Analysis I
(Examination of GAPDH and HPRT1 as control genes for lung clinical samples)

Set A
Paired samples of lung cancer patients
n=15 normal; n=15 tumor

Analysis II
(Test of expression stability of 16 reference genes in a pilot study of lung clinical samples and cell lines)

Set B
Normal lung and tumor paired samples of 6 NSCLC patients
n=6 normal; n=6 tumor

Lung cultured cells
n=2 primary cells; n=7 NSCLC cell lines

Best genes for lung clinical samples
Best common genes
Best genes for lung cultured cells

Analysis III
(Clinical validation of the selected optimally performing endogenous genes)

Set C
Paired samples of patients:
n=16 normal; n=16 tumor

Paired samples of patients
n=31 normal; n=31 tumor