

FIG. 1. Jump-diffusion foraging model parametrized **b**ye probability of jumping. (a) The forager moves to nonadjacent sites with probability and to adjacent sites with probability $\frac{1}{2}$, accounting for the possibility of diffusion. (b)—(d) Example forager paths (blue lines and dots) for Q, 1, QO3. Green represents sites with food, while white represents empty sites.

jumps with probability, or diffuses with probability β_j [Fig. 1(d)]. Providing our forager with both types of movement allows us to consider how much time the forager should spend exploiting a given location, and how frequently the forager should move to other locations. We demonstrate that the mean lifetime of the forager varies nonmonotonically with respect p_0 , and the forager $\tilde{\mathbf{G}}$ lifetime is maximized through a mixture of jumping and diffusion.

This work extends the recent studies90f [

FIG. 3. (a), (b) Forager lifetime computed from E(a) (n the case of pure diffusional

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drops as we change 1 tos = 2 since in the case of pure diffusion (and = 1) the forager will live at least two time steps, whereas the pure jumper may not. Asseo2, dtais effect becomes negligible. Furthermore, this drop in the ratio becomes less severe for larger valuesince the jumper will almost always live at least two time steps.3(dFig. we display the ratio as a surface plot along boahdthe axes. Increasingclearly expands the region (outlined) of values, for which diffusion is a better strategy. Note that for very small values of ($n \le 5$) the cover time for diffusion is less than the cover time for jumping, leading to an advantage of jumping over diffusion at those parameter valuesis. When large relative to the diffusive forager benet to form a larger



when using a mixture of these modes of movement. This explanation is further validated by the qualitative similarities of the jump-diffusion and jump-wait models. In either model, makings larger orn smaller lead to situations in which the

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