

On Fisher's geometrical model

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Fisher's geometrical model considers evolution in a high-dimensional phenotypic space. Let us consider a situation in which fitness is a function of n quantitative traits, $x = (x_1, \dots, x_n)$, where n is a large number. We assume that the optimal phenotype lies at a point O in the space of phenotypes, and that the fitness decreases as the Euclidean distance from the maximum increases. Without loss of generality, we can choose O as the origin of the coordinates. Then the surfaces of equal fitness are spheres centered in O , and the larger the radius of the sphere, the smaller the fitness value.

Let us now consider an organism whose phenotype is not optimal, and is described by a point P at some distance from O . We would like to evaluate the probability that a random mutation, that generates a phenotype at a distance r from P , produces a phenotype which is better adapted than P . Without loss of generality, we can choose the unit of length to be equal to twice the distance of P from O , so that P lies on a sphere Σ , centered in O , of radius equal to $\frac{1}{2}$. The mutated phenotype lies on a sphere S centered in P of radius r . Its fitness will be larger than that of P if it lies in the interior of Σ . We wish therefore to evaluate the fraction of the surface of S that lies in the interior of Σ (cf. Figure 1).

Let us first remind that the surface of a sphere of radius r in n dimensions is given by $S_{n-1} r^{n-1}$, where the constant S_{n-1} is given by

$$S_{n-1} = \frac{2\pi^{n/2}}{\Gamma(n/2)}, \quad (1)$$

where $\Gamma(z)$ is Euler's gamma function.

The condition that a point Q belonging to S lies in the interior of Σ is equivalent to the condition that its projection on the OP radius lies at a distance $(\frac{1}{2} - w)$ from O , where $w \geq u$, and u is the distance from P of the projection Q' on OP of a point Q lying at the intersection of Σ and S . To evaluate u let us consider a point Q belonging to both Σ and S , and let us draw the plane containing O , P and Q (Figure 1), with the x -axis along OP and the y -axis normal to it. Let the coordinates of Q be $(\frac{1}{2} - u, y)$. We then have

$$u^2 + y^2 = r^2; \quad (2a)$$

$$\left(\frac{1}{2} - u\right)^2 + y^2 = \frac{1}{4}. \quad (2b)$$

This implies

$$u = r^2. \quad (3)$$

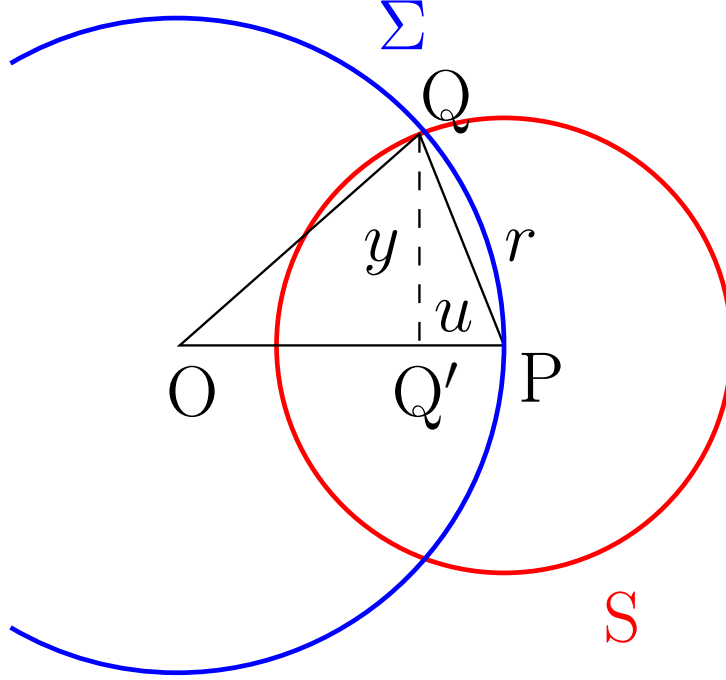


Figure 1: Fisher geometrical model. The sphere Σ is a sphere of radius $\frac{1}{2}$ in an n -dimensional space, with $n \gg 1$, centered in O . The sphere S is an analogous sphere of radius r , centered in P , where P belongs to Σ . One wishes to evaluate the fraction of the surface of S that lies in the interior of Σ . This corresponds to all points of S whose distance from P along the radius OP are larger than u , where u is the distance from P of the projection Q' on OP of the points (like Q) that belong to both Σ and S .

We must now evaluate the surface of the spherical cap corresponding to $w > u$. This is given by the surface of S , divided by 2, minus the surface of the zone (region of the sphere between two parallels) determined by the condition $0 \leq w \leq u$. Let us consider a zone determined by the parallels w and $w + dw$. The sum y^2 of all coordinates but the first one must satisfy

$$y^2 + w^2 = r^2. \quad (4)$$

Thus, for a given value of w , the region of space that satisfies this condition is a sphere in $(n - 1)$ dimensions of radius $\sqrt{r^2 - w^2}$, which has a total area equal to $S_{n-1}(r^2 - w^2)^{n/2-1}$. On the other hand, the area of the zone intercepted by dw will be proportional to $\sqrt{dw^2 + dy^2}$, which is readily seen to be equal to $dw r / \sqrt{r^2 - w^2}$. Thus the area of the zone we are interested in is given by

$$\mathcal{A} = r \int_0^{r^2} dw S_{n-2} (r^2 - w^2)^{n/2-1}. \quad (5)$$

One can check that

$$\int_{-r}^r dw r S_{n-2} (r^2 - w^2)^{n/2-1} = r^{n-1} S_{n-2} \int_{-1}^1 du (1 - u^2)^{n/2-1} = S_{n-1} r^{n-1}. \quad (6)$$

Let us now evaluate \mathcal{A} for $n \gg 1$. We have

$$\begin{aligned}\mathcal{A} &= r \int_0^{r^2} dw S_{n-2} (r^2 - w^2)^{n/2-1} \\ &= r^{n-1} S_{n-2} \int_0^r du (1 - u^2)^{n/2-1}.\end{aligned}\quad (7)$$

Assuming that $r \ll 1$ (whose consistency can be checked later) we can express the integrand as follows:

$$(1 - u^2)^{n/2-1} \approx e^{-nu^2/2}, \quad (8)$$

where we have also neglected $1/2$ with respect to n . Define the function

$$\phi(x) = \sqrt{\frac{2}{\pi}} \int_0^x dt e^{-t^2/2}. \quad (9)$$

This function vanishes for $x = 0$ and approaches 1 as $x \rightarrow \infty$. We then have

$$\mathcal{A} = r^{n-1} S_{n-2} \sqrt{\frac{\pi}{2n}} \phi(r\sqrt{n}). \quad (10)$$

The probability of being in the considered zone is given by $\mathcal{A}/(S_{n-1}r^{n-1})$, i.e.,

$$P_1 = \frac{S_{n-2}}{S_{n-1}} \sqrt{\frac{\pi}{2n}} \phi(r\sqrt{n}) \simeq \frac{1}{\sqrt{2}} \phi(r\sqrt{n}). \quad (11)$$

Now we have

$$\lim_{x \rightarrow \infty} \frac{\Gamma(x + 3/2)}{\sqrt{x} \Gamma(x + 1)} = 1, \quad (12)$$

as can be checked from Stirling's formula, and therefore

$$\lim_{n \rightarrow \infty} \frac{S_{n-2}}{S_{n-1}} \sqrt{\frac{\pi}{2n}} = \lim_{n \rightarrow \infty} \frac{\Gamma(n/2)}{\sqrt{\pi} \Gamma((n-1)/2)} \sqrt{\frac{\pi}{2n}} = \frac{1}{2}. \quad (13)$$

Thus the probability of being in the cap is given by

$$P = \frac{1}{2} - P_1 = \frac{1}{2} (1 - \phi(r\sqrt{n})). \quad (14)$$

Since $\phi(u)$ approaches 1 rapidly when its argument grows beyond 1, the probability of being in the cap, i.e., that the mutation is beneficial, vanishes rapidly when $r \gtrsim 1/\sqrt{n}$, which, for large n , is a small number. Thus, Fisher concludes, the probability that a mutation is beneficial does not substantially vanish only when their phenotypic effect is very small. The behavior of this function is shown in Figure 2.

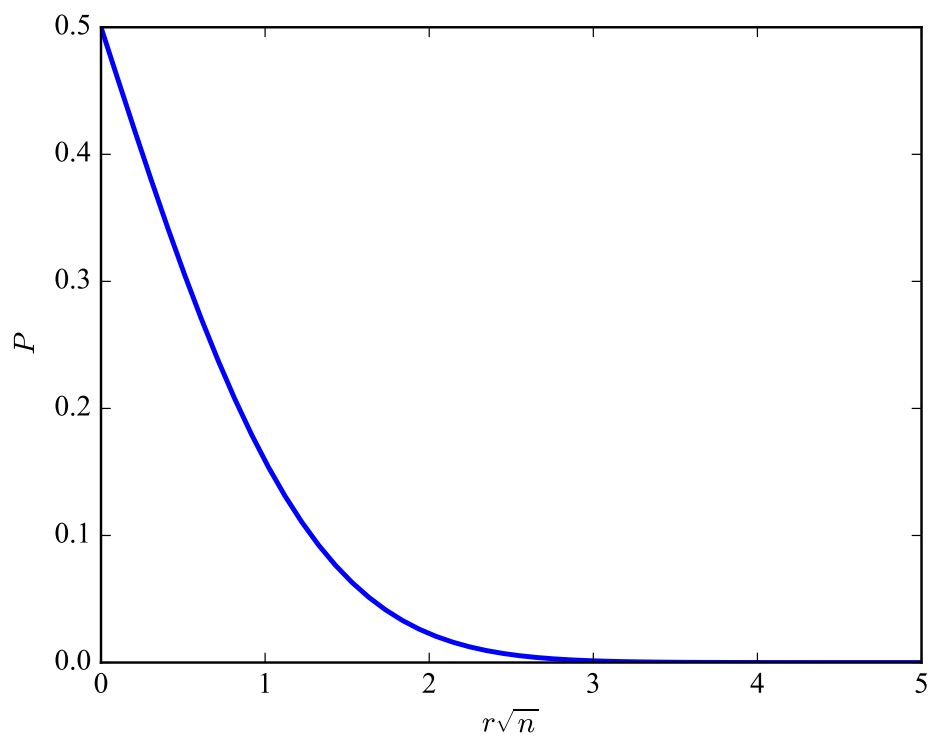


Figure 2: Probability of a beneficial mutation as a function of its size r , according to equation (14).