CV vs. mean after EigenMS normalisation, hc = 2, con_sp.hsa.canonical.isoforms_junc.exons, protein groups 0.3 0.2 - \lesssim 0.0 0.0e+00 5.0e+07 1.0e+08 1.5e+08 2.0e+08 Mean Intensity (normalised)