### SUPPLEMENTAL TABLE 2
Ross et al. 2015

| Gene sym. | Protein description | Mean Fragment intensity value | Basal | A-769662 treated | Background |
|-----------|---------------------|-------------------------------|-------|----------------|------------|
| FAT1      | FAT tumor suppressor 1 | 0.728 ± 0.024 | n.d. | n.d. | n.d. |
| FAT2      | FAT tumor suppressor 2 | 0.795 ± 0.035 | n.d. | n.d. | n.d. |
| COL6A3    | collagen, type VI, alpha 3 | 0.855 ± 0.009 | n.d. | n.d. | n.d. |
| COL14A1   | collagen, type XIV, alpha 1 | 0.856 ± 0.028 | n.d. | n.d. | n.d. |
| ITGA11    | integrin, alpha 11  | 0.869 ± 0.013 | n.d. | n.d. | n.d. |
| ITGA4     | integrin, alpha 4   | 0.938 ± 0.041 | n.d. | n.d. | n.d. |
| ICAM5     | intercellular adhesion molecule 5 | 0.860 ± 0.011 | n.d. | n.d. | n.d. |
| MYH13     | Myosin heavy chain 13 | n.d. | 0.772 ± 0.022 | n.d. | n.d. |
| PDZD2     | PDZ domain containing 2 | n.d. | 0.731 ± 0.037 | n.d. | n.d. |
| DNAH5     | Dynein heavy chain 5 | 0.547 ± 0.030 | 0.710 ± 0.037 | n.d. | n.d. |
| VAC14     | ArPIKfyve           | 1.295 ± 0.029 | 1.047 ± 0.017 | n.d. | n.d. |
| TLN1      | Talin 1             | 0.694 ± 0.013 | 1.235 ± 0.048 | 0.887 ± 0.012 | 0.642 ± 0.004 |
| MCAF1     | Act. Trans. Fact. 7 Interact. Protein | 0.661 ± 0.005 | 0.647 ± 0.006 | 0.642 ± 0.004 | 0.642 ± 0.004 |