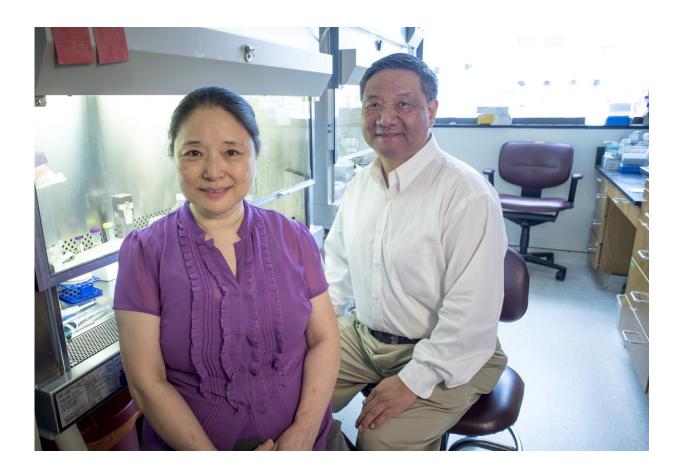


Researchers describe mechanism that underlies age-associated bone loss

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Wei Chen, left, and Yi-Ping Li. Credit: UAB

A major health problem in older people is age-associated osteoporosis—the thinning of bone and the loss of bone density that increases the risk of fractures. Often this is accompanied by an increase



in fat cells in the bone marrow.

University of Alabama at Birmingham researchers have now detailed an underlying mechanism leading to that osteoporosis. When this mechanism malfunctions, <u>progenitor cells</u> stop creating <u>bone</u>-producing <u>cells</u>, and instead create fat cells. Knowledge of this mechanism can provide targets in the search for novel bone-loss therapeutics to treat human osteoporosis with minimal side effects.

The UAB researchers found that a protein called Cbf-beta plays a critical role in maintaining the bone-producing cells. Furthermore, examination of aged mice showed dramatically reduced levels of Cbf-beta in bone marrow cells, as compared to younger mice.

Thus, they propose, maintaining Cbf-beta may be essential to preventing human age-associated osteoporosis that is due to elevated creation of fat cells.

Bone is a living tissue that constantly rebuilds. Bones need a constant new creation of cells specific to their tissue, including the boneproducing cells called osteoblasts. Osteoblasts live only about three months and do not divide.

The progenitor cells for osteoblasts are bone marrow mesenchymal stem cells. Besides osteoblasts, mesenchymal stem cells can also differentiate into the chondrocyte cells that make cartilage, the myocyte cells that help form muscles and the adipocytes, or fat cells. Thus, the same progenitor cell has four possible tracks of differentiation.

UAB researchers and colleagues focused on the molecular mechanism that controls the lineage commitment switch between the osteoblast and adipocyte tracks. Led by Yi-Ping Li, Ph.D., UAB professor of pathology, and Wei Chen, M.D., UAB associate professor of pathology,



they investigated the key role played by Cbf-beta, or core-binding factor subunit beta.

Study details

The team led by Li and Chen generated three mouse models by deleting Cbf-beta at various stages of the osteoblast lineage. All three mouse models showed severe osteoporosis with accumulation of <u>fat cells</u> in the bone marrow, a pathology that resembles aged bone from enhanced adipocyte creation.

Bone marrow <u>mesenchymal stem cells</u> and <u>bone cells</u> from the skulls of Cbf-beta-deficient mice showed increased expression of adipocyte genes.

Looking at the mechanism downstream, the researchers found that the loss of Cbf-beta impeded the canonical Wnt signaling pathway, particularly through decreased Wnt10b expression. In nonmutant mice, they found that the protein complex composed of Cbf-beta and the Runx2 transcription factor binds to the Wnt10b promoter to drive Wnt10b expression. The Cbf-beta/Runx2 complex also inhibited expression of the enhancer protein C/EBP-alpha that promotes differentiation of adipocytes.

In addition, the researchers showed that Cbf-beta maintains the osteoblast lineage commitment in two ways—through the Wnt paracrine pathway to affect nearby cells and through endogenous signaling within the cell to suppress adipogenesis gene expression.

Altogether, this knowledge of the mechanism driven by Cbf-beta can help explain the imbalance in bone maintenance seen in older people.

More information: Mengrui Wu et al, Cbfβ governs



osteoblast–adipocyte lineage commitment through enhancing β-catenin signaling and suppressing adipogenesis gene expression, *Proceedings of the National Academy of Sciences* (2017). DOI: 10.1073/pnas.1619294114

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