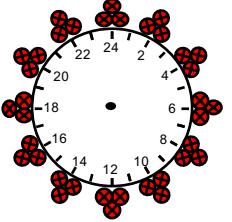


Dataset: **1 data per time-point**

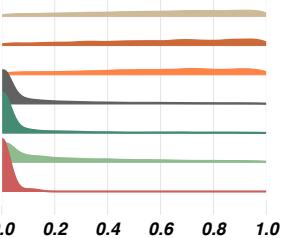
Density distribution of p-values



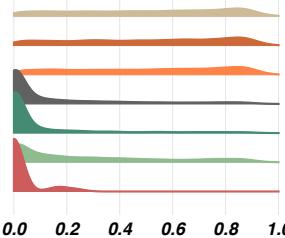
*every 2h over 72h
[36 time-points]*



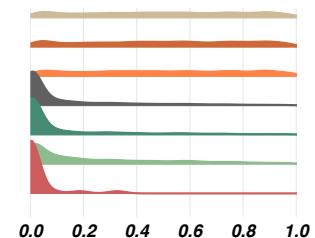
ARS



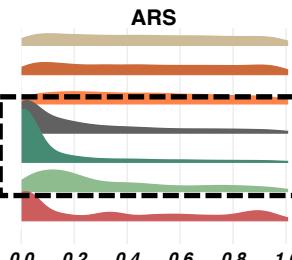
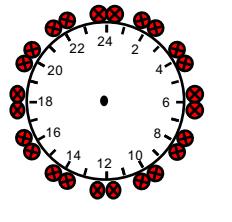
GeneCycle



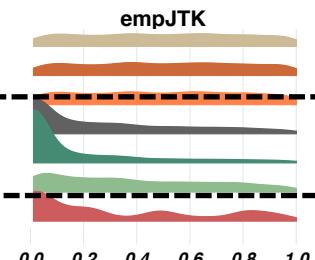
empJTK



*every 2h over 48h
[24 time-points]
(microarray)*



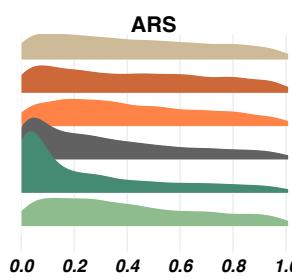
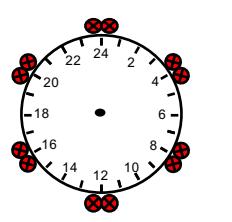
GeneCycle



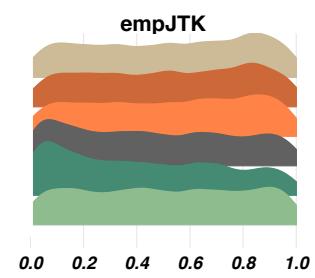
empJTK



*every 4h over 48h
[12 time-points]*



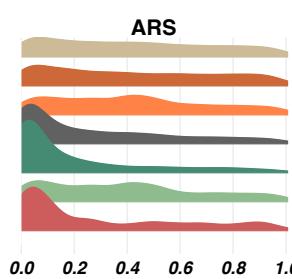
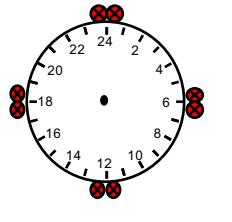
GeneCycle



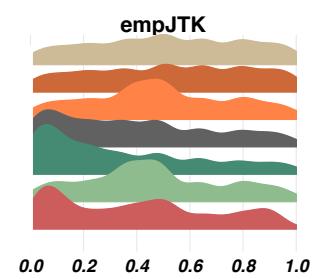
empJTK



*every 6h over 48h
[8 time-points]
(RNAseq)*



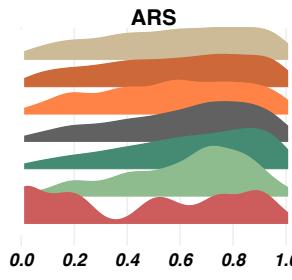
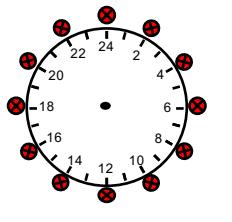
GeneCycle



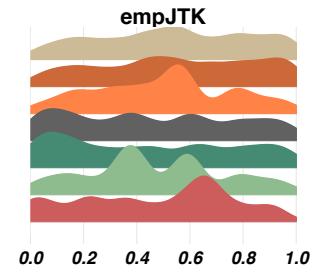
empJTK



*every 2h over 24h
[12 time-points]*



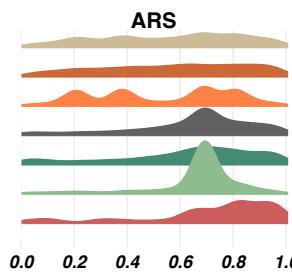
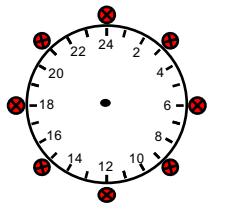
GeneCycle



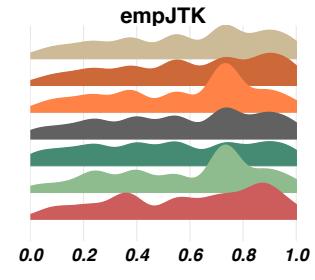
empJTK



*every 3h over 24h
[8 time-points]*



GeneCycle



empJTK

a

default p-value