PCA standard deviations based on gene expected count with RUVr treament 0.4 timepoint ud 6h 12h 0.2 24h 3d PC2 (2.05%) 9d 12d replicatenumber r1 r2 0.0 r3 size 2 -0.2 **-**0.10 0.15 0.20 0.25 PC1 (96.2%)