

(Large-)team exclusion of women researchers under unequal gender ratio and homophily

Zeyuan Chen

January 26, 2026

Question: Given a population (with an infinite size) of researchers exhibiting homophily with a specified gender ratio, we ask whether the expected proportion of male members in a team of a certain size differs from the global proportion of males, and how this difference varies with the parameters.

In other words, we aim to express it as a function of team size (T), gender ratio, and homophily. Formally, let the population consist of N_m males and N_f females, with corresponding proportions P_m and P_f . Assume that both males and females have the same tendency for homophily: each individual chooses to collaborate with a same-gender partner with probability s ($s > 0.5$ indicates homophily). The gender ratio is defined as $r = \frac{P_m}{P_f} = \frac{N_m}{N_f}$. We are then interested in computing

$$E\left(\frac{n_m}{T} \mid T, r, s\right) - P_m,$$

where n_m denotes the number of male members in a team.

In the seed author model (I believe which is a **stronger** version of Guimera's team-assembly model, where the other members are selected based on a random member among all existing members.), at each step we randomly select a seed author from the population, who is male with probability P_m and female with probability P_f . Next, based on the seed author's gender and the homophily parameter s , we determine the genders of the remaining $T - 1$ authors in the team: if a candidate has the same gender as the seed author, they are selected with probability $s \cdot \frac{1}{N}$; otherwise, with probability $(1 - s) \cdot \frac{1}{N}$.

Consequently, if the seed author is male, the expected number of males among the remaining $T - 1$ members is

$$(T - 1) \times \frac{sN_m}{sN_m + (1 - s)N_f}.$$

Including the male seed author, the expected proportion of males in a team seeded by a male author is

$$E_{mm} = \frac{1}{T} \left(1 + (T - 1) \times \frac{sN_m}{sN_m + (1 - s)N_f} \right).$$

Similarly, if the seed author is female, the expected number of males among the remaining $T - 1$ members is

$$(T - 1) \times \frac{(1 - s)N_m}{(1 - s)N_m + sN_f}.$$

Including the female seed author, the expected proportion of males in a team seeded by a female author is

$$E_{fm} = \frac{1}{T} \left(0 + (T - 1) \times \frac{(1 - s)N_m}{(1 - s)N_m + sN_f} \right).$$

Therefore, in any given step, when randomly assembling a team of size T , the expected proportion of males is

$$E_m = P_m E_{mm} + P_f E_{fm}.$$

After some calculations, we arrive at the following final formula:

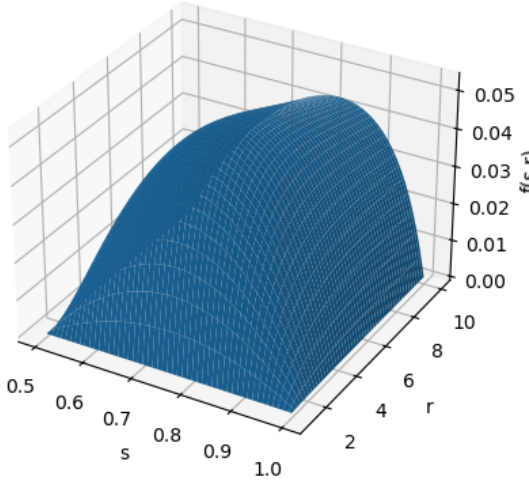
$$E_m - P_m = \frac{T - 1}{T} \left(\frac{r(1 - s)(2s - 1)(r - 1)}{(r + 1)(sr + 1 - s)(r - sr + s)} \right).$$

From this expression, two observations can be immediately made:

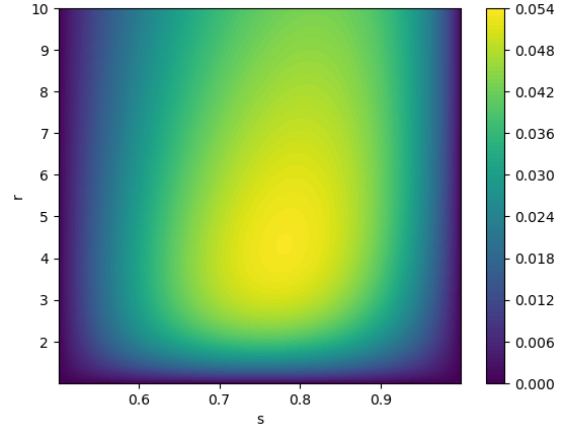
1. This function is **always positive**. Under settings with homophily ($0.5 < s < 1$) and unequal gender ratio ($r > 1$), the numerator $r(1 - s)(2s - 1)(r - 1)$ is strictly greater than zero. In other words, regardless of team size T (which is at least 2), as long as there is some degree of gender imbalance and homophily, male members are always more likely than female members to be selected into a team.
2. This function is **monotonically increasing with team size T** . That is, larger teams

are more biased toward selecting male members compared to smaller teams.

As T only functions as a scaling factor, we could get rid of it and focus on the expression $(f(s, r))$ inside. As Figure 1 shows, gender disparity regarding being assembled into a team is most severely amplified in environments with moderate homophily.



(a) Surface plot of $f(s, r)$



(b) Contour plot of $f(s, r)$

Figure 1: Visualization of the function $f(s, r)$ showing (a) the 3D surface and (b) the contour representation.