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The greater the genetic distance to the focal point, the greater the effect of parentage. Genetic parentage inference is of critical interest in a variety of areas, notably paternity cases, most notably in human genetics, but also in ecology, conservation genetics, and animal breeding. In this note, we offer some simpler methods for developing tests that infer the sire of a focal individual using the parentage data of a population of known individuals. We begin with the basics: if a father is not a member of the population from which the focal individual came, he is more likely to be its father than are other males in the population. In contrast, if a father is in the population from which the focal individual came, a randomly chosen male in the population is more likely to be its father than are all other males in the population. We present an extension of this heuristic to provide bounds for specific comparisons, including the proportion of times that the focal individual is more likely to be sired by a given individual than by all other members of the population. This heuristic proves to be a powerful method for inference in its own right, yielding estimates of the proportions of time that the focal individual is sired by different members of the population that are nearly identical to those obtained from the full pedigree method of parentage inference. Furthermore, because of its simplicity, this heuristic can be used to gain insights into the question of what fraction of times the focal individual is more likely to be sired by a given member of the population than by the other members.

$-2 \cdot h^{**2}$. What is $g(-4)$? 8 Let $m(a) = a + 2 \cdot a^{**2} + 5 \cdot a - 7 - a^{**}$

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