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## Bisphosphonates pathway

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### Keywords

alendronate; bisphosphonate; bone diseases; bone mineral density; bone resorption; farnesyl pyrophosphate synthase; mevalonate pathway; pharmacogenomics; osteoporosis; riseridronate

### Description

Bisphosphonates (BPs) are potent inhibitors of osteoclast-mediated bone resorption. They are widely used in the management of osteoporosis and other diseases of high bone turnover [1]. BPs' pronounced affinity for bone, but not other tissues, makes them the ideal candidates for treatment of bone diseases. BPs have been shown to increase bone mineral density (BMD), reduce bone turnover markers, and reduce the risk of osteoporotic fractures. They are currently the most important and effective class of drugs for metabolic bone disorders such as osteoporosis, Paget's disease, and osteogenesis imperfecta [1–3]. Intravenous BPs are also used in the treatment of malignant hypercalcemia, osteogenesis imperfecta, and prevention of skeletal-related events in patients with multiple myeloma or breast and prostate cancers [3–7].

Chemically, BPs are synthetic analogs of pyrophosphate, an endogenous regulator of bone mineralization that contains a nonhydrolysable P-C-P backbone with two side chains (R<sup>1</sup> and R<sup>2</sup>) [8]. The two phosphonate groups are required for both binding to bone mineral and for antiresorptive potency. Modification to one or both phosphonate groups significantly reduced the binding affinity as well as the antiresorptive potency of BPs [9]. The R<sup>1</sup> and R<sup>2</sup> side chains are responsible for a wide range of activities. Acting together with the two phosphonate groups, the presence of a hydroxyl group (-OH) or an amino group (-NH<sub>2</sub>) rather than an H group in the R<sup>1</sup> chain enhances binding to calcium minerals. The presence of a nitrogen or amino group in the R<sup>2</sup> side chain significantly increases the antiresorptive potency of BPs and also affects binding to hydroxyapatite [9–11].

BPs can be broadly classified into two major classes with distinct mechanisms of action: the nonnitrogen containing class and the nitrogen-containing class. The earlier nonnitrogen containing BPs (e.g. clodronate, tiludronate, and etidronate) act by incorporation into ATP, whereas the newer, more potent nitrogen-containing BPs (N-BPs, e.g. pamidronate, alendronate, ibandronate, riseridronate, and zoledronate) act by inhibiting farnesyl diphosphate synthase (FDPS) in the mevalonate pathway [12].

## Pharmacokinetics

Pharmacokinetic studies showed that all BPs have high affinity for bone mineral and poor intestinal absorption [13]. The amount of BPs taken up by the skeleton depends on several factors including renal function, rate of bone turnover and on the affinity for bone mineral. The skeleton has a high capacity to retain BPs and the major route of elimination is through kidney [8]. Renal transporters for BPs have not been elucidated. Potential candidate transporters for BPs are the sulfate transporters and phosphate transporters that belong to SLC22A, SLC17A, and SLC13A families [14,15].

The two classes of BPs are metabolized differently. The simple, nonnitrogen-containing BPs are metabolized to cytotoxic, nonhydrolysable analogs of ATP [16,17]. Accumulation of these toxic byproducts interferes with mitochondrial function and ultimately leads to apoptosis of osteoclasts. However, the more potent nitrogen-containing BPs are not metabolized and are excreted unchanged through the kidney. BPs are typically given when fasting, as food reduces the bioavailability of oral nitrogen-containing BPs [18].

## Pharmacodynamics

BPs preferentially bind to the surface of bone at sites of active remodeling and become internalized into osteoclasts through endocytosis [19]. Two distinct mechanisms of action have been revealed. The nonnitrogen containing BPs inhibit bone resorption by generating the cytotoxic analogs of ATP which interfere with mitochondrial function and induce apoptosis of osteoclasts [16,17]. In contrast, the more potent nitrogen-containing BPs bind to and inhibit key enzymes of the intracellular mevalonate pathway, thereby preventing the prenylation and activation of small GTPases that are essential for the bone-resorbing activity and survival of osteoclasts (Fig. 1, BP pathway, <https://www.pharmgkb.org/do/serve?objId=PA154423660&objCls=Pathway>) [1,20–22]. The mevalonate pathway is a biosynthetic pathway responsible for production of cholesterol, other sterols and isoprenoid lipids [23]. Some of these isoprenoid lipids (e.g. farnesyl pyrophosphate and geranyl-geranyl pyrophosphate) are essential for the prenylation and activation of the small GTPases such as Ras, Rho, Rac, Rab, and Cdc42 [23–25]. The small GTPases are important signaling proteins regulating osteoclast morphology, cytoskeleton arrangement, membrane ruffling, trafficking, and cell survival [26,27]. Inhibition of enzymes along the mevalonate pathway may impair the prenylation process and lead to loss of function of the small GTPases. Several enzymes along the mevalonate pathway have been studied as potential molecular targets for nitrogen-containing BPs. Earlier studies showed that incadronate and ibandronate, but not other BPs, are inhibitors for squalene synthase (FDFT1), an enzyme required for cholesterol biosynthesis in the mevalonate pathway. Inhibition of FDFT1, however, does not lead to loss of protein prenylation [28,29], suggesting that other enzymes upstream in the pathway may be inhibited by N-BPs. The main target protein of the N-BPs is currently considered to be FDPS, a key regulatory enzyme catalyzing the production of isoprenoid lipids. Multiple studies demonstrated that FDPS is inhibited by all the N-BPs and the antiresorptive potency of various N-BPs correlates with their ability to inhibit FDPS [23–25]. X-ray crystallography study also confirmed that the potent N-BPs (risedronate and zoledronate respectively) bind and inhibit the active site of FDPS through the nitrogen atoms [30]. Inhibition of FDPS by N-BP blocks the synthesis of farnesyl pyrophosphate and geranyl-geranyl pyrophosphate, which in turn prevents prenylation of small GTPases and disrupts normal osteoclast function. These studies clearly indicate that inhibition of FDPS is a central mechanism by which N-BPs inhibit bone resorption.

Bisphosphonates are currently the most widely used and effective antiresorptive therapy for osteoporosis. They are well tolerated by most; however, the efficacy and safety of BPs vary among patients [31]. It is estimated that around 5–10% patients fail to respond to BP therapy [32,33]. In addition, some patients with intravenous BP treatment experienced adverse events such as atrial fibrillation [34,35], acute phase response [36–38], renal insufficiency, and osteonecrosis of the jaw (ONJ) [39,40]. Despite the large amount of genetic evidence elucidated for the susceptibility to osteoporosis, the understanding about genetic factors contributing to osteoporosis treatment response is still limited. The majority of published pharmacogenetic studies for BP response to date have focused on candidate genes related to BMD, bone turnover and osteoporotic fracture risk [8,41–44]. These include the vitamin D receptor gene (*VDR*), the estrogen receptor beta gene (*ESR2*), the type 1 collagen gene (*COL1A1* and *COL1A2*), and the low-density lipoprotein receptor-related protein 5 gene (*LRP5*). Common polymorphic variation in the *VDR* gene (*BsmI* genotype, *rs1544410*) has been reported to contribute to individual response to etidronate [45] and alendronate [46,47] in Caucasian postmenopausal women, with the *bb* genotype associated with lower increase in BMD after BP treatment. It has also been observed that the *BsmI* (*rs1544410*) and *TaqI* (*rs731236*) polymorphisms of *VDR* and – 511 C/T polymorphism (*rs16944*) of *IL1B* were significantly associated with acquired resistance to BP treatment in Caucasian patients with Paget's disease of the bone [48,49]. The *COL1A1* Sp1 polymorphism (*rs1800012*) has been associated with decrease in bone mass and osteoporotic fractures. The *SS* genotype of this polymorphism was associated with a better response to etidronate treatment in terms of femoral neck BMD but not lumbar spine BMD, indicating site-specific BMD response influenced by *COL1A1* genotype to cyclical etidronate therapy [50]. Variations in *ESR2* (*RsaI*) and *LRP5* (*rs3736228* and *rs4988321*) were also evaluated for their influence on response to alendronate and risedronate respectively, however, no association was found [51,52]. As BPs modulate the mevalonate pathway, polymorphisms in genes encoding target enzymes of the mevalonate pathway may modulate the response to N-BP treatments. Marini *et al.* [53] has demonstrated that a polymorphism in *FDPS*, a target of N-BPs, impacted the response to long-term N-BP treatment in Danish postmenopausal women. Patients with the *CC* genotype for *rs2297480* showed a decreased response of bone turnover markers to N-BP therapy. Moreover, a recent genome-wide association study identified a polymorphism of *CYP2C8* (*rs1934951*) to be associated with BP-related ONJ [54]. Individuals homozygous for the *T* allele showed an increased risk for developing ONJ. Results from all the studies described above highlight the possibility of using genetic testing to tailor BP treatments for optimal response. However, further population analysis involving larger patient cohorts, different ethnic backgrounds and examining all functional gene variants simultaneously is required to validate the existing findings and to fully elucidate the mechanisms underlying the variation in the treatment response to BPs.

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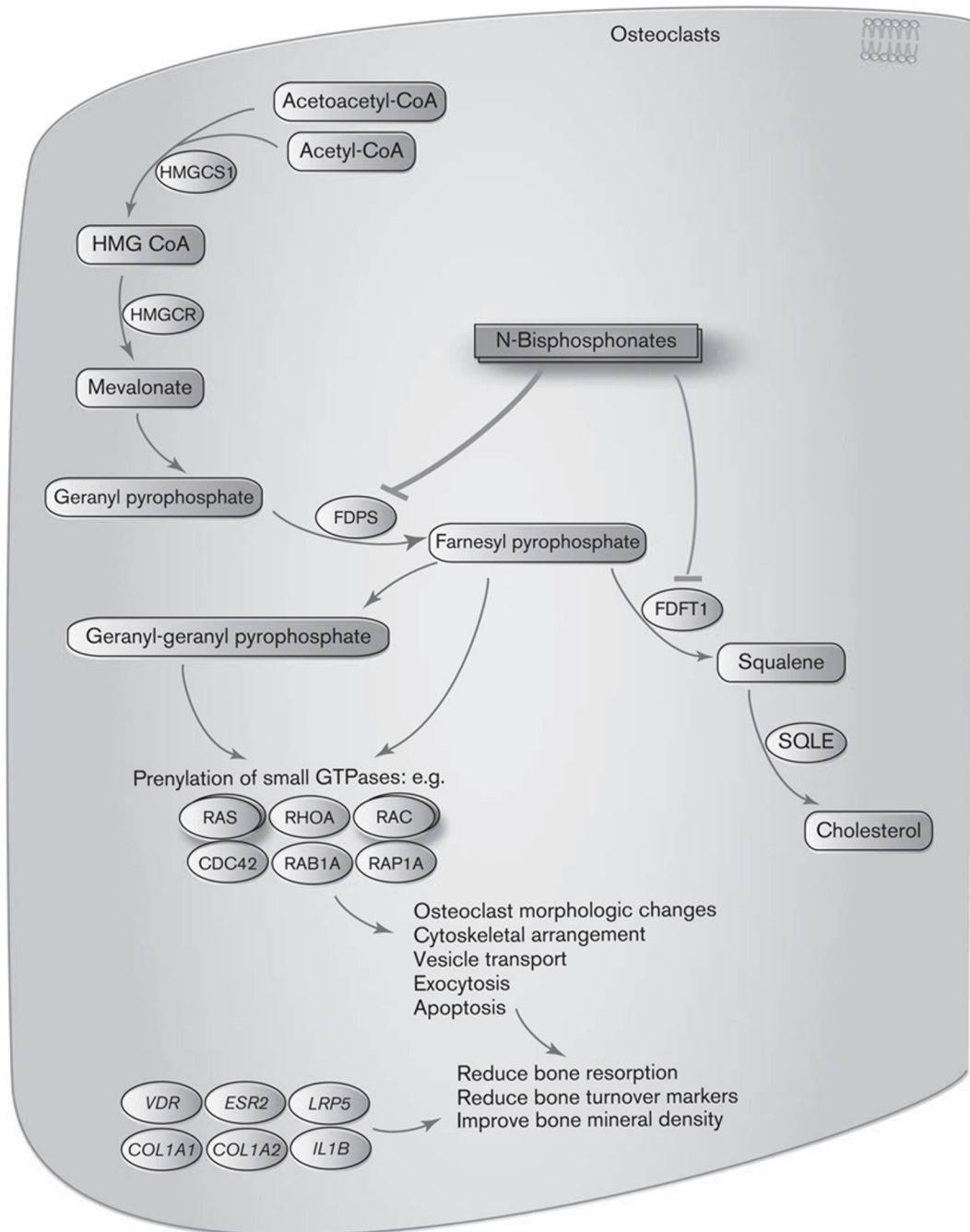
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**Fig. 1.**  
Bisphosphonates pathway: Mechanism of action of N-bisphosphonates in osteoclasts.