0.5 Cross validated R² 0.4 0.3 sRRR, r=13 0.2 sRRR, r=2 0.1 0.0 ò 20 40 80 60 Number of selected genes E_{leak} 9м g_{Na} au_{max} R_{input} g_L Multivariate $R^2 = 0.20$ τ g_{Nat} g_{Kv31} g_{Kd} r_{SS} 0.0 0.1 0.2 0.3 0.4 0.5 0.6 R^2