FIG. S14. Time evolution of the pairwise mean Hamming distance in the $f_{sm}$. Parameters are $N = 100$, $L = 10$, $p = 0.5$ with four different values of the mutation rate $\mu$. Each panel compares obligately recombining ($r = 1$) and non-recombining ($r = 0$) populations. Thick lines represent the mean over 5000 landscape realizations and the shaded areas the corresponding standard deviation.