McKinlay et al. Suppl Figure 1

GENE	$Cp_{biotin\text{-}UTP}$	Cp_{UTP}	Fold enrichment = $2^{(Cp_{UTP}-Cp_{biotin\text{-}UTP})}$
5'ETS (45S rRNA)	20.2	26.9	100.43
LHCB1.3	26.3	33.5	144.01
PSAD1	28.8	39.6	1820.35
LHCB6	30.1	38.3	292.71
ACT2	33.1	39.6	478.82
RBSC2B	33.3	37.2	15.93
AT2G45180.1	33.3	40.0	106.4
DRT112	34.1	38.8	24.76

Average Cp values correspond to the average cross-point values from triplicate NRO reactions