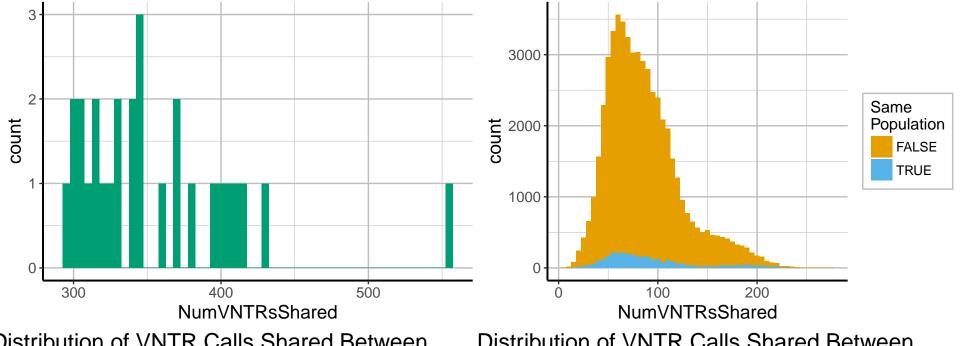
Distribution of VNTR Calls Shared Between Distribution of VNTR Calls Shared Between Two Genomes, all pairs (Source == "WG\$\forallow 00Genomes, all pairs (Source == "1000 Genomes")



Distribution of VNTR Calls Shared Between Distribution of VNTR Calls Shared Between nomes, all pairs (Source == "Platinum **Toem Greeo**mes, all pairs (Source == "HapMap Yoruban Far 1.00 10 0.75 Same Same count 0.50 conut Population **Population** TRUE **TRUE** 0.25 0.00 500 550 2000 450 600 1800 1900 2100 2200 NumVNTRsShared NumVNTRsShared