Retroposon breakdown by by eight-clusters and Family 0.5 1.00 -0.75 repName SVA_A SVA_B conut SVA_C SVA_D SVA_E SVA_F 0.25 -0.00 NR EAR_close ESR_close ESR_OC LR_open LR_close all_peaks EAR_open ESR_open mrc