Potential of human fetal chorionic stem cells for the treatment of osteogenesis imperfecta

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Abstract

Osteogenesis Imperfecta (OI) is a genetic bone pathology with prenatal onset, characterised by brittle bones in response to abnormal collagen composition. There is presently no cure for OI. We previously showed that human first trimester fetal blood mesenchymal stem cells (hfMSC) transplanted into a murine OI model (*oim* mice) improved the phenotype. However, the clinical use of fetal MSC is constrained by their limited number and low availability. In contrast, human fetal chorionic stem cells (e-CSC) can be used without ethical restrictions and isolated in high numbers from the placenta during ongoing pregnancy. Here we show that intra-peritoneal injection of e-CSC in *oim* neonates reduced fractures, increased bone ductility and bone volume, increased the numbers of hypertrophic chondrocytes, and upregulated endogenous genes involved in endochondral and intramembranous ossification. Exogenous cells preferentially homed to long bone epiphyses, expressed osteoblast genes and produced collagen COL1A2. Together, our data suggests that exogenous cells decrease bone brittleness and bone volume by directly differentiating to osteoblasts and indirectly stimulating host chondrogenesis and osteogenesis. In conclusion, the placenta is a practical source of stem cells for the treatment of OI.

Introduction

Osteogenesis Imperfecta (OI), or brittle bone disease, is a debilitating inherited skeletal dysplasia with prenatal onset that affects 1 in 15,000-20,000 births. OI is characterised by short stature, osteopenia and multiple fractures. The severity of the disease ranges across the 11 known types depending on the causative mutation in collagen type I or genes involved in its biosynthesis, with type III being the most severe that survive the neonatal period [1,2,3,4]. Existing treatments largely provide symptomatic relief, but there is currently no cure. The 'gold standard' bisphosphonates temporarily improve bone strength by inhibiting bone resorption, but do not improve growth or bone pain beyond a year [5] and do not reduce fracture incidence long term [6].

Cell therapy in OI aims to prevent morbidity and deformity as well as mortality, by introducing healthy cells, early in development, with the aim that exogenous cells will home to bones and contribute to bone formation to decrease the severity of the disease [7]. Cell therapy for OI holds much promise, with most studies showing beneficial effects. In humans, whole bone marrow and bone marrow mesenchymal stem cells (MSC) have been transplanted in OI children with gains in body length and bone mineralization [8,9], whilst allogeneic fetal liver derived stem cells transplanted *in utero* led to apparent phenotypic improvement in an OI fetus, although confounded by concomitant bisphosphonate use [10]. In rodent OI models, transplantation of whole bone marrow/ bone marrow MSC led to increased collagen content [11], improved bone strength, reduced perinatal lethality [12] and increased osteoblast differentiation [13,14]. Marked therapeutic benefits were shown following transplantation of fetal MSC from human first-trimester blood in a mouse model of human type III OI (*oim)* including improved bone plasticity and a two-third reduction in long bone fractures [15,16].

However, there are a number of hurdles to overcome before fetal stem cell therapy can be translated to the clinic. For example, it is essential to have a source of stem cells that have high therapeutic potential and are easily accessible for clinical use. Extra-embryonic fetal tissues, such as the placenta, are readily available either from termination of pregnancy or surplus tissue at routine prenatal diagnostic procedures [17,18], or at term delivery [19,20,21,22]. Recently we have shown early gestation chorionic stem cells (e-CSC) isolated from human placental tissue accelerated tissue repair in dermal excision skin wounds and improved bone quality and plasticity in *oim* mice. This tissue repair capacity of e-CSC was greater than its late gestation counterparts *in vivo*, as was the osteogenic differentiation and cell expansion potential *in vitro* [23]. This may be due to the more primitive characteristics of e-CSC compared to term isolated CSC, which showed an intermediate phenotype between human embryonic stem cells (hESC) and MSC [23].

We hypothesised that transplantation of stem cells derived from first trimester placenta would have therapeutic benefits in a mouse model of OI. Here, we show that exogenous e-CSC engrafted at sites of bone growth and repair in the *oim* model, differentiated to osteoblasts that produced COL1A2 and mediated changes in endogenous ossification genes, which resulted in reduced fractures and increased bone flexibility.

Materials and Methods

Cells

Collection of human early chorionic stem cells (e-CSC) was as previously described [23] from first trimester chorionic villous tissue sampled during pregnancy termination (9-10 weeks

gestation age) as approved by the Research Ethics Committee of Hammersmith & Queen Charlotte's Hospital. Isolated cells were plastic adherent and cultured in Dulbecco's modified Eagle's medium high glucose (DMEM-LG) (Sigma) supplemented with 10% fetal bovine serum (BioSera), 2mmol/l L-glutamine, 50IU/ml penicillin, 50mg/ml streptomycin (Gibco-BRL) (D10 medium). Cells were expanded at 70-80% confluence on plastic dishes and used at passage 6-8.

The chondrogenic ATDC5 cells (generous gift from G.H.D Bassett and G.R. Williams) were expanded in D10 medium. Differentiation was induced chemically by culturing the cells with 10 ng/ml TGF-beta3, 1X ITS (insulin, transferrin, selenium), 10 nanomolar dexamethasone and 100 micromolar ascorbate-2-phosphate for 7 days.

Fluorescence immunostaining and confocal microscopy

Human e-CSC were fixed in 4% then 8% PFA in 125mM HEPES (pH7.6), then permeabilized in 0.5% Triton X-100 (Sigma), incubated with 20µM glycine (Sigma) and blocked in PBS supplemented with 1% bovine serum albumin (BSA), 0.2% gelatine and 0.1% casein (pH7.6). Cells were stained with primary antibodies (listed in Supplementary Table S1) then secondary antibody; donkey anti-mouse or anti-rabbit IgG (Jackson ImmunoResearch Laboratories), before being mounted in VectaShield labelled with DAPI (Vector Labs) [24]. Fluorescence confocal laser scanning microscopy images were collected on a Leica TCS SP5 (X1000 PL APO oil objective). Positive controls were hESC and negative controls differentiated cells.

Flow cytometry

Cells were detached, blocked with PBS supplemented with 1% BSA (Sigma) and either fixed in 0.01% PFA and permeabilized with 0.5% Triton X-100 for intracellular staining, or immediately stained with primary antibodies for cell surface staining (see Supplementary Table S1). For unconjugated antibodies, cells were subsequently washed with 1% BSA and incubated with

secondary goat anti-murine IgM PE (Santa Cruz) [23]. Otherwise cells were analyzed by FACS calibur flow cytometry (Becton Dickinson) using hESC as positive and antibody-specific isotypes as negative controls.

Cell Differentiation

Cells were differentiated along the osteoblast lineage for 2 weeks in DMEM-LG supplemented with 10 mM β-glycerophosphate, 0.2 mM ascorbic acid and 10^{-8} M dexamethasone, then fixed in 10% formalin and stained with von Kossa (2% silver nitrate) or 2% alizarin red. Cells were differentiated along the adipocyte lineage over 2 weeks in DMEM supplemented with 0.5 mM hydrocortisone, 0.5 mM isobutyl methylxanthine and 60 mM indomethacin, then fixed and stained with oil red O [23]. Cells were differentiated along the chondrocyte lineage over 2 weeks in DMEM-LG supplemented with 0.01µg/ml TGF-β3, 0.1µM dexamethasone, 0.17mM ascorbic acid, 1mM sodium pyruvate, 0.35mM L-proline, 1% ITSS, 50µg/ml Linoleic Acid (reagents from Sigma), then cells were fixed in and stained with alcian blue (2%).

Animals

All experimental protocols complied with Home Office guidelines (PPL 70/6857). Heterozygous male and female (B6C3Fe a/a-Col1a2^{oim}/Col1a2^{oim}) mice (Jackson Laboratory) were housed in individual ventilated cages in 12:12-hour light dark cycle (21^{o}C) with water and chow. Offspring were genotyped by sequencing the *oim* fragment then homozygous and wild type colonies established. Progeny were weaned at 30±1 day and culled at 8 weeks of age. Human e-CSC (10^6 cells resuspended in 20 µl of cold PBS) were injected intra-peritoneally (i.p.) into 3-4 day-old *oim* neonates (n=11 males, n=11 females) and mice were culled for analysis when they were 8 week old. We noted no variability between different isolated placenta specimens in terms of e-CSC

phenotype (data not shown) and donor cells injected in *oim* mice were from a single donor. Controls comprised age-matched non-transplanted *oim* and wild type mice.

Immunohistochemistry

Dissected tibias were decalcified in 10% EDTA pH7.4 and subsequently embedded in paraffin. Four micron sections were cut, deparaffinised in xylene and rehydrated. Heat-induced epitope retrieval was performed in a steamer (Dako), followed by incubation with peroxidise block (Dako). The presence of donor cells in transplanted 8 week old *oim* mice was determined in 3 different regions of the non-fractured tibia (epiphysis, diaphysis and bone marrow) as well as in fracture callus. Donor cells were visualised using human-specific mouse monoclonal vimentin (Dako) primary antibody (Supplementary Table S1) and incubation with HRP-labelled anti-mouse polymer followed by DAB+ substrate-chromogen staining. Positive cells were counted in bone marrow (n=4 samples and n=4 sections for each). Staining specificity was verified using non-transplanted negative controls.

Detection of Collagen type X and Osteopontin (Supplementary Table S1) was performed on four micron sagittal sections of tibia from 8 week old mice, using HRP-labelled polymer followed by DAB+ substrate-chromogen staining.

Engraftment measured by quantitative real time PCR

Femurs of the same mice were dissected and separated into callus if present (n=8), epiphysis (n=6) and diaphysis (n=6). Liver (n=6) was also used. RNA was then extracted using TRIzol (Invitrogen) followed by cDNA synthesis with M-MLV reverse transcriptase (Promega). To calculate donor cell engraftment quantitative real time PCR (qRT-PCR) was performed using SYBR green dye (Applied Biosystem) and the ABI Prism 7700 Sequence Detection System with human specific and human-mouse non-specific *β-actin* primers (Supplementary Table S2).

Human:mouse chimerism was estimated as the ratio of human *β-actin* to total human and mouse *β-actin* in the total cDNA sample to give the $2^{-\Delta Ct}$ value. Samples were considered positive with a human specific *β-actin* Ct above 36 at a threshold of 0.13ΔRn. Negative controls were non-transplanted *oim* [15].

Quantitative real time RT-PCT

Osteoblast gene expression was performed by quantitative real time RT-PCR (QRT-PCR) using SYBR green dye (Qiagen) and the MJ-Opticon with human specific *Osteopontin* and *Osteocalcin* primers (Supplementary Table S2). Results with a Ct below 36 were normalised to human *β-actin* to give the $2^{-\Delta Ct}$ value. Expression in transplanted *oim* femurs (n=6) was compared to e-CSC (n=3) undifferentiated and grown in osteogenic permissive media for 2 weeks. Negative controls were non-transplanted *oim* [15]. Sox9 expression in ATDC5 cells was measured by QRT-PCR using the $2^{-\Delta Ct}$ method. Manufactured mouse-specific primers were from SABiosciences (Qiagen).

Western blot

Collagen was extracted from ground bone over 72hrs at 4ºC in a lysis buffer of 6M guanidine HCl and 100mM Tris pH7.4 containing protease inhibitor cocktail. Proteins were precipitated with 10% TCA, re-suspended in RIPAE buffer (1xTBS, 1% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% SDS, 0.004% sodium azide; Sigma) containing PMSF (Sigma) and protease inhibitor cocktail, run on an 8% SDS-PAGE, transferred to nitrocellulose, blocked with milk and stained with a COL1A2 (129kDa) primary antibody (Abcam), then with an HRP-linked anti-rabbit IgG secondary antibody (GE healthcare), followed by enhanced chemiluminescence detection (Thermo scientific). The loading control used was β-ACTIN (43kDa) (Santa Cruz) [23]. Detection of COL1A2 in transplanted *oim* bones was confirmed using wild type positive controls and specificity confirmed using non-transplanted *oim* negative controls (n=4 per group).

Mechanical testing

Three-point bending tests were performed as described [23] using a materials testing machine (5866 Instron) on 8 week old unfractured frozen and thawed femurs (n=22 transplanted *oim*, n=34 *oim* controls, n=14 wild type). Bones were bent mid-diaphysis to fracture on two supports 9 mm apart at a loading rate of 50 µm/s. Force deflection curves were analyzed (Matlab, MathWorks) to measure bending stiffness (slope of the linear elastic deformation; N/mm), load to fracture (maximum force sustained prior to fracture; N), maximum deflection (deflection at fracture in mm).

X-ray microradiography

Tibias from 8 week-old mice (n=19 transplanted, n=20 *oim*, n=11 wild type) were fixed in formalin for 24 hours and stored in 70% ethanol prior to removal of soft tissues. Digital X-ray images were obtained at a 10µm pixel resolution using a Faxitron MX20 variable kV point projection X-ray source and digital image system (Qados, Cross Technologies plc, Sandhurst, Berkshire, UK). An X-ray image of a digital micrometer was used to calibrate ImageJ 1.41 software (http://rsb.info.nih.gov/ij/) prior to determination of cortical bone thickness and diameter at 5 locations along the mid shaft, and bone length. Relative bone mineral content (BMC) was determined by comparison with 1 mm diameter steel, aluminium and polyester standards included in each frame. 16 bit DICOM images were converted to 8 bit Tiff images using ImageJ and the image histogram stretched between the polyester (grey level 0) to steel (grey level 255) standards. Bone mineralization densities were represented by a pseudocolour scheme representing 16 equal intervals [25].

Counting of fractures

Fractures in both femurs, tibias and humeri were assessed at 8 weeks of age by determination of callus formation (n=120 transplanted, n=78 *oim* control). The number of mice with at least one long bone fracture and the fracture incidence (number of fractured bones/total bones assessed) were calculated by two independent observers blinded to transplantation status. Deformities and callus formation in the caudal vertebrae (n=16 transplanted, n=10 *oim* control) were counted on digital X-ray images and the fracture rate was calculated as above. The presence of vertebral deformity and callus formation detected by X-ray microradiography was verified by micro computerised tomography (μCT40 Scanco Medical) at 10μm voxel resolution (45kV, 177μA, 200ms integration time). Unfractured *oim* vertebrae did not differ in shape from wild type vertebrae (**Supplementary Fig.S1A**) and had normal morphology (**Supplementary Fig.S1B**), while deformed vertebra had evidence of callus formation (**Supplementary Fig.S1C and Fig.S1D**) [16].

Dynamic histomorphometry

Animals (n=7 transplanted, n=6 wild type, n=5 *oim*) were injected 10 days and 3 days before sacrifice with 20mg/kg of calcein (Sigma). Tibias were then fixed in formalin for 24hrs and transferred to 70% ethanol, before being dehydrated in acetone for 48hrs, infiltrated over 6-9 days at -20^{o}C and embedded in methylmethacrylate (MMA) [26]. Embedded samples were imaged on a Leica TCS SP5 confocal laser scanning microscope and analysed using ImageJ. Fluorescent images of calcein labels were taken 500μm and 1000μm below the proximal growth plate of the trabecular and endo-cortical regions respectively The amount of mineralizing surface per total bone surface (MS/BS;%), the daily mineral apposition rate (MAR;μm/day) and the bone formation rate (BFR;$\mu m^3/\mu m^2$/day) were calculated.

Static histomorphometry

MMA embedded samples were cut into 8µm sections and stained using the Leucognost AP kit (Merck), according to the manufacturer's instructions (n=6 transplanted, n=6 *oim*, n=5 wild type). Sections were analysed on a light microscope using the Osteomeasure system (OsteoMetrics Inc). Histomorphometric measurements of the secondary spongiosa were performed on stained sections 500µm from the end of the hypertrophic zone of the growth plate; % trabecular bone volume per total tissue volume (BV/TV) was quantified. For growth plate analysis dissected tibia were decalcified, paraffin embedded and 5µm sections were cut and stained with alcian blue 8GX (2%), Weigert's haematoxylin and van Gieson, mounted and growth plate morphology analysed using ImageJ (n=14 transplanted *oim*, n=5 *oim*, n=5 wild type).

Osteogenesis PCR array

Total RNA was extracted from femoral epiphysis of 8 week-old mice using TRIzol (Invitrogen), followed by RNA clean up (RNeasy Qiagen) and cDNA synthesis using an RT^2 First Strand Kit (Qiagen). Gene expression was investigated using an RT^2 Profiler mouse osteogenesis PCR array (Qiagen) and analysed according to the manufacturer's instructions (n=3 mice per group). To verify results, quantitative real time PCR was performed using RT^2 qPCR Master Mix and primers (Supplementary Table S2) and analysed with MJ-opticon (Biorad). Data were normalised to 2 housekeeping genes (β-Actin and Hsp90ab1) and the $2^{-\Delta Ct}$ of each sample calculated (n=8 transplanted *oim*, n=5 *oim* controls).

Protein measurement

Cells were cultured either in D10 medium (non-primed) or in co-culture without cell contact with ATDC5 cells (primed with ATDC5) or in the presence of blood serum from *oim* or wild type mice (primed with *oim* or WT serum) for 7 days. The mouse cell line ATCD5 is chondrogenic and goes through a sequential process analogy to chondrocyte differentiation, constituting an

excellent *in vitro* model cell line for analysing skeletal development and studying the factors involved in chondrogenesis [27]. The presence of protein was measured in the medium using Mini ELISA development kits (Peprotech, London, UK) for the detection of human-specific basic fibroblast growth factor (bFGF), platelet-derived growth factor-BB (PDGF-BB) and connective tissue growth factor (CTGF), following the manufacturer's protocol. Briefly, ELISA microplates (Corning) were incubated overnight with capture antibody, washed with 0.05% Tween-20 in PBS (Sigma), incubated with standards or samples for 2 hours, washed and incubated with detection antibody for 2 hours, followed by washing and incubation with Avidin-HRP conjugate for 30 minutes. Finally, substrate was added to the wells and colour development was monitored at 405nm with wavelength correction set at 650nm. For the detection of human factor IX in blood serum of transplanted *oim* mice, a kit containing microplates pre-coated with antibody was used (Abcam), using the protocol recommended by the manufacturer. Detection was carried out at 450nm.

Statistical analysis

Data were expressed as mean ± s.e.m (standard error). Normally distributed data were analyzed by unpaired two-tailed Student's *t*-test or one-way ANOVA followed by a Tukey's multiple comparison post-hoc test. $P < 0.05$ was considered significant. Two-tailed 2x2 Fisher exact was used for categorical comparisons. Cumulative frequency distributions of bone mineral densities were compared using the Kolomogorov-Smirnov test. Chi-squared with Yates correction and to one degree of freedom was used to compare fracture incidence.

Results

Characterisation of e-CSC

The e-CSC transplanted in neonatal *oim* have a pre-pluripotent phenotype as previously described [23], showing some characteristics of both MSC [28] and human embryonic stem cells (hESC) [29,30]. MSC traits were demonstrated by positive expression of adhesion molecules CD29 and CD44, and the MSC-associated markers CD73, CD90 and CD105, and absent expression of the endothelial or hematopoietic markers CD14, CD34 and CD45, while presenting low levels of intracellular HLA I and no expression of HLA II, similar to fetal liver MSC [31] (**Fig.1A**). A sub fraction of cells expressed key hESC markers required for the maintenance of pluripotency; OCT4A, SOX2, TRA-1-60 and SSEA4 (**Fig.1B** and **Fig.1C**) As expected, e-CSC showed tri-lineage differentiation capability; osteogenic differentiation by alizarin red staining of calcium deposits and von kossa staining of mineralisation, chondrogenic differentiation by Safranin O staining of cartilage matrix, and adipogenic differentiation by oil red O staining of lipid droplets (**Fig.1D**).

Bones of transplanted mice are less liable to fracture

Eight weeks after e-CSC were transplanted, 11/20 *oim* mice (55%) had no long bone (femur, tibia and humerus) fractures, whereas all non-transplanted *oim* controls (100%; n = 13) had at least one or more long bone fracture (**Fig.2A**). The fracture incidence in long bones, calculated as the number of fractured tibia, femur and humeri over the total number of these bones, was reduced from 29.5% (23/78 total bones) in non-transplanted *oim* to 10.0% (12/120 total bones) in e-CSC transplanted *oim*. This corresponds to a 66% decrease in fracture rate (X^2=21, P<0.001) (**Fig.2B**).

We next counted the number of caudal vertebra fractures on digital X-Ray, with a fracture classified as any vertebra having a callus or evidence of bone remodelling (see **Supplementary Fig.1** and Methods section for classification of normal and fractured vertebra by microCT). Compared to non-transplanted *oim* (n=10), which showed an average 46.6% ± 4.2 incidence of fractured vertebra per mouse, transplanted mice (n=16) had 29.2% fewer vertebral fractures at an average incidence of 33.0% ± 2.2 per mouse (X^2=33.8, P<0.001) (**Fig.2C**). The reduction of vertebral fractures from the non-transplanted control group was widespread, with overall numbers of vertebrae with callus reduced across the majority of vertebral positions (**Fig.2D**). Most fractures were found in proximal caudal vertebrae, where force is exerted when rearing up to feed.

Transplanted mice have bones with reduced stiffness and increased ductility

We previously reported 3-point bending data from femurs of *oim* mice transplanted with either early (e-CSC) or late (l-CSC) gestation CSC [23]. Mice transplanted with e-CSC had greater plasticity and overall bone quality than l-CSC transplanted or control *oim* due to an increased post-yield strain. Here we further analysed the 3-point bending load-displacement curves to show that e-CSC transplanted *oim* were also more ductile due to a reduction in stiffness in the pre-yield region and an increase in maximum deflection before fracture (**Fig.2E**). Bone stiffness of e-CSC transplanted *oim* was reduced by an average of 16% compared to non-transplanted *oim* femurs (29.7 N/mm ± 2.1 s.e.m vs. 35.3 N/mm ± 1.7 s.e.m respectively, P<0.05) (**Fig.2F**), while the maximum deflection was increased in transplanted compared to control *oim* by an average of 24% (0.26 mm ± 0.02 s.e.m vs. 0.21 mm ± 0.01, P<0.05 respectively) (**Fig.2G**). However, the maximal load sustained by transplanted and non-transplanted *oim* femurs prior to fracture was similar (4.5 N ± 0.3 vs. 4.8 N ± 0.3) (**Fig.2H**). In contrast, wild type compared to *oim* bones were stiffer (73.6 N/mm ± 2.0, P<0.001) (**Fig.2F**), with greater maximum deflection (0.88 mm ± 0.07,

P<0.001) (**Fig.2G**) and sustained higher loads before fracture (14.0 N ± 0.5, P<0.001) (**Fig.2H**). Thus the material properties of bones from transplanted *oim* were not intermediate between the properties of wild type and *oim* bone. Instead bones from transplanted *oim* were of similar strength to non-transplanted *oim,* but displayed greater plasticity and ductility which may explain their reduced fracture susceptibility.

Transplanted e-CSC preferentially home to oim epiphysis

We first performed an ELISA for human factor IX in blood serum of transplanted mice. Results showed absence of mouse anti-human antibodies, indicating an absence of immune reaction of the neonatal murine immune system (data not shown).

Donor cells were visualised by immunohistochemistry in 8 week-old e-CSC transplanted *oim* using a rabbit monoclonal to human vimentin. Staining was localized at the epiphysis, diaphysis, and sites of fracture callus, with some cells present in the primary spongiosa below the growth plate (**Fig.3A**). Quantitative real time PCR (qRT-PCR) showed that donor cell engraftment in transplanted *oim* was highest in the epiphysis, the site of active bone formation. This was 7.1 fold (P<0.001) higher than in the non-fractured diaphysis where bone formation is less active, and 11.7 fold (P<0.01) higher than engraftment in the liver (**Fig.3B**). Donor cells also preferentially homed to sites of bone repair, where engraftment was 4 fold (P<0.01) higher than engraftment in the liver. Engraftment in fractured and non-fractured diaphysis were not significantly different, but interestingly more mice were positive for human cDNA in the diaphysis if a fracture callus was present; 90% compared to 60%. Engraftment within the femoral epiphysis was inversely correlated (R^2=0.64, y= -0.96x + 41.43, P<0.01 deviation from zero) with bone stiffness, indicating that bone flexibility increases with increasing numbers of donor cells (**Fig.3C**).

Exogenous cells undergo osteogenic differentiation in vivo

To determine if transplanted cells underwent osteogenic differentiation *in vivo* expression using qRT-PCR was determined for human specific *Osteopontin (OP)*; a major interfacial non-collagenous extracellular matrix proteins found in bone and secreted by osteoblasts [32] and also for *Osteocalcin (OC)*; an osteoblast specific gene [33] with an important role in osteoblast differentiation [34]. Results showed expression of human *OP* and *OC* in the transplanted mouse bones (0.38 $2^{-\Delta Ct}$ ± 0.08 s.e.m and 0.85 $2^{-\Delta Ct}$ ± 0.18, respectively), which was greater than expression in e-CSC after growth in osteogenic permissive media for 2 weeks (0.05 $2^{-\Delta Ct}$ ± 0.03, $P<0.05$ and 0.06 $2^{-\Delta Ct}$ ± 0.04, $P<0.05$, respectively) and greater than the low/null basal expression level of the undifferentiated cells (**Fig.3D**). Western blot analysis showed the COL1A2 protein, missing in non-transplanted *oim* [35], was present in the femoral bones of *oim* transplanted with e-CSC (**Fig.3E**), which demonstrates osteogenic differentiation of donor cells to functional osteoblasts.

Transplantation of e-CSC did not affect bone length or cortical bone formation

Tibial length was unaffected by transplantation being similar in transplanted *oim* compared to control *oim* (15.1mm ± 0.1 and 15.2mm ± 0.1 respectively), with both being shorter than wild type tibia (17.0mm ± 0.1, $P<0.001$) (**Fig.4A**). The diameter of the tibia at the mid-diaphysis was also similar in transplanted and control *oim* (1.01 mm ± 0.01 and 1.05mm respectively) but less than in wild types (1.28 mm ± 0.02, $P<0.001$) (**Fig.4B**). The cortical bone thickness was decreased in *oim* compared to wild type (17.9% ± 0.5 and 20.4% ± 0.3, $P<0.001$ respectively), and was similar in untransplanted and transplanted *oim* mice (17.9% ± 0.5 vs 17.2% ± 0.5) (**Fig.4C**).

Cortico-endosteal bone formation rate (BFR) was greater in wild type mice compared to *oim* (2.4 $\mu m^3/\mu m^2$/day ± 0.21 s.e.m vs. 1.54 $\mu m^3/\mu m^2$/day ± 0.23, $P<0.05$ respectively) (**Fig.4D**). This difference resulted from an increased mineral apposition rate (MAR) at the cortico-endosteal interface (2.6 µm/day ± 0.3 vs. 1.7 µm/day ± 0.3, P<0.05 respectively) (**Supplementary Fig.S2A**), because there was no difference in mineralizing surface (92% ± 2 vs. 90% ± 2, respectively) (**Supplementary Fig.S2B**). The MAR and mineralizing surface, however, did not differ between transplanted and non-transplanted *oim* mice (Fig.4D and **Supplementary Fig.S2A and Fig.S2B**).

Transplantation increases trabecular bone volume, but not BMC

The total bone mineral content (BMC) of combined trabecular and cortical bone compartments did not differ between transplanted and non-transplanted *oim* mice, both of which had markedly reduced BMC compared to wild-type ($P<0.001$) (**Fig.4E and Supplementary Fig.S3**). Trabecular bone volume per total tissue volume (BV/TV), however, was increased in transplanted compared to non-transplanted *oim* mice (4.1% BV/TV ± 0.6 vs. 2.0% ± 0.4 respectively, $P<0.05$), but remained lower than in wild type mice (11.9% ± 0.9) (**Fig.4F**). Nevertheless, trabecular BFR did not differ between transplanted and non-transplanted *oim* mice (0.58 $\mu m^3/\mu m^2$/day ± 0.03 vs. 0.52 $\mu m^3/\mu m^2$/day ± 0.03 respectively) and was reduced compared to wild-type (1.10 $\mu m^3/\mu m^2$/day ± 0.1, $P<0.001$) (**Fig.4G**). Furthermore, transplanted and non-transplanted *oim* had similar trabecular bone MAR (0.86 µm/day ± 0.04 vs. 0.89 µm/day ± 0.04, $P<0.05$ respectively) that was reduced compared to wild-type (1.62 µm/day ± 0.09, $P<0.001$) (**Supplementary Fig.S4A**). MS/BS did not differ between transplanted *oim*, non-transplanted *oim* and wild type mice (**Supplementary Fig.S4B**).

Transplantation reduces endogenous Smad3 expression and increases expression of genes activated during endochondral ossification.

The mouse osteogenesis array (SABiosciences) was used to analyse changes in endogenous gene expression within the femoral epiphysis, and showed a global increase in expression of cartilage gene groups in e-CSC transplanted mice compared to non-transplanted *oim* controls (**Fig.5A**). This included up regulation of genes involved in the early stages of endochondral ossification: 2.7 fold for *Sox9* ($P<0.01$), 1.6 fold for *Twist1* ($P<0.05$), 6.8 fold for *Col2α1* ($P<0.01$) and 3.3 fold for *Col11α1* ($P<0.05$). In addition, late hypertrophic chondrocyte differentiation genes were up-regulated 4.4 fold for *Col10α1* ($P<0.05$) and 2.6 fold for alkaline phosphatase ($P<0.05$) [36], while chondrocyte assembly gene *Comp* was also up-regulated 1.7 fold ($P<0.05$) [37].

The array results were confirmed (**Fig.5B**) by qRT-PCR for the key chondrogenesis transcription factor *Sox9* ($5.6 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.1 \times 10^{-2}$ in transplanted *oim* vs. $2.1 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.01 \times 10^{-2}$ in *oim* controls, $P<0.05$) [38]. Downstream up-regulation of the *Sox9* transactivation target *Col2α1* [39,40] was also confirmed in transplanted *oim* compared to *oim* controls ($0.11\ 2^{-\Delta Ct} \pm 0.01$ vs. $0.04\ 2^{-\Delta Ct} \pm 0.01$ respectively, $P<0.05$) and of the key cartilage matrix component aggrecan [41,42] ($4.8\ 2^{-\Delta Ct} \pm 0.8$ vs. $1.1\ 2^{-\Delta Ct} \pm 0.4$ respectively, $P<0.01$), found in proliferating chondrocytes. However, expression of the *Sox9* target gene *Pthrp*, which inhibits chondrocyte maturation [43,44,45] was similar in transplanted and non-transplanted *oim* ($5.2 \times 10^{-4}\ 2^{-\Delta Ct} \pm 0.7 \times 10^{-4}$ vs. $4.8 \times 10^{-4}\ 2^{-\Delta Ct} \pm 0.5 \times 10^{-4}$). Importantly, *Smad3*, which inhibits maturation of chondrocytes by mediating TGF-β signalling [46] was down-regulated in e-CSC transplanted mice compared to *oim* controls ($1.8 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.1 \times 10^{-2}$ vs. $4.4 \times 10^{-2}\ 2^{-\Delta Ct} \pm 1.1 \times 10^{-2}$ respectively, $P<0.05$). This correlated with increased expression of *Col10α1*, a marker of chondrocyte maturation [47], in transplanted mice compared to non-transplanted *oim*, ($1.0\ 2^{-\Delta Ct} \pm 0.2$ vs. $0.2\ 2^{-\Delta Ct} \pm 0.1$ respectively, $P<0.01$).

Up-regulation of genes activated during intramembraneous ossification in transplanted mice is associated with increased endogenous expression of Runx2

The PCR array also identified genes involved in intramembraneous ossification that were up-regulated in transplanted mice compared to *oim* controls. There was a 2.8 fold ($P<0.05$) increase in *Phex* and a 2.9 fold ($P<0.05$) increase in *Dmp1*, genes that are co-expressed by osteoblasts and osteocytes and which regulate osteoblast maturation as well as bone mineralization via FGFR signalling pathways [48,49]. Also up-regulated in transplanted *oim* was *Bgn, which* has a role in osteoblast differentiation and matrix mineralisation [50], and *Serpinh1*, which acts as a molecular chaperone in collagen biosynthesis [51] (2.6 fold, $P<0.05$, and 3.5 fold, $P<0.05$ respectively). These findings correlated with higher expression of extracellular matrix proteins in transplanted mice, including a 3.2 fold increase in *Col1α1* ($P<0.01$), involved in fibril formation of the abundant collagen type I [1,52] (**Fig.5A**).

Array results were confirmed by qRT-PCR and showed *Runx2* expression, essential for osteoblast differentiation [53,54,55], was also increased in transplanted mice compared to non-transplanted *oim* ($2.1 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.2 \times 10^{-2}$ vs. $1.3 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.3 \times 10^{-2}$ respectively, $P<0.05$) (**Fig.5C**). However, expression of the downstream transcription factor osterix, also required for osteoblast differentiation [56,57] was similar in both e-CSC transplanted *oim* and non-transplanted controls ($3.7 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.5 \times 10^{-2}$ vs. $3.3 \times 10^{-2}\ 2^{-\Delta Ct} \pm 0.7 \times 10^{-2}$ respectively). In contrast *Igf1*, which regulates both osteoblasts [58] and osteoclastogenesis via induction of RANK-L synthesis [59,60] and stimulates linear growth [61] was up-regulated in transplanted mice ($0.43\ 2^{-\Delta Ct} \pm 0.05$ vs. $0.21\ 2^{-\Delta Ct} \pm 0.02$ for *oim* controls, $P<0.05$).

Expression of endogenous ossification genes correlated linearly with their co-activators and trans-activation targets as expected when mice were analysed on an individual basis. For example, *Sox9* expression was positively correlated with expression of its transactivation target *Col2α1* (R^2=0.84) (*P*<0.001) (**Supplementary Fig.S5A**). Likewise expression of the co-activators *Dmp1* and *Phex* were strongly correlated (R^2=0.91, *P*<0.001) (**Supplementary Fig.S5B**), as was expression of the ECM genes *Col1α1* and *Bgn* (R^2=0.93, *P*<0.001) (**Supplementary Fig.S5C**). Protein evaluation was performed *in situ* by immunohistochemistry, confirming increased expression of cartilage hypertrophic marker Collagen Type X and increased expression of the osteoblastic marker osteopontin in mice treated with e-CSC compared to non-treated mice (**Fig.5D and Fig.5E**)

Growth plate height is increased in e-CSC transplanted oim

Analysis of growth plate height confirmed previous findings [15] that *oim* have larger growth plates than wild type mice (153μm ± 7 vs. 137μm ± 9 respectively, *P*<0.05). Interestingly e-CSC transplanted *oim* had a substantially wider growth plate (184μm ± 6) than *oim* controls (*P*<0.05), primarily the result of a larger hypertrophic zone (84μm ± 4 for e-CSC transplanted *oim* vs. 60μm ± 7 for *oim* controls, *P*<0.01) (**Fig.6A and Fig.6B**). When the relative proportions of the growth plate zones were calculated, the hypertrophic zone formed a larger proportion of the total growth plate in the transplanted mice than in non-transplanted *oim* mice (46.4% ± 1.1 vs. 39.5% ± 3.2 respectively, *P*<0.05), in which the hypertrophic zone was similar to wild type (41.7% ±1.2) (**Fig.6C**). This finding was consistent with gene expression studies demonstrating that transplanted *oim* had a 10:1 ratio of expression of the late hypertrophic chondrocyte marker *Col10α1* in the epiphysis compared to expression of the proliferating chondrocyte marker *Col2α1*, whereas in non-transplanted *oim* the ratio was 4:1 (*P*<0.05).

There was a positive correlation (R^2=0.75, y=20.13x – 0.84, P<0.01) between endogenous *Col10α1* expression and the size of the hypertrophic zone of chondrocytes in the growth plate (**Fig.6D**). Whilst *Runx2* expression was also correlated with the expression of *Col10α1* (R^2=0.76, y=65.56x – 0.41, P<0.001) (**Fig.6E**) and may therefore be involved in mediating the larger hypertrophic zone in the growth plate. We also show a strong correlation between *Igf1* and *Bgn* expression (R^2=0.89, y=8.64x – 0.15, P<0.001), highlighting the importance of *Igf1* in regulating genes activated during intramembraneous ossification (**Fig.6F**).

We next wanted to provide mechanistic clues as to how e-CSC transplantation induces up-regulation of endogenous genes involved in skeletogenesis. We hypothesized that donor cells produce growth factors that stimulate maturation of endogenous chondrocyte progenitors. To test this hypothesis, we cultured e-CSC with the chondrogenic cell line ATDC5 to investigate whether e-CSC would produce factors that stimulate chondrogenic differentiation and maturation of ATDC5 cells. Although expression of the chondrogenic marker sox9 was higher in ATDC5 cells cultured in chondrogenic differentiated medium compared to levels found in non-induced cells, sox9 levels were not up-regulated when ATDC5 cells were co-cultured without cell contact with e-CSC, indicating that e-CSC do not produce soluble factors that induce chondrocyte maturation *in vitro* (**Fig.6G**). Interestingly, although ELISA analysis showed e-CSC did not produce bFGF, CTGF and PDGF-BB when cultured in D10 medium, they produced CTGF, but not bFGF or PDGF-BB, when co-cultured with ATDC5 (**Fig.6H**). When primed with *oim* or wild type seri, e-CSC produced both bFGF and CTGF, but not PDGF-BB, indicating the cells might respond to *in vivo* signals present in blood serum (**Fig.6H**).

Discussion

This study demonstrates that fetal stem cells derived from human first trimester chorionic placental tissue (e-CSC), have therapeutic benefits in the OI mouse model (*oim*) as evidenced by a two-third decrease in long bone fracture incidence and decreased bone brittleness compared to non-transplanted controls. These results are in line with our previous studies [15,16,62]. Fracture reduction in e-CSC transplanted mice was attributed to an increase in bone plasticity, as previously demonstrated [23], as well as greater bone ductility. Changes to the bone mechanical properties of transplanted *oim* were most likely mediated by the exogenous cells since higher engraftment levels in bones correlated with decreased bone stiffness. This is in agreement with recent work from our group showing that up regulation of *CXCR4* in transplanted fetal blood MSC increased cell homing to sites of injury via the CXCR4-SDF1 pathway [62,63], which subsequently increased donor cell engraftment as well as bone plasticity and bone quality [62]. Transplanted e-CSC homed to areas of bone growth and fracture repair and expressed osteoblast differentiation genes *Osteopontin* and *Osteocalcin* as well as the COL1A2 protein, indicating their differentiation to functional osteoblasts. These findings are in agreement with previous studies in the *oim* model that demonstrated the direct differentiation of transplanted cells to osteoblasts [13,14,15,16] and subsequent improvements in disease pathology. In addition, we used the detection of human factor IX as an immunoassay to detect the presence of mouse anti-human antibodies in the serum of mice transplanted with human cells and we were able to show the absence of immune reaction against allogeneic cells.

The trabecular bone volume (BV/TV) of *oim* is lower than wild type mice due to the impaired osteoblast differentiation of *oim* [64,65], which results in a high numbers of preosteoblasts that

support greater osteoclast bone resorption [64]. *Oim* mice transplanted with e-CSC had a higher BV/TV than non-transplanted *oim*, despite bone formation rate remaining the same, which could be due to differentiation of exogenous cells to normal osteoblasts that better regulate bone remodelling. Increased BV/TV may also result from an indirect effect of the transplanted cells on osteoblast differentiation as demonstrated by the upregulated expression of endogenous genes in transplanted *oim* that were associated with osteoblast differentiation, including *Dmp1*, *Phex* and *Bgn* [48,49,50]. Others have also shown an effect of transplantation on endogenous osteoblast activity, for example transplantation of osteogenic differentiated MSC in SCID mice resulted in increased bone being produced by host cells [66], and endogenous osteoblast numbers were increased after transplantation of term placental stem cells in a SCID-rab mouse model of medullary myeloma-associated bone loss [67]. We also showed upregulation in transplanted *oim* of endogenous chondrogenesis genes including chondrogenesis regulator *Sox9* [38] and *Runx2*, implicated in chondrocyte maturation through *Col10α1* transactivation [68]. Whilst expression of chondrocyte maturation inhibitor *Smad3* [46] was downregulated in transplanted mice. These changes were associated with a larger zone of hypertrophic chondrocytes within the growth plate, and indicate transplantation may have increased endogenous endochondral ossification.

The larger growth plate of e-CSC transplanted *oim* compared to *oim* controls is in contrast to previous data with hfMSC that instead showed normalisation of growth plate height in prenatally-transplanted *oim* compared to controls [15]. This may suggest different mechanisms of action between different transplanted cell sources. For example, recent work by Horwitz *et al*. in *oim* mice suggested different sources of cells contributed through different mechanisms when used in cell therapy, with non-adherent bone marrow cells differentiating to osteoblasts that produced normal collagen, whilst bone marrow MSC increased lumbar vertebrae length via paracrine mechanisms on chondrocyte proliferation at the growth plate, possibly through release of soluble growth factors [69].

BMC measured at the whole bone scale did not increase in *oim* after transplantation of e-CSC, despite changes in bone mechanical properties. Others have shown the importance of the collagen matrix organization on bone mechanical properties [70,71] and recently we have shown using nanoindentation that compared to wild type mice, *oim* have greater mineralization of a poorly organized matrix [72]. Therefore exogenous cells may have affected the bone mineralization or improved organization of bone matrix collagen fibres in the *oim* bones at the microscopic matrix scale, potentially in response to production of normal COL1A2.

To test the hypothesis that e-CSC produce growth factors which promote endogenous chondrocyte progenitor maturation and differentiation, we co-cultured e-CSC with ATDC5 in vitro. Interestingly, co-culture with ATDC5 cells induce e-CSC to produce connective tissue growth factor CTGF, which is known to induce chondrocytic proliferation, maturation and hypertrophy *in vitro* [73]. Interestingly, when primed with blood serum, e-CSC produced both CTGF and bFGF. CTGF is also known for stimulating proliferation and differentiation of cultured osteoblastic cells, and bFGF, which stimulates proliferation in the perichondrium [72]. Together, these results suggest e-CSC respond to *in vivo* signals to produce CTGF and bFGF, which may stimulate endogenous osteogenesis and chondrogenesis.

In summary, our study demonstrates that fetal stem cells derived from first trimester chorionic tissue have the potential to treat osteogenesis imperfecta.

List of Abbreviations

e-CSC: early fetal placental chorionic stem cells, hESC: human embryonic stem cells, HZ: hypertrophic zone, i.p.: intraperitoneal, MSC: mesenchymal stem/stromal cells, OI: osteogenesis imperfecta, *oim*: osteogenesis imperfecta murine, PZ: proliferative zone, RZ: reserve zone

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Figure Legends

Figure 1

e-CSC express MSC and hESC markers and differentiate down mesenchymal lineages (**A**) Representative confocal immunofluorescence images for expression (green) of adhesion molecules (CD29 and CD44), MSC-associated markers (CD73, CD90 and CD105), endothelial marker (CD14), hematopoietic markers (CD34 and CD45) and MHC antigens (HLA I and II). Nuclei stained with DAPI (blue). (**B**) Flow cytometry for percent of e-CSC population positive for OCT4A, SOX2, TRA-1-60 and SSEA4 (isotype control in black). (**C**) Confocal images for expression of OCT4A, SOX2, TRA-1-60 and SSEA4 in the e-CSC whole population. (**D**) Von Kossa staining of calcium mineralisation and alizarin red staining of mineralising nodules following osteogenic differentiation of e-CSC. Safranin O staining of cartilage matrix following chondrogenic differentiation. Oil red O staining of lipid droplets following adipogenic differentiation. Samples were either cultured in the presence of differentiation medium (induced) or in growth medium (un-induced, negative controls). All scale bars 100µm.

Figure 2

Transplanted *oim* show improvement in disease pathology (**A**) Percentage of mice with any long bone fracture. (**B**) Fracture rate; total proportion of fractured femurs, tibias and humeri over total number of these bones per mouse. (**C**) Percentage caudal vertebral fractures calculated over total number of vertebrae per mouse. (**D**) Percentage of mice with vertebral fractures shown per caudal vertebra from the base of the tail (vertebra number 1) to the tip of the tail (vertebra number 30). (**E**) Three-point bending load (N) displacement (mm) curves shown up to the critical fracture point (**F**) Bending stiffness of femurs (slope of the linear elastic deformation; N/mm). (**G**) Maximum deflection at fracture (displacement extension to the point of fracture; mm). (**H**) Load to fracture (maximum force sustained by femur prior to fracture; N). All mice were 8 weeks old

and either wild type (WT; red), non-transplanted *oim* (OIM; blue) or e-CSC transplanted *oim* (e-CSC; black). n.s. not significant, * $P<0.05$, *** $P<0.001$; Student's t-test or Fisher exact test. Error bars +/- s.e.m.

Figure 3

Transplanted e-CSC engraft and differentiate to osteoblasts in *oim* bone (**A**) Visualisation of human donor cells with DAB staining (brown) of human specific vimentin in the tibial epiphysis, growth plate, diaphysis and fracture callus of *oim* neonatally transplanted with e-CSC, compared to age-matched non-transplanted *oim* controls. Zoomed in human MSC in the fracture callus shown. Nuclei counter stained with Haematoxylin (blue). AC; articular cartilage, SOS; secondary ossification site, HC; hyaline cartilage, M; metaphysic, PS; primary spongosia, P; periosteum, CB; cortical bone, E; endosteum. Scale bars all 100μm. (**B**) Quantitative real time PCR of donor cell engraftment calculated as the $2^{-\Delta Ct}$ of human specific *β-actin* normalised to human-mouse non-specific *β-actin* in the femoral epiphysis (Epi), diaphysis (Dia); with (+Fr) and without fracture callus (-Fr), and in the liver (Liv). * $P<0.05$, ** $P<0.01$, ***$P<0.001$; One-way ANOVA followed by Tukey's post hoc test. Error bars are s.e.m. (**C**) Linear correlation and regression equation for donor cell engraftment ($2^{-\Delta Ct}$) in the femoral epiphysis per mouse against femur stiffness (N/mm) calculated from the three point bending test. Linear line of best fit given. (**D**) Quantitative real time PCR of expression of human specific *Osteopontin* (OP) and *Osteocalcin* (OC) normalised to human specific *β-actin* ($2^{-\Delta Ct}$) in the femurs of e-CSC transplanted oim (e-CSC Tx bone; black) and compared to the basal expression level of e-CSC (basal cells; stripes) and expression level of cells grown in osteogenic permissive media for 2 weeks (differentiated cells; checks). * $P<0.05$; One-way ANOVA followed by Tukey's post hoc test. Error bars are s.e.m. (**E**) Western blot of expression of COL1A2 protein in the femurs of e-CSC transplanted

oim (e-CSC) compared to age matched wild type (WT) control and non-transplanted *oim* (OIM). Loading control is GAPDH.

Figure 4

Increased BV/TV but not BMC (**A**) Tibial length (mm) for 8 week old wild type (WT; red), non-transplanted *oim* (OIM; blue) and e-CSC transplanted *oim* (e-CSC; black). (**B**) Tibial periosteal diameter (mm) at the mid shaft for each group shown medullary bone marrow cavity (light grey) and cortical bone (dark grey). (**C**) Relative thickness of tibial cortex compared to periosteal diameter. (**D**) Bone formation rate (BFR; $\mu m^3/\mu m^2$/day) for the endosteal cortex calculated from dual calcein labelling. (**E**) Bone mineral content shown as relative frequency (%) across 16 equal intervals of mineralisation density (Displayed as a pseudocolour scheme where 0 grey level (black): low mineralisation; 256 grey level (white): maximum mineralisation). (**F**) Percentage of trabecular bone volume in the tibial metaphysis per total tissue volume (BV/TV). (**G**) Bone formation rate (BFR; $\mu m^3/\mu m^2$/days) for trabecular bone calculated from dual calcein labelling. n.s. not significant, * $P<0.05$, *** $P<0.001$; Student's t-test or Kolmogorov-Smirnov test. Error bars are s.e.m.

Figure 5

Transplanted *oim* have increased expression of genes involved in endogenous osteogenesis and chondrogenesis. (**A**) Fold changes in gene expression in the femoral epiphysis of *oim* transplanted with e-CSC when compared to non-transplanted *oim* controls, generated from a mouse osteogenesis PCR array (SABiosciences). (**B**) Expression of genes involved in endochondral ossification; *Sox9*, *Col2α1*, *Aggrecan* (*Acan*), *Col10α1*, *Smad3* and *Pthrp*. (**C**) Expression of genes involved in intramembranous ossification; *Dmp1*, *Phex*, *Biglycan* (*Bgn*), *Col1α1*, *Runx2*, *Osterix* (*Osx*) and *Igf1*. Results are given as $2^{-\Delta Ct}$ normalised to mouse *β-actin* and

hsp90ab1 for non-transplanted *oim* (*OIM*; blue) and *oim* transplanted with e-CSC (e-CSC; black). * P<0.05, ** *P*<0.01; Student's *t*-test. Error bars are s.e.m. (**D**) Visualisation of Collagen Type X with DAB staining (brown) in the tibial epiphysis of *oim* neonatally transplanted with e-CSC, compared to age-matched non-transplanted *oim* controls. Scale bar 100μm. (**E**) Visualisation of Osteopontin with DAB staining (brown) in the tibial epiphysis of *oim* neonatally transplanted with e-CSC, compared to age-matched non-transplanted *oim* controls. Scale bar 100μm.

Figure 6

Transplantation increases hypertrophic chondrocytes (**A**) Tibial growth plate architecture of 8 week old wild type (WT), non-transplanted *oim* (OIM) and *oim* transplanted with e-CSC (e-CSC). Chondrocyte matrix is stained blue and contains a reserve zone (RZ) of cells undergoing clonal expansion, a proliferative zone (PZ) containing columns of proliferating chondrocytes, and a hypertrophic zone (HZ) of differentiated hypertrophic chondrocytes. Scale bar is 20μm. (**B**) Mean widths of the tibial growth plate and growth plate zones (RZ, PZ, HZ). Significance shown for total growth plate height. n.s. not significant, * P<0.05, ** P<0.01, *** P<0.001; Student's t-test. Error bars are s.e.m. (**C**) Ratio of zones (RZ, PZ, HZ) within the growth plate. (**D**) Correlation between *Col10α1* gene expression (given as $2^{-\Delta Ct}$ normalised to mouse *β-actin* and *hsp90ab1*) and width of the hypertrophic zone (HZ) of the growth plate. (**E**) Correlation of *Runx2* and *Col10α1* gene expression. (**F**) Correlation of *Igf1* and *Bgn* gene expression. Linear line of best fit given. (**G**) Sox9 expression in ATDC5 cells was measured by quantitative real time RT-PCR using the $2^{-\Delta Ct}$ method. ATDC5 cells were either cultured in D10 medium alone (non-induced), induced to differentiate chemically, or co-cultured without cell contact with e-CSC for 7 days. Samples were tested in triplicates, and results are shown as mean±stdev. ** indicates P<0.01, when compared to non-induced ATDC5 cells; ANOVA variance analysis. (**H**) Measurement of bFGF, CTGF and PDGF-BB by e-CSC either cultured in D10 medium alone (non-primed), in co-culture with

ATDC5 cells without cell contact (primed with ATDC5), or the presence of blood serum from *oim* (primed with *oim* serum) or wild type mice (primed with WT serum). Samples were tested in triplicates, and results are shown as mean±stdev. *** indicates P<0.001 when compared to non-primed e-CSC; ANOVA variance analysis.

Supplementary Figure Legends

Supplementary Figure S1

X-ray image of tail vertebra and the corresponding MicroCT images shown at 4 different points of rotation for (**A**) a normal wild type vertebra, (**B**) a normal *oim* vertebra, (**C**) an *oim* vertebra with an obvious callus, (**D**) an *oim* vertebra with evidence of bone remodelling suggesting previous callus.

Supplementary Figure S2

Parameters of bone formation for the endosteal region of the tibial cortex, analysed from dual calcein labelling for 8 week old wild type (WT; red), non-transplanted *oim* (OIM; blue) and e-CSC transplanted *oim* (e-CSC; black). (**A**) Mineral apposition rate (MAR) in μm per day. (**B**) Percentage mineralizing surface per total bone surface (MS/BS). n.s. not significant, * $P<0.05$; Student's *t*-test. Error bars are s.e.m.

Supplementary Figure S3

Digital X-ray images of tibia from 8 week old wild type (WT), non-transplanted *oim* (OIM) and *oim* transplanted with e-CSC (e-CSC), where grey level has been pseudocoloured according to a 16-colour palette; ranging from black (low mineralization density) to white (high mineralization density).

Supplementary Figure S4

Parameters of bone formation for the tibial trabecular bone, analysed from dual calcein labelling for 8 week old wild type (WT; red), non-transplanted *oim* (OIM; blue) and e-CSC transplanted *oim* (e-CSC; black). (**A**) Mineral apposition rate (MAR) in µm per day. (**B**) Percentage mineralizing surface per total bone surface (MS/BS). n.s. not significant, *** $P<0.001$; Student's *t*-test. Error bars are s.e.m.

Supplementary Figure S5

Linear correlations of gene expression (given as $2^{-\Delta Ct}$ normalised to mouse *β-actin* and *hsp90ab1*) (**A**) *Sox9* and *Col2α1*, (**B**) *Dmp1* and *Phex*, (**C**) *Col1α1* and *Bgn*.

Supplementary Table S1

List of antibodies used for immuno-fluorescence (IF), immuno-histochemistry (IH) and western blot (WB).

Supplementary Table S2

List of primers used for RT-PCR and quantitative real time PCR.