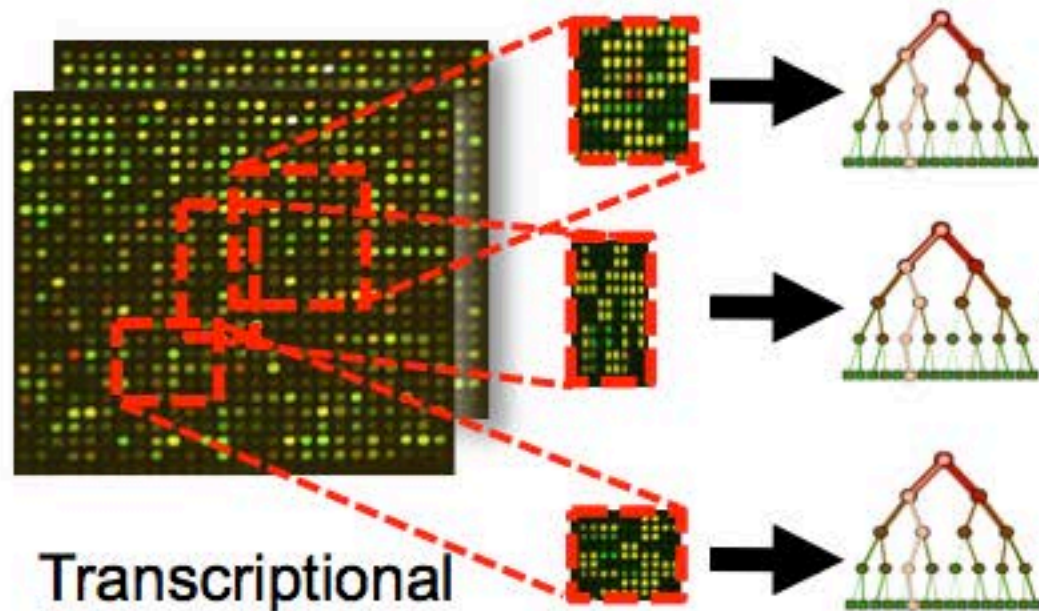


Decision trees over regulator subset
and expression subsamples



Transcriptional
Datasets

Random forest regression
for each target gene g

