

Supporting Text

Clonal Expansion of Hepatocyte Variants in Chronic Hepatitis. This analysis examines the quantitative relationship between the expansion of variant hepatocyte clones during liver turnover, based on a selective advantage of the variant. Two types of selective advantage are considered together: resistance to killing and preferential cell regeneration.

Assume a liver consisting of two cell types, designated c_0 and c_1 , where c_0 cells are normal hepatocytes, and c_1 cells are variants. $c_0 = 1$ liver equivalent, and $c_1 \ll c_0$ at all times. First-order kill rate constants for the two cell types are d_0 and d_1 . Normal hepatocytes, c_0 , because they are in a vast majority, can be considered to regenerate at the same absolute rate as they are killed, and, with a first-order growth rate constant $k_0 = d_0$. Variant hepatocytes, c_1 , may die and regenerate with different probabilities from that of c_0 , so that k_1 and d_1 might not be equal to k_0 or d_0 . Because $c_0(t) \gg c_1(t)$, the extra mass added to the liver by the addition or depletion of c_1 variants is neglected.

$$\frac{dc_0(t)}{dt} = c_0(t)(k_0 - d_0) = 0, \text{ and} \quad \frac{dc_1(t)}{dt} = c_1(t) \cdot (k_1 - d_1).$$

If $a = \frac{k_1}{k_0}$ and $b = \frac{d_1}{d_0}$, where $a, b \geq 0$. The expansion of the variant is as follows:

$$\frac{c_1(t)}{c_0(t)} = e^{(a \cdot k_0 - b \cdot d_0)t}, \text{ and} \quad \ln\left(\frac{c_1(t)}{c_1(0)}\right) = (a \cdot k_0 - b \cdot d_0) \cdot t, \text{ or since } k_0 = d_0,$$

$$\ln\left(\frac{c_1(t)}{c_1(0)}\right) = (a - b) \cdot d_0 \cdot t.$$

Because $d_0 \cdot t$ is the total hepatocyte killing, or the total turnover, $T(t)$, the relationship between clonal expansion of c_1 and turnover is as follows:

$$\ln\left(\frac{c_1(t)}{c_1(0)}\right) = (a - b) \cdot T(t)$$

Fig. 5, which is published as supporting information on the PNAS web site, illustrates this relationship for five values of $a - b$, where $a - b = 0$ represents no selective advantage for the variant.

Clonal Outgrowth due to Random Turnover. We previously published an analysis of the relationship between random turnover of a population of cells and the clonal distribution of cells in that population (1). Briefly, the distribution $F(n, k)$ of clones, where n is the number of members in a clone and k is the number of cycles of random cell killing followed by random cell division, is given by recursive computation of the two equations:

$$f(n, k) = F(n, k - 1) / c - n \cdot F(n, k - 1) / c + (n + 1) \cdot F(n + 1, k - 1) / c$$

$$F(n, k) = f(n, k) - n \cdot f(n, k) / (c - 1) + (n - 1) \cdot f(n - 1, k) / (c - 1)$$

where $F(n = 1, 0) = c$, $F(n > 1, 0) = 0$, $f(0, k) = 0$, and $n = 0 \rightarrow c$. To calculate how many cycles of killing would be required to expand one cell out of 5×10^5 cells to a size of 1,000 or more cells, the equations were computed until

$$\sum_{n=1,000}^{n=c} F(n, k) = \frac{c}{5 \cdot 10^5},$$

where $c = 10,000$. The expression represents the number of clones of size >1000 cells when c cells undergo k cycles of killing and regeneration. When $c = 10,000$ the number of clones is 0.02. On average, one clone will expand to a size of 1,000 cells or more when $10,000/c$, or 5×10^5 cells, undergo k/c turnovers. The number of turnovers that satisfied this condition was 123.54.

1. Summers, J., Jilbert, A. R., Yang, W., Aldrich, C. E., Saputelli, J., Litwin, S., Toll, E. & Mason, W. S. (2003) *Proc. Natl. Acad. Sci. USA* **100**, 11652–11659.