

Supplementary Table 7. Function of the top 16 hub genes of module 1 in bone metabolism.

| Gene symbol | Full name | Function |
|----------------|---|---|
| <i>PRSS23</i> | <i>serine protease 23</i> | NA |
| <i>TNC</i> | <i>tenascin C</i> | An extracellular matrix glycoprotein involved in osteogenesis and bone mineralization [1]. |
| <i>FSTL1</i> | <i>follistatin like 1</i> | Promotes chondrocyte apoptosis [2] and osteoclast formation [3]. |
| <i>FBN1</i> | <i>fibrillin 1</i> | Limits osteoclast formation and function [4]; a negative regulators of bone resorption [5]. |
| <i>APOE</i> | <i>apolipoprotein E</i> | Plays crucial roles in maintaining bone mass by promoting osteoblast differentiation and suppressing osteoclast differentiation [6]. |
| <i>LGALS1</i> | <i>galectin 1</i> | Relates to osteoblast maturation [7], and plays a role in cell-cell and cell-matrix interactions of osteoblastic cells [8]. |
| <i>SPARCL1</i> | <i>SPARC like 1</i> | An extracellular matrix remodel gene [9]; a member of the osteonectin family of proteins [10]; suppresses osteosarcoma metastasis [11]. |
| <i>IGFBP4</i> | <i>insulin like growth factor binding protein 4</i> | Highly expressed in adipocytes and osteoblasts [12]; regulates bone metabolism [13–15]. |
| <i>MXRA8</i> | <i>matrix remodeling associated protein 8</i> | NA |
| <i>MFGE8</i> | <i>milk fat globule EGF and factor V/VIII domain containing</i> | Regulates osteoclast homeostasis and inflammatory bone loss [16]. |
| <i>GAS6</i> | <i>growth arrest specific 6</i> | Enhances the bone resorbing activity of mature osteoclasts [17]; induces osteoclast differentiation [18]. |
| <i>TIMP1</i> | <i>TIMP metallopeptidase inhibitor 1</i> | Inhibits the activity of MMPs and then regulate the degradation of bone extracellular matrix molecules [19]. |
| <i>IGFBP7</i> | <i>insulin like growth factor binding protein 7</i> | Inhibits osteoclastogenesis and osteoclast activity [20]; enhanced osteogenic differentiation of BM-MSCs <i>in vitro</i> and promoted new bone formation <i>in vivo</i> [21]. |
| <i>CYR61</i> | <i>cysteine-rich protein 61</i> | Modulates mature osteoblast and osteocyte function to regulate bone mass [22]; stimulates proliferation and differentiation of osteoblasts <i>in vitro</i> and contribute to bone remodeling <i>in vivo</i> in myeloma bone disease [23]; regulates adipocyte differentiation from mesenchymal stem cells [24]. |
| <i>CP</i> | <i>ceruloplasmin</i> | Inhibits osteoblast activity, mineralization [25, 26]. |
| <i>IGFBP5</i> | <i>insulin like growth factor binding protein 5</i> | The IGFBP5 produced by osteoblasts stimulates osteoclastogenesis and bone resorption, and as an osteoblast-osteoclast coupling factor [27]. |

SPARC: secreted protein acidic and cysteine rich. TIMP: tissue inhibitor of metalloproteases; MMPs: matrix metalloproteinases; BM-MSCs: bone marrow-derived mesenchymal stem cells.

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