Supplementary Table 1: Results of the semiquantitative PCR analysis of EGFR amplification. We calculated the $E G F R / C F T R$ ratios $[x]$ from peripheral blood DNA of 20 healthy Caucasian adults. For tumor samples, a value $[x+3 \mathrm{SD}=2.3]$ was considered as the cut-off level for the normal gene copy number. Ratios higher than $[x+3 \mathrm{SD}]$ were regarded as evidence of more than 2 copies of the $E G F R$ gene.

|  | EGFR/ CFTR <br> ratios |
| :--- | :---: |
| G1p | 5.9 |
| G1r | 6.9 |
| G2p | 2.2 |
| G2r | 19.3 |
| G3p | 1.5 |
| G3r | 1.6 |
| G4p | 14.2 |
| G4r | 15.9 |
| G5p | 3.5 |
| G5r | 4.1 |
| G6p | 1.5 |
| G6r | 8.7 |
| G7p | 1.1 |
| G7r | 1.5 |
| G8p | 1.6 |
| G8r | 9.1 |
| G9p | 1.6 |
| G9r | 9.9 |
| G10p | 6.5 |
| G10r | 9.1 |
| G11p | 3.8 |
| G11r | 8.3 |
| G12p | 14.4 |
| G12r | 20.1 |
| G13p | 3.6 |
| G13r | 4.7 |
| G14p | 1.3 |
| G14r | 1.4 |
| G15p | 9.6 |
| G15r | 9.9 |
| G16p | 1.3 |
| G16r | 1.6 |
| G17p | 4.4 |
| G17r | 5.1 |
| G18p | 3.2 |
| G18r | 3.7 |
| G19p | 8.3 |
| G19r | 8.6 |
| G20p | 10.2 |
| G20r | 12.2 |
|  |  |

