Differential expression analysis for individual cancer samples based on robust within-sample relative gene expression orderings across multiple profiling platforms

SUPPLEMENTARY TABLE

Supplementary Table S1: Reproducibility evaluation of significantly stable REOs detected in small samples for each platform

	Platform	Normal	Number	POG
Lung				
GSE31210	Affymetrix	20	191,721,331	0.8923
GSE31267	Illumina	24	247,667,868	0.9168
GSE15197	Agilent	13	156,176,364	0.7923
Colorectal				
GSE21510	Affymetrix	25	190,854,700	0.9475
GSE31279	Illumina	32	135,225,886	0.8578
GSE46271	Agilent	22	137,856,117	0.8453

Note: POG denotes the percentage of the significantly stable REOs for the combined data for each platform are included in the corresponding small sample dataset.