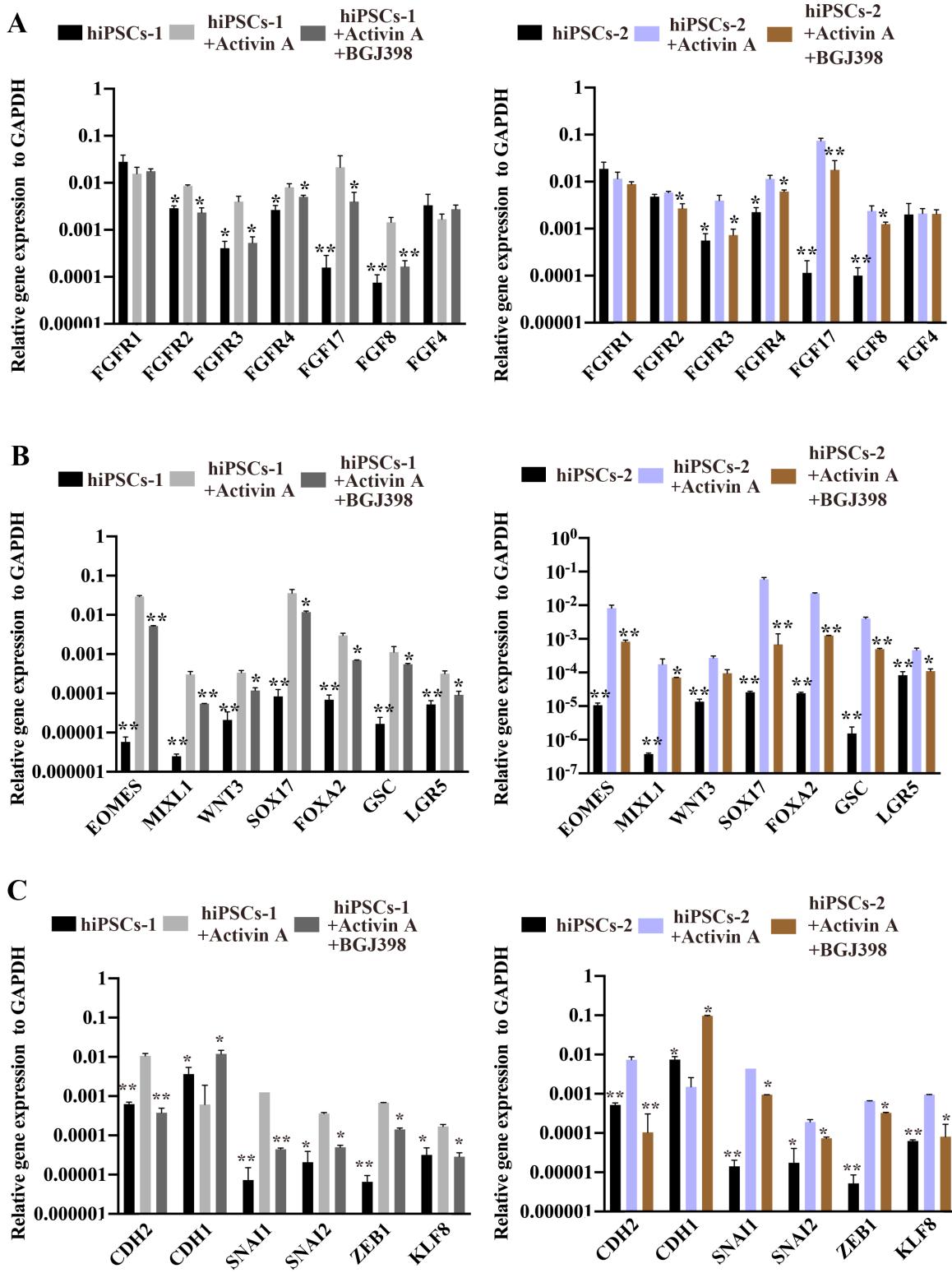


## SUPPLEMENTARY MATERIAL

**corresponding to:**

# **FGF signaling mediates definitive endoderm formation by regulating epithelial-to-mesenchymal transition and cell proliferation**

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**Supplementary Fig. S1.** The mRNA expression of FGF family members, EMT associated genes and DE markers are disturbed by BGJ398 in two hiPSCs models. **(A, B, C)** RT-qPCR analyses for the indicated genes expressed in Activin A-induced cells in the absence and presence of BGJ398 in hiPSC-1 and hiPSC-2 models. Data represent means  $\pm$  s.d. from three biological repeats. \* $p<0.05$ ; \*\* $p<0.01$ , compared to Activin A-induced cells in the absence of BGJ398.

**SUPPLEMENTARY DATA 1: PRIMER SEQUENCES FOR RT-QPCR**

| Human Gene Name | Forward (5' to 3')     | Reverse (5' to 3')      |
|-----------------|------------------------|-------------------------|
| EOMES           | CCGCCACCAAACGTGAGATGA  | ACATTGGCTGCCTGCATGT     |
| MIXL1           | CCGAGTCCAGGATCCAGGTAA  | CTCTGACGCCGAGACTTGG     |
| WNT3            | AGGGCACCTCCACCATTG     | GACACTAACACGCCGAAGTC    |
| SOX17           | CGCACGGAATTGAAACAGTA   | GGATCAGGGACCTGTCACAC    |
| FOXA2           | ACTACCCCCGGTACGGTTC    | AGGCCGTTTGTGCGTGA       |
| GSC             | AACGCGGAGAAGTGGAAACAAG | CTGTCGAGTCCAATCGC       |
| LGR5            | CTCCCAGGTCTGGTGTGTTG   | GAGGTCTAGGTAGGAGGTGAAG  |
| FGFR1           | CCCGTAGCTCCATATTGGACA  | TTGCCATTTCAACCAGCG      |
| FGFR2           | GGTGGCTGAAAAACGGGAAG   | AGATGGGACCAACTTCCATA    |
| FGFR3           | CCCAATGGGAGCTGTCG      | CCCGGTCTTGCAATGCC       |
| FGFR4           | CCATAGGGACCCCTCGAATAG  | CAGCGGAACCTGACGGTGT     |
| FGF17           | CTGCTGATTCTGCTGTCAA    | GTAAGTTGGTACTCGCGGAT    |
| FGF8            | GACCCCTTCGCAAAGCTCAT   | CCGTTGCTTGGCGATCA       |
| FGF4            | CTCGCCCTTCTCACCGATG    | GTAAGACTCGTAGGCCTGTA    |
| CDH2            | TGGGAATCCGACGAATGG     | TGCAGATCGGACCGGATACT    |
| CDH1            | TGCCAGAAAATGAAAAAGG    | GTGTATGGCAATGCGTC       |
| SNAI1           | ACTGCAACAAGGAATACCTCAG | GCACTGGTACTCTTGACATCTG  |
| SNAI2           | CATGCCCTGTCATACCACAAAC | GGTGTAGATGGAGGAGGG      |
| ZEB1            | AGCAGTGAAGAGAAGGGATGC  | GGTCCTCTCAGGTGCCTCAG    |
| KLF8            | CCCAAGTGAACCAGTTGACC   | GACGTGGACACCAAGGG       |
| SOX2            | CCCAGCAGACTTCACATGT    | CCTCCCATTCCCTCGTTT      |
| PAX6            | ATGTGTGAGTAAAATTCTGGCA | GCTTACAACCTCTGGAGTCGCTA |
| NESTIN          | CTGGAGCAGGAGAACAGG     | TGGGAGCAAAGATCCAAGAC    |
| GAPDH           | AGGGCTGCTTTAACCTCGGT   | CCCCACTTGTATTGGAGGGA    |