GOEAs recovering HR GO IDs; propagate_counts=True 1.00 **B**1 100% Null 0.075 Genes 0.75 0.050 0.50 0.025 0.25 0.000 0.00 1.00 0.075 0.050 0.025 **General Population** 0.75 0.50 9% 0.25 0.000 0.00 1.00 **B**3 0.075 50% Null 0.75 0.050 0.50 0.025 0.25 0.000 0.00 1.00 34%64%83% 90% 0.075 25% Null 0.75 0.050 0.50 of 0.025 0.25 0.000 0.00 Percentage 1.00 0.075 0.050 0.025 0.075 0.75 0.50 0.25 0.000 0.00 16 8 FDR Sensitivity Specificity Number of Genes in a Study Group