



Assessment of pluripotency in control and DCM iPSC clones.

A, Quantitative RT-PCR (qRT-PCR) analysis of endogenous genes associated with pluripotency (c-MYC, KLF4, OCT3/4, SOX2, NANOG, REX1, and TDGF1) in two control (1 and 2, black bars) and two DCM iPSC clones (1 and 2, grey bars). Expression values are relative to corresponding primary skin fibroblasts (PSF, red bars), normalized to GAPDH, and presented as mean \pm s.e.m., $n=3$.