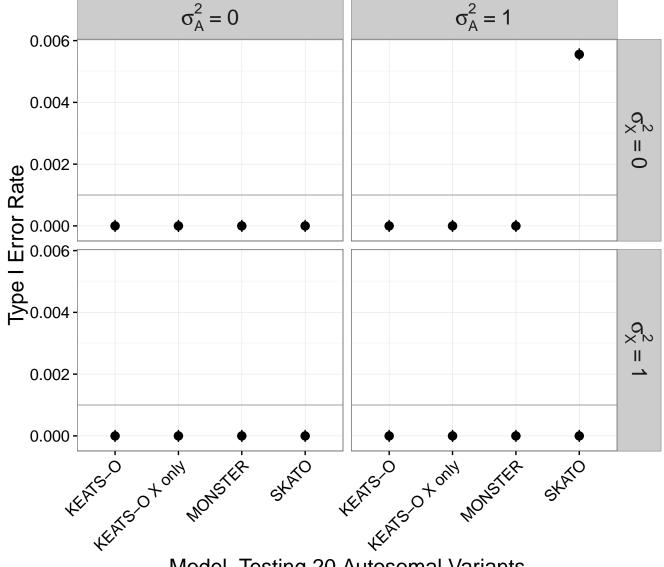
Type I Error Rate, α =0.001 rarecommon variants, AF diff in 0.2, 0.4



Model, Testing 20 Autosomal Variants