## **Supplementary Figures**

- *Journal*: Cells
- *Manuscript* #: cells-1018009
- *Title of Paper*: Characterization of early-onset finger osteoarthritis-like condition using patient-derived induced pluripotent stem cells
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**Figure S1.** Gene expression confirmation using three different reference genes; *GAPDH*,  $\beta$ -actin (ACTB), and 18S RNA. (a) Relative expression of *SOX9*. (b) Relative expression of *ACAN*. (c) Relative expression of *COL2A1*. (d) Relative expression of *COL1A1*. (e) Relative expression of *COL10A1*. (f) Relative expression of *RUNX2*. (g) Relative expression of *MMP13*. (h) Relative expression of *VEGFA*. (i) Relative expression of *AQP1*. Significant differences between the healthy control (HC) and early finger osteoarthritis (efOA) on each time point (day 7 and 21; D7 and D21) are each indicated by the asterisk symbol (\* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001).



**Figure S2.** Confirmation of chondrogenic gene expression in human induced pluripotent stem cells (hiPSCs) and derived chondrogenic pellets (CPs). (a) Relative expression of *LIN28*. (b) Relative expression of *SOX9*. (c) Relative expression of *ACAN*. (d) Relative expression of *COL2A1*. (e) Relative expression of *COL1A1*. (f) Relative expression of *COL10A1*. Statistically significant differences between the hiPSC control and CPs on each time point are indicated by the hash symbol (# p < 0.05, ## p < 0.01, ### p < 0.001). The significance between each group analyzed using ANOVA was indicated using the at sign (@ p < 0.05, @@ p < 0.01, @@@ p < 0.001).



**Figure S3.** Further characterization of day 7 (D7) and 21 (D21) chondrogenic pellets (CPs). (**a**) Pellet size measurement of CPs. (**b**) Vacuole count in CPs. Statistical significance between healthy control (HC) and early finger osteoarthritis (efOA) CPs are indicated by the asterisk symbol (\* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001). (**c**) Measurement of vacuole size in CPs. The statistical significance was measured between D7 HC-CPs and other groups. The expression of each gene was normalized to that of GAPDH.



**Figure S4.** Gene expression confirmation using three different reference genes; *GAPDH*,  $\beta$ -*actin* (*ACTB*), and 18S RNA. (a) Relative expression of *IL6*. (b) Relative expression of *MMP1*. (c) Relative expression of *MMP10*. (d) Relative expression of *IL1B*. (e) Relative expression of *TNFA*. Statistical significance between the healthy control (HC) and early finger

osteoarthritis (efOA) on each time point (day 7 and 21; D7 and D21) are each indicated by the asterisk symbol (\* p < 0.05, \*\* p < 0.01, \*\*\* p < 0.001).



**Figure S5.** Confirmation of *RUNX2*, *VEGFA*, *IL1B*, *IL6*, *MMP1*, and *MMP10* expression using three different reference genes; *GAPDH*,  $\beta$ -actin (ACTB), and 18S RNA. (a) Relative expression of *RUNX2*. (b) Relative expression of *VEGFA*. (c) Relative expression of *IL1B*. (d) Relative expression of *IL6*. (e) Relative expression of *MMP1*. (f) Relative expression of *MMP10*. Statistically significant differences between the normal control (NC) and the treatment groups are each indicated by the +, @, and & (One-way ANOVA, Dunnett's test, +,@,& p < 0.05, ++,@@,&& p < 0.01, +++,@@@,&&& p < 0.001).