left[\frac{\left(\frac{X}{10^0}\right)^2}{1+\left(\frac{X}{10^0}\right)^2}\right]\cdot\left[\frac{1}{1+\frac{X}{10^1}}\right]+\left[\frac{B\frac{X}{10^2}}{1+\frac{X}{10^2}}\right]\left[\frac{\frac{X}{10^1}}{1+\frac{X}{10^1}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^2}}\right]+\left[\frac{C\frac{X}{10^3}}{1+\frac{X}{10^3}}\right]\left[\frac{\frac{X}{10^2}}{1+\frac{X}{10^2}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^3}}\right]$$

$$+\left[\frac{D\frac{X}{10^4}}{1+\frac{X}{10^4}}\right]\cdot\left[\frac{\frac{X}{10^3}}{1+\frac{X}{10^3}}\right]\cdot\left[\frac{1}{1+\frac{X}{10^4}}\right]+\left[\frac{E\frac{X}{10^5}}{1+\frac{X}{10^5}}\right]\cdot\left[\frac{\frac{X}{10^4}}{1+\frac{X}{10^4}}\right] \quad (59)$$

Figure 11: (**A-D**) The series which computes $\log_{10}$. The total output (blue) of the first five receptors for various range of signal plotted on $\log_{10}$-transformed abscissa. The straight line (red) is drawn for comparison. The last receptor lacks the negative arm. The first receptor possesses the positive arm, with m=2, in order to dampen the output when signal is below one. β_1/α_1 of the five receptors were: 2.89, 6.299, 8.587, 12.36, 8.317, respectively. The R^2 was 0.9989.

Figure 12: (**A-D**) The series which computes $\log_e$. The total output (blue) of the first five receptors for various range of signal plotted on $\log_e$-transformed abscissa. As earlier, the last receptor lacks the negative arm and the first receptor possesses the positive arm with m=2. β_1/α_1 of the five receptors were: 6.654, 14.5, 19.77, 28.47, 19.15, respectively. The R^2 was 0.9989.

3. Concluding Remarks

Sensory systems need to respond to stimuli which vary over several orders of magnitude. One way to achieve this is by "resetting" the system, that is, by employing the mechanism of exact adaptation. The sensory mechanism presented in this work differs fundamentally from the one which uses exact adaptation. In the latter mechanism it's the transient response that provides information about the signal, as the system's output returns back to its resting value. On the other hand, in the former mechanism the equilibrium state of the system possesses information about the signal.

It should be noted that there has to be a physical limit on the maximum production rate (β_1). However, because the expression 2 (and hence also 17) is linear in β_1, the overall output can be increased by having the signal engender the production from multiple Z producers. There is also a physical limit to the value of the equilibrium dissociation constants, otherwise the free energy of binding will approach the thermal energy. Thus, some mechanism has to exist such that the apparent value of the equilibrium dissociation constants increase even when the equilibrium dissociation constants for the actual binding reach a limit. This can be achieved, for example, by having a mechanism in which the signal has to occupy multiple producer elements of the producer Z at the same time in order to engender production of Z. An implication of these two considerations is that the four-node motif is likely to be present in a hidden form in biological networks.

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