Figure S2.—Diversity in recombining and non-recombining steady-state populations. Shown as a function of population size are a) number of different genotypes in the population, b) mean average genotypic distance (see Methods), c) genotype diversity index, for populations in mutation-selection balance (open circles) and mutation-selection-recombination balance (closed circles). Data are based on circuits with \( N=12 \) genes, number of regulatory interactions per circuit in the interval \((N,3N)\), \( \mu=0.1 \) mutations per circuit and generation, and orthogonal vectors \( S(0) \) and \( S(\infty) \). Note that all relevant circuit properties depend only on the angle between these vectors (Ciliberti et al. 2007).