

Figure S11. Number of potentially problematic bivalent domain-containing genes expressed in different human iPSC lines.

The iPSCs (different lines, different clones or different passages of the same line) from the available human datasets are represented on the X axis. For each iPSC, the number of bivalent domain-containing genes expressed in the given iPSC whereas silent in 100% (blue ■) or in at least 90% (red ■) of the human ESC lines analyzed, is represented on the Y axes.

