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Supplemental Information

Pluri-IQ: Quantification of Embryonic Stem Cell Pluripotency through

an Image-Based Analysis Software

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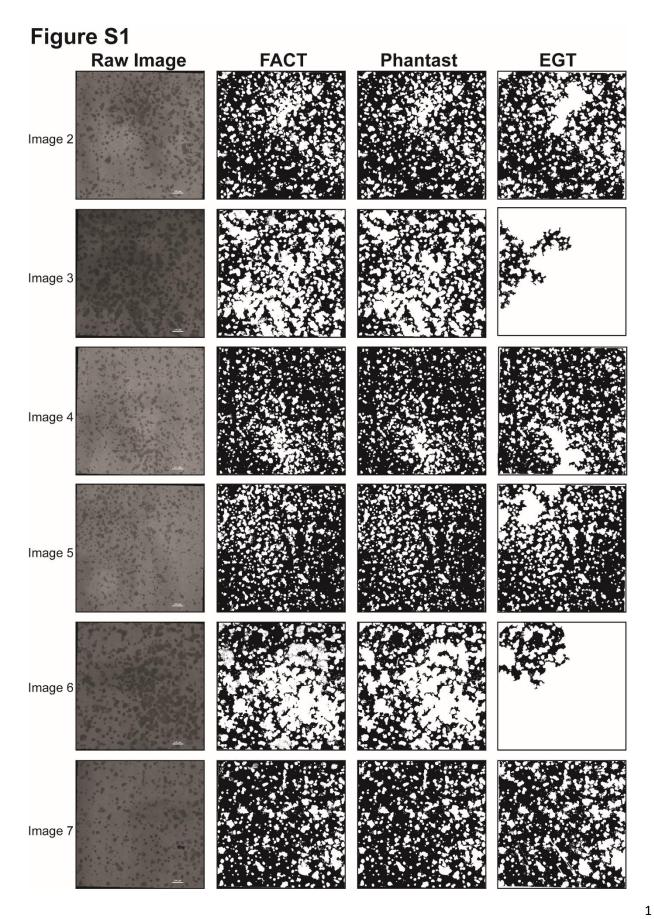


Figure S2

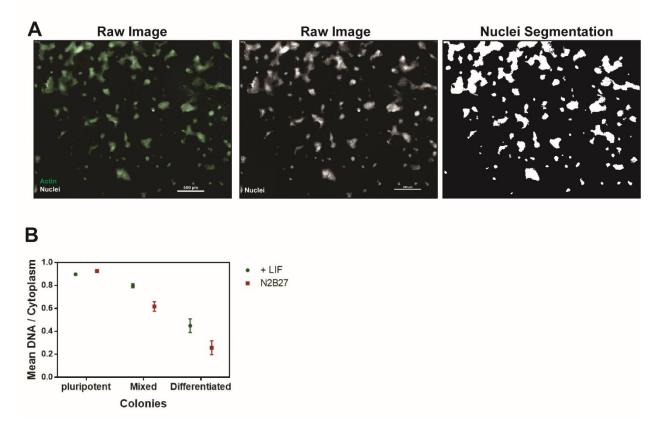


Table S1: Features used to train the random forest classifier and discriminate betweenpluripotent, mixed and differentiated colonies.Related to Figure 5.

Feature	Feature type
Colony circularity	Morphological
Intensity mode	Intensity
Intensity mean	Intensity
Intensity variance	Texture
Intensity Histogram Skew	Intensity
Variation of intensity histogram peaks	Intensity

Table S2 - Sequences of forward and n	everse primer (5' to 3	') sets used in this study, rela	ted to Figure 6.

Target Gene	Forward	Reverse	Primer Bank ID
Dppa3	GACCCAATGAAGGACCCTGAA	GCTTGACACCGGGGTTTAG	21218416a1
Esrrb	GCACCTGGGCTCTAGTTGC	TACAGTCCTCGTAGCTCTTGC	31542617a1
Fgf5	AAGTAGCGCGACGTTTTCTTC	CTGGAAACTGCTATGTTCCGAG	3721900a1
Klf4	GTGCCCCGACTAACCGTTG	GTCGTTGAACTCCTCGGTCT	6754456a1
Oct4	CGGAAGAGAAAGCGAACTAGC	ATTGGCGATGTGAGTGATCTG	356995852c3
Rex1	CCCTCGACAGACTGACCCTAA	TCGGGGCTAATCTCACTTTCAT	7110739a1
Rplp0	AGATTCGGGATATGCTGTTGGC	TCGGGTCCTAGACCAGTGTTC	6671569a1
\overline{T}	GGATTCACATCGTGAGAGTTGG	GTCACAGCTATGAACTGGGTC	118130357c3

Figure S1: Comparison between FACT algorithm segmentation and Phantast and EGT segmentation algorithms. Related to Figure 4.

FACT, Phantast and EGT segmentation results of large images with different degrees of colony density. Binary images, where white is representative of presence of a colony. Scale bar: 500 µm.

Figure S2: Nuclei cytoplasm ratio decreases upon differentiation. Related to Figure 7.

(A) Representative image of the nuclei segmentation through FACT algorithm. Raw image color code: green – actin; white – nuclei. Scale bar: 500 μ m. (B) Average nuclei cytoplasm ratio, calculated by Pluri-IQ, of pluripotent, mixed and differentiated cells when cultured in a pluripotent medium (+LIF) and in a neuronal differentiation medium (N2B27). Result is expressed as mean \pm SEM and are derived from two replicates.