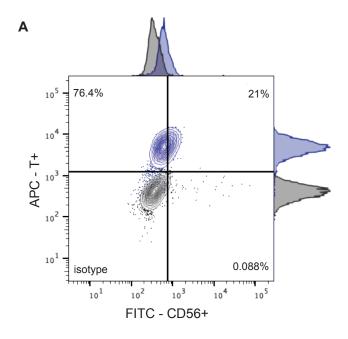
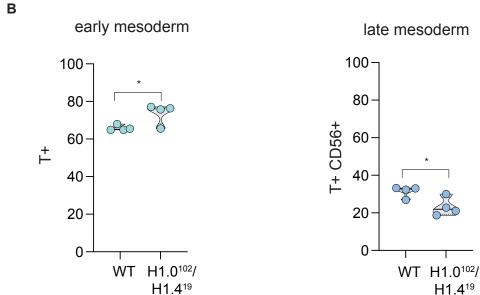
## H1.0<sup>102</sup>/H1.4<sup>19</sup> mesoderm FACS analysis





Flow cytometry analysis of direct mesodermal differentiation of linker histones H1.0/H1.4 hESC double knockout cells (clone 102/19: H1.0<sup>102</sup>/H1.4<sup>19</sup>).

- (A) Contour plot of the population distribution and percentage of cells positive for brachyury (T+) and CD56+. Isotype staining was performed as control. Result of a representative experiment is shown.
- (B) Quantification of cell populations from (A) with T+ cells representing early mesoderm (left) and T+/CD56+ cells representing late mesoderm (right). Truncated violin plot is shown. n=4 corresponding to two independent biological replicates (independent cell differentiation) and two technical replicates (FACS analysis). Black line denotes the mean; the p-values were determined by two tailed unpaired t-test \* p<0.05, \*\* p<0.01, \*\*\* p<0.001.