

Suppl. Table 5. Numbers of genes that show significantly differential expression (false discovery rate (FDR) < 0.05 after correction for multiple testing) between UKF-NB-3 and 1 or more of its sub-lines adapted to RITA (3 sub-lines investigated: UKF-NB-3^rRITA^{10 μ M}II, UKF-NB-3^rRITA^{10 μ M}III, UKF-NB-3^rRITA^{10 μ M}IV) or nutlin-3 (4 sub-lines investigated: UKF-NB-3^rNutlin^{10 μ M}II, UKF-NB-3^rNutlin^{10 μ M}V, UKF-NB-3^rNutlin^{10 μ M}VI, UKF-NB-3^rNutlin^{10 μ M}VIII).

Comparison	number of differentially expressed genes
UKF-NB-3 vs. at least 1 RITA-adapted sub-line	5363
UKF-NB-3 vs. at least 2 RITA-adapted sub-lines	2030
UKF-NB-3 vs. 3 RITA-adapted sub-lines	663
UKF-NB-3 vs. at least 1 nutlin-3-adapted sub-line	8891
UKF-NB-3 vs. at least 2 nutlin-3-adapted sub-lines	5685
UKF-NB-3 vs. at least 3 nutlin-3-adapted sub-lines	3592
UKF-NB-3 vs. 4 nutlin-3-adapted sub-lines	1597