Supplemental Figure S4

Fig. S4. ROC analysis to compare the ability of ChIP sites in variably sized windows to predict induced genes: cancer expression compendium and primary targets. Positives and negatives are taken as in Fig. 3. Format is identical as for Fig. 4, but different sites are used: stringent ChIP-chip sites (t>16) in the upper row and ChIP-pet sites in the bottom row.