

Riege et al., Figure 7 – figure supplement 1

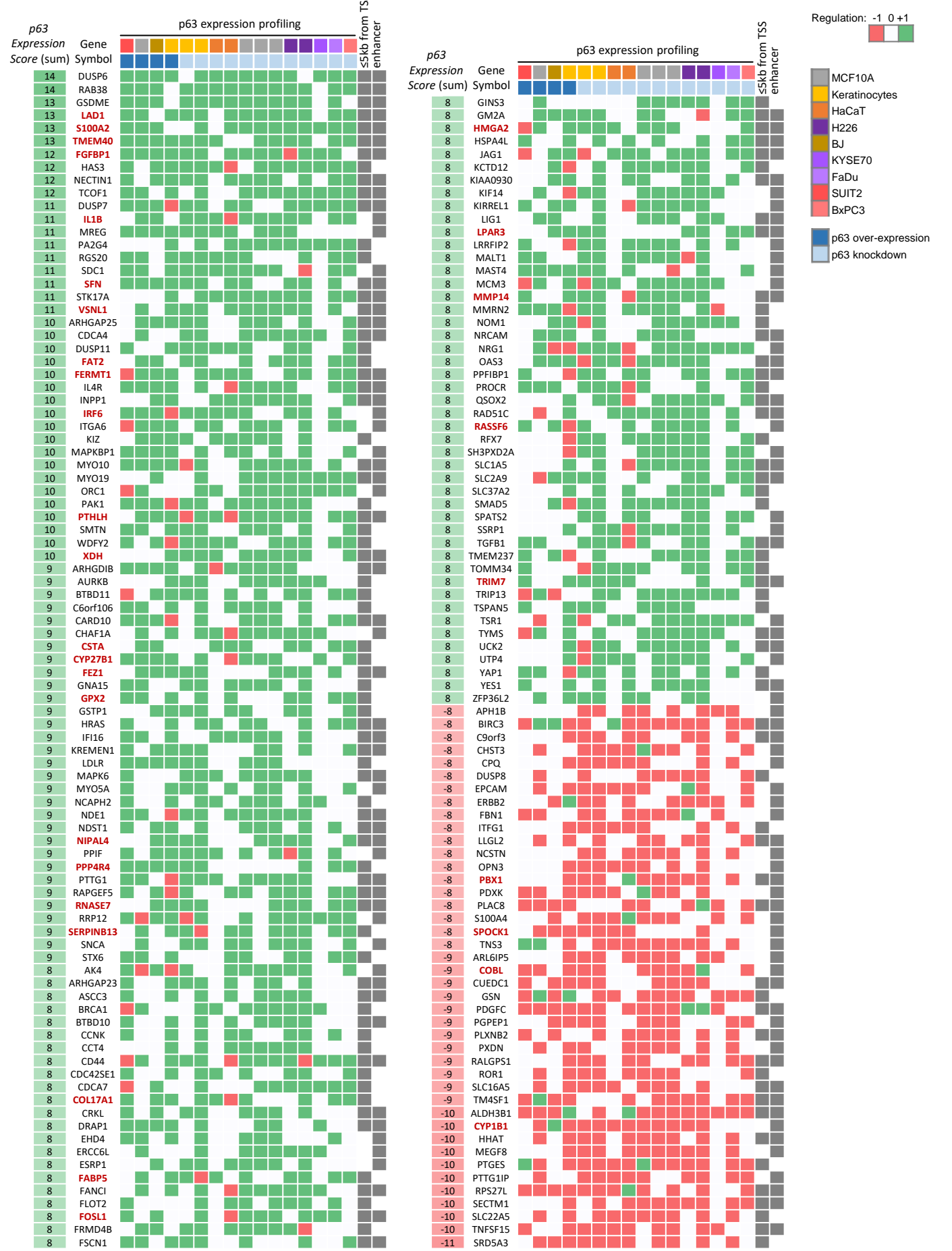


Figure 7 – figure supplement 1

Complement to Table 1. Genes identified as significantly up- or down-regulated in at least the half of all datasets ($|p63 \text{ Expression Score}| \geq 8$) that are linked to p63 binding sites supported by at least half of all datasets (≥ 10) through binding within 5 kb from their TSS or through double-elite enhancer:gene associations (Fishilevich et al., 2017). Using these thresholds we identified 138 and 42 high-probability candidates as directly up- and down-regulated by p63, respectively. Gene names marked in red are also up- or down-regulated across SCCs (Campbell et al., 2018).