Osteoclasts control sensory neurons axonal growth through epidermal growth factor

receptor signaling

Estrela Neto[1, 2, 3], Daniela M. Sousa[1, 2], Francisco Conceição [1, 2, 4], Luís Leitão[1, 2, 4], Cecília J. Alves[1, 2],

Inês S. Alencastre[1, 2], Jonathan West[5], Richard O. C. Oreffo[6] and Meriem Lamghari[1, 2, 4]*

[1]i3S - Instituto de Investigação e Inovação em Saúde, Universidade do Porto, Rua Alfredo Allen

280, 4200-135 Porto, Portugal

[2]INEB - Instituto de Engenharia Biomédica, Universidade do Porto, Rua Alfredo Allen 280, 4200-

135 Porto, Portugal

[3]FMUP - Faculdade de Medicina da Universidade do Porto, Alameda Prof. Hernâni Monteiro,

4200-319 Porto, Portugal

[4]ICBAS - Instituto de Ciências Biomédicas Abel Salazar, Universidade do Porto, Rua de Jorge

Viterbo Ferreira n.º 228, 4050-313 Porto, Portugal

[5]Institute for Life Sciences, University of Southampton, University of Southampton Highfield

Campus, Southampton SO17 1BJ, United Kingdom

[6]Centre for Human Development, Stem Cells and Regeneration, Institute of Developmental

Sciences, Tremona Rd, Southampton SO16 6YD, United Kingdom

*Correspondence:

Meriem Lamghari,

Rua Alfredo Allen 280, 4200-135 Porto, Portugal; Phone: +351 220408800; lamghari@ineb.up.pt

Abstract

Dense ectopic sprouting of nerve fibers was reported in several bone pathologies featuring high osteoclast number and/or activity. Osteoclasts play a critical role on nociceptors activation; however, their contribution to nerve fibers outgrowth is unknown. This study provides compelling evidence that osteoclastic lineage has an intrinsic capacity to promote axonal outgrowth of dorsal root ganglia (DRG), surpassing the neurotrophic potential of bone marrow stromal cells. Using microfluidic devices, we showed that osteoclast-secreted molecules induced nerve growth via local action on nerve terminals. Interestingly, the axonal outgrowth mediated by osteoclast is neurotrophin-independent but implicate epidermal growth factor receptor (EGFR)/ErbB2 signaling. Ligand search studies showed lack of EGFR agonists in osteoclast secretome, however, demonstrated an increase of endogenous EGF in DRG under osteoclast stimulation. Moreover, proteomic analysis revealed molecules able to trigger EGFR signaling to induce osteoclast-mediated axonal outgrowth, as fibronectin, low-density lipoprotein receptor-related protein 1 and periostin.

Keywords: osteoclasts; axonal growth; sensory neurons; secretome; epidermal growth factor receptor (EGFR)/ErbB2 signaling; microfluidics

Introduction

The presence of nerve fibers in the bone microenvironment has been directly implicated in bone homeostasis and bone regeneration [1–5]. Anatomical mapping of innervation during skeletal development shows co-localization of nerve fibers with a rich capillary network in areas with high osteogenic activity [6–10]. Moreover, intense sprouting of nerve fibers has been observed in injured or diseased bones [11–15]. In normal bone formation, the pattern of neuronal dynamics observed during development is typically repeated [7,16–18], highlighting the critical role of neuronal regulation. Furthermore, several denervation studies have revealed an impairment on fracture healing, emphasizing the impact of the peripheral nervous system on bone turnover [19–23].

Sensory innervation has attracted significant interest concerning bone formation. Beyond their role in bone nociception, evidence demonstrates that the lack of sensory nerves within the bone microenvironment significantly impairs bone mass accrual. Thus, sensory nerves play a key role in bone homeostasis [24,25].

Bone cells have been shown to actively respond to neural factors [26], and neural key molecules participate in bone turnover [27]. Nevertheless, it remains unclear how bone cells create or maintain a permissive environment for the presence and sprouting of nerve fibers within the skeleton microenvironment. The high neurotrophic potential of bone marrow cells is well known [28–35], although the role of osteoclasts in the modulation of nerve growth remains far from clear. Interestingly, in pathological conditions such as bone cancer where ectopic sprouting of nerve fibers in the bone marrow, mineralized bone, and periosteum occurs [11,12,14,36], osteoclasts play a major role in the activation of nociceptors. The matrix resorption in the bone compartment results in a decrease in the pH milieu as a consequence of active proton secretion into the extracellular space [37–39], stimulating the acid-sensing ions channels in nerve endings leading to pain. Notably, clinical data showed that osteoclast inhibitors significantly reduced cancer-associated bone pain [40,41].

While such studies provide support for the role of osteoclasts in the activation of nociceptors in diseased bone [42], the extent to which osteoclasts contribute to the changes in the innervation pattern in normal or diseased bone is unknown. The current study has examined the effects of the osteoclast on the growth of sensory nerve fibers. We demonstrate the paracrine effect of osteoclasts at different maturation stages on axonal outgrowth and have investigated the main signaling pathways and possible molecules involved in intercellular crosstalk.

Results

Osteoclast secretome promotes sensory neurons neurite outgrowth

Osteoclasts were differentiated from the hematopoietic lineage of bone marrow cells and the

phenotype confirmed by increased expression of tartrate-resistant acid phosphatase (TRAP),

cathepsin K (CTSK) and osteoclast associated immunoglobulin-like receptor (OscAR) (Figure 1A).

Immunostaining revealed a large population of multinucleated osteoclasts (Figure 1B) with the

low contribution of positive F4/80 macrophages present in the culture. Reactivity to TRAP also

confirmed the mature osteoclast phenotype (Figure 1C).

Figure 1: Characterization of osteoclasts differentiation *in vitro*. Osteoclasts were differentiated
from mouse bone marrow hematopoietic lineage. **A.** Differentiation markers (tartrate-resistant
acid phosphatase (TRAP), cathepsin K (CTSK) and osteoclast associated immunoglobulin-like
receptor (OscAR)) were evaluated by qPCR. Data represented as mean±S.E.M. **B.** Image of
multinucleated osteoclasts stained for F-actin (red), nuclei (blue) and macrophages stained with
F4/80 (green). **C.** TRAP staining of mature osteoclasts differentiated *in vitro*. Scale bar 100 μm.
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The secretome derived from osteoclast precursors (pre-OC) and mature osteoclasts (OC) populations was tested on explants of embryonic dorsal root ganglia (DRG) cultures (Figure 2A). Outgrowth from sensory neurons was used to assess the neurotrophic potential of differentiating osteoclasts. After treatment, axonal outgrowth was calculated using a Matlab-based algorithm [43]. DRG treated with osteoclastic lineage conditioned media (pre-OC and mature OC), showed significantly higher axonal outgrowth when compared to the neurobasal-nerve growth factor (neurobasal (+NGF)) supplemented control (Figure 2B-C). The mature osteoclast secretome demonstrated an approximate 3-fold stronger influence on sensory neurons growth than the secretome from bone marrow stromal cells (BMSC), previously demonstrated to have a positive neurotrophic potential [33,44]. Additionally, there was a notable effect on axonal outgrowth induced by the mature osteoclasts indicating an intrinsic capacity magnified with the maturation stage (Figure 2B-C). Explant DRG cultures were also treated with osteoclast differentiation medium (OCm) to rule out the effect of receptor activator of nuclear factor kappa-B ligand (RANKL) and macrophage colony-stimulating factor (M-CSF) present in the medium known to modulate axonal outgrowth [45,46]. No significant differences were observed when compared to neurobasal control media (Figure 2B).

Figure 2: **Effect of osteoclasts on axonal outgrowth of sensory neurons. A.** Schematic representation of experimental setup. Pre-osteoclast conditioned media (Pre-OC CM) and mature osteoclast conditioned media (OC CM) was collected and added to embryonic DRG culture. **B.** Representative images of DRG outgrowth after treatment (stained for βIII-tubulin, scale bar 500 μm). **C.** Automatic axonal outgrowth quantification of DRG treated with Neurobasal (+nerve growth factor (NGF)), bone marrow stromal cells conditioned medium (BMSC CM), Pre-OC CM, OC CM and osteoclast medium (OCm, dashed line). Data represented as box & whiskers (line=median, +=mean). Statistical analysis was performed *vs.* controls – Neurobasal and OCm (*) and between groups (#).

In physiological conditions, the cell soma of sensory neurons is confined to the DRG and only the

axonal terminals are present in the bone microenvironment. Therefore, microfluidic platforms

were used to recapitulate the physical separation of soma/axonal structures with the possibility

to exclusively stimulate the axonal compartment (Figure 3A). To understand the direct effect of

the soluble factors released by the mature osteoclasts in the activation of receptors present in

the axonal terminals, the secretome of mature osteoclasts was specifically applied to axon

compartment. Neurite outgrowth was measured with AxoFluidic, an algorithm designed to

quantify neurite projection within microfluidic platforms [47]. The Axofluidic calculates the

exponential fit in the axonal compartment given by the spatial dependence decay function

$f(x) = A\,e^{-x/\lambda}$, where χ represents the spatial variable and λ represents the scale of spatial

decay (associated with the length of the neurite). The secretome of mature osteoclasts

reproduced the effects observed with NGF stimulation through axonal terminals (Figure 3B-C).

These data demonstrate that local administration of osteoclast secretome on the axonal

terminals of sensory neurons induced neurite outgrowth. Similar to the effect of NGF, mediated

by the activation of TrkA receptors present in axonal growth cones, the effect of the osteoclast

secretome could be mediated through neurotrophins present in the conditioned media and their

corresponding receptors. Therefore, the neurotrophins present in the osteoclast secretome were

quantified.

Figure 3: Axonal outgrowth in microfluidic platforms. DRG were cultured in microfluidic devices to
isolate the axonal terminals from cell soma. **A.** Schematic representation of a DRG cultured on a
microfluidic platform (somal side). Osteoclast secretome was specifically added to the axonal side
of the chamber. **B.** Quantification of axonal projection in the axonal side using Axofluidic. λ
constant, retrieved from the exponential decay function, represents the distance of the neurites
projection. Data represented as scatter plot (mean±SD) **C.** Representative images of neurite

projection on axonal compartment (NB (+NGF) left and OC CM right). β III tubulin in black, scale
bar = 500 μm.

Osteoclast-induced axonal outgrowth is neurotrophin-independent

Neurotrophins are classified as promoters for neuronal survival and differentiation. Four
neurotrophins are known to be expressed in mammals: NGF, brain-derived neurotrophic factor
(BDNF), Neurotrophin-3 (NT-3) and NT-4 [48]. Therefore, neurotrophins in the osteoclast
secretome were measured to understand the contribution of these proteins in the crosstalk
between osteoclasts and sensory neurons.

NGF, BDNF, NT-3 and NT-4/5 were not detected in the mature osteoclast secretome, despite our
observation of higher neurite outgrowth on embryonic DRG explant cultures, while NGF and
BDNF were detected in the BMSC secretome (Figure 4), as previously described in other reports
[33,44]. Although previous studies by Togari *et al.* demonstrated the mRNA expression of NGF
and BDNF by osteoclasts, these were not detected by ELISA or Western Blot [49].

Figure 4: Analysis of neurotrophins in secretome. Quantification of NGF, BDNF, NT-3 and NT-4/5
on bone marrow stromal cells conditioned medium (BMSC CM), pre-osteoclast conditioned
medium (Pre-OC) and mature osteoclast conditioned medium (OC CM). NT-3 was not detected
in any condition (data not shown). The dashed line represents the limit of detection for the ELISA
assay. Data represented as scatter plot (mean±SD).

In our work, we performed the screening of receptor tyrosine kinases phosphorylation,
implicated in neuronal development, growth, survival and axonal regeneration [50]. Low levels of
TrkA, TrkB and TrkC activation were observed further confirming the low contribution of NGF,
BDNF, NT-3 and NT-4/5 neurotrophins.

Among the different phosphorylated receptors, epidermal growth factor receptor (EGFR), ErbB2,
ErbB3 and platelet-derived growth factor receptor alpha (PDGFRα) displayed raised activation
levels in the condition where high axonal outgrowth was observed (Figure 5A). Interestingly,

under osteoclast stimulation, there was an increased activation of the ErbB2 receptor in DRG

neurons when compared to treatment with BMSC (Figure 5B). ErbB2, an orphan receptor with no

characterized ligand, can be activated by spontaneous homodimer formation or by

heterodimerization with another ligand-bound or epidermal growth factor (EGF) family

transactivated receptor (Figure 5C) [51,52]. The high levels of EGFR phosphorylation indicate

heterodimerization of EGFR/ErbB2 leading to the pathway activation. These results are in

agreement with the literature where both EGFR and ErbB2 were shown to be increased in DRG

upon lesion, in the context of nerve repair [53]. Therefore, osteoclasts may promote axonal

outgrowth through EGFR family signaling, described to be involved in the neuronal repair.

Figure 5: Analysis of activated receptor tyrosine kinases in DRG cultures. A. The graphic represents
the mean spot intensity of the most phosphorylated receptors in embryonic DRG treated with
different secretomes, showing the major activation of two different families: epidermal-growth

factor receptors (EGFR), platelet-derived growth factor (PDGF) receptor alpha. **B.** Graphic
highlighting the phosphorylation levels of EGFR family. **C.** Schematic representation of possible
EGFR family dimers formation and the correspondent ligands (adapted by permission from
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EGFR/ErbB2 signaling is involved in osteoclast-mediated axonal outgrowth

Up to date, the effect of EGFR family activation on axonal outgrowth is not clear yet [54]. While

in vivo and *in vitro* studies showed that the inhibition of EGFR promotes axonal regeneration over

inhibitory environments [55–58], other studies reported that EGFR activation enhances neurite

outgrowth in DRG neurons [59,60].

To understand the contribution of EGFR and ErbB2 signaling in the osteoclast-induced axonal

outgrowth, two complementary approaches have been applied, one by receptor-mediated

inhibition using pharmacological inhibition and other by ligand searching.

Erlotinib is an EGFR inhibitor that reversibly binds to the intracellular tyrosine kinase domain of

the receptor. It is described to selectivity block the EGFR over ErbB2 but at high concentrations,

inhibits both receptor kinases [52,61]. Our results show that the neurotrophic effect of

osteoclasts was reduced in the presence of the higher tested Erlotinib concentration (100 μM)

(Figure 6A-B), suggesting that both EGFR/EGFR homodimers and EGFR/ErbB2 heterodimers might

contribute the osteoclast-mediated effect on axonal outgrowth.

To unravel the input of ErbB2 in osteoclast neurotrophic effect the monoclonal antibody

Trastuzumab® (also known as Herceptin®) was used against the extracellular domain of the

receptor [62]. Upon treatment with Trastuzumab®, no differences on the outgrowth area of the

sensory neurons were observed (Figure 6C-D), indicating that sole inhibition of ErbB2 signaling

fails to circumvent the osteoclast secretome stimuli. These results show that osteoclast-mediated

axonal outgrowth required both EGFR- and ErbB2 receptor-dependent pathways.

Figure 6: Pharmacological inhibition of EGFR and ErbB2. **A.** Representative images of DRG treated with different concentrations of Erlotinib (βIII tubulin in green and nuclei in blue, scale bar 500 μm). **B.** Quantification of axonal outgrowth of sensory neurons blocked with epidermal growth factor receptor (EGFR) inhibitor – Erlotinib. **C.** Quantification of axonal outgrowth of sensory neurons blocked with ErbB2 receptor inhibitor – Trastuzumab®. **D.** Representative images of DRG treated with different concentrations of Trastuzumab® (βIII tubulin in green and nuclei in blue, scale bar 500 μm). Data represented as box&whiskers (median).

Endogenous expression of EGF is triggered upon osteoclast stimulation

Among EGFR agonists, EGF was previously showed to mediate axonal extension in several studies [59,63,64]. Thus, EGF released by osteoclasts was quantified in their secretome, and endogenous EGF expression by stimulated DRG was performed by immunohistochemistry. The results showed that EGF was not present in osteoclast secretome at detectable concentrations (data not shown). However, immunocytochemistry analysis revealed spread localization of EGF positive staining in the DRG stimulated with osteoclast conditioned medium in comparison to the control DRG (neurobasal+NGF). EGF was expressed by cell soma and neurite projections of sensory neurons (Figure 7B'-D', white arrows) as well as in the cells migrating from explant ganglion (Figure 7E', white arrows) while the control DRG, EGF expression was confined to the core of the ganglion (Figure 7A').

Together, these results indicate that under osteoclast stimulation, the endogenous EGF expression in DRG cells can mediate the activation of the EGF signaling cascade through the EGFR/EGFR and EGFR/ErbB2 dimers.

Figure 7: Expression of epidermal growth factor (EGF) in dorsal root ganglia (DRG). A. EGF staining on DRG cultured in control conditions – neurobasal + NGF. **A'.** Isolated red channel from image A showing the positive staining of EGF. **B.** EGF staining on DRG cultured with mature osteoclasts conditioned medium. **C.** Zoom in from image B, showing in detail the expression of EGF in cell soma of sensory neurons. **D.** Detailed image of neurites stained for EGF. **E.** Detailed image of cells

migrating from DRG positively stained for EGF. White arrows show positive staining (EGF in red/white, βIII tubulin in green and nuclei in blue, scale bar 100 μm (A, B) and 50 μm (C-E)).

Osteoclasts soluble proteins targeting axonal outgrowth

To identify potential molecules mediating the osteoclast-mediated axonal outgrowth, directly or indirectly, the total murine osteoclast secretome was analyzed.

Liquid chromatography and mass spectrometry of the osteoclast secretome identified more than two hundred proteins against the *Mus musculus* database. Pro Group Algorithm software calculates an unused score of 2 for a peptide with 99% identity confidence, and an unused score of 1.3 for a peptide with 95% confidence level. The unused protein score, representing the sum of the log confidence contributions of unique peptides used exclusively in the identification of the given protein, was applied. Proteins identified with an unused score above 15 were afterwards analyzed with STRING database v.10.0 and enriched for different cellular components (Table S1).

Most of the identified proteins can be assigned to the extracellular region, mainly as components of the extracellular matrix (ECM) but also from the plasma membrane. According to Gene Ontology biological function terms, extracellular proteins (both ECM and secreted proteins) are related to cell adhesion and five proteins appear related to nervous system development (bold type in Table S1) suggesting a possible contribution of these proteins in neurite growth.

Among the different identified proteins, fibronectin, low-density lipoprotein receptor-related protein 1 (LRP1) and periostin have been described as regulators of axon growth [65–68]. Analysis of the first shell of interaction of fibronectin, LRP1 and periostin with STRING software (Figure S1) identified molecular interactions with PDGF, EGFR, and TGF, which can be important for osteoclasts - sensory neurons crosstalk. In fact, adhesion to fibronectin stimulates tyrosine phosphorylation of the EGFR in the absence of receptor ligands [69]. Interestingly, periostin also binds to the integrins αvβ3, αvβ5, and α6β4, promoting the recruitment of EGFR and the activation of the Akt/PKB and FAK-mediated signaling pathways [70,71]. These results indicate the presence of other molecules in the osteoclast secretome capable of activating this signaling pathway.

Discussion

The present study provides compelling evidence that osteoclast lineage has a remarkable intrinsic
capacity to stimulate axonal outgrowth. Furthermore, this potential is concomitant with the
differentiation stage of osteoclasts and surpasses the effect of BMSC on axonal growth. Axonal
terminals integrate the cues secreted by differentiating osteoclasts after local administration of
osteoclast secretome using microfluidic devices. Remarkably, proteomic analysis revealed that
the observed osteoclast-mediated axonal growth was not derived from the action of the classical
neurotrophins such as NGF, BDNF, NT-3 and NT4/5. In addition, receptors tyrosine kinase
phosphorylation analysis confirmed low levels of TrkA, TrkB and TrkC activation. Interestingly,
EGFR and ErbB2 presented high levels of activation, highlighting their possible involvement in the
integration of instructive cues secreted by the osteoclasts. Furthermore, inhibition of EGFR/EGFR
and EGFR/ErbB2 dimers confirmed the contribution of this signaling cascade to the osteoclast-
mediated axonal outgrowth. The analysis of total osteoclast secretome indicated different
proteins that can be triggering the axonal outgrowth of sensory neurons, such as fibronectin,
LRP1 and periostin.

Different studies point to a parallelism between osteoclast activity and sensory neurons
outgrowth in physiological and pathological conditions. Several reports have described that
during embryonic long bone development osteoclasts are found in the process of endochondral
ossification, degrading the new formed trabecular bone, creating the bone marrow space [72]. In
another hand, longitudinal studies on the innervation process showed nerves appearing in areas
with high osteogenic and chondrogenic activity during the embryonic development [6,73,74].
Thus, the presence of osteoclasts in stages where innervation process occurs suggests a latent
role of these cells in the innervation pattern. In bone pathologies with increased osteoclasts
number: Paget's disease [75], fibrous dysplasia [76], *osteogenesis imperfecta* [77], and with
increased osteoclast activity: osteoporosis and metastatic bone disease, are frequently
associated with activation of sensory nerves [78–80]. Although the pattern of innervation in these
pathological conditions has been only described in metastatic bone disease, where an exuberant
sprouting of sensory nerve fibers has been reported, several studies indicate osteoclasts play a
fundamental role in bone pain [38,42,79,81]. Our finding that osteoclasts hold notable effect on
axonal outgrowth strength the reported parallelism between osteoclast lineage and the bone
innervation process and support the possibility that osteoclasts might contribute actively to the
dynamics of bone innervation pattern in physiological or pathological conditions.

Immunohistological studies described osteoclasts in close contact with nerve fibers within bone

microenvironment though typical synapses were not detected in the nerve endings [9,82,83]. Yet,

osteoclasts are described to activate nociceptors in nerve endings through acid-sensitive

nociceptors that innervate the marrow, mineralized bone, and periosteum [14,38]. These

channels are sensitized and excited by a decrease in pH, which is generated by osteoclasts activity

[81,84]. Using compartmentalized microfluidic devices resembling the compartmentalized bone

microenvironment, where only the nerve terminals are present, we show that the osteoclast-

mediated axonal growth was achieved by the ability of neurite terminals to integrate, through

receptors, the cues secreted by differentiating osteoclasts. The most probable candidate would

be TrkA receptors, well known to be activated, in permissive axonal growth and regeneration

[48,85]. Surprisingly, tyrosine kinase receptors phosphorylation analysis performed on DRG under

osteoclast secretome stimulation showed that neurotrophins receptor family (TrkA/ TrkB/ TrkC)

were not activated. These data corroborate with the lack of their universal ligands (NGF, BDNF,

NT-3 or NT4/5) in osteoclast secretome demonstrated by ELISA and further confirmed by mass

spectrometry.

In the bone microenvironment, BMSC and cells of the hematopoietic lineage have been described

to enhance axonal outgrowth through different signaling pathways. It is known that BMSC

stimulate the neuron survival and neurite outgrowth of spinal or DRG neurons [34,44,86–92], and

were reported to produce NGF, BDNF, basic fibroblast growth factor (bFGF), glial cell line-derived

neurotrophic factor (GDNF), vascular endothelial growth factor (VEGF) and ciliary neurotrophic

factor (CNTF) to stimulate neurite outgrowth *in vitro* [33,44,86]. The macrophages, which share

common progenitors with osteoclasts, were shown to play a supportive role in axonal

regeneration and outgrowth [93,94]. M1 and M2 macrophages have divergent secretory

repertoires and induce different responses on axonal behavior, being the M2 macrophages

predominantly neuroprotective and promoters of axon growth [95,96]. The macrophages were

shown to enhance neurite extension through the action of granulocyte-colony stimulating factor

[97] or by activating calcium/calmodulin kinase-dependent pathways [98]. Herein we show that

the osteoclasts operate via different secreted factors and target different signaling, mainly EGFR

family, to promote axonal outgrowth.

EGFR activation is involved in neuron differentiation and survival and has been implicated in

responses to neuronal damage [53,59,60,99]. Nevertheless, there is an extensive discussion in

the literature about the direct involvement of EGFR activation/inhibition on axonal regeneration

[54,57,100,101]. EGFR phosphorylation has been implicated in signaling inhibition of axonal

growth in central nervous system [54,55,57]. Koprivica *et al.* showed that EGFR inhibitors

effectively promoted neurite outgrowth from cultured DRG [55]. Nevertheless, others reported

that EGFR knockdown did not induce neurite outgrowth of retinal ganglion cells and inhibitors of

EGFR showed to enhance neurite outgrowth in the absence of the receptor [54]. We and others

previously demonstrated an increased expression of EGFR family in DRG after lesion [53,99,102].

To understand the contribution of EGFR and ErbB2 signaling in the osteoclast-induced axonal

outgrowth, two complementary approaches were applied, one by receptor-mediated inhibition

using pharmacological inhibition and other by ligand searching. Our data indicate that the EGFR

signaling has a role in axonal outgrowth promoted by osteoclast secretome. Upon EGFR inhibition

with Erlotinib, described to inhibit both EGFR and ErbB2 receptor kinases at high concentrations

[52], we observed a significant reduction in the axonal outgrowth area. These findings are

consistent with prior observations that the phosphorylation of EGFR was showed to enhance

neurite outgrowth [59,60,63,103–105]. The EGFR-mediated axonal outgrowth, in the absence of

EGFR ligands from the osteoclast secretome, leads us to hypothesize that endogenous expression

of EGF, by the neurons and glial cells from DRG explant cultures, could occur. In fact, we showed

that EGF was endogenously expressed by DRG cells in culture upon exposure to osteoclast

secretome and, therefore, having an autocrine/paracrine effect on the activation of EGFR

signaling pathway. It remains to be deciphered which molecules present in the osteoclast

secretome can directly influence either the endogenous expression of EGF and the activation of

EGFR family.

Notably, from the total secretome analysis, other proteins described to promote axonal

outgrowth and activate EGFR signaling pathway were identified such as fibronectin, LRP1 and

periostin. Adhesion to fibronectin stimulates tyrosine phosphorylation of the EGFR in the absence

of receptor ligands [69]. Periostin, also detected in the osteoclast secretome, was shown to

promote neurite outgrowth in cerebellar granule neuron and DRG neurons [67,68] and to recruit

the EGFR, supporting its possible role on EGFR modulation [70]. LRP1 was hypothesized to

promote axonal sprouting and regeneration in the adult central nervous system by activation of

canonical signaling pathways associated with neurotrophin receptors [66]. These molecules

present in the osteoclast secretome can directly influence the activation of EGFR signaling

cascade. Moreover, others can act directly on neurons or glial cells to induce the endogenous

expression of EGF. Some authors claimed that the effect of EGF on axonal outgrowth might be an

off-target effect. EGF can act on glial cells to increase the expression of neurotrophins and,

consequently, intensify the axonal outgrowth [100,106]. In line with these observations, the glial

cells at the explant culture can be producing neurotrophins upon osteoclast secretome

stimulation to improve the neurite outgrowth. Nevertheless, the TrkA, TrkB and TrkC receptors
did not show high phosphorylation levels.

In conclusion, our data provide robust evidence of the neurotrophic potential of the osteoclastic
lineage on axonal outgrowth, delivering new insights into the crosstalk between osteoclasts and
sensory neurons. Clearly, besides the activation of nociceptors through microenvironment
acidification, osteoclasts and their secreted molecules can be implicated in cancer-associated
bone pain possibly through the control of nerve sprouting and consequently remodeling of the
pathological innervation pattern. This is a subject that deserves further investigation to i)
control/reduce the formation of neuroma-like structures and ii) specify new candidates as
potential targets for bone pain therapies.

383 Materials and methods

All animal procedures were approved by the IBMC/INEB ethics committee and by the Portuguese
Agency for Animal Welfare (*Direção-Geral de Alimentação e Veterinária*) in accordance with the
EU Directive (2010/63/EU) and Portuguese law (DL 113/2013). Mice were housed at 22 °C with a
12 h light/dark cycle with *ad libitum* access to water and food. Adult C57Bl/6 male mice (6-8
weeks-old) were sacrificed in a carbon dioxide chamber to obtain bone marrow cells, while
pregnant females were sacrificed by cervical dislocation.

391 Cell culture

392 Bone marrow cells cultures

Bone marrow cells were isolated from tibiae and femur by flushing the bone marrow with α-MEM
(Gibco, Thermo Fisher Scientific, Waltham, MA, USA) containing 10% (v/v) heat inactivated (30
min at 56 °C) foetal bovine serum (FBS, Gibco, Thermo Fisher Scientific) and 1% (v/v)
penicillin/streptomycin (P/S, Gibco, Thermo Fisher Scientific). Non-adherent cells were removed
after 3 days and fresh medium was added. After three days the conditioned medium was
collected, centrifuged at 140 g, 4 °C, 5 min and stored at -80 °C. The conditioned medium was
divided into small aliquots (500 μL - 1 mL) before freezing to avoid repeated freeze/thaw cycles.

Osteoclast cultures

Bone marrow cells were isolated from tibiae and femur by flushing the bone marrow with α-MEM containing 10% (v/v) FBS and 1% (v/v) P/S. To generate primary osteoclast precursors, the bone marrow mononuclear cell suspension was treated with red blood cells lysis buffer (ACK lysing buffer, #A1049201, Gibco, Thermo Fisher Scientific) for 1 min at room temperature (RT) and, after centrifugation, cells were plated in 10-cm-diameter Petri dishes with 10 ng/mL macrophage colony-stimulating factor (M-CSF, PeproTech, London, UK) for 24 h. Afterwards the M-CSF concentration was increased to 30 ng/mL for an additional three days (pre-osteoclasts). Conditioned medium from osteoclasts precursors was collected, centrifuged at 140 g, 4 °C, 5 min and stored at -80 °C. Adherent cells were then detached with a cell scrapper and seeded at a density of 5×10^4 cells/cm^2 in the presence of 30 ng/mL M-CSF and 100 ng/mL receptor activator of nuclear factor kappa-B ligand (RANKL, PeproTech) [107,108]. Cells were maintained at 37 °C in a humidified atmosphere of 5% CO_2 for 3-4 days and then osteoclast conditioned medium was collected and stored at -80 °C.

Evaluation of osteoclast differentiation

Evaluation of osteoclast differentiation was performed by qRT-PCR analysis. RNA was isolated and purified as previously described in [102] and the mRNA expression of differentiation markers (tartrate-resistant acid phosphatase (TRAP, Fw: CGACCATTGTTAGCCACATACG; Rv: TCGTCCTGAAGATACTGCAGGTT, Invitrogen, Carlsbad, CA, USA), cathepsin K (CTSK, Fw: ATATGTGGGCCAGGATGAAAGTT; Rv: TCGTTCCCCACAGGAATCTCT, Invitrogen), osteoclast associated immunoglobulin-like receptor (OscAR, Fw: TGGCGGTTTGCACTCTTCA; Rv: GATCCGTTACCAGCAGTTCCAGA, Invitrogen) and glyceraldehyde 3-phosphate dehydrogenase (GAPDH, Fw: GCCTTCCGTGTTCCTACC; Rv: AGAGTGGGAGTTGCTGTTG, Invitrogen) was performed.

Immunocytochemistry was performed to assess the morphology of the differentiated osteoclasts and the possible presence of macrophages in culture. Briefly, cells were fixed with 4% paraformaldehyde (PFA, Merck Millipore, Kenilworth, NJ, USA) in phosphate-buffered saline (PBS) during 10 min followed by permeabilization with 0.25% (v/v) Triton X-100 (Sigma-Aldrich, St. Louis, MO, USA) in PBS and incubated, for 30 min at RT, with blocking solution composed of 1% (w/v) Bovine Serum Albumin (BSA, Sigma-Aldrich) in PBS for 30 min at RT to block antibody unspecific binding. The cells were incubated with macrophages specific marker – anti-F4/80 (#14-4801-85, eBioscience, Thermo Fisher Scientific), 1:250 in blocking solution overnight at 4 °C. Afterwards cells were washed with PBS and incubated with Alexafluor 488-Phalloidin (Life

Technologies, Thermo Fisher Scientific) 1:100 in PBS for 30 minutes at RT in the dark. At the end of the incubation time, the cells were washed with PBS and mounted using Vectashield with DAPI (Vector Labs, Burlingame, California, USA). Images were captured using the Inverted Fluorescence Microscope (Zeiss AxioVert, Carl Zeiss, Oberkochen, Germany), equipped with AxioVision SE64 Rel. 4.8 software. TRAP staining was also performed using an Acid phosphatase, Leukocyte (TRAP) kit (Sigma-Aldrich) according to manufacturer's instructions. Images were acquired using a stereomicroscope (SZX10, Olympus, Shinjuku, Tokyo, Japan) coupled to a digital camera (DP21, Olympus).

Dorsal root ganglia (DRG) culture

Embryonic DRG were obtained from 16 to 18 days-old (E16-18) murine embryos. Pregnant female mice were exposed to isoflurane by inhalation. The embryos were harvested and maintained on ice-cold Hank's balanced salt solution (HBSS, Invitrogen). Embryonic ganglia were accessed through the dorsal side of the embryo after spinal cord removal. The meninges were cleaned from the isolated DRG and the roots were cut. The ganglia were kept in cold HBSS until seeding. DRG were isolated under the stereoscopic magnifier and seeded into the lower wells of a 15-well µ-Slide Angiogenesis plate (#81506, Ibidi, Martinsried, Germany). Fibrin hydrogels were formed in the plates by applying equal volumes of a solution of plasminogen-free fibrinogen, pooled from human plasma, and a thrombin solution containing $CaCl_2$ and aprotinin (final concentration of fibrin components: 6 mg/mL fibrinogen (Sigma-Aldrich); 2 NIH U/mL thrombin from human plasma (Sigma-Aldrich); 2.5 mM $CaCl_2$ (Sigma-Aldrich); 10 µg/mL aprotinin (Sigma-Aldrich)). Before being used in the preparation of fibrin gels, fibrinogen was dissolved in ultrapure water, dialyzed against tris-buffered saline (TBS, pH 7.4), sterile-filtered and diluted to 12 mg/mL with sterile TBS. The fibrin gel was allowed to polymerize for 30 min at 37 °C in a 5% CO_2 humidified incubator, before the addition of culture media. DRG were cultured with neurobasal medium supplemented with 2% v/v B-27 Serum-Free Supplement® (B-27, Invitrogen), 60 µM 5-fluoro-2'-deoxyuridine (FDU, Sigma-Aldrich), 25 mM glucose (Sigma-Aldrich), 1 mM pyruvate (Sigma-Aldrich), 50 ng/mL 7S Nerve Growth Factor (NGF, Calbiochem, Merck Millipore), 2 mM glutamine (BioWittacker, Lonza, Basel, Switzerland) and 1% penicillin/streptomycin (P/S). Embryonic DRG explant cultures were left undisturbed for 24 h. Afterwards, neurobasal media was replaced by the conditioned media collected from BMSC and osteoclast cultures at different stages of differentiation for additional three days. Controls with neurobasal supplemented with NGF and α-MEM supplemented with M-CSF and RANKL (no contact with cells) were performed.

Osteoclast conditioned medium was also tested on axonal terminals of embryonic DRG cultured

using microfluidic devices. Briefly, microfluidic devices acquired from Merck Millipore were

modified and assembled as described in [47], to allow the alignment of the ganglia with the

microgrooves. Upon DRG plating, the somal side of the microfluidic chamber containing the DRG,

was cultured with neurobasal medium supplemented with NGF whereas the axonal compartment

was treated with mature osteoclast conditioned medium. Fluidic isolation was achieved by

maintaining the volume difference between the two compartments during 48 h, after which

axonal outgrowth was quantified.

Quantification of axonal growth

Axonal outgrowth was quantified after 72 h of treatment with conditioned medium and controls.

The embryonic DRG samples were fixed with 2% PFA in PBS during 10 min followed by 10 min at

37 °C with 4% of PFA in PBS with 4% sucrose. Ganglia were permeabilized with 0.25% (v/v) Triton

X-100 in PBS and incubated, for 30 min at RT, with blocking solution composed of 5% v/v normal

goat serum (Invitrogen) and 5% v/v FBS in PBS. DRG cultures were incubated with an antibody

directed against the neuronal specific marker βIII tubulin (Promega, Madison, WI, United States)

diluted 1:2000 in blocking solution, overnight at 4 °C. Afterwards, cells were washed and

incubated 1 h at RT with secondary antibody (Alexafluor 488/568, Invitrogen). Images were

captured either with a confocal laser scanning microscope (CLSM) Leica SP2 AOBS SE equipped

with LCS 2.61 software (Leica Microsystems, Wetzlar, Germany) at the i3S Advanced Light

Microscopy Unit or with an IN Cell Analyzer 2000 equipped with IN Cell Investigator software (GE

Healthcare, Little Chalfont, UK) at the i3S BioSciences Screening Unit. Radial outgrowth, defined

as the area between the ganglion edge and the outgrowth front, was determined. The outgrowth

area was computed according to Bessa *et al.* [43]. To quantify the axonal outgrowth in the

microfluidic platforms, neurite outgrowth was measured with AxoFluidic, an algorithm designed

to quantify neurite projection within these platforms [47].

Quantification of neurotrophins in secretome by enzyme-linked immunosorbent assay (ELISA)

496 Conditioned media from osteoclasts and BMSC were concentrated 10 times by centrifugation at

497 4000 g for 45 min, using 3 kDa MW cut-off filter units (Merck Millipore). In order to detect and

498 quantify the amount of NGF, brain-derived neurotrophic factor (BDNF), neurotrophin-3 (NT-3)

499 and NT4/5 in the conditioned medium, a multi-neurotrophin rapid screening ELISA kit (#BEK-

500 2231, Tebu-bio, France) was used according to the manufacturers' protocol.

Analysis of activation of Phospho-Receptor Tyrosine Kinase (RTK)

A proteome profiler mouse phospho-RTK array kit (#ARY014, R&D system, Minneapolis, MN, USA)

was used to quantify the phosphorylation level of 39 tyrosine kinase receptors, according to the

manufacturer's instructions. Protein lysate of DRG treated with conditioned medium was

analyzed. Each array membrane was exposed to X-ray film using a chemiluminescence detection

system (Amersham, GE Healthcare). The film was scanned using Molecular Imager GS800

calibrated densitometer (Bio-Rad, Hercules, CA, USA) and pixel density was quantified using

Quantity One 1-D Analysis Software, v 4.6 (Bio-Rad).

Pharmacological inhibition of epidermal growth factor receptor (EGFR) and ErbB2

Embryonic DRG were cultured in 15-well slides as previously described. Erlotinib, an EGFR

inhibitor, was added to the conditioned medium at 10 nM, 100 nM, 1 µM, 10 µM and 100 µM

and tested on DRG cultures during 72 h. Trastuzumab (Herceptin®), a monoclonal antibody

against the ErbB2 receptor, was kindly donated by the Instituto Português de Oncologia do Porto

(IPO, Porto, Portugal). Trastuzumab was tested in DRG cultures at a concentration of 10 µg/mL in

conditioned medium. Afterwards, axonal outgrowth was measured as previously described.

Immunostaining of EGF in explant DRG upon osteoclast secretome treatment

DRG exposed to osteoclast secretome were stained for EGF expression. Explant cultures were

fixed and permeabilized as above-mentioned. DRG were incubated with EGF primary antibody

(#06-102, Merck Millipore) diluted 1:400 and βIII tubulin diluted 1:2000 in blocking solution,

overnight at 4 °C. Afterwards, cells were washed and incubated 1 h at RT with secondary antibody

(Alexafluor 568/488, Invitrogen). Images were captured with a CLSM Leica SP5 AOBS SE equipped

with LCS 2.61 software.

Analysis of total secretome of mature osteoclast by information dependent acquisition (IDA) / liquid chromatography – mass spectrometry (LC-MS)

To reduce the amount of abundant proteins (such as albumin from bovine serum) and increase the signal from low abundance proteins (such as cytokines and growth factors), ProteoMiner kit (BioRad) was used. Conditioned media collected from mature osteoclasts cultures, was concentrated 10 times by centrifugation at 4000 g for 15 min, using 30 kDa MW cut-off filter units (Merck Millipore). Protein was quantified and then digested with trypsin. After reduction (DTT, 10 mM, 40 min, 56 °C) and alkylation (IAA, 20 mM, 30 min in the dark, RT, 18–20 °C), overnight digestion with trypsin (trypsin:sample = 1:50) was performed followed by a second in-solution trypsin digestion (3 h) in 80% (v/v) acetonitrile (trypsin:sample = 1:100).

Protein samples were analyzed by Nano LC–MS using Triple TOF 6600 (ABSciex, Framingham, MA, USA). External calibration was undertaken using a beta-galactosidase digest (ABSciex). The mass spectrometer was set for IDA scanning full spectra (400–2000 m/z) for 250 ms. The top 40 ions were selected for subsequent MS/MS scans (150–1800 m/z for 50 ms each) using a total cycle time of 2.3 s. The selection criteria for parent ions included a charge state between +2 and +5, and counts above a minimum threshold of 125 counts per second. Ions were excluded from further MS/MS analysis for 12 s. Fragmentation was performed using rolling collision energy with an energy spread of 5. Data were acquired with the Analyst software TF 1.7 (ABSciex, Framingham, MA, USA). Proteins were identified from the raw MS and MS/MS data using Protein Pilot Software v. 5.0 (ABSciex). A search was performed against the SwissProt protein database with taxonomic restriction to *Mus musculus* (release 2017_03). Protein identification was considered with an unused score greater than 1.3 (95% confidence). An independent false discovery rate (FDR) analysis using the target-decoy approach provided with the Protein Pilot™ software was used to assess the quality of the identifications.

LC-Triple TOF MS protein identification was performed at UniMS – Mass Spectrometry Unit, iBET/ITQB-UNL, Oeiras, Portugal.

Statistical analysis

All experiments were run in triplicate and repeated at least three times. Data analysis was performed using the Mann-Whitney U test in GraphPad Prism 6.00 for Windows (GraphPad Software, San Diego CA, USA). Differences between groups were considered statistically significant when $*p<0.05$, $**p<0.01$, $***p<0.001$.

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Author contributions

Conceptualization, E.N. and M.L.; Methodology, E.N., D.M.S and F.C.; Investigation, E.N., D.M.S,

F.C. and L.L.; Writing – Original Draft, E.N. and M.L.; Writing – Review & Editing, all authors;

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