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TEMPLE BAR SES

STRAND, W. C. 2.

2nd July, 1930.

Dear Fisher:

I gathered from a passage in your book that you probably regarded all mutations as having begun first as recessives, and later become dominants, but evidently I was mistaken in that. I, like you, am inclined to doubt if any mutation is strictly completely dominant in the quantitative sense.

As regards <u>rubricalyx</u>, it has the same chromosome linkage (four pairs and a ring of six) as <u>rubrinervis</u>, from which it arose, and all the evidence shows that it was originally a simple gene mutation.

with regard to blood groups, of course Bernstein and Furuhata have proposed a multiple allelomorph series, 0, A, B, but from later evidence I am rather inclined to favour the view of two pairs of factors with linkage and crossing-over. 106 generations certainly seems a long time, and it seems impossible that the gene A arose as much as 25,000,000 years ago. In your book, if I remember, you postulated mutation rates of the order of one in one million. I do not quite understand the basis of this estimate, which seems to me low,

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if Drosophila is any sort of criterion, but perhaps my mathematics are astray !

I take it that as regards the dispersal of A and B, crossing and migration of races alone without repeated mutations would not be adequate to account for the present frequencies.

Forgive me for troubling you, but it seems to me that if anything can contrinute to analysis of this kind, your mathematics can.

Yours sincerely,