

**LETTER TO THE EDITOR**

Dear Editor,

*An intuitive insight into a result of Cowan*

Motivated by a problem in DNA replication, Cowan (2001) considered a sequence of water springs distributed along a straight road out of town according to a Poisson process of rate  $\lambda$ . An infinite team of workers leaves the town at constant speed  $r$ ; on reaching any spring, one worker peels off and builds a pipe back towards town at rate  $c$ . The first worker stops when he reaches town, later ones stop when they reach the previous spring.

Given  $t > 0$ , let  $N_t$  be the number of workers building at time  $t$ , and let  $g_j(t) = P(N_t = j)$ ,  $g_j = \lim_{t \rightarrow \infty} g_j(t)$ , and  $\phi(s) = \sum_{j \geq 0} g_j s^j$ , the corresponding probability generating function of the asymptotic number actually building. Cowan showed that  $\phi(s) = \prod_{n \geq 1} (1 + (s - 1)b^n)$ , where  $b = r/(c + r)$ , and noted that, if  $\{I_n\}$  are independent Bernoulli( $b^n$ ) variables, then  $\phi$  is the probability generating function of  $\sum_{n \geq 1} I_n$ . We give here an intuitive explanation of this elegant representation.

Suppose that  $\{X_i : i = 1, 2, \dots\}$  are independent  $\text{Exp}(\lambda)$  variables, i.e. exponential with mean  $1/\lambda$ , representing the distance between successive springs, and let  $S_n = \sum_{i=1}^n X_i$ , so that  $S_1 < S_2 < \dots$  denote the positions of the springs. Let  $A_t = \max\{n : S_n < rt\}$  be the label of the last spring found before time  $t$ . When  $A_t \geq 1$ , define  $Y_0(t) = rt - S_{A_t}$  and write  $Y_n(t) = X_{A_t+1-n}$  for each  $n = 1, 2, \dots, A_t$ . Then  $N_t = \sum_{n \geq 1} I_n(t)$ , where

$$I_n(t) = \begin{cases} 1 & \text{if } Y_n(t) > \frac{c(Y_0(t) + Y_1(t) + \dots + Y_{n-1}(t))}{r}, \\ 0 & \text{otherwise.} \end{cases}$$

We seek the asymptotic distribution of  $N_t$ . Note that the variates  $\{Y_i(t) : i \geq 0\}$  are not independent and have a complicated joint distribution. As  $t \rightarrow \infty$ , however, it is intuitively clear that these variates are asymptotically independent and  $\text{Exp}(\lambda)$  distributed. Formal passage to the limit of  $I_n(t)$  is difficult but, by using the intuitively clear asymptotics for the  $Y$  variates, we can describe the limiting *properties* of  $\{I_n(t)\}$ : let  $Y_0, Y_1, \dots$  be independent  $\text{Exp}(\lambda)$  variables, and write  $N = \sum_{n \geq 1} I_n$ , where

$$I_n = \begin{cases} 1 & \text{if } Y_n > \frac{c(Y_0 + Y_1 + \dots + Y_{n-1})}{r}, \\ 0 & \text{otherwise.} \end{cases}$$

It is straightforward to show that  $P(I_n = 1) = b^n$ , as  $Y_n$  is independent of  $Y_0 + Y_1 + \dots + Y_{n-1}$ , which has a gamma density. For independence, it is sufficient to show that  $P(I_1 = 1, I_2 = 1, \dots, I_n = 1) = P(I_1 = 1)P(I_2 = 1) \dots P(I_n = 1)$  for all  $n \geq 1$ . We do this explicitly for the case  $n = 3$ : the general case follows by the same method. Plainly, we may take  $\lambda = 1$ . Now,  $P(I_1 = 1, I_2 = 1, I_3 = 1)$  is the same as

$$P\left(Y_1 > \frac{cY_0}{r}, Y_2 > \frac{c(Y_0 + Y_1)}{r}, Y_3 > \frac{c(Y_0 + Y_1 + Y_2)}{r}\right),$$

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which can be written as an integral and then evaluated as

$$\begin{aligned}
 & \int_0^\infty e^{-y_0} \int_{cy_0/r}^\infty e^{-y_1} \int_{c(y_0+y_1)/r}^\infty e^{-y_2} \int_{c(y_0+y_1+y_2)/r}^\infty e^{-y_3} dy_3 dy_2 dy_1 dy_0 \\
 &= \int_0^\infty e^{-y_0/b} \int_{cy_0/r}^\infty e^{-y_1/b} \int_{c(y_0+y_1)/r}^\infty e^{-y_2/b} dy_2 dy_1 dy_0 \\
 &= b \int_0^\infty e^{-y_0/b^2} \int_{cy_0/r}^\infty e^{-y_1/b^2} dy_1 dy_0 \\
 &= b \cdot b^2 \int_0^\infty e^{-y_0/b^3} dy_0 \\
 &= b \cdot b^2 \cdot b^3 \\
 &= P(I_1 = 1) P(I_2 = 1) P(I_3 = 1).
 \end{aligned}$$

### Reference

COWAN, R. (2001). A new discrete distribution arising in a model of DNA replication. *J. Appl. Prob.* **38**, 754–760.

Yours sincerely,

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