

Molecular mechanisms of bone resorption

An update

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The osteoclast is morphologically and functionally polarized (Figure 1), with a pole facing the bone matrix, where attachment occurs and towards which most of the secretion is targeted (the apical pole), and a pole facing the soft tissues in the local microenvironment (bone marrow or periosteum) and which provides mostly, but not exclusively, regulatory functions (the basolateral pole).

Attachment apparatus

The most striking and unique feature of the osteoclast actin cytoskeleton is at the site of cell contact with the substratum, where there is a prominent peripheral ring of F-actin which contains actin filaments oriented parallel to the plane of the underlying substrate around the cell periphery as well as numerous punctate structures where the actin filaments are organized in bundles perpendicular to the plane of the substratum, the podosomes. When osteoclasts are cultured on bone, the ring circumscribes the area of active bone resorption.

The punctate actin structures in the peripheral band, termed "podosomes" (Marchisio et al. 1984; Teti et al. 1991), apparently occur only in cells of monocytic origin (osteoclasts and monocytes) and in cells which have been transformed by the *src*, *fps* and *abl* oncogenes (Marchisio et al. 1987). In addition to the bundles of actin filaments, the podosomes contain a number of other proteins which have been reported to occur at sites of cell-substratum or cell-cell interaction (see Teti et al. 1991 for review). These include fimbrin, α -actinin and gelsolin, which are closely associated with the actin filaments in the core of the podosome, as well as vinculin and talin, which appear to form rosette structures surrounding the podosome cores (Marchisio et al. 1984, Teti et al. 1991). Furthermore, several tyrosine kinases and/or their substrates have also been localized to focal adhesions and to the sealing zone in osteoclasts.

Recent observations by Ali et al. (1984), Lakkakorpi et al. (1989; Lakkakorpi and Vaananen 1991) and Kanehisa et al. (Kanehisa and Heersche

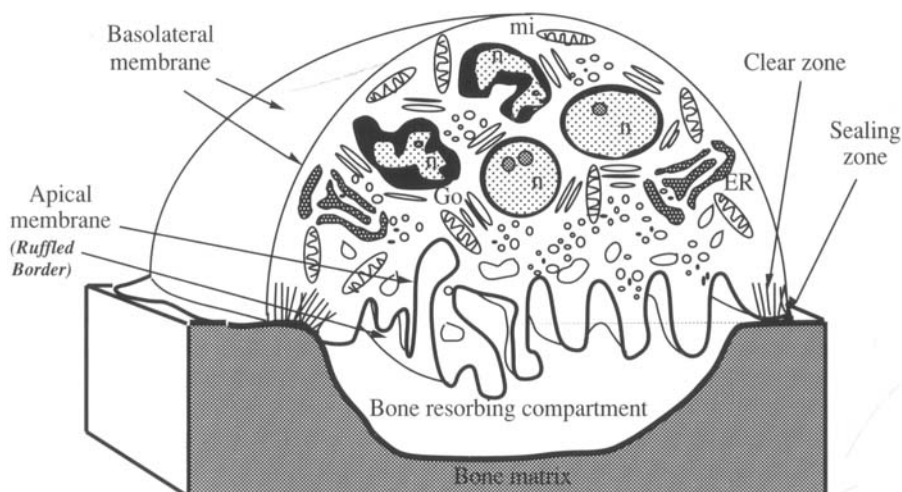


Figure 1. Functional organization and polarity of the osteoclast; n= nuclei; Go= Golgi complexes; mi= mitochondria; ER= endoplasmic reticulum.

1988, Kanehisa et al. 1990) provide us with a dynamic view of the attachment of the osteoclast to the bone matrix (Figure 2). In highly motile ("walking") osteoclasts, few podosomes are observed and seem to be confined to the irregularly shaped leading edge of the cell, or lamellipodium. Upon arrest and attachment, much more numerous podosomes are formed and are organized in a peripheral ring, as described in cells attached on glass. Rapidly thereafter, the seal is established: then the punctate "podosome" structures are either replaced by two concentric rings of vinculin and talin circumscribing a broad central zone of F-actin (Kanehisa et al. 1990, Lakkakorpi and Vaananen 1991) or reach such density that they cannot be individually observed when they arrange into two concentric rings. These observations therefore suggest a distinction in time and in specific cell-matrix interactions between the motile cell ("walking"), the cell recently arrested at a future resorbing site ("sitting"), forming a first ring-like structure of punctate attachment sites, and the resorbing cell, with a functionally tight sealing zone.

Transmembrane proteins of the integrin family of adhesion molecules, mediate cell-substratum interactions at the interface between the osteoclast and bone. Integrins are heterodimeric molecular assemblies of an α subunit and a β subunit with specific, receptor-like, extracellular binding sites which recognize the Arg-Gly-Asp (RGD) sequence, a motif known to represent the core ligand for all members of the integrin family (Ruoslahti and Pierschbacher 1987). The amino acid sequence surrounding the RGD motif determines which integrin will recognize and bind a specific matrix protein (Ruoslahti and Pierschbacher 1987, Horton and Davies, 1989).

Osteoclasts express multiple integrins ($\alpha\beta 3$, $\alpha 2\beta 1$ and $\alpha\beta 1$; Nesbitt et al. 1993) some of which are involved in osteoclast adhesion to the bone matrix. Several RGD-containing matrix proteins have been identified (Teti et al. 1991). Of these, collagen-type I, osteopontin and bone sialoprotein II are the most likely candidates to fill the role of integrin-binding proteins in bone (Teti et al. 1991). Most interestingly, data is now accumulating that suggests that the osteoclast synthesizes and secretes both osteopontin and BSP. II.

That the integrins play an important role in the osteoclast function and, despite the controversy regarding their presence at the sealing-zone membrane, most likely mediate the attachment of the osteoclast to the bone surface, is demonstrated by the fact that several RGD-containing proteins have been shown to inhibit bone resorption *in vitro* and *in vivo*.

The osteoclast secretes lysosomal enzymes as well as matrix-metalloproteinases in the bone resorbing compartment

The morphological polarity of the osteoclast is paralleled by a functional polarity, with the osteoclasts synthesizing and vectorially secreting several classes of enzymes towards the apical ruffled border (Baron et al. 1985, Baron et al. 1988). It is well established that these enzymes comprise acid phosphatases, aryl-sulfatases, beta-glucuronidase, beta-glycerophosphatase as well as several cysteine-proteinases (Baron et al. 1985, Baron et al. 1988, Vaes 1988), including cathepsins B, C, D, L and a novel cathepsin K, which are capable of fully digesting collagen at low pH. Recent evidence from several laboratories however suggests that osteoclasts also secrete metalloproteinases (collagenase type I and type IV, stromelysin) and that these are important in the process of bone resorption (4–7). Thus, it is the cooperative action of a set of enzymes with acidic and neutral optimal pH that leads to a complete degradation of the extracellular matrix at the resorbing site.

The apical ruffled-border membrane and the process of acidification

The apical membrane of the osteoclast, or ruffled-border, is directly involved in the molecular mechanisms of bone resorption. It is the target for the specific delivery of the newly synthesized secretory enzymes (Baron et al. 1985, 1988) and it is the site of proton extrusion for acidification of the bone resorbing compartment (Baron et al. 1985, Blair et al. 1989). The H^+ ATPase present in osteoclast membranes is of the Vacuolar-type, closely resembling the pumps present in lysosomes, coated vesicles or kidney membranes (Blair et al. 1989, Bekker and Gay 1990a, Vaananen et al. 1990).

There are however subtle differences in the properties and structure of mammalian V-ATPases (Nelson, 1991, Forgac 1989, Wang and Gluck 1990) and recent results from our laboratory indeed show that the osteoclast proton pumps, at least in chicken, differ in some respects from kidney vacuolar ATPases (Chatterjee et al. 1992). Briefly, proton transport by osteoclast-derived vesicles is more sensitive to inhibition by vanadate, nitrate and Tiludronate than similar preparations derived from the kidney in the same animals. Furthermore, cloning of the catalytic domain of the enzyme showed that the B subunit is of the ubiquitous type, different from the one expressed in kidney and that there are two isoforms of the catalytic subunit A, although their exact distribution in cells and tissues is not clearly elucidated.

The osteoclast proton pump may therefore have a

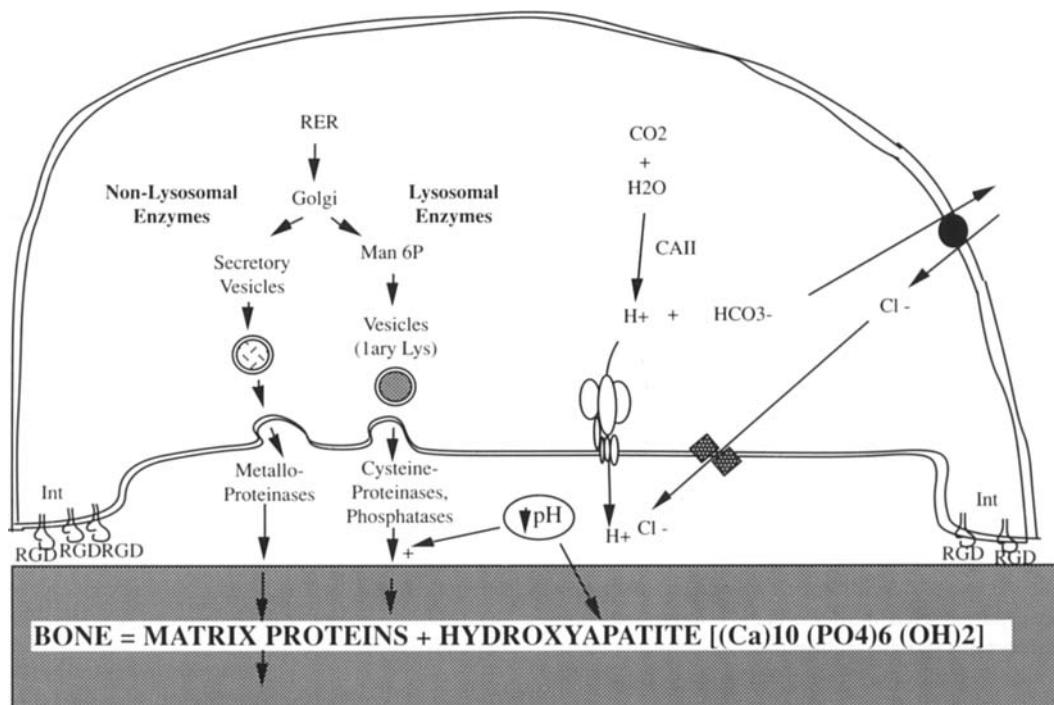


Figure 2. Schematic representation of the relationship of enzyme secretion, attachment and acidification in the mechanisms of bone resorption. Sealing at the periphery of the resorbing compartment is assured by binding of integrin receptors to RGD-containing extracellular proteins; cysteine-proteinases and metalloproteinases are secreted into this sealed-off compartment, possibly transported in two separate secretory vesicles; acidification is ensured by vacuolar-like proton ATPase in the ruffled-border membrane; the protons are generated intracellularly by carbonic anhydrase and the by-product of the reaction, bicarbonate, is secreted basolaterally in exchange for chloride by a band 3-like anion exchanger; the chloride is then transported into the resorbing compartment by chloride channels, thereby avoiding hyperpolarization of the osteoclast's membrane that would result from proton transport. The low pH favors the action of lysosomal enzymes and dissolves the hydroxyapatite, and the presence of metalloproteinases broadens the pH range at which resorption can occur.

unique pharmacology and the assembly of specific isoforms of two subunits in the catalytic portion of the enzyme may confer specificities to this ATPase.

In parallel with proton pumping, the plasma membrane of the osteoclast in the apical ruffled border domain also expresses chloride channels, measured as a chloride conductance, and the extrusion of this anion is absolutely required for the acidification process.

Basolateral ion transport systems are tightly coupled to the process of apical acidification

The apparently simple process of pumping protons across the ruffled border membrane to acidify the resorption lacuna imposes, in fact, a complex and demanding set of ionic requirements to the cell, essentially designed to maintain the electrochemical balance of the osteoclast during bone resorption. This ionic balancing act requires the coordinated activity of electrogenic ion pumps, ion channels and electro-neutral ion exchangers in order to maintain the cytoplasmic pH and the transmembrane electrical poten-

tial within narrow physiological ranges.

Both an acid extruder (the Na⁺/H⁺ exchanger) and an acid-loader (i.e. base-extruder, the HCO₃⁻/Cl⁻ exchanger) are present in the osteoclast (Teti et al. 1989, Ravesloot et al. 1995). The Na⁺/H⁺ exchanger is driven by a Na⁺ gradient established across the basolateral membrane by the activity of the sodium pumps (Baron et al. 1986).

The hyperpolarization resulting from the activities of the H⁺ pump and the Na⁺/K⁺ ATPase, to which the activity of the Ca²⁺ ATPase also contributes (Bekker and Gay 1990b), is alleviated by the activity of voltage-sensitive ion channels, which transport positively charged ions into the cell (K channels) and/or negatively charged ions out of the cell (Cl channels), thus discharging the electrical potential (Ravesloot et al. 1989b, Ravesloot et al. 1989a, Sims and Dixon 1989, Sims et al. 1991, Schoppa et al. 1990, Blair et al. 1991). Studies performed by our laboratory, mostly in kidney cells, suggest that the main mode of action of calcitonin may be via the regulation of several of these basolateral transporters.

The proto-oncogene c-src plays a critical functional role while the proto-oncogene c-fos is required for the differentiation of osteoclasts

The discovery that deletion of the genes encoding either *c-src* or *c-fos* lead to osteopetrosis, directly and unequivocally demonstrated that these proteins, one a cytoplasmic tyrosine kinase and the other a nuclear transcription factor, play critical and unique roles in bone resorption (Soriano et al. 1991, Grigoriadis et al. 1994).

The *c-fos* deletion has been demonstrated to result in the absence of osteoclast formation from hematopoietic precursors while macrophages differentiate normally and are even markedly increased (Grigoriadis et al. 1994), thereby demonstrating that *c-fos* is required for the branching from monocytes to osteoclasts but not to macrophages.

In contrast, *c-src* is playing its critical role within the osteoclast itself, i.e. the osteopetrotic phenotype in the *c-src* mutants is cell autonomous (Lowe et al. 1993). *C-Src* is expressed at high levels in osteoclasts, where it is associated with both the plasma membrane and intracellular organelles, but not in other bone cells (Horne et al. 1992).

Recent studies in our laboratory have indicated that *c-src* is involved in several critical signaling pathways in the osteoclast, downstream of integrin and CSF-1 receptors. Integrin binding leads to a rapid and transient wave of tyrosine phosphorylation in isolated osteoclasts. *c-src*-kinase gets phosphorylated in osteoclast-like cells. Such responses are blunted in *src*-deficient cells. In contrast, calcium signaling from integrins is not affected. Furthermore, *src* substrates are found in association with several intracellular vesicles in the osteoclast as well as with both the sealing zone and the ruffled border membrane. *c-src* may therefore play several roles in osteoclast function, one or more of which cannot be compensated when the gene for *c-src* is deleted.

Acknowledgements

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