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SELENIUM NANOPARTICLES PREPARED BY MYCO-FABRICATION: A NOVEL BONE-FORMING AGENT FOR MANAGING/PREVENTING POSTMENOPAUSAL OSTEOPOROSIS

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Selenium Nanoparticles Prepared by Myco-fabrication: A Novel Bone-forming Agent for Managing/Preventing Postmenopausal Osteoporosis

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A thesis submitted in partial fulfillment of the requirements for

the degree of Doctor of Philosophy

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ABSTRACT

Osteoporosis is a skeletal disease characterized by low bone mass and density as well as deterioration in bone microarchitecture, leading to high risk of bone fracture. With an increase in aging population, osteoporosis has become one of the major public health issues nowadays, causing significant medical and socioeconomic burdens.

Selenium is an essential trace mineral to human health. Substantial evidences have demonstrated that selenium deficiency is detrimental to bone microarchitecture and even associated with osteopenia, Kashin-Beck osteoarthropathy and osteoporosis, suggesting its crucial role in bone metabolism. Recently, selenium nanoparticles (SeNPs) have become the new research target, since they were found to possess relatively low toxicity and remarkable anti-tumor efficacy compared to other organic and inorganic selenocompounds. Nevertheless, scientific research concerning their effects on bone health is currently very limited.

By using the mushroom polysaccharide-protein complex (PSP) isolated from *Pleurotus tuber-regium*, our research team has successfully prepared novel SeNPs (PTR-SeNPs) under a simple redox system [US patent no.: 9,072,669]. We

discovered that PTR-SeNPs existed as well-dispensed spherical particles in water with an average diameter of 91.3 \pm 1.53 nm and was highly stable without significant increase in size after 13 weeks. Further characterization using TEM also showed that the particle size of SeNPs in PTR-SeNPs was around 20 nm. For individual SeNP, the clear lattice fringes (3.34Å), SAED pattern as well as EDX spectrum obtained by HR-TEM-EDX collectively indicated that the resulting nanoparticle possessed a polycrystalline structure with high level of Se (81.2%), implying a successfully fabrication of SeNPs using mushroom PSP.

Endocytosis has been widely reported as the major cellular uptake mechanism for nanoparticles. Our study found that coumarin-6 labelled PTR-SeNPs mainly localized in lysosomes of the murine preosteoblast MC3T3-E1 subclone 4 cells (bone forming progenitor cells) as early as 5 min after cellular internalization. It is the first study of its kind to report the intracellular localization of SeNPs by MC3T3-E1 cells via endocytosis. Besides, substantial previous findings have demonstrated that promoting bone formation is one of the effective strategies to prevent and/or manage postmenopausal osteoporosis. Interestingly, our study indicated that PTR-SeNPs was found to exhibit a significant dose-dependent proliferation effect (range from 1.33 – 1.73 folds) on the MC3T3-E1 cells for 24, 48 and 72 hours with the most effect dosage of 10μ M. More importantly, PTR-SeNPs (10μ M) were found to markedly induce both osteoblast differentiation and bone mineral formation of the MC3T3-E1 cells as evidenced by a significant increase in ALP activity (1.36 ± 0.10 folds), as well as an enhancement of bone nodule formation (Von Kossa and Alizarin Red S staining). Further investigation on their possible interaction with osteoclasts (bone resorption cells) also showed that PTR-SeNPs could significantly up-regulate the gene expression ratio of the bone remodeling markers OPG/RANKL (2.11 ± 0.33 folds), suggesting their indirect inhibition effect on osteoclastogenesis in addition to promoting osteoblastogenesis.

Osteoblast differentiation is a crucial step of bone formation. Previous studies have demonstrated that BMP-2 signaling via Smad-dependent and/or Smad-independent pathway(s) is one of the most important signaling transduction cascades for regulating osteoblast differentiation. Interestingly, our study discovered that PTR-SeNPs could trigger both BMP-2/Smad dependent and independent pathways simultaneously as evidenced by a significant upregulation of BMP-2 gene expression (1.64 \pm 0.24 folds) as well as Smad 1/5/8 (1.31 \pm 0.15 folds), ERK (3.67 \pm 1.48 folds), p38 (2.86 \pm 0.58 folds) and JNK (1.36 \pm 0.07 folds) proteins phosphorylation in the MC3T3-E1 cells after 4-day treatment. Besides, PTR-SeNPs were found to significantly up-regulate the gene expression of those major downstream biomarkers in bone remodeling such as Dlx5 (1.54 ± 0.06 folds), Runx2 (2.11 ± 0.1 folds), Osx (1.87 ± 0.60 folds), ALP (2.09 ± 0.10 folds), and OCN (9.69 ± 1.07 folds), further supporting their active role in promotion of osteoblast differentiation and bone mineralization.

Last but not least, by using Oryzias latipes (medaka) larvae, we have successfully established a brand new in vivo model to investigate the simulating effect of PTR-SeNPs on bone formation. And we found that PTR-SeNPs (10ppm) could significantly enhance the vertebrate development (both bone and cartilage) of medaka larvae after feeding for 11 days as visualized by Alizarin Red S and Alcian Blue double staining. Consistent with the findings of in vitro study, the gene expression level of those major osteogenic markers such as ALP (1.53 ± 0.1 folds), Runx2 (1.45 \pm 0.10 folds), Osx (1.41 \pm 0.15 folds) as well as OPG/RANKL (4.38 \pm 1.31 folds) were all significantly upregulated. Findings of this study could provide significant insights into the in vitro and in vivo effects of PTR-SeNPs on bone formation. Besides, medaka larvae, as a novel non-mammalian animal model for bone formation study, could be further developed into a rapid and economical in vivo screening platform for bone formation prior to conducting traditional animal study using mice or rats. Our long-term goal is to develop an evidence-based bone-forming agent for promoting the bone health of postmenopausal patients in our community.

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LIST OF ABBREVIATION

ALP	Alkaline phosphatase
AR	Androgen receptor
ASTM	American Society for Testing and Materials
BCA	Bicinchoninic acid
BMD	Bone mineral density
BMP	Bone morphogenetic protein
BMSC	Bone marrow stromal cell
BSA	Bovine serum albumin
BSI	British standards institution
DDI	Double deionized water
DEPC	Diethylpyrocarbonate
DLS	Dynamic light scattering
DNA	Deoxyribonucleic acid
EDTA	Ethylenediaminetetraacetic acid
EDX	Energy-dispersive X-ray spectroscopy
ER	Estrogen receptor
ERK	Extracellular receptor kinase
FBS	Fetal bovine serum

GAPDH	Glyceraldehyde 3-phosphate dehydrogenase
GSK	Glycogen synthase kinase
HRP	Horseadish peroxidase
HRT	Hormonal replacement therapy
HR-TEM	High resolution transmission electron microscopy
ICP-MS	Inductively coupled plasma mass spectrometry
ICP-OES	Inductively coupled plasma optical emission spectrometry
ICSBP	Interferon consensus sequence binding protein
IgG	Immunoglobulin G
Ihh	Indian hedgehog homolog
IOF	International osteoporosis foundation
IR	Infrared spectroscopy
ISO	International organization for standardization
ITF-2	Immunoglobulin transcription factor 2
LDL	Low density lipoprotein
Lef	Lymphoid enhancer-binding factor
LRP	Low-density lipoprotein receptor-related protein
МАРК	Mitogen-activated protein kinase
Mdm2	Mouse double minute 2 homolog

MicroCT	Micro computed tomography
MTS	3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-
	(4-sulfophenyl)-2H-tetrazolium
MWCO	Molecular weight cut off
NTA	Nanoparticles tracking analysis
OCN	Osteocalcin
OPG	osteoprotegerin
Osx	Osterix
PAGE	Polyacrylamide gel electrophoresis
PCR	Polymerase chain reaction
РКС	Protein kinase C
PKD	Protein kinase D
PMS	Phenazine methosulfate
PMSF	phenylmethylsulfonyl fluoride
PSP	Polysaccharide saccharide protein complex
РТН	Parathyroid hormone
PTR	Pleurotus tuber-regium
qPCR	Real time polymerase chain reaction
RANKL	Receptor activator of nuclear factor kappa- β ligand

Rho	Ras homolog
RIPA	Radioimmunoprecipitation assay buffer
RNA	Ribonucleic acid
RT-PCR	Reverse transcription polymerase chain reaction
Runx2	Runt-related transcription factor 2
SAED	Selected area electron diffraction
SD rats	Sprague Dawley rats
SDS	Sodium dodecyl sulfate
Se0	Elemental selenium
SEM	Standard error mean
SeNPs	Selenium nanoparticles
SERM	Selective estrogen receptor modulator
TBST	Tris-buffer saline with tween 20
Tcf	T-cell factor
TEM	Transmission electron microscopy
TGF	Transforming growth factor
XRD	X-ray diffraction

1. Introduction and Background

1.1. Osteoporosis

Osteoporosis is medically defined as a skeleton decrease with low bone mineral density or deterioration in bone micro-architecture (Figure 1), leading to high risk of bone fractures in different parts of human body. Since its only observable symptom is bone fracture, osteoporosis is commonly regarded as a kind of "silent disease". In addition to elderly, postmenopausal woman is the major high-risk group of osteoporosis due to their loss of estrogen production. As reported by International Osteoporosis Foundation¹, osteoporosis influenced 200 million women worldwide, and bone fracture commonly occurred at body parts like hip, forearm and vertebrate. Subject to the increase of world aging population, osteoporosis has become one of the major public health issues nowadays, causing significant medical & socio-economical burdens. According to IOF, the incident rate of hip fracture in men and women is projected to be increased by 310% and 240%, respectively. Thus, there is a pressing need to develop effective approaches for managing and/or preventing osteoporosis.

Normal Bone



Figure 1. Micro-architecture between normal and osteoporotic bone²

1.2. Local Situation of Bone Health and Osteoporosis

According to the statistics released from the IOF in 2013, the life expectancy of Hong Kong population will increase from 82 to 84 years old by 2050 (**Figure 2**). Although the total population of Hong Kong will be decreased, people aged over 50 and 70 years are projected to be increased for 28% and 165% by 2050, respectively (**Figure 3**).



Figure 2. Life expectancy of Hong Kong population¹



Figure 3. Projection of Hong Kong population¹

Recently, Lau and her co-investigators 3 also indicated that despite the trend of hip fracture is decreasing, the incidence of hip fracture of last 50 years was dramatically increased by 300% from 1960's to 1990's 4 .

In response to the need of health-conscious population in Hong Kong, numerous bone health products such as calcium and vitamin D supplement have been launched to local market in the past two decades. Nevertheless, a community-based study showed that the lifestyle of Hong Kong women such as insufficient physical activity and limited exposure to sunlight, significantly increased their risk to osteoporosis, resulting in a higher % of postmenopausal osteoporosis cases than that of the Western countries ⁵. As reported by the Hospital Authority of Hong Kong, approximately 4,500 cases of hip fracture were received annually, causing hospital expenditure of about 52 billion USD every year. More importantly, the cost is expected to be significantly increased in the next 10 years due to the expansion of aging population in Hong Kong.

1.3. <u>Current Medications for Osteoporosis</u>

In order to treat osteoporosis, different medications have been released and studied. The most commonly used medications are summarized

as follow:

Name of treatment	Examples	Mechanism(s) of action	Side effect(s)	Eligible for reimbursement
				in Hong Kong
Bisphosphonates	Alendronate	Inhibit bone resorption with increased	Increase cardiovascular risks	100% with prior fracture
	Risedronate	BMD and reduction of bone fractures ⁶ , ⁷	8-11	
Calcitonin	Salmon	Inhibit osteoclast with mild increase in	Very rare allergic reaction;	Only employed for pain
	calcitonin	BMD ¹²	Low efficacy ¹²	treatment in acute vertebral
				fracture
Denosumab	Denosumab	Inhibit bone resorption by preventing	Serious allergic reaction;	100% with prior fracture and
		RANKL bind to osteoclast ¹³	Hypocalcemia ^{14, 15}	intolerance to oral alendronate
Selective Estrogen	Raloxifene	Act as estrogen agonist on bone only;	Myocardial infaraction;	No
Receptor	Bazedoxifene	Do not stimulate breast cancer in	Increase risk of fatal stroke ¹⁸	
Modulators		postmenopausal women		
(SERM)		16, 17		

Table 1. Summary of current medications commonly used in osteoporosis.

Strontium Ranelate	Strontium	Slightly inhibit bone resorption and	Nausea and diarrhea;	100% with prior fracture and
	Ranelate	promote bone formation ^{19, 20}	Heart attack ²¹	intolerance to oral alendronate
Teriparatide	Teriparatide	Stimulate bone formation and inhibit	Osteosarcoma ²⁴	No
		bone resorption ^{22, 23}		
Hormone	Estrogen	Slow down bone turnover and increase	Increase risk of cardiovascular	Yes
Replacement	Progestin	BMD in postmenopausal women ²⁵⁻²⁷	diseases; Breast cancer	
Therapy (HRT)			26, 28	

Table 1 showed a list of medication currently used for treating osteoporosis. In particular, hormone replacement therapy (HRT) has been widely used to alleviate the physical symptoms and to prevent clinical consequences of postmenopausal osteoporosis ²⁹. However, recent studies by Women's Health Initiative as well as the Million Women reported that HRT would significantly increase the risk of postmenopausal women in developing breast cancer, stroke, thrombosis and cardiovascular disease ^{28, 30}. Hence, there is a pressing need to develop alternative approach for managing and/or preventing postmenopausal osteoporosis.

1.4. Bone Remodeling

Bone remodeling is a lifelong process of regulating "new bone formation" and "old bone resorption" (**Figure 4**). These two sub-processes are governed by specialized cell types named "osteoblast" as well as "osteoclast" and an imbalance of which results in metabolic bone diseases like osteoporosis ³¹. Substantial evidences have demonstrated that promoting bone formation is one of effective strategies to reduce the risk of first and recurrent fractures. Hence, there is a clear clinical need to develop new bone forming agents for the prevention and treatment of postmenopausal osteoporosis.



Figure 4. Bone remodeling system ³²

1.5. Osteoblastic Differentiation

Osteoblasts are derived from mesenchymal stem cells, which are also progenitors of myocytes, chondrocytes and adipocytes ³³. During osteoblastogenesis, osteoblast differentiation is one of the crucial steps (**Figure 5**). Substantial evidences have demonstrated that BMP-2 (bone morphogenetic protein-2) signaling via Smad-dependent and/or Smad-independent pathway(s) is one the most important signaling transduction cascades for regulating osteoblast differentiation ³⁴.



Figure 5. Osteoblastogenesis [Adopted from ³⁵]

BMPs are a group of phylogenetically conserved signaling molecules showed potent osteogenic effects and were initially identified by their capacity to induce endochondral bone formation ³⁶⁻³⁸. BMP-2, -4 and -6 are the most readily detectable BMPs in osteoblast cultures ^{36, 39}. Different BMPs have their own unique functions in osteoblast differentiation. BMP-2, BMP-6, BMP-7 and BMP-9 have been found to promote bone formation, while BMP-3 acts as an inhibitor ⁴⁰.

As shown in **Figure 6**, on receptor activation, BMP-2 transmit signals through Smad-dependent and/or Smad-independent pathways⁴¹. In Smad-dependent pathways, three classes of Smads are involved: (i) Receptor-regulated Smads (R-Smads), which are activated by Smad 1, 5 and 8; (ii) Common partner BMP mediator Smads (Co-Smads) such as Smad 4; and (iii) Inhibitory Smads (I-Smads) such as Smad 6 and 7. Upon ligand stimulation and activation by type II receptors, type I receptors phosphorylate R-Smad followed by forming complexes with Co-Smads⁴². The R-Smad/Co-Smads complexes will then translocate into the nucleus, regulating the transcription of target genes by interacting with various transcription factors and transcriptional coactivators or co-repressors. I-Smads will negatively regulate signaling by the R-Smads and Co-Smads. Upon BMP-2 signaling activation, R-Smads and Runx2 physically interact with each other and co-operatively regulate the transcription of target genes, leading to osteoblast differentiation of mesenchymal progenitor cells ⁴³⁻⁴⁵. Interestingly, through the action of R-Smads, BMP does not induce the Runx2 expression in the mesenchymal progenitor cells directly. Instead, it facilitates the expression of Dlx5 ^{46, 47}, which will then indirectly promote the Runx2 expression in osteoprogenitor cells. Previous studies also discovered that transcription factors including Hey1 (also termed HesR1 and Herp2), Tcf7, ITF-2, and ICSBP were specifically expressed in osteoblast cells by BMP-2 treatment, involving both Notch and Wnt signaling ^{48, 49}.

In Smad-independent pathway, BMP-2 has been reported to activate MAP kinases such as ERK, JNK and p38 in osteoblastic cells, regulating alkaline phosphatase (ALP) and osteocalcin (OCN) expressions ^{50, 51}. It is worth noting that BMP-2 was found to activate JNK and p38 via protein kinase D (PKD) during osteoblastic differentiation, independent of protein kinase C (PKC) activity ⁵². Previous studies also reported that Smad and p38 MAPK pathways converged at the Runx2 to regulate cell differentiation ⁵³ and Runx2 plays a central role in the BMP-2-induced differentiation of C2C12 cells, diverting them from myogenic to

osteogenic pathway ^{54, 55}. Similar to Smad-dependent pathway, the homeobox gene Dlx5 is an upstream target of Smad-independent pathway, stimulating the downstream transcription factor, Runx2 followed by upregulating bone-specific genes during osteoblastic differentiation ⁴⁷. In addition to Runx2, BMP-2 has been found to promote osterix (Osx) expression in mouse progenitor cells and chondrocytes mediated by Dlx5 but is independent of Runx2 ⁵⁶⁻⁵⁸. Interestingly, this BMP-2 inducing effect was mediated via p38 but not via ERK. Nevertheless, under osteogenic culture conditions, both ERK and p38 were involved in mediating the Osx expression ⁵⁹.



Figure 6. BMP-2 signaling pathway ^{60, 61}

1.6. Experimental Design

Selenium (Se) is an essential trace mineral to human health with a recommended daily allowance of $55\mu g/day$ for adults ⁶². Physiologically, Se is an important constituent of selenoproteins (e.g. thioredoxin reductase), which are important anti-oxidant enzymes to help protect the body from cellular damage by free radicals, supporting our immune system, cell proliferation and differentiation.⁶³⁻⁶⁶. In the past decades, substantial evidences have demonstrated that Se deficiency is detrimental to bone micro-architecture and associated with osteopenia, osteoporosis as well as Kashin-Beck osteoarthropathy ^{64, 67, 68}, suggesting its crucial role in bone metabolism. Previous studies also indicated that Se supplementation could not only restore the anti-oxidative capacity and prevent cell damage in bone marrow stromal cells, but also protect their osteoblastic differentiation inhibited by H₂O₂ via suppressing oxidative stress and ERK signaling pathway ⁶⁹.

The efficacy and toxicity of Se are highly dependent on its chemical form and dosage. Recently, Se nanoparticles (SeNPs) have become the new research target, since they were found to possess remarkable anti-cancer efficacy and low toxicity ^{70,} ⁷¹ compared to those organic and inorganic selenocompounds. Apart from toxicity evaluation ^{72, 73} as well as chemical and bio-synthesis ^{74, 75}, the *in vitro* anti-tumor

efficacy of SeNPs have been widely reported in literature for the past ten years ⁷⁶⁻⁷⁸. Although SeNPs stabilized with glutathione and bovine serum albumin have been recently found to promote the growth of primary human calvarial osteoblasts and alleviate the bone loss in ovariectomized rats, scientific research concerning the effects of SeNPs on bone health is still very limited. Besides, SeNPs aggregate easily and their health-promoting effects will be significantly reduced, once their nano size could not be maintained. This limiting factor has attracted many scientists to search for novel biomolecules that could stabilize the SeNPs without compromising their health-promoting effects.

Pleurotus tuber-regium (Fr.) Singer (also known as "tiger milk mushroom" in China), is an edible and medicinal mushroom mainly distributed in tropical and subtropical regions such as China, Australia and Nigeria ⁷⁹. This basidiomycete is the only *Pleurotus* species that can produce mushroom sclerotium ⁸⁰. Although this white-rot fungus is widely used for food and medicinal purposes in Nigeria, it is under-utilized in Asia and its large-scale cultivation is not currently available ⁸¹. By using its water-soluble polysaccharide-protein complex (PSP), we have successfully prepared novel SeNPs (PTR-SeNPs) under a simple redox system. In contrast to normal cells, PTR-SeNPs were found to significantly inhibit the growth of human

breast carcinoma MCF-7 cells (IC₅₀= 3.7μ M) by apoptosis induction via activating a ROS-mediated mitochondrial pathway ⁸². Comparing with the IC₅₀ value of native SeNPs (200 μ M) and mushroom PSP (400 μ g/mL), interestingly, the mushroom PSP surface decoration did not only stabilize the SeNPs, but also significantly enhance their cellular uptake and anti-proliferative effect on the MCF-7 cells ⁸².

Thus, in order to figure out whether PTR-SeNPs could be developed into a novel bone forming agent for prevention and treatment of osteoporosis, in this project, the *in vitro* and *in vivo* effects of PTR-SeNPs on bone formation were further investigated using the following approaches and a project summary is shown in **Figure 7**:

1.6.1. To investigate the direct effects of PTR-SeNPs on bone formation in vitro

Preosteoblast murine MC3T3-E1 subclone 4 cells (ATCC; CRL-2593) is one of *in vitro* models widely used for studying bone formation ⁸³. In this study, a dose- and time-dependent study of the osteogenic effect of PTR-SeNPs (including cell proliferation, osteoblast differentiation and bone mineral formation) on MC3T3-E1 cells was performed. The effective dosage of PTR-SeNPs for the MC3T3-E1 cells was firstly determined and then applied in the time-dependent studies.

1.6.2. To elucidate the mechanism of action of PTR-SeNPs on bone formation in vitro

Substantial evidences have demonstrated that BMP-2 signaling via Smad-dependent and/or Smad-independent pathway(s) is one the most important signaling cascades in regulating osteoblastogenesis (bone formation) ³⁴. Upon cell surface ligand binding, BMP-2 forms a hetero-tetrameric receptor complex with the dimers of both type II and type I transmembrane serine/threonine kinases, triggering phosphorylation of Smad1/5/8 in the cytoplasm. Phosphorylated Smad1/5/8 then forms a trimetric complex with Smad4 followed by translocation to nucleus, where they regulate the transcription of target genes for osteoblast differentiation. In particular, homeobox gene Dlx5, which is an upstream target of both BMP-2/Smad
dependent and BMP-2/Smad independent pathways, plays a pivotal role in stimulating the downstream transcription factors such as Runx2 and Osx³⁴. MAPKs (including ERK, p38 and JNK) signaling, which are upstream regulators of Dlx5 in the BMP-2/Smad independent pathway, has been found to play important role in osteoblast differentiation, regulating the ALP, Runx2 and OCN expressions ³⁴. Interestingly, the involvement of BMP-2/Smad dependent pathway in SeNPs mediated osteoblast differentiation has been recently reported ⁸⁴. Thus, in this study, we investigated whether PTR-SeNPs mediated osteoblastogenesis is related to BMP-2 signaling like other SeNPs using the MC3T3-E1 cells. In addition to BMP-2, a time-dependent study on the gene expression of major osteogenic markers (such as Dlx5, ALP, Runx2, Osx, OCN and OPG/RANKL) as well as protein expression and phosphorylation of Smad1/5/8 triggered by the PTR-SeNPs was performed in the MC3T3-E1 cells by qPCR and Western blotting, respectively. Besides, in order to determine the role of MAPKs signaling in the PTR-SeNPs mediated osteoblast differentiation, a time-dependent study on the protein expression and phosphorylation of ERK, p38 and JNK triggered by the PTR-SeNPs was also conducted in the MC3T3-E1 cells.

1.6.3. To study the cellular uptake behavior of PTR-SeNPs

Endocytosis has been widely reported as the major cellular uptake mechanism for nanoparticles ⁸⁵. Interestingly, our preliminary study found that after cellular internalization, coumarin-6 labelled PTR-SeNPs (120µM) co-localized with lysosomes in the MCF-7 human breast carcinoma cells round 10 min ⁸². Similarly, in this study, the cellular uptake behavior of PTR-SeNPs by the MC3T3-E1 cells was traced intracellularly by fluorescence microscopy.

1.6.4. To evaluate the effects of PTR-SeNPs on bone formation in vivo

Rat and mouse are commonly used animal models for bone formation study ⁸⁶⁻⁸⁹. Due to space requirement and time consuming, teleost fish such as zebrafish and medaka have been suggested previously to study bone metabolism in limited space ⁹⁰. Similar to mammals, the bone matrix of teleost fish is synthesized by osteoblasts in the presence of osteocyte ^{91, 92} and undergoes resorption by multinucleated osteoclasts ⁹³. More importantly, mammalian's bone remodeling markers and transcription factors such as alkaline phosphatase, tartrate-resistant phosphatase, runx2, twist, col10a1 and osterix were all found in teleost fish with similar structure and function^{90, 92, 94-96}. Compared to the rat and mouse models, advantages of using fish model include short experimental period, less space requirement and sample use as well as large sample size and cost effective.

Research studies on aging and bone metabolism using Japanese medaka (*Oryzias latipes*) as animal model have been reported previously ⁴. In this project, we established a brand new *in vivo* screening model to investigate the effect of PTR-SeNPs on bone formation using medaka larvae. In brief, a time dependent study on the vertebrae development (both bone and cartilage) of medaka larvae after feeding PTR-SeNPs for 11 days was visualized by Alizarin Red S and Alcian Blue double staining. Besides, the gene expression of major osteogenic markers such as ALP, Runx2, Osx and OPG/RANKL induced by PTR-SeNPs was also being determined.

We anticipate that findings of this study could provide significant insights into the *in vitro* and *in vivo* effects of PTR-SeNPs on bone formation. Our long-term goal is to develop an evidence-based bone-forming agent for promoting the bone health of postmenopausal patients in our community.

1.7. Hypothesis

- a) PTR-SeNPs could induce bone formation on preosteoblast MC3T3-E1 cells via BMP-2 signaling cascades
- b) PTR-SeNPs would enter the MC3T3-E1 cells via endocytosis
- c) PTR-SeNPs could stimulate bone formation in Japanese medaka larvae

1.8. Objectives

- a) To characterize the direct effects of PTR-SeNPs on bone formation using preosteoblast MC3T3-E1 cells
- b) To investigate the BMP signaling cascades modulated by PTR-SeNPs during osteoblast differentiation using MC3T3-E1 cells
- c) To study the cellular uptake behavior of PTR-SeNPs by the MC3T3-E1 cells
- d) To investigate the stimulatory effect of PTR-SeNPs on bone formation using Japanese medaka larvae





2. <u>Materials and Methods</u>

2.1. Chemicals and Reagents

All chemicals were at least analytical grade and purchased from Sigma-Aldrich Co. (St. Louis, MA, USA) unless specifically stated.

2.2. <u>Mushroom Samples</u>

Mushroom sclerotia of *P. tuber regium* were obtained from Sanming Mycological Institute in the Fujian Province of China. All sclerotia were cleaned with minimum amount of ultra-pure water to remove the dust and soils adhered on their surfaces before air-drying. The rinds (outer covering) of sclerotia were then removed in order to avoid any contamination that might come from the compost as well as the wrapping plastics bag during cultivation. All sclerotia were pulverized to pass through a screen with an aperture of 0.5 mm using a cyclotech mill (Tecator, Höganäs, Sweden). The milled mushroom powders were then transferred to airtight plastic bags and stored in a desiccator at room temperature (24 °C) prior to further polysaccharide protein complex extraction.

2.3. <u>Hot Water Extraction of Mushroom Polysaccharide Protein Complex</u>

About 500g of sclerotial powders were used for hot water extraction at 95-100°C for 2 hours with a sample-to-solvent ratio of 1:20 (w/v). After cooling, the extract was centrifuged at 18,600 x g for 1 hour at 4°C and the supernatant was collected. The whole extraction process was repeated twice and all supernatants were pooled together followed by rotary evaporation (60° C) and centrifugation under the same conditions. Subsequently, the concentrated supernatant was dialyzed using Spectra/Por® Dialysis Membrane (*M*w cutoff: 8,000) until the total dissolved solute (TDS) was similar to that of Milli-Q water. Dialyzate, which is the hot water soluble polysaccharide protein complex (PSP) was then freeze dried into powder by FreeZone 6L Console Freeze Dry Systems with Stoppering Tray Dryers (Labconco, USA) and stored in desiccator prior to selenium nanoparticles preparation.

2.4. <u>Preparation of PTR-SeNPs</u>

According to our award-winning and patented nanotechnology, 10mL of PTR-SeNPs were prepared by standardized procedures using the hot water-soluble PSP. In brief, 4mL of aqueous PSP (1.5mg/mL) was mixed with 1mL of aqueous sodium selenite (25mM) under magnetic stirring for 2 hours before drop-wise addition of 1mL of freshly prepared ascorbic acid (100mM). After reconstituting with ultrapure water to

10mL, the mixture was stirred for 10 min under room temperature and allowed to react for 24 hours at 4° C before extensive dialysis (*M*w cutoff: 8,000) until the TDS was similar to that of Milli-Q water. The resulting dialysate, which is the PTR-SeNPs was stored at 4° C prior to further composition analysis and structure characterization.

2.5. Composition Analysis and Structure Characterization of PTR-SeNPs

2.5.1. Selenium concentration of PTR-SeNPs

In brief, 0.5mL of PTR-SeNPs was mixed with 10mL of trace metal grade 70% concentrated nitric acid followed by microwave digestion with ETHOS One Advanced Microwave Digestion System (Milestone S.r.l., Italy), using a temperature program of initial temperature 180°C with a hold of 20 min followed by a temperature rise to 250°C with a final hold of 30 min. Digestate was then transferred into a volumetric flask and marked up to 25mL using Milli-Q water. After filtered with 0.45µm cellulose acetate syringe filter, the Se concentration of filtrate was determined by Agilent 710 ICP-OES (Agilent Technologies Inc., USA) using 196.026 as emission wavelength and 203.985 as reference. For quantification, a standard curve of selenium ranging from 0.1ppm to 5ppm was prepared.

2.5.2. Particles size and stability of PTR-SeNPs

In brief, PTR-SeNPs was diluted 1,000 folds followed by injection into Nanosight NS300 (Malvern Instruments Ltd., UK) for particles size determination using nanoparticles tracking analysis (NTA). To investigate its stability, PTR-SeNPs, nanoparticles was stored at 4°C and samples were collected for every 2 weeks prior to particles size determination using the Nanosight NS300.

2.5.3. Structure characterization of PTR-SeNPs using HR-TEM-EDX

In short, 30μ L of PTR-SeNPs was firstly dropped onto a hollow carbon grid and air-dried. The whole procedure was repeated for 5 times. The particle size of specimen was then visualized by JEM-2100F Field Emission Electron Microscope (JEOL Ltd., Japan) with magnification of $20,000\times$, $40,000\times$ and $150,000\times$ using 200kV electrons beam. Subsequently, single nanoparticle was selected under magnification of $150,000\times$ and its elemental composition and crystal structure were further investigated by using HR-TEM-EDX (JEOL JEM-2100F + Horiba EX-250). All images obtained were analyzed by software ImageJ.

2.5.4. Structure interaction between SeNPs and PTR-PSP

In brief, PTR-PSP and freeze-dried PTR-SeNPs were individually ground into powder together with potassium bromide under humidity-controlled environment. Subsequently, pellet discs of each powder mixture were prepared and their Infrared spectra were determined and compared by using NicoletTM iSTM 50 Fourier Transform Infrared Spectrometer (FT-IR) in order to figure out the major functional groups that are responsible for the association between PTR-PSP and SeNPs.

2.6. <u>Cell Culture</u>

MC3T3-E1 subclone 4 preosteoblast cells were purchased from American Type Culture Collection (ATCC ® CRL-2593, Rockville, MD, USA) and maintained in ascorbic acid free α -Minimum Essential Medium (α -MEM; Invitrogen, San Diego, CA, USA) supplemented with 10% fetal bovine serum (FBS; Invitrogen, San Diego, CA, USA), 1% 100× antibiotics and anti-mycotics (Invitrogen, San Diego, CA, USA) in a 37°C incubator with 95% relative humidity and 5% CO₂. Cells were routinely passaged using 0.25% trypsin/0.1% EDTA at 70 – 90% confluence.

2.7. Direct Effects of PTR-SeNPs on Bone Formation

2.7.1. Cell proliferation assay

In brief, MC3T3-E1 cells were firstly seeded into 96-well tissue culture plates at 15,000 cells/well. Cells pretreated with different concentrations of PTR-SeNPs (10nM - 10µM), 6µg/mL PTR-PSP or 10µM sodium selenite were then incubated for 24, 48 and 72 hour followed by determining cell proliferation using MTS [3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetr azolium] assay (Promega). To perform MTS assay, 4.6 mg of phenazine methosulfate (PMS) was dissolved in 5mL of PBS, while 42mg of MTS powder was dissolved in 21mL of PBS with pH adjustment to 6.0 - 6.5. Subsequently, 4.2 mL of MTS solution was mixed with 1mL of PMS solution. After filtered with 0.22µM filter, cells were added into the MTS/PMS mixture ($20\mu L$ /well) and incubated for 1-2 hours at $37^{\circ}C$ in a humidified incubator with 5% CO₂. Absorbance was determined at 490nm using a microplate reader (Clariostar, USA).

To study osteoblastic differentiation, MC3T3-E1 preosteoblast cells cultured in osteogenic medium containing 10nM β -glycerophoshpate and 50 µg/mL ascorbic acid were pretreated with 10µM of PTR-SeNPs for 7 days. ALP activity in the MC3T3-E1 cells was then determined by spectrophotometry at 410 nm, while ALP-containing cells (osteoblasts) were visualized by ALP staining kit. Fresh culture media were replenished every 3 – 4 days.

2.7.2.1. ALP activity

In brief, MC3T3-E1 cells pretreated with 10 μ M of PTR-SeNPs for 7 days were washed with PBS followed by lysed with 50mM Tris-HCl (200 μ L/well, pH 8.0) and 0.5% Triton X-100. Cell lysates were then harvested and transferred into 1.5mL centrifuge tubes before storage at -80°C. After defrosted, all lysates were centrifuged (21,000 × g; 20 min; 4°C) and 10 μ L of supernatant was added into 96 wells in triplicate followed by incubation with 200 μ L of 10mM *p*-nitrophenyl phosphate (*p*NPP) in 1M diethanolamine and 0.5mM MgCl₂ (pH 9.8) for 15 – 30 min. Consequently, the reaction was quenched by adding 50 μ L of 3M NaOH and the resulting absorbance was measured at 410nm using a microplate reader. To normalize the ALP activity, the total protein content was also determined by BCA assay. In short, 200 μ L of BCA solution (mixing BCA reagent A with B as indicated in manufacture's manual) was added into a well containing 2μ L of protein sample and 23μ L of water followed by incubation at 37°C for 30 min. Similarly, the resulting absorbance was measured at 595nm using a microplate reader.

2.7.2.1. ALP staining

In brief, MC3T3-E1 cells pretreated with 10µM of PTR-SeNPs for 7 days were washed with PBS and fixed with 10% formalin for 15 min. The ALP in cells were then stained by Fast Blue RR salt and Naphthol AS-MX phosphate alkaline solution for 60 min followed by rinsing thoroughly with deionized water.

2.7.3. Mineralization assay

To study bone mineral formation, MC3T3-E1 cells cultured in osteogenic medium containing 10nM β -glycerophoshpate and 50 μ g/mL ascorbic acid were pretreated with different concentration of PTR-SeNPs (10 μ M) for 16 days. Bone nodules formed in the MC3T3-E1 cells were then visualized and compared by Von Kossa and Alizarin Red S staining.

2.7.3.1. Von Kossa staining

To detect calcium deposit, cells were pre-washed with PBS and fixed with 10% formalin for 15 min. Fixed cells were then treated with 5% silver nitrate for 30 min

under light following by sequential washing with deionized water, 5% sodium carbonate in 10% formalin (1 min), deionized water and 5% sodium thiosulfate (5 min). Consequently, stained well was captured and analyzed by ImageJ to quantify calcium deposition.

2.7.3.2. Alizarin Red S staining

Similarly, cells were pre-washed with PBS and fixed with 10% formalin for 15 min. Fixed cells were then stained with 40nM Alizarin Red S solution (pH 4.1) for 20 min with gently shaking followed by heavy washing 4 times with 4mL of deionized water for 5 min per well. Excessive water in the wells was removed and the resulting image was captured. To quantify the calcium deposit, 800µL of 10% (v/v) acetic acid was added into each well followed by incubation with shaking for further 30 min. All stained cells were then scraped with cell scraper and transferred into a 1.5mL centrifuge tube using acetic acid. After vortex-mixing for 30 sec, 500µL of mineral oil was added and the slurry was heated up to 85°C for 10 min followed by cooling down in ice for 5 min. All tubes were then centrifuged at $20,000 \times g$ for 15 min and 500μ L of the supernatant was transferred into new Eppendorf tubes. After adding 200μ L of 10% (v/v) ammonium hydroxide, the mixture was aliquot in a 96 well plate in triplicate followed by measuring the absorbance with a microplate reader at 405nm.

To further characterize the direct effects of PTR-SeNPs on osteoblastogenesis, mRNA expression of major osteogenic markers (such as ALP, Dlx5, Runx2, Osx, OCN and OPG/RANKL) in the MC3T3-E1 cells pretreated with 10µM of PTR-SeNPs for 3, 7 and 16 days were quantified by real-time PCR. For total RNA extraction, cells were washed twice with 1mL of ice cold PBS and total RNA was isolated with 1mL of Trizol reagent per 6 wells according to manufacturer's protocol (Life technologies, USA). After homogenization, 200µL of chloroform was added with gentle shaking and the mixture was allowed to stand for 3 min at room temperature. All samples were then centrifuged (12,000 \times g; 10 min; 4°C) and the top aqueous layers were transferred into new tubes prior to adding 500µL of 2-propanol. Subsequently, the whole reaction mixture was centrifuged under the same conditions and the 2-propanol was removed followed by washing with 1mL of 75% ethanol in DEPC H₂O. After centrifugation $(7,500 \times g; 5 \text{ min}; 4^{\circ}\text{C})$, the 75% ethanol was removed and the RNA pellet was air-dried. Once the RNA pellets turned into milky, 10 - 20µL of DEPC H₂O was added to re-dissolve the RNA and stored at -80°C until further qPCR analysis.

2.7.5. Real-time polymerase chain reaction (qPCR)

In brief, 20ng of RNA was firstly prepared by One-Step SYBR Green RT-qPCR according to manufacturer's instructions. In a 20μ L reaction, the reaction mixture

contained 10µL of 2X One Step SYBR RT-PCR Buffer 4, 0.8µL of PrimeScript One-Step Enzyme Mix 2, 0.4µM of Forward PCR primers, 0.4µM of Reverse PCR primers and 20ng of RNA sample. qPCR analysis of BMP-2 and major osteogenic markers (such as ALP, DLx5, Runx2, Osx, OCN and OPG/RANKL) was performed by a Pikoreal thermal cycler system (Thermo, USA) using thermal cycling program and primers shown in **Table 2** and **3**, respectively. Relative mRNA expression of each osteogenic marker gene was normalized to that of GAPDH.

Tuble 2. Thermal eyening program of the encland york (mouse)				
Stage	Temperature	Hold Time		
Reverse transcription	42 °C	5 min		
(1 cycle)	95 °C	10 sec		
PCR reaction	95 °C	5 sec		
(40 cycle)	60 °C	20 sec		
Melting curve analysis	95 °C	0 sec		
	65 °C	15 sec		
	95°C	0 sec		

 Table 2. Thermal cycling program of qPCR analysis (mouse)

 Table 3. Primer list for qPCR analysis (mouse)

Gene	Forward Primer (5' – 3')	Backward Primer (5' – 3')
Runx2	AAGTGCGGTGCAAACTTTTCT	TCTCGGTGGCTGGTAGTGA
ALP	AACCCAGACACAAGCATTCC	GAGAGCGAAGGGTCAGTCAG
Dlx5	GTTTCAGAAGACTCAGTACCT	TGACTGTGGCGAGTTACAC
BMP2	CCCCAAGACACAGTTCCCTA	GAGACCGCAGTCCGTCTAAG
Osx	AGAGGTTCACTCGCTCTGACGA	TTGCTCAAGTGGTCGCTTCTG
OCN	TAGTGAACAGACTCCGGCGCTA	TGTAGGCGGTCTTCAAGCCAT
OPG	ACGAACTGCAGCACATTTGG	TGTTTCCGGAACACACGTTG
RANKL	ACGCCAACATTTGCTTTCGG	ATTGCCCGACCAGTTTTTCG
GAPDH	CATGGCCTTCCGTGTTCCTA	CCTGCTTCACCACCTTCTTGAT

2.7.6. Total protein extraction

With a purpose to dissect the BMP-2 signaling (both Smad dependent and independent pathways) mediated by PTR-SeNPs during osteoblast differentiation, protein expression and phosphorylation of Smad1/5/8 ERK, p38 and JNK in the MC3T3-E1 cells pretreated with 10 μ M of PTR-SeNPs for 4 days were determined by Western blot analysis. For total protein extraction, cells were washed twice with PBS and lysed with 50 μ L of RIPA Lysis buffer containing PMSF and 1X protease and phosphatase inhibitor cocktail. Cells were then scraped and transferred into 1.5mL centrifuge tubes followed by sonication (1 min) and centrifugation (22,000 × g; 20 min; 4°C). Protein content of the supernatant was determined by BCA protein assay. After mixed with 5X sample buffer, the reaction mixture was heated up to 95°C for 5 min and stored at -20°C prior to Western blot analysis.

2.7.7. Western blot analysis

For Western blotting, 40µg of protein sample was loaded and separated by 10% SDS-PAGE using gel electrophoresis. Separated proteins were then transferred to PVDF membranes using iBlot 2 Dry Blotting System. After blocking with 5% BSA (w/v) or 5% blocking grade non-fat dry milk powder (Bio-rad, USA) in Tris-buffered saline containing 0.1% (v/v) Tween-20 (TBST) for 1 hour at room temperature, all

membranes were incubated with specific primary antibodies including p-Smad 1/5/8, Smad 1/5/8, p-ERK, ERK, p-p38, p-38, p-JNK, JNK and GAPDH [dilution: 1:1000; 5% BSA in TBST] at 4°C overnight. Subsequently, all membranes were extensively washed with TBST and incubated with the secondary anti-mouse IgG HRP-linked antibodies at 1:2000 dilution for 1 hour at room temperature. Finally, the membranes were further washed with TBST, while chemiluminescence for detection was developed using EMD Millipore Immobilon Western Chemiluminescent HRP substrate (Millipore, USA) according to the manufacturer's instructions. Image acquisition was performed using AzureTM c600 digital imager (Azure Biosystem Inc., USA) followed by quantification with ImageJ.

2.7.8. Cellular uptake behavior of PTR-SeNPs

In brief, MC3T3-E1 cells were firstly seeded in a 35mm confocal dish for 24 hours followed by sequential staining with Hochest 33342 (1µg/mL; nucleus; Invitrogen, CA, USA) and Lysotracker Deep Red (75µM; lysosomes; Invitrogen, CA, USA) for 20 and 30min, respectively. After further incubation with 10µM of coumarin-6-loaded PTR-SeNPs at 37°C for 10 min, the cellular uptake behavior of PTR-SeNPs by MC3T3-E1 cells was intracellularly traced by fluorescence microscopy.

2.8. Japanese Medaka

Orange-red, outbred line of Japanese medaka (Oryzia latipes) was originated from the Molecular Aquatic Toxicology Laboratory of Duke University. A brood colony was maintained at 100-200 individuals (60-80 fish \times 3 tanks) at any given time. This brood colony produced all the embryos for the project. In brief, adult fish in brood colony was maintained at 25°C at 14:10 light/dark cycle to provide favorable conditions for egg laying. Under this condition medaka fish produce eggs daily. Fifty percent of tank water was changed daily with de-chlorinated tap water and was maintained at pH 8.0 and 30% salinity. Water pH and temperature was monitored daily; nitrate, nitrite, ammonia and water hardness were checked every month. Fish were fed twice daily with commercial fish feed, Otohime B1 (Nisshin Co, Japan), and supplemented with hatched brine shrimp Artemia nauplii (Lucky Brand, O.S.I., USA) for 3 days per week. Embryo less than 4 hours post-fertilization was collected and transferred to a petri dishes containing de-chlorinated water for culturing.

2.8.1. Japanese medaka embryo culture

All collected medaka embryos were firstly cultured in a petri dish using saturated sea salt water for 1 day-disinfection followed by culturing in de-chlorinated water. Embryos were cleaned and dead eggs were pick out every day during change of water in order to prevent contamination. Embryos were cultured at $25 - 30^{\circ}$ C until hatched and larvae were transferred to new containers before performing PTR-SeNPs exposure experiment.

2.8.2. Fish diet preparation

In brief, 10 g of commercial fish feed (Otohime β 1) was firstly mixed with 10mL of PTR-SeNPs to make a PTR-SeNPs based diet containing 10 ppm Se. After freeze-drying, the PTR-SeNPs based diet was grinded by using mortar and pestle to pass through a 100µm sieve. For control diet, 10mL of MilliQ water was used instead. Final Se concentration of both control and SeNP-based diets were measured by ICP-MS before use.

2.8.3. PTR-SeNPs exposure experiment using medaka larvae

After collecting the 5 days post-hatched medaka larva, both control and PTR-SeNPs treatment groups were cultured in 6-well cell-culture plates (one medaka larvae per well) and fed the control or PTR-SeNPs based diet (10ppm) twice daily for a total of 11 days. During fish sampling, medaka larvae were anaesthetized by immersing in ice-cold aquarium water for 30 sec.

Similar to the *in vitro* study, the mRNA expression of major osteogenic markers (such as ALP, Runx2, Osx and OPG/RANKL) in the medaka larvae fed with 10ppm of PTR-SeNPs for 2 days were quantified by qPCR. As mentioned in 2.7.4 and 2.7.5, total RNA isolation from medaka larvae (n=3) were performed using Trizol reagent followed by One-Step SYBR Green RT-qPCR analysis using Pikoreal thermal cycler system (Thermo, USA). Thermal cycling program and primers used are shown in **Table 2** and

Table 5, respectively. Relative mRNA expression of each osteogenic marker gene was normalized to that of 18S.

2.8.4. Bone mineralization of medaka larvae

Mineralization is the end point of bone formation. Bone and cartilage in medaka larvae after treatment with 10ppm of PTR-SeNPs for 11 days (n = 6) were double-stained by Alizarin Red S and Alcian Blue. In brief, medaka larvae were firstly fixed in 10% formalin (2 hours) followed by dehydration using 50% ethanol (10 min). After washing three times with PBS (10 min), medaka larvae were stained overnight with 0.02% Alcian blue 8GX in 70% ethanol containing 0.06M MgCl₂. Subsequently, medaka larvae were washed three times with absolute ethanol (10 min) followed by sequential washing with 90%, 75% 50%, 25% ethanol and finally PBS (20 min each). After bleaching with 1.5% H₂O₂ in 1% KOH (40 min) and washing three time with PBS (10 min), medaka larvae were further digested with 1% trypsin solution in 30% saturated sodium borate solution (40 min). For bone staining, digested larvae were washed three times with PBS (10 min) and stained with saturated Alizarin Red S in 0.5% KOH overnight followed by de-staining three times with 0.5% KOH (10 min). After sequential washing with 20% glycerol/0.25% KOH and 50% glycerol/0.25% KOH, all medaka larvae specimens were stored in 100% glycerol at 4°C prior to examination using stereomicroscope Nikon SMZ18 and upright microscope Eclipse Ci-E.

2.8.5. Total RNA isolation

To further characterize the direct effects of PTR-SeNPs on osteogenesis *in vivo*, mRNA expression of major osteogenic markers (such as ALP, Runx2, Osx and OPG/RANKL) in medaka larvae pretreated with 10ppm of PTR-SeNPs for 2 days were quantified by real-time PCR. For total RNA extraction, total RNA of 3 larvae were isolated with 1mL of Trizol reagent according to manufacturer's protocol (Life technologies, USA). After homogenization, 200µL of chloroform was added with gentle shaking and the mixture was allowed to stand for 3 min at room temperature. All samples were then centrifuged (12,000 × g; 10 min; 4°C) and the top aqueous layers were transferred into new tubes prior to adding 500µL of 2-propanol. Subsequently, the whole reaction mixture was centrifuged under the same conditions and the 2-propanol was removed followed by washing with 1mL of 75% ethanol in DEPC H₂O. After centrifugation (7,500 \times g; 5 min; 4°C), the 75% ethanol was removed and the RNA pellet was air-dried. Once the RNA pellets turned into milky, 10 - 20µL of DEPC H₂O was added to re-dissolve the RNA and stored at -80°C until further qPCR analysis.

2.8.6. Real-time polymerase chain reaction (qPCR)

In brief, 50ng of RNA was firstly prepared by One-Step SYBR Green RT-qPCR according to manufacturer's instructions. In a 20 μ L reaction, the reaction mixture contained 10 μ L of 2X One Step SYBR RT-PCR Buffer 4, 0.8 μ L of PrimeScript One-Step Enzyme Mix 2, 0.4 μ M of Forward PCR primers, 0.4 μ M of Reverse PCR primers and 20ng of RNA sample. All qPCR analysis of those major osteogenic markers (such as ALP, Runx2, Osx and OPG/RANKL) was performed by a Pikoreal thermal cycler system (Thermo, USA) using thermal cycling program and primers shown in **Table 2** and **5**, respectively. Relative mRNA expression of each osteogenic marker gene was normalized to that of 18S.

Stage	Temperature	Hold Time
Reverse Transcription	$42^{\circ}C$	5 minutes
(1 cycle)	95°C	10 seconds
PCR reaction	95°C	5 seconds
(40 cycle)	60°C	20 seconds
Melting Curve analysis	95°C	0 seconds
	65°C	15 seconds
	95°C	0 seconds

Table 4. Thermal cycling program of qPCR analysis (medaka larvae)

Table 5. Primer list for qPCR analysis (medaka larvae)

Gene	Forward Primer (5' – 3')	Backward Primer (5' – 3')
Runx2	TCAGACGCAGTCACAGA	ACTCAGGTAGGACGGATA
ALP	AGGGAAGCCGATGTTTCCTG	TCCTGCATCTTTGGCCCATT
Osx	TCTCCCCTCAGCTTCCTTAG	CTGTGTTGCAGACAGCCAGT
OPG	GCTGGTTATGTGCGTAAA	AAGAATGCTGAGAGGAA
RANKL	CTGCTTCCGCCACTACAA	GGGTTGCCAATATCCACAG
18S	CATGGCCTTCCGTGTTCCTA	CCTGCTTCACCACCTTCTTGAT

2.9. Statistical Analysis

All experiments were conducted in triplicate and results were presented as mean \pm standard deviation. Except gene expression and mineralization results, which were analyzed by two-tailed Student's *t*-test, all mean values were analyzed by parametric one-way ANOVA followed by multiple comparisons using Turkey-HSD to detect significant differences at *p* < 0.05, *p* < 0.01 *p* < 0.001 (GraphPad Prism, San Diego, CA, USA).

3. <u>Results and Discussion</u>

3.1. Preparation of PTR-SeNPs

In this study, PTR-SeNPs was prepared according to our previously reported method ⁸² with some modifications, in which sodium selenite and ascorbic acid were used as the Se source and reducing agent, respectively (

Figure 8). The structure and composition PTR-SeNPs were then characterized

by Nanosight NS300, HR-TEM-EDX (JEOL JEM-2100F + Horiba EX-250), and FT-IR (NicoletTM iSTM 50) currently used in our laboratories.

During PTR-SeNPs preparation, selenite (SeO_3^{-2}) was firstly reduced to elemental selenium (Se^0) by ascorbic acid according to the following equations ⁹⁷.

 $\text{SeO}_3^{-2} + 2\text{H}^+ \rightarrow \text{H}_2\text{SeO}_3$

$$H_2SeO_3 + 2C_6H_8O_6 \rightarrow Se^0 + 2C_6H_6O_6 + 3H_2O$$

Se⁰ were then aggregated to form a "Se nucleus" and continuously grew bigger. Fabricated by suitable concentration of PTR-PSP, further aggregation of "Se nucleus" was prevented within a nano range to form Se nanoparticles.



Figure 8. Schematic illustration of PTR-SeNPs preparation

3.2. Particles size and stability of PTR-SeNPs

Nanoparticle tracking analysis (NTA) measures particle size using laser to track the movement of nanoparticles within a period of time, calculating their particles size according to the rate of Brownian motion and Stokes-Einstein equation. Based on the NTA using Nanosight NS300, size distribution of the PTR-Se nanocomposite was found to has a mode of 82nm (**Figure 9A**) with an average particle size of 91.3 \pm 1.53nm. As shown in **Figure 9B**, the average particle size of PTR-Se nanocomposite was highly stable without significant change in size (\pm 10nm) after 13 weeks' storage, suggesting its relatively high stability compared to other SeNPs prepared by using bovine serum albumin ⁷³. Besides, the tiny peak detected by NTA at 275nm might be due to the presence of mushroom PSP or other impurities (**Figure 9A**).

For HR-TEM, particle images are produced by differential amount of electron beam passing through the specimen and background, creating a darker areas for more electron dense regions (i.e. the particles) ⁹⁸. In this study, further characterization of PTR-SeNPs using HR-TEM (JEOL JEM-2100F; 200kV; magnification 40000×) also indicated that SeNPs itself (without mushroom PSP decoration) existed as well-dispensed spherical particles in water with an average diameter of 17.3 ± 15.0 nm (Figure 9C & D). It is also worth to note that majority of the resulting SeNPs was ranged from 5 to 35nm (Figure 9D). Among those existing international standards for "nanoparticles" shown in **Table 6**, the most widely accepted definition is that at least two dimensions of the particle are less than 100nm in size ⁹⁹. For NTA, the particles size is determined by tracking the x and y axis Brownian motion of particles, involving 2 dimension calculation of hydrodynamic diameter ¹⁰⁰. For ImageJ analysis of HR-TEM image, the Feret diameter measured is the longest distance of 2 parallel planes of an object with any shape ¹⁰¹. In the present study, as the average particle size of PTR-SeNPs determined by both NTA and HR-TEM are less than < 100nm (Figure 9), PTR-SeNPs met the international standards and could be regarded as "nanoparticles".







Figure 9. Particles size and stability of PTR-SeNPs.

D

(A) Particles size distribution of PTR-SeNPs measured by Nanosight NS300. PTR-SeNPs were diluted and its particle size distribution was determined by nanoparticle tracking analysis using Nanosight NS300. (B) Time dependent study on stability of PTR-SeNPs. PTR-SeNPs were prepared and stored at 4°C for 13 weeks. Samples of PTR-SeNPs were collected every 2 weeks and their particles size was measured by Nanosight NS300. Particle size was measured 5 times for each independent determination and results were presented as mean value of three independent measurements ± S.D. (C) Representative TEM image of PTR-SeNPs. PTR-SeNPs were seeded on a hollow carbon grid and examined under HR-TEM (JEOL JEM-2100F; 200kV; magnification 40000×; scale bar: 100nm). (D) Particles size distribution of SeNPs measured by HR-TEM. HR-TEM images obtained were analyzed with ImageJ to calculate particle size of SeNPs. Results were based on three independent measurements (600 particles/ measurement).

	1	
Standard	Nanoparticle	
International Organization for	A particle spanning 1 – 100nm (diameter)	
Standardization (ISO)		
American Society of Testing Materials	An ultrafine particle whose length in 2 or 3	
(ASTM)	places is 1 – 100nm	
National Institute of Occupational	A particle with diameter between 1 and	
Safety and Health (NIOSH)	100nm, or a fiber spanning the range 1 -	
	100nm	
British Standards Institution (BSI)	All the fields or diameters are in the	
	nanoscale range	

Table 6. Existing international standards for nanoparticles

3.3. Characterization of PTR-SeNPs using HR-TEM-EDX and FT-IR

For individual PTR-SeNP, in addition to having clear lattice fringes/d-spacing (0.33nm; Figure 10A), the SAED pattern of PTR-SeNPs exhibited diffuse rings constructed with white dots (Figure 10B), collectively indicating that the nanoparticle possessed a polycrystalline structure ⁹⁸. Besides, PTR-SeNPs shared a very similar 6-ring SAED pattern to that of Se-nanocrystallites in SiO₂, suggesting a cubic crystal structure as previously reported ¹⁰². Further investigation on its elemental composition by EDX analysis also showed that PTR-SeNPs contained high level of Se (81.2%; Figure 10C), indicating a successfully fabrication of SeNPs using mushroom PSP. Table 7 showed that protein (50.3%) and polysaccharide (48.8%) were the major components of PTR-PSP, suggesting a rich source of hydroxyl (-OH-) and imino (-CO-NH-) groups. In this study, the chemical binding of mushroom PSP to the surface of SeNPs was further investigated by FT-IR analysis. As shown in Figure 10D, the absorption peaks at 1538.7 and 3390 – 3400cm⁻¹ in the IR spectrum of PTR-PSP corresponded to the secondary imino (-CO-NH-) groups of its proteins and the stretching vibrations of hydroxyl (-OH-) groups of its polysaccharides. Interestingly, PTR-PSP shared very a similar FT-IR spectrum to that of PTR-SeNPs, indicating that PTR-PSP formed part of the PTR-SeNPs nanocomposite. Nevertheless, both absorption peaks of hydroxyl groups and imino groups shifted from 1538.7 to 1537.2 and from 3397.23 to 3393.47, respectively, indicating the occurrence of interaction

between the hydroxyl groups and imino groups of PTR-PSP and the Se atoms as previously reported ⁸². These results suggest that the SeNPs were capped by mushroom PSP mainly through the formation of Se-O and Se-N bonds, leading to a stable spherical structure of SeNPs.





Figure 10. Representative HR-TEM image of (A) lattice fringes and (B) corresponding selected area electron diffraction (SAED) pattern of PTR-SeNPs. PTR-SeNPs was seeded on a hollow carbon grid and examined under HR-TEM (20keV; 150000×). Lattice fringes as well as diameter, radius and d-spacing of diffraction rings were measured using ImageJ. (C) Representative EDX analysis of PTR-SeNPs. (D) FT-IR analysis of PTR-SeNPs and PTR-PSP. Freeze-dried PTR-SeNPs and PTR-PSP were mixed with KBr to form sample discs followed by analysis using NicoletTM iSTM 50 FT-IR Spectrometer.
Table 7. Composition of PTR-PSP

Composition	Percentage
Polysaccharide	48.8%
Protein	50.3%

3.4. Direct Effects of PTR-SeNPs on Bone Formation Using MC3T3 E-1 Cells

3.4.1. Cell proliferation

In general, PTR-SeNPs was found to exhibit a significant dose-dependent proliferation effect (range from 1.33 – 1.73 folds) on the MC3T3-E1 cells for 24, 48 and 72 hours with the most effect dosage of 10µM (Figure 11A-C). In addition to 10µM of sodium selenite (common food source of inorganic Se), we further compared the proliferation effect of PTR-SeNPs (10µM) and PTR-PSP alone at concentration used for preparing 10μ M of PTR-SeNPs (i.e. 6μ g/mL) in order to investigate whether mushroom PSP decoration would contribute the proliferation effect of PTR-SeNPs on MC3T3-E1 cells. Interestingly, except for 48-hour treatment, PTR-PSP had no effect on the proliferation of MC3T3-E1 cells, and sodium selenite (10µM) was even found to significantly inhibit the growth of MC3T3-E1 cells for all time intervals (Figure **12A-C**). In contrast, the significantly highest proliferation effect of PTR-SeNPs on the MC3T3-E1 cells (Figure 12A-C) suggested that SeNPs itself, instead of PSP, was mainly responsible for the proliferation effect of whole PTR-Se nanocomposite, having relatively lower toxicity than that of sodium selenite under the same dosage. Besides, it is worth noting that, compared with the effective dosage (640µM) of other SeNPs on bone cell proliferation ¹⁰³, PTR-SeNP exhibited strong proliferation effect on bone cells with remarkable low concentration (10µM only).



Figure 11. Dose and time dependent study of the proliferation effect of PTR-SeNPs on MC3T3-E1 cells. Cells were treated with different concentrations of PTR-SeNPs (10nM - 10 μ M) in normal medium for (A) 24 (B) 48 and (C) 72 hours. Cell proliferation was determined by MTS assay and results were normalized and compared with that of negative control group (1% deionized water; 1% DDI). Results are mean values of three determinations ± SEM. Means with "***" are significantly different (*p* < 0.001, one-way ANOVA).



Figure 12. Time-dependent study of proliferation effect of PTR-SeNPs, PTR-PSP and sodium selenite on MC3T3-E1 cells. Cells were treated with PTR-SeNPs (10 μ M), PTR-PSP (6 μ g/mL) and Na₂SeO₃ (10 μ M) in normal medium for (A) 24, (B) 48 and (C) 72 hours. Cell proliferation was determined by MTS assay and results were normalized and compared with negative control group (1% deionized water; 1% DDI). Results are mean values of three determinations ± SEM. Means with letters (a-d) are significantly different (*p* < 0.05, one-way ANOVA).

ALP is a phosphor-hydrolytic enzyme that plays a major role in bone formation, regarding as a phenotypic marker for osteoblasts during differentiation. As shown in **Figure 13A** and **15A**, PTR-SeNPs (10 μ M) markedly induced the osteoblastic differentiation of MC3T3-E1 cells after a 7-day treatment as evidenced by a significant increase in ALP activity (1.36 ± 0.10 folds) and up-regulation of ALP gene expression (2.09-folds). The ALP containing MC3T3-E1 cells pre-treated with 10 μ M of PTR-SeNPs (purple-blue area) were further visualized by ALP staining (**Figure 13B**).

Mineralization is the final stage of osteogenesis and osteocalcin (OCN) which expresses in differentiating cells of intermediate and mature nodules, is a specific marker gene for mineralization. As shown in **Figure 14A&B**, PTR-SeNPs (10µM) significantly promoted the bone nodule formation in MC3T3-E1 cells as visualized by both Von Kossa (20 folds) or alizarin red S staining (3 folds) followed by semi-quantification of corresponding calcium deposition. Further investigation on the gene expression of OCN also indicated that PTR-SeNPs (10µM) could significantly up-regulate this mineralization maker gene for 9.69 folds (**Figure 15B**), demonstrating its strong stimulating effect on bone mineralization. Runx2 and osterix (Osx) are the two main osteoblastic transcription factors in osteogenesis ^{104, 105}. Activation of Runx2 has been found to upregulate several osteoblast-specific genes (such as OCN), promoting bone formation in both cell and animal models ¹⁰⁶⁻¹⁰⁸. Similarly, Osx is another critical transcriptional factor in osteoblast differentiation and bone formation previously identified in a Osx-null mutant mice study ⁵⁶. As shown in **Figure 15B & C**, the gene expression of both Runx2 (2.11 folds) and Osx (1.87 folds) in MC3T3-E1 cells pre-treated with 10μM of PTR-SeNPs were significantly up-regulated, indicating that PTR-SeNPs (10μM) enhanced the osteoblastic differentiation and bone formation in MC3T3-E1 cells mediated by these two transcription factors.

Bone remodeling is a lifelong process of regulating "new bone formation" and "old bone resorption". These two sub-processes are governed by specialized cell types named "osteoblast" as well as "osteoclast" and an imbalance of which results in metabolic bone diseases like osteoporosis ³¹. OPG and RANKL are bone remodeling markers secreted by osteoblasts for inhibiting and activating osteoclasts, respectively ^{109, 110}. Thus, OPG/RANKL ratio is a widely used parameter to figure out the possible interaction between osteoblasts and osteoclasts. Interestingly, our study discovered that PTR-SeNPs (10µM) significantly up-regulate the expression of both OPG and RANKL

to different extent, however, resulting in significant increase of the OPG/RANKL ratio $(2.11 \pm 0.33 \text{ folds}; \text{Figure 15D})$. In addition to direct promoting osteoblastogenesis, this finding suggested that PTR-SeNPs (10 μ M) also exhibited indirect inhibition effect on osteoclastogenesis.



Figure 13. PTR-SeNPs induced osteoblastic differentiation in MC3T3-E1 cells. (A) ALP activity in MC3T3 cells pre-treated with different concentrations of PTR-SeNPs (0.1nM-10 μ M) in osteogenic medium for 7 days was determined by spectrophotometry at 410nm and normalized relative to the total cellular protein content. Normalized results were compared with that of the negative control group (1% deionized water; 1%DDI). (B) ALP containing MC3T3-E1 cells pre-treated with 10 μ M of PTR-SeNPs or 1% (v/v) DDI were visualized by ALP staining. Results are mean values of three determinations ± SEM. Means with "***" are significantly different (p < 0.001, one-way ANOVA or Student's *t* test).



Figure 14. PTR-SeNPs simulated bone mineral formation in MC3T3-E1 cells. Bone nodules formation in MC3T3 cells pre-treated with 10 μ M of PTR-SeNPs in osteogenic medium for 16 days was visualized by (A) Von Kossa staining and (B) Alizarin Red S staining. (C) Semi-quantification of Von Kossa staining was obtained by ImageJ. (D) Semi-quantification of Alizarin Red S staining was determined by spectrophotometry. Results are mean values of three determinations ± SEM. Means with "***" are significantly different (*p* < 0.001; one-way ANOVA or Student's *t* test).



Figure 15. PTR-SeNPs upregulated the major osteogenic markers in MC3T3-E1 cells during osteoblastic differentiation and mineralization. Gene expression of (A) ALP, (B) Runx2, (C) Osx, (D) OCN and (E) OPG/RANKL in MC3T3-E1 cells pre-treated with 10μM of PTR-SeNPs or 1% (v/v) DDI for 3 (Runx2), 7 (ALP & Osx), 11

(OPG/RANKL) and 16 days (OCN) was quantified by qPCR analysis and normalized to that of GAPDH. Results are mean values of three determinations \pm SEM. Means with "*" or "***" are significantly different (p < 0.05 and p < 0.001; Student's *t* test).

3.5. <u>BMP-2 Signaling Cascades Modulated by PTR-SeNPs in the MC3T3 E-1</u>

<u>Cells During Osteoblast Differentiation</u>

evidences Substantial have demonstrated that BMP-2 signaling via Smad-dependent and/or Smad-independent pathway(s) is one the most important signaling cascades in regulating osteoblastogenesis ³⁶⁻³⁸. In Smad-dependent pathway, upon cell surface ligand binding, BMP-2 forms a hetero-tetrameric receptor complex with the dimers of both type II and type I transmembrane serine/threonine kinases, triggering phosphorylation of Smad1/5/8 in the cytoplasm. Phosphorylated Smad1/5/8 then forms a trimetric complex with Smad4 followed by translocation to nucleus, where they regulate the transcription of target genes for osteoblast differentiation. In particular, homeobox gene Dlx5, which is an upstream target of both BMP-2/Smad dependent and BMP-2/Smad independent pathways, plays a pivotal role in stimulating the downstream transcription factors such as Runx2 and Osx³⁴. MAPKs (including p38, ERK and JNK) signaling, which are upstream regulators of Dlx5 in the BMP-2/Smad independent pathway, has been found to play important role in osteoblast differentiation, regulating the ALP, Runx2 and OCN expressions ³⁴. Interestingly, we discovered that PTR-SeNPs (10µM) not only significantly upregulated the gene expression of BMP-2 (1.64 \pm 0.24 folds) and Dx15 (1.54 \pm 0.06 folds) but also markedly increased the protein phosphorylation of Smad1/5/8 (1.31 \pm 0.06 folds) and

all MAPKs such as p38 (2.86 \pm 0.58 folds), ERK (3.67 \pm 1.48 folds) and JNK (1.36 \pm 0.07 folds) (**Figure 19**). The involvement of BMP-2/Smad dependent pathway in SeNPs (functionalized by ruthenium and citrate) mediated osteoblast differentiation has been reported recently ⁸⁴. Surprisingly, our findings suggest that PTR-SeNPs could trigger both BMP-2/Smad-dependent and Smad-independent pathways in the MC3T3-E1 cells during osteoblast differentiation. The proposed BMP-2 signaling cascades modulated by PTR-SeNPs in the MC3T3-E1 cells during osteoblast differentiation, the specific role of PTR-SeNPs in regulating osteoblastogenesis of MC3T3 E-1 cells via BMP-2/Smad dependent and independent pathways is not clear at this stage and will be further discussed later.



Figure 16. PTR-SeNPs upregulated the BMP-2 and Dlx5 in MC3T3-E1 cells during osteoblastic differentiation. Gene expression of (A) BMP-2 and (B) Dlx5 in MC3T3-E1 cells pre-treated with 10 μ M of PTR-SeNPs or 1% (v/v) DDI for 3 (BMP-2) and 7 days (Dlx5) was quantified by qPCR analysis and normalized to that of GAPDH. Results are mean values of three determinations ± SEM. Means with "*" and "**" are significantly different (*p* < 0.05 and *p* < 0.01; Student's *t* test)



Figure 17. PTR-SeNPs induced the BMP-2/Smad dependent pathway in MC3T3-E1 cells. Protein expression and phosphorylation of Smad1/5/9 in MC3T3-E1 cells pre-treated with 10µM of PTR-SeNPs or 1% (v/v) DDI for 4 days was quantified by Western blot analysis and normalized to that of GAPDH. Semi-quantification of images was obtained by ImageJ. Results are mean values of three determinations \pm SEM. Means with "***" are significantly different (p < 0.001; Student's t test).



68 | P a g e



Figure 18. PTR-SeNPs induced the BMP-2/Smad independent pathway in MC3T3-E1 cells. Protein expression and phosphorylation of (A) p38, (B) ERK and (C) JNK in MC3T3-E1 cells pre-treated with 10 μ M of PTR-SeNPs or 1% (v/v) DDI for 4 days was quantified by Western blot analysis and normalized to that of GAPDH. Semi-quantification of images was obtained by ImageJ. Results are mean values of three determinations ± SEM. Means with "*" and "**" are significantly different (*p* < 0.05 and *p* < 0.001; Student's *t* test).



Figure 19. Proposed BMP-2 signaling mediated by PTR-SeNPs in MC3T3-E1 cells during osteoblast differentiation

3.6. <u>Cellular Uptake Behavior of PTR-SeNPs by the MC3T3-E1 Cells</u>

Endocytosis has been widely reported as the major cellular uptake mechanism for nanoparticles ⁸⁵. Our preliminary study found that after cellular internalization, coumarin-6 labelled PTR-SeNPs (120μM) co-localized with lysosomes in the MCF-7 human breast carcinoma cells around 10 min ⁸². Similarly, as shown in **Figure 20**, the overlay of the blue (nucleus), green (PTR-SeNPs) and red (lysosomes) fluorescence clearly indicated the co-localization of PTR-SeNPs (10μM) and lysosomes in the MC3T3-E1 cells as early as 5 min, suggesting that lysosome is the main target organelle of PTR-SeNPs. Subsequently, some PTR-SeNPs were found to escape from lysosomes and released into cytosol from 6 min onwards. It is the first study of its kind to report the intracellular localization of SeNPs by MC3T3-E1 cells via endocytosis.



Figure 20. PTR-SeNPs co-localized with lysosomes in the MC3T3-E1 cells after cellular internalization. Cells were pre-treated with Hoechst 33342 (blue; nucleus), LysoTracker Deep Red (Red; lysosomes) and courmarin-6-loaded PTR-SeNPs (10μM; green) at 37 °C for different period of time followed by visualization under fluorescence microscope (Scale bar: 250μm).

3.7. Stimulatory Effect of PTR-SeNPs on Bone Formation Using Medaka

<u>Larvae</u>

Japanese medaka (*Oryzias latipes*) is a teleost fish, which has been found to possess a very high similarity in bone structure and metabolism to those of mammals ⁹¹⁻⁹³. Because of having a similar estrogen profile to humans, medaka has been recently suggested to serve as a non-mammalian animal model to study sex and age-related bone metabolism *in vivo* ⁴. In this study, we have successfully established a brand new *in vivo* model to investigate the simulating effect of PTR-SeNPs on bone formation using medaka larvae. By comparing the mineralization level (area and intensity of purple-red color) at different regions of vertebrae column (such as anal fin, dorsal fin and caudal fin), PTR-SeNPs (10ppm) was found to significantly enhance the vertebrate development of medaka larvae after feeding for 11 days as visualized by Alizarin Red S (bone) and Alcian Blue (cartilage) double staining (

Figure 21)

Expression of mammalian osteogenic markers such as ALP, Runx2, Osx and OPG/RANKL have been previously found in medaka fish during their bone developmental stage $^{94-96, 111}$. Similarly, the gene expression of ALP (1.53 ± 0.1 folds), Runx2 (1.45 ± 0.10 folds), Osx (1.41 ± 0.15 folds) as well as OPG/RANKL (4.38 ± 1.31 folds) in the medaka larvae fed with PTR-SeNPs based diet (10ppm) for 2 days were all significantly up-regulated (**Figure 22**). In consistent with the findings of our *in vitro* study, PTR-SeNPs (10ppm) not only promoted osteoblastogenesis, but also indirectly inhibited osteoclastogenesis, significantly enhancing the vertebrae development in medaka fish larvae.

Control diet



<u>PTR-SeNPs</u> based diet (10 ppm)





Figure 21. PTR-SeNPs enhanced the vertebrae development of medaka larvae. Medaka larvae were cultured in 6-well cell-culture plates (one medaka larvae per well) and fed either the control or PTR-SeNPs based diet (10ppm) twice daily for 11 days. Bone (pink) and cartilage (blue) were double-stained by Alizarin Red S and Alcian Blue, respectively. Images were captured by Nikon Elicpse Ci-E upright microscope.



Figure 22. PTR-SeNPs upregulated the major osteogenic markers in medaka larvae during vertebrae development. Gene expression of (A) ALP, (B) Runx2, (C) Osx and (D) OPG/RANKL in medaka larvae (n = 3) fed with 10ppm of PTR-SeNPs based diet or control diet for 2 days was quantified by qPCR analysis and normalized to that of 18S. Results are mean values of three determinations \pm SEM. Means with "*" are significantly different (p < 0.05; Student's *t* test).

3.8. <u>Comparison between PTR-SeNPs and Other SeNPs</u>

Table 8 briefly summarized different types of SeNPs reported in previous literatures. Interestingly, SeNPs fabricated with different stabilizers showed different physcio-chemical properties and health promoting functions. Further comparison with the osteogenic Ru@Se also indicated that PTR-SeNPs did not only possess a remarkably smaller average particle size, but also exhibited a significantly higher stability and cellular uptake efficiency (

Table 9). More importantly, in contrast to the Ru@Se, the major component of PTR-SeNPs is edible mushroom PSP instead of highly toxic and carcinogenic Ru(II) complex, suggesting its high potent to be developed as a novel agent for bone health.

Table 8. Brief summary of different types of SeNPs

Particles	Preparation	Stabilizer	Average	Effect
	method		particles size	
SeNPs fabricated with Undarin pinnatifida	Redox	Polysaccharide from	59nm	Induced apoptosis of A375 human
polysaccharide ⁷⁸		Undarin pinnatifida		melanoma cells
SeNPs stabilized by Pseudomonas	Biosynthesis	Pseudomonas	200nm	Unknown
alcapliphilia ⁷⁴		alcaliphilia		
Transferrin-chitosan-	Redox	Transferrin	130nm	Enhanced cellular uptake of doxorubicin
doxorubicin-SeNPs 112		Chitosan		and induced apoptosis of breast cancer
				cells
Nano-Se ¹¹³	Redox	Bovine serum	36nm	Chemoprevention via increasing
		albumin		glutathione-S-transferase activity
Ruthenium (II) functional SeNPs (Ru@Se)	Redox	Ruthenium (II)	170nm	Enhanced osteoblastic differentiation in
84		complex		stem cell via BMP signaling pathway

 Table 9. Comparison between PTR-SeNPs and Ru@Se

	PTR-SeNPs	Ru@Se ⁸⁴	
Composition of SeNPs	PTR-PSP, Se	Ruthenium, Se	
Average particles size	< 100nm	> 100nm	
Stability	> 3 months (well-dispensed spherical particles in water)	Unknown; (spherical particles aggregated in water)	
Safety	PTR-PSP: edible mushroom	Ru: highly toxic and carcinogenic metal	
Cell line	Preosteoblasts	Stem cells	
Mechanism	Via both Smad dependent and independent pathways	Via Smad dependent pathway only	
Cellular uptake efficiency	5 min	48 hrs	
In vivo study	Medaka larvae	Unknown	

3.9. Comparison between PTR-SeNPs and Other Osteogenic Nanoparticles

Table 10 briefly summarized different osteogenic nanoparticles and their possible mechanism of actions. Similar to the silica-based nanoparticles ¹¹⁴, PTR-SeNPs were found to enter the MC3T3-E1 cells via endocytosis followed by activating the MAPK pathways (including ERK1/2) and stimulating osteoblast differentiation and mineralization. Thus, it would be very interesting to find out whether PTR-SeNPs mediated osteoblastogenesis is related to autophagy like the silica-based nanoparticles in the MC3T3-E1 cells ¹¹⁴. As proposed in Figure 23, PTR-SeNPs might enter the MC3T3-E1 cells through endocytosis followed by stimulating a MAPK pathway dependent autophagy and upregulating the expression of major osteogenic marker genes. Besides, the increased transcription of BMP-2 would further react with BMP-2 receptor to activate the BMP-2/Smad dependent pathway, hereby collectively promoting the osteoblastogenesis.

Nanoparticles	Composition	Safety	Average particles size	Cell line	Mechanism of action
Calcium phosphate nanoparticles	Na ₂ HPO ₄ , CaCl _{2,} CTAB	CTAB: toxic	20nm	Bone marrow mesenchymal stem cells	Unknown
Hydroxyapatite-coated iron oxide nanoparticles ^{116,} ¹¹⁷	Fe ₃ O _{4.} Hydroxyapatite	Fe ₃ O ₄ : inedible	60nm long, 20nm wide	hFOB 1.1 human osteoblast-like cells	Enhanced fibronectin adsorption
Zinc oxide nanoparticles	ZnO	Safe	30nm	Mesenchymal stem cells	Unknown
Silica-based nanoparticles	SiO ₂	inedible	50nm	MC3T3-E1 cells	Stimulated osteoblastic differentiation and mineralization through activating the ERK1/2 pathway dependent autophagy
Gold nanoparticles	Au, Sodium citrate	Safe	20nm	Mesenchymal stem cells	Promoted osteogenic differentiation via p38 pathway by causing mechanical stress

 Table 10. Brief summary of different osteogenic nanoparticles



Figure 23. Possible mechanism of PTR-SeNPs stimulated osteoblastogenesis via BMP-2 signaling cascade

4. <u>Conclusions</u>

4.1. <u>Conclusions</u>

By using the mushroom polysaccharide-protein complex (PSP) isolated from *Pleurotus tuber-regium*, we have successfully prepared novel SeNPs (PTR-SeNPs) under a simple, food grade redox system. PTR-SeNPs existed as well-dispensed spherical particles in water with an average diameter of 91.3 \pm 1.53 nm, possessing a polycrystalline structure with high level of Se (81.2%). X-ray Diffraction (XRD) is a versatile technique that can investigate more degree of diffraction than conventional SAED-TEM ¹²¹ and is commonly used in crystallography analysis for studying lattice structure and orientation of inorganic nano-compounds ^{122, 123}. Further characterization using XRD would allow providing a clearer crystal structure of the PTR-SeNPs.

Substantial previous findings have demonstrated that promoting bone formation is one of the effective strategies to prevent and/or manage postmenopausal osteoporosis. Interestingly, our study indicated that PTR-SeNPs (10µM) not only promoted osteoblastogenesis (including cell proliferation, osteogenic differentiation and bone mineral formation), but also indirectly inhibited osteoclastogenesis in the murine preosteoblast MC3T3-E1 cells mediated by both BMP-2/Smad dependent and independent pathways. Thus, it would be interesting to further characterize their direct effects on osteogenesis using different cell stages of human osteoblasts such as human bone marrow-derived mesenchymal stem cells and hFOB 1.19 human fetal osteoblast cells.

Endocytosis has been widely reported as the major cellular uptake mechanism for nanoparticles. PTR-SeNPs (10µM) was found to co-localize with the lysosomes in the MC3T3-E1 cells as early as 5 min after cellular internalization. It is the first study of its kind to report the intracellular localization of SeNPs by MC3T3-E1 cells via endocytosis. Further elucidation of the molecular mechanism of PTR-SeNPs induced endocytosis is highly valuable.

Last but not least, we discovered that PTR-SeNPs (10ppm) could significantly enhance the vertebrate development (both bone and cartilage) of medaka larvae after feeding for 11 days together with up-regulating of those major osteogenic markers in bone metabolism. In order to develop medaka larvae as a brand new *in vivo* model for bone research, it is crucial to further determine the effect of PTR-SeNPs on the medaka fish at different life stages in terms of bone mineral density and micro-architecture using microCT as well as gene and protein levels of major osteogenic markers (both osteoblasts and osteoclasts) using qPCR and Western blotting. Besides, in order to understand the possibility of over mineralization in the PTR-SeNPs fed medaka fish, the bone mineral density and calcification level in medaka fish at different life stages after feeding with very high concentration of PTR-SeNPs based diet (e.g. 50ppm) should be comprehensively investigated. Furthermore, verification and validation of the findings of fish model by using traditional mice or rats (mammals) are also essential.

Findings of this study could provide significant insights into the *in vitro* and *in vivo* effects of PTR-SeNPs on bone formation. Besides, medaka larvae, as a novel non-mammalian animal model for studying bone formation, could be further developed into a rapid and economical *in vivo* screening platform prior to conducting traditional animal study using mice or rats, which is relatively expensive, time-consuming, labor intensive and required plenty of space. Our long-term goal is to develop PTR-SeNPs into an evidence-based bone-forming agent for promoting the bone health of postmenopausal patients in local community.
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