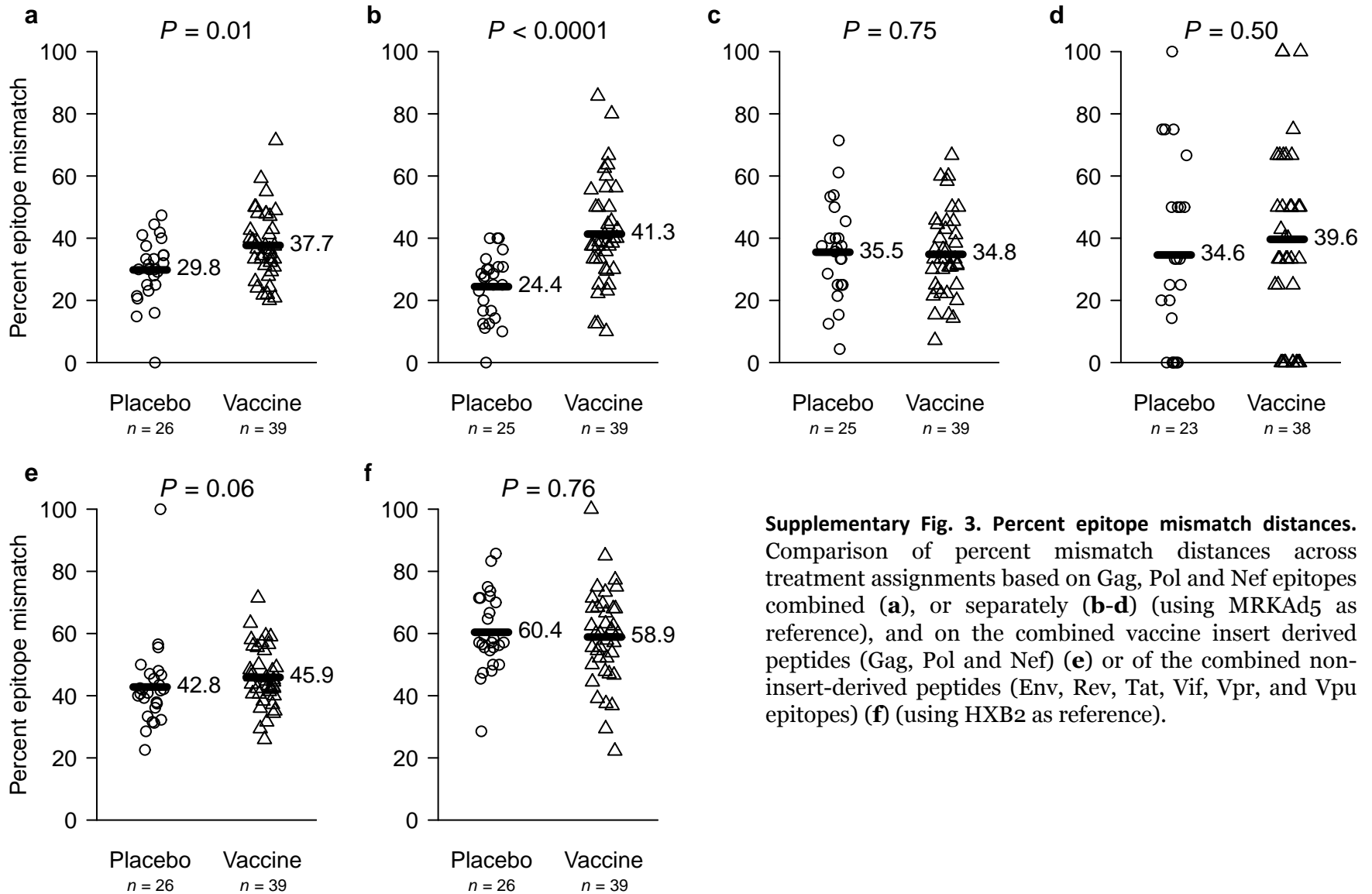Supplementary Fig. 3. Percent epitope mismatch distances. Comparison of percent mismatch distances across treatment assignments based on Gag, Pol and Nef epitopes combined (**a**), or separately (**b-d**) (using MRKAd5 as reference), and on the combined vaccine insert derived peptides (Gag, Pol and Nef) (**e**) or of the combined non-insert-derived peptides (Env, Rev, Tat, Vif, Vpr, and Vpu epitopes) (**f**) (using HXB2 as reference).