



Osteogenesis Is Improved by Low Tumor Necrosis Factor Alpha Concentration through the Modulation of Gs-Coupled Receptor Signals

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ABSTRACT In the early phase of bone damage, low concentrations of the cytokine tumor necrosis factor alpha (TNF- α) favor osteoblast differentiation. In contrast, chronic high doses of the same cytokine contribute to bone loss, demonstrating opposite effects depending on its concentration and on the time of exposure. In the bone microenvironment, TNF- α modulates the expression/function of different G protein-coupled receptors (GPCRs) and of their regulatory proteins, GPCR-regulated kinases (GRKs), thus dictating their final biological outcome in controlling bone anabolic processes. Here, the effects of TNF- α were investigated on the expression/responsiveness of the A_{2B} adenosine receptor (A_{2B}AR), a Gs-coupled receptor that promotes mesenchymal stem cell (MSC) differentiation into osteoblasts. Low TNF- α concentrations exerted a prodifferentiating effect on MSCs, pushing them toward an osteoblast phenotype. By regulating GRK2 turnover and expression, the cytokine impaired A_{2B}AR desensitization, accelerating receptor-mediated osteoblast differentiation. These data supported the anabolic effect of TNF- α submaximal concentration and demonstrated that the cytokine regulates GPCR responses by interfering with the receptor desensitization machinery, thereby enhancing the anabolic responses evoked by A_{2B}AR ligands. Overall, these results indicated that GPCR desensitization plays a pivotal role in osteogenesis and that its manipulation is an effective strategy to favor bone remodeling.

KEYWORDS A2B adenosine receptor, G protein-coupled receptor kinases, tumor necrosis factor alpha, mesenchymal stem cells, osteoblasts, proteasome

Bone diseases, characterized by a progressive reduction of bone strength and susceptibility to fracture, are increasingly frequent pathologies, especially due to the remarkable increase in the average age of the population. Currently, no Food and Drug Administration-approved therapy is available to effectively promote bone regeneration (1, 2). For this reason, understanding molecular pathways involved in fracture reparative processes is pivotal to develop new, simple, and cost-effective therapeutic agents and strategies.

In a healthy body, injury is usually followed by both inflammation and immunologic reaction that is caused by local necrosis and bacterial infection. Inflammatory cells and factors released in the injured site influence the microenvironment of bone tissue in order to favor osteogenic reparative processes (3–5). Among the inflammatory players, tumor necrosis factor alpha (TNF- α) assumes an important role in bone healing, affecting mesenchymal stem cell (MSC) behavior in a dose- and time-dependent manner. In the early inflammatory phase, immediately after fracture, TNF- α is released at low levels and favors MSC migration, survival, and differentiation, thus promoting

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bone repair (6). In contrast, in the late and chronic uncontrolled phase of inflammation, higher doses of the same cytokine have destructive effects on bone and contributed to bone loss. However, the intracellular pathways and mechanisms involved in bone remodeling and responsible for the paradoxical effects of TNF- α remain unclear (7–9). For that matter, it is uncertain whether or not TNF- α dialogues with different G protein-coupled receptors (GPCRs) and, through the modulation of the expression and/or functional activity of these receptors, may interfere with the biological responses evoked by receptor stimulation under physiological and pathological conditions.

Several GPCRs, such as the parathyroid hormone receptor (PTH1R), E series prostaglandin receptors (EP2 and EP4), P1 and P2Y purinergic receptors, and many others, are directly involved in bone remodeling (10, 11). Clinical trials with PTH R-targeted molecules demonstrated that receptor desensitization, subsequent to the continuous administration of receptor agonists, may explain the switch from anabolic to catabolic effects of PTH on bone (12–14). Based on these pieces of evidence, the investigation of the molecular mechanisms underlying desensitization processes of GPCRs may be crucial to understanding the final biological outcome of these receptors to controlling bone remodeling and to propose new and alternative therapeutic strategies for bone diseases.

Among GPCRs, the adenosine A_{2B} receptors (A_{2B} ARs) have been demonstrated to promote MSC differentiation and increase osteogenesis both *in vitro* and *in vivo*, suggesting this receptor is an innovative target for bone diseases (15–19). A_{2B} ARs are functionally activated by endogenous adenosine only in damaged and inflamed tissues in which adenosine is massively released following ATP degradation. Moreover, several studies have demonstrated that the responses evoked by A_{2B} ARs are selectively regulated by proinflammatory cytokines in both chronic and acute inflammation (20, 21). In particular, TNF- α has been shown to modulate the A_{2B} AR desensitization process through a direct effect on GPCR receptor kinase 2 (GRK2) in human astroglial cells, thus modifying the receptor-mediated responses (22, 23). Indeed, GPCR desensitization is largely mediated by direct phosphorylation of receptor serine and threonine residues by a family of kinases, termed GRKs. Among the different isoforms, GRK2 and GRK3 are the most common kinases involved in the regulation of GPCR responsiveness in bone (24). The expression of GRK2 is temporally regulated during osteoblast development, and this kinase has been shown to modulate GPCR responsiveness during the osteogenic process (25). It is worth noting that the activity of GRKs is regulated by the inflammatory microenvironment. Several studies have demonstrated that TNF- α decreases the cellular levels of GRK2 in *in vitro* and *in vivo* models of inflammatory diseases and may contribute to the regulation of the desensitization processes of different GPCRs, thereby prolonging agonist-mediated receptor activation over time (26–28).

To date, no data are available on the regulation of A_{2B} AR desensitization in bone under inflammatory conditions. Based on this evidence, the aim of this work was to investigate the effect of TNF- α on the kinetics of A_{2B} AR desensitization (taken as a model of Gs protein-coupled receptors) in the regulation of osteoblast differentiation from MSCs. We demonstrated that a low TNF- α concentration, mimicking a submaximal inflammatory state, induced a significant inhibition of GRK2 activity, thereby impairing agonist-induced A_{2B} AR desensitization. Therefore, the inflammatory cytokine increased the osteogenic effects elicited by the purinergic system, contributing to osteoblast differentiation. These data indicate that local inflammatory environment manipulation is a simple and effective way to enhance bone formation and accelerate fracture repair.

RESULTS

MSC proliferation and differentiation to osteoblasts: effects of TNF- α . MSCs were cultured in complete growth medium. To induce differentiation into osteoblasts, a specific osteogenic medium was used (15). The time course and complete cell osteogenic process were monitored as described previously (15).

The effects of different concentrations of TNF- α (0.1 ng/ml to 10 ng/ml) on MSC proliferation were evaluated. TNF- α did not induce any significant effect on the proliferation rate of both undifferentiated MSCs (Fig. 1A) and cells differentiated for 5 days in osteogenic medium (Fig. 1B). These data demonstrated that the cytokine, in our culture experimental medium, did not have a toxic effect up to a 10 ng/ml concentration.

We then investigated the effects of TNF- α on MSC differentiation to osteoblasts. MSCs were subjected to the proinflammatory stimulus during their differentiation in osteogenic medium. The effects of TNF- α (1 ng/ml) on the expression of Runx2, a pivotal transcription factor that drives cells toward an osteoblast phenotype, and ALP, an early marker of osteoblast differentiation, were evaluated at different stages of differentiation. TNF- α induced a time-dependent increase in the expression levels of both Runx2 (19.69- \pm 2.69-fold versus 12.44- \pm 0.91-fold in the control after 15 days of treatment) (Fig. 1C) and ALP (20.37- \pm 1.34-fold versus 13.26- \pm 1.09-fold in the control after 15 days of treatment) (Fig. 1D).

In addition to the induction of osteogenic markers, calcium deposition after 15 days of cell incubation with the cytokine in osteogenic medium was measured as a late indicator of osteogenic differentiation. TNF- α significantly increased calcium deposition, confirming it favored matrix mineralization (Fig. 1E and F). In contrast, any significant effects were observed in MSCs cultured in growth medium (data not shown). These results suggested the proinflammatory cytokine contributes to the early osteogenic differentiation of MSCs, potentiating the effects of prodifferentiating factors.

Osteoblast differentiation: interplay between TNF- α and A_{2B}AR. It is known that proinflammatory cytokines regulate the expression and activity of several intracellular proteins; therefore, it contributes to the modulation of the responses evoked by GPCRs (22, 29). Among these, TNF- α has been shown to modulate expression and functionality of the A_{2B}AR (21, 23), a GPCR involved in MSC differentiation (17).

Based on this evidence, the modulation of TNF- α on A_{2B}AR-induced MSC differentiation was evaluated. Consistent with literature data, the selective A_{2B}AR agonist BAY60-6583 caused a time-dependent increase in both ALP and Runx2 expression after 5 and 15 days of cell treatment (Fig. 2A and B). These results confirmed that A_{2B}AR stimulation induced a significant enhancement of osteogenic processes. Moreover, the treatment with BAY60-6583 for 15 or 21 days produced an evident increase of mineralization (Fig. 2C and D), as previously reported (15).

When BAY60-6583 was used in combination with TNF- α , the pro-osteogenic effects of A_{2B}AR agonist were significantly increased. Indeed, the expression levels of Runx2 and ALP induced by BAY60-6583 in the proinflammatory medium were 1.28- and 1.25-fold higher than those obtained with the agonist alone (Fig. 2A and B).

These results were confirmed by the mineralization assay (Fig. 2C and D). Indeed, a significant increase in matrix mineralization was detected when TNF- α and BAY60-6583 were used in combination. The percentage of calcium deposit induced by the A_{2B}AR stimulation in the presence of cytokine was 144.6% \pm 5.1% and 155.3% \pm 4.8% after 15 and 21 days of differentiation, respectively, and it was 127.1% \pm 6.0% and 136.5% \pm 7.6% in the absence of TNF- α .

The effect of TNF- α on the responses evoked by BAY60-6583 was completely abrogated by the selective A_{2B}AR antagonist MRS1706, confirming it is selectively ascribed to the activation of the A_{2B}AR subtype. Notably, the effect of TNF- α alone was slightly decreased by the treatment with MRS1706, pointing out that the A_{2B}AR activation could be one of several mechanisms at the basis of TNF- α osteoblastogenesis induction.

These data, taken together, suggested that the MSC inflammatory microenvironment modulates the mineralization process, favoring the responses evoked by A_{2B}AR stimulation.

Agonist-induced A_{2B}AR desensitization at different stages of MSC differentiation: effect of TNF- α . The putative molecular mechanisms involved in the effects of

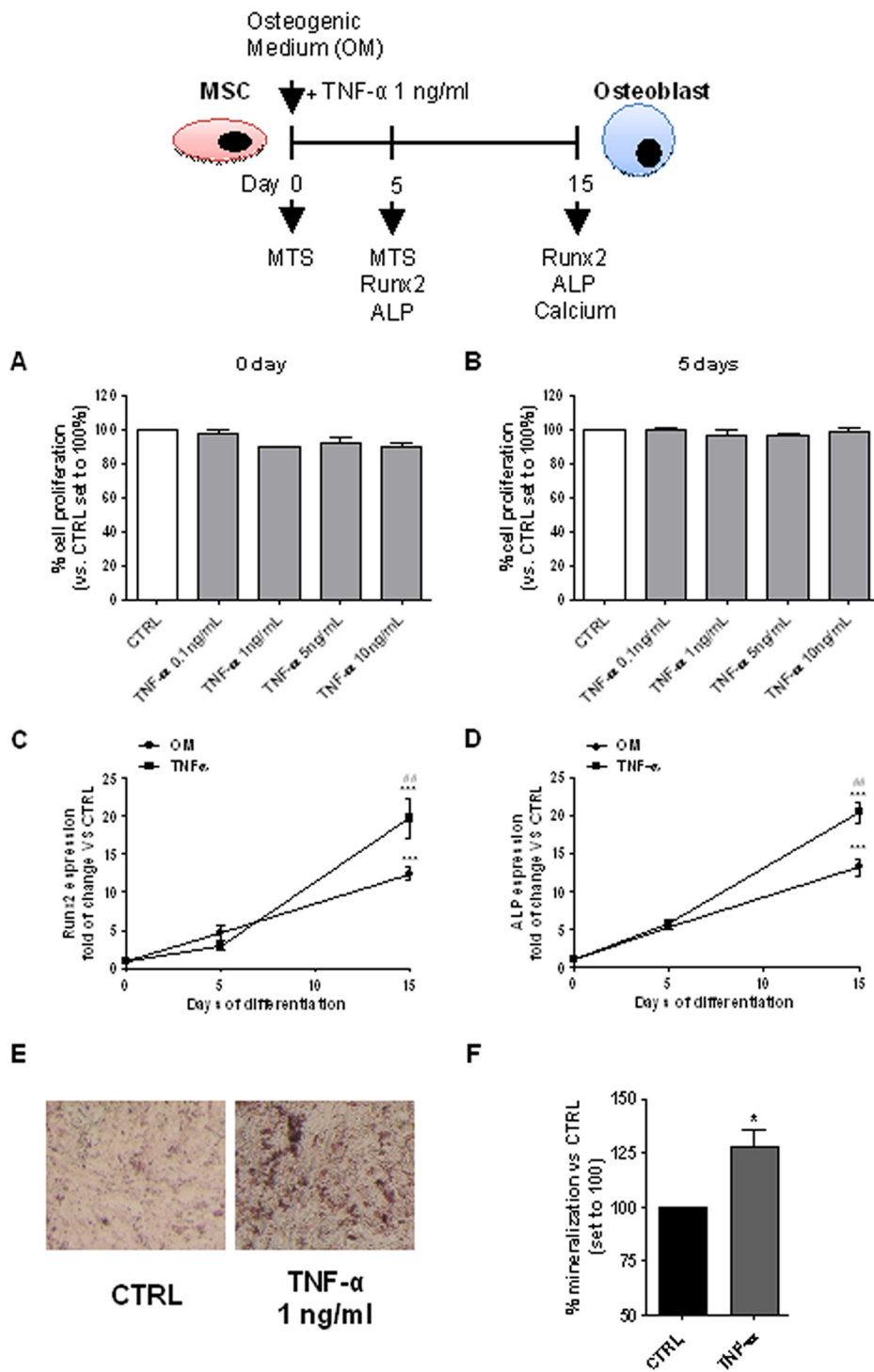


FIG 1 Effect of TNF- α on MSCs and osteoblast viability and differentiation. (A and B) MSCs were cultured in proliferation medium for 72 h (A) or in osteogenic medium for 5 days (B) in the presence of TNF- α (0.1 ng/ml to 10 ng/ml). After treatments, cell viability was detected using the MTS assay. The data were expressed as percentages of cell viability with respect to the untreated cells (control, OM), which was set to 100%, and they were presented as the mean values \pm SEM from three independent experiments, each performed in triplicate. (C and D) MSCs were cultured in osteogenic medium for different times (0 to 15 days), in the absence (control, OM) or in the presence of 1 ng/ml TNF- α . At each time, mRNA expression levels of transcription factors Runx2 (C) and ALP (D) were quantified by real-time RT-PCR. The data were expressed as fold changes with respect to the basal value, which was set to 1 (mean values \pm SEM; $n = 3$). ***, $P < 0.001$ versus basal; ##, $P < 0.01$ versus untreated cells. (E and F) MSC mineralization was evaluated after 15 days of differentiation in the absence (control) or presence of TNF- α (1 ng/ml). After treatments, cells were stained with alizarin red S, representative images were taken (E), and absorbance was counted using a plate reader (530 nm) (F). The data were expressed as a percentage of MSC mineralization with respect to that for the untreated cells (control), which was set to 100% (mean values \pm SEM; $n = 3$). *, $P < 0.05$ versus the control.

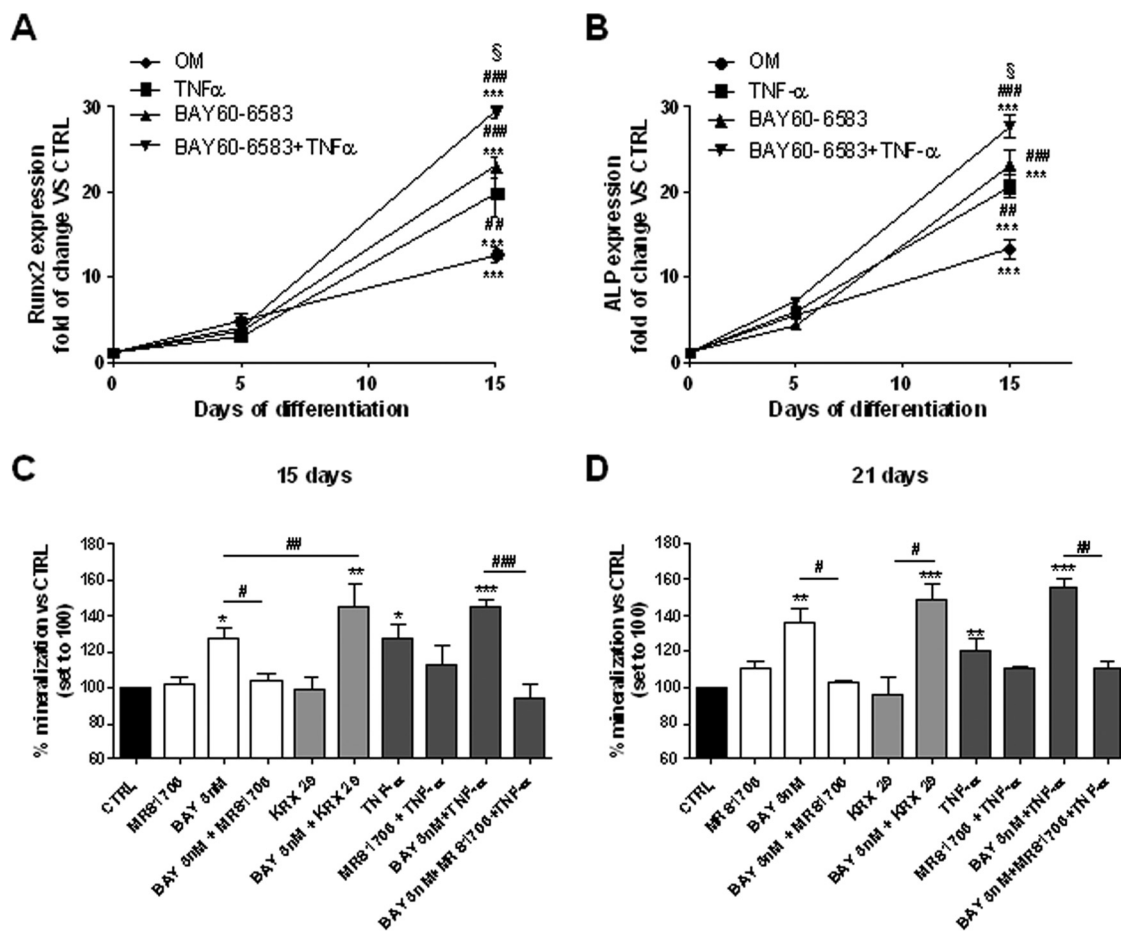


FIG 2 TNF- α modulation of A_{2B}AR-mediated MSC mineralization. (A and B) MSCs were cultured in osteogenic medium for different times (0 to 15 days), in the absence (control) or in the presence of 5 nM BAY60-6583 or 1 ng/ml TNF- α , alone or in combination. At each time, mRNA expression levels of transcription factors Runx2 (A) and ALP (B) were quantified by real-time RT-PCR. The data were expressed as fold changes with respect to the basal value (set to 1), and they were presented as the mean values \pm SEM from three independent experiments. The significance of the differences was determined by one-way ANOVA, followed by Bonferroni's *post hoc* test. ***, $P < 0.001$ versus basal; ##, $P < 0.01$; ###, $P < 0.001$ versus untreated cells; \$, $P < 0.05$ versus BAY60-6583 alone. (C and D) The MSC mineralizations were evaluated after 15 (C) or 21 (D) days of differentiation in the absence (control, OM) or presence of BAY60-6583 (5 nM), TNF- α (1 ng/ml), MRS1706 (1 μ M), and KRX29 (1 μ M), alone or in combination. After treatments, cells were stained with alizarin red S and absorbance was counted using a plate reader (530 nm). The data were expressed as a percentage of MSC mineralization with respect to that for the untreated cells (control), which was set to 100% (mean values \pm SEM; $n = 3$). *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$ versus the control; #, $P < 0.05$; ##, $P < 0.01$; ###, $P < 0.001$.

TNF- α on A_{2B}AR responses were investigated. First, the modulation of A_{2B}AR gene and protein expression evoked by the cytokine was investigated. As shown in Fig. 3, a significant upregulation of A_{2B}AR gene (2.03- \pm 0.17-fold versus the control) and protein (147% \pm 8% versus the control) expression was detected after 48 h of MSC treatment with the cytokine (Fig. 3A, C, and D). In contrast, challenging MSCs for 5 days in osteogenic medium did not affect A_{2B}AR content at either gene or protein levels (Fig. 3B, C, and D).

The effects of TNF- α on A_{2B}AR functional responsiveness then were investigated. A_{2B}ARs are mainly coupled to Gs proteins and activated adenylyl cyclase, thereby increasing intracellular cyclic AMP (cAMP) levels. Desensitization is one of the main processes that regulate GPCR functionality. On this basis, TNF- α modulation on agonist-mediated A_{2B}AR desensitization was evaluated at different stages of MSC differentiation (0 to 5 and 15 days). For this purpose, control and TNF- α -treated cells were preincubated with A_{2B}AR agonist (BAY60-6583) for different times (5 to 120 min), washed, and stimulated with the same agonist for 15 min. As depicted in Fig. 4, A_{2B}AR functional responses were quickly impaired by cell prechallenge with the agonist BAY60-6583 (Fig. 4) (30, 31). The desensitization kinetics appeared to be faster in

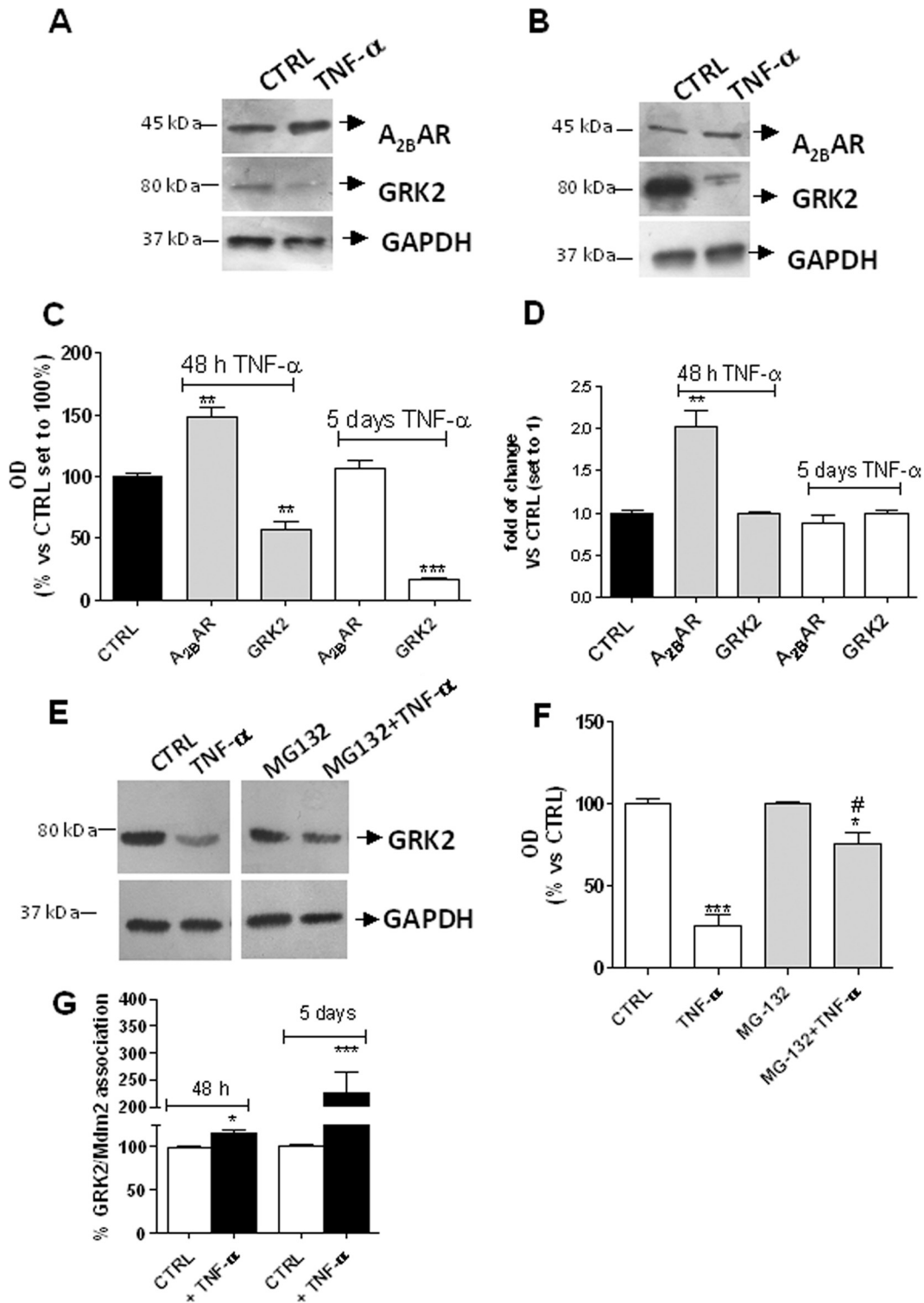


FIG 3 Effect of TNF- α on A_{2B}AR and GRK2 expression. (A to D) MSCs were cultured in osteogenic medium for 48 h (A, C, and D) or for 5 days (B, C, and D) in the absence (control) or in the presence of TNF- α (1 ng/ml). (A to C) At the end of the treatment, cells were lysed and the A_{2B}AR and GRK2 protein (A to C) or mRNA (D) levels were evaluated using Western blotting or real-time RT-PCR analyses, respectively. (A and B) Representative Western blots; GAPDH was the loading control. (C) Densitometric analysis of the immunoreactive bands, performed using the ImageJ program. The data were expressed as optical density (OD) percentage with respect to the untreated cells (control; set to 100%) and are mean values \pm SEM ($n = 3$). (D) Real-time PCR data, expressed as fold changes with respect to the control value, set to 1 (mean values \pm SEM; $n = 3$). (E and F) MSCs were pretreated with saline or 10 μ M MG-132 and then incubated in osteogenic medium for 48 h in the absence (control) or in the presence of TNF- α (1 ng/ml). At the end of the treatment, GRK2 protein levels were evaluated using Western blot analysis as reported for panels A to C. (G) MSCs were treated as described for panel A. At the end of the treatment

(Continued on next page)

undifferentiated MSCs (half-life [$t_{1/2}$], $1.46 \pm 0.008 \text{ min}^{-1}$) and in the early phase of differentiation ($t_{1/2}$, $4.17 \pm 0.021 \text{ min}^{-1}$) (Fig. 4A and B). In contrast, in the late phase of the differentiation program, A_{2B}AR functional responses remained preserved for a longer time, showing slower desensitization kinetics ($t_{1/2}$, $16.32 \pm 1.12 \text{ min}^{-1}$) (Fig. 4C). These differences may be ascribed to the different expression of A_{2B}ARs along with MSC maturation.

TNF- α did not significantly alter the basal response of A_{2B}AR to agonist but substantially impaired the degree and the rate of receptor desensitization. These effects were already evident in undifferentiated cells ($t_{1/2}$, $17.61 \pm 1.22 \text{ min}^{-1}$) and became more significant at different stages of the differentiation process (5 days, $t_{1/2}$ of $36.75 \pm 2.15 \text{ min}^{-1}$; 15 days, $t_{1/2}$ of $50.77 \pm 3.81 \text{ min}^{-1}$). The diverse effect of TNF- α during the differentiation process could reflect the presence of different cell phenotypes that are characterized by the expression of peculiar intracellular proteins involved in the receptor regulatory machinery.

Effects of TNF- α on GRK2 expression. GRKs, particularly the isoform 2, are the most relevant kinases involved in the phosphorylation and desensitization of GPCRs, including A_{2B}ARs (32, 33). To elucidate the mechanism through which TNF- α impaired A_{2B}AR functionality, we investigated the effects of the inflammatory cytokine on GRK2 expression and on its association with A_{2B}ARs upon agonist-mediated receptor stimulation. Notably, the cytokine caused a significant and time-dependent inhibition of GRK2 protein levels (Fig. 3A to C).

In order to dissect if a TNF- α -mediated decrease in GRK2 levels could involve a transcriptional mechanism, a real-time PCR analysis was performed. The results showed that the cytokine did not affect the expression of GRK2 mRNA after 48 h or after 5 days of differentiation (Fig. 3D). Based on such data, posttranscriptional regulation of GRK2 by TNF- α was investigated, considering, in particular, the involvement of the proteasome pathway, which has been identified as a major mechanism for modulating GRK2 expression levels (34–36). For this purpose, MSCs were challenged with the proteasome inhibitor MG-132 (37) in the absence or presence of the inflammatory cytokine for 48 h. As depicted in Fig. 3E and F, MG-132 significantly, but not completely, prevented TNF- α -mediated decrease of GRK2 expression.

Because several proteins are targeted to the proteasome pathway go through polyubiquitination (34, 38, 39), the potential role of ubiquitination in GRK2 degradation was examined by assessing the kinase association with the mouse double minute 2 (Mdm2) ubiquitin ligase (38, 40). The results showed that the cytokine induced a significant GRK2 association with Mdm2 following both 48 h and 5 days of treatment (Fig. 3G). Altogether, our data indicate that TNF- α regulation of GRK2 involves, at least partially, the ubiquitin-dependent proteasome pathway.

Effects of TNF- α on GRK2 association with A_{2B}ARs. The stimulation with BAY60-6583 caused a significant increase of A_{2B}AR-GRK2 association ($148.6\% \pm 4.8\%$; P value of <0.001 versus basal expression) (Fig. 4D). Moreover, A_{2B}AR-GRK2 association decreased during different time points of MSC differentiation to osteoblasts. The degree of this association paralleled the rate of desensitization induced by the agonist (Fig. 4A to C). In addition, a reduction in GRK2 association with A_{2B}ARs was detected when the MSCs were maintained in the presence of TNF- α , particularly in the early phase of the MSC differentiation process (Fig. 4D). These data suggest the cytokine modulates the desensitization machinery of different Gs-coupled receptors in MSCs, thereby favoring the functional activity of these receptors in response to agonist stimulation, as was demonstrated for the A_{2B}AR.

Effect of GRK2 inhibition/overexpression on the kinetics of A_{2B}AR desensitization. In order to corroborate the hypothesis that TNF- α affected the GRK2 activity

FIG 3 Legend (Continued)

period, the interaction between Mdm2 and GRK2 was quantified by an ELISA method as reported in Materials and Methods. The data were expressed as percentages of GRK2 association with respect to the untreated cells (mean values \pm SEM; $n = 3$). *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$ versus the control; #, $P < 0.05$ versus cells treated with TNF- α .

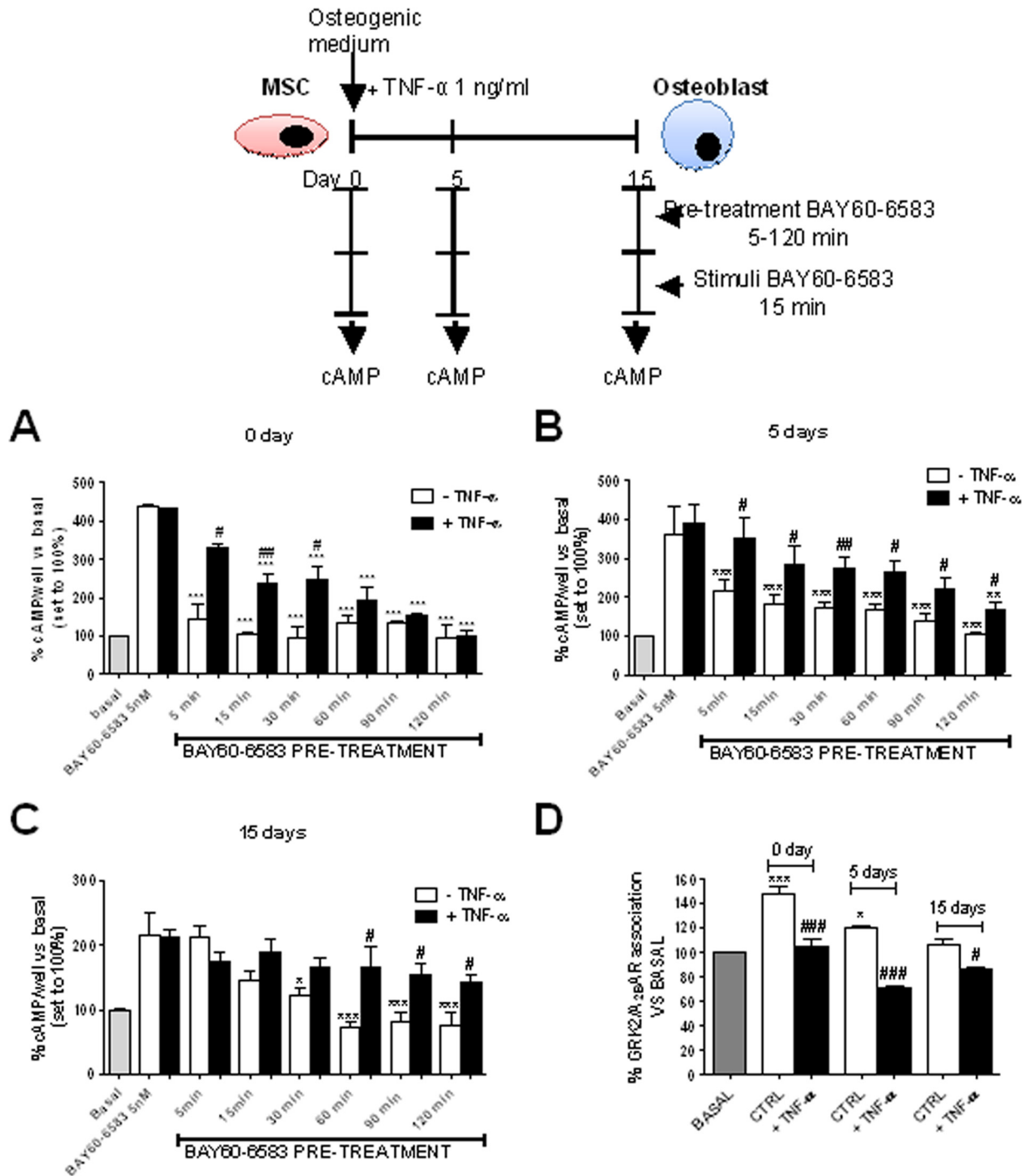


FIG 4 Effect of TNF- α on A_{2B}AR functional response during MSC differentiation. (A to C) MSCs were cultured in osteogenic medium in the absence or presence of 1 ng/ml TNF- α for 0 (A), 5 (B), or 15 (C) days. At each time point, MSCs were incubated with 5 nM BAY60-6583 for different times (5 to 120 min). After extensive washing, cells were treated for 15 min with 5 nM BAY60-6583. Intracellular cAMP levels were evaluated as reported in Materials and Methods. The data were expressed as cAMP percentages with respect to the level for untreated cells (basal), which was set to 100% (mean values \pm SEM; n = 3). **, P < 0.01; ***, P < 0.001 versus respective BAY60-65383 treatments; #, P < 0.05; ##, P < 0.01 versus TNF- α . (D) MSCs were differentiated for 0, 5, or 15 days in the absence or in the presence of TNF- α (1 ng/ml) and then were stimulated for 5 min with 5 nM BAY60-6583. At the end of the treatment period, cells were collected and the interaction between A_{2B}AR and GRK2 was quantified by an ELISA method as reported in Materials and Methods. The data were expressed as a percentage of GRK2 association with respect to that for the untreated cells (basal), which was set to 100% (mean values \pm SEM; n = 3). *, P < 0.05; ***, P < 0.001 versus the basal level; #, P < 0.05; ###, P < 0.001 versus the respective control.

leading to the impairment of Gs-coupled receptor desensitization, the effects of selective GRK2 inhibition and overexpression on A_{2B}AR desensitization were evaluated. Challenging MSCs with the synthetic GRK2 inhibitor KRX29 (33, 41) completely prevented A_{2B}AR desensitization at all of the osteoblast differentiation stages (Fig. 5A to C).

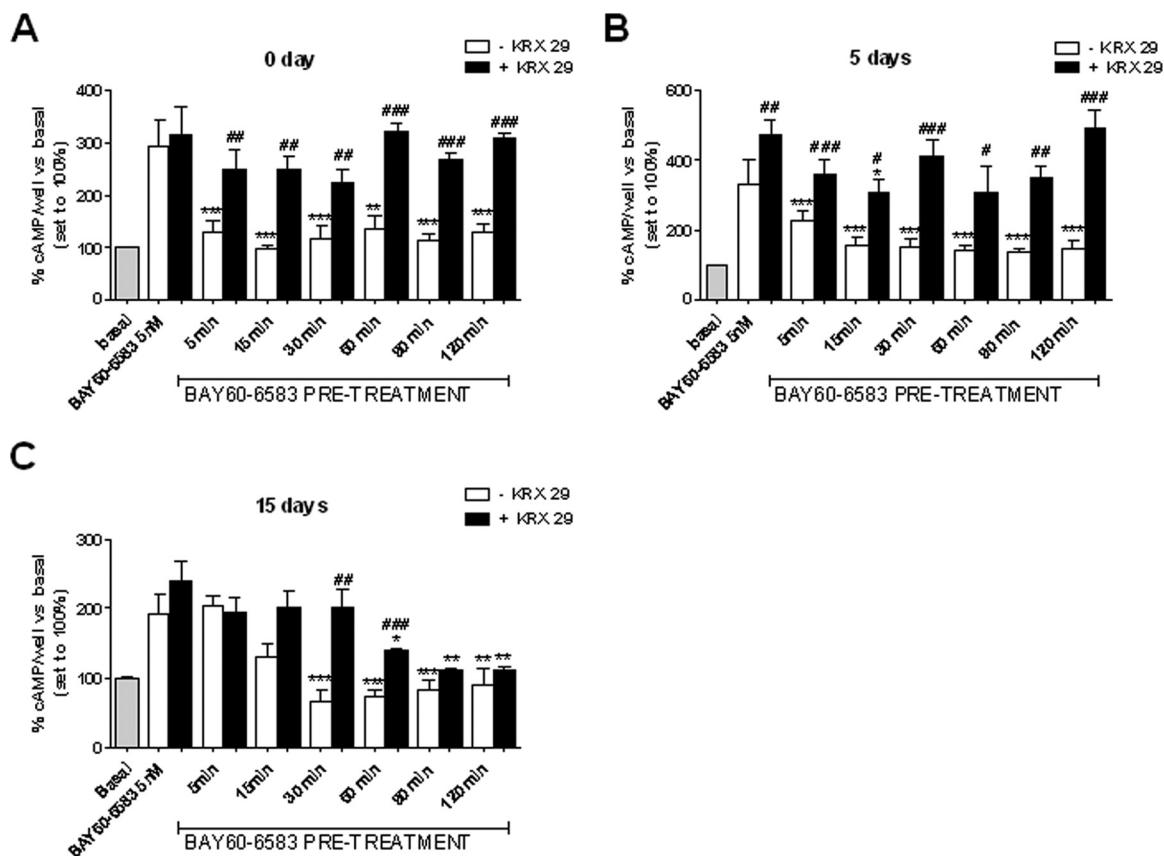


FIG 5 Effect of the GRK2 inhibitor on A_{2B}AR functional response during MSC differentiation. MSCs, cultured in osteogenic medium for 0 (A), 5 (B), or 15 (C) days, were treated with 5 nM BAY60-6583 for different times (5 to 120 min) in the absence or presence of 1 μ M KRX29. After extensive washing, cells were treated for 15 min with 5 nM BAY60-6583. Intracellular cAMP levels were evaluated as reported in Materials and Methods. The data were expressed as cAMP percentages with respect to the level for the untreated cells (basal), which was set to 100% (mean values \pm SEM; $n = 3$). *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$ versus respective BAY60-65383 treatments; #, $P < 0.05$; ##, $P < 0.01$; ###, $P < 0.001$ versus KRX29.

The data are in accordance with the effect of KRX29 on mineralization (Fig. 2C and D). In fact, the inhibitor slightly increased the mineralization induced by the agonist BAY60-6583, producing an effect comparable to that exerted by TNF- α . The similar effect exerted by the cytokine and the kinase inhibitor highlighted the involvement of GRK2 in agonist-mediated A_{2B}AR desensitization.

In parallel, we also evaluated the effect of GRK2 overexpression on TNF- α -mediated impairment of A_{2B}AR functional responses. MSC transfection with GRK2 plasmid induced a significant increase in protein expression (2.5-fold versus empty vector-transfected cells; $P < 0.001$), as demonstrated by Western blotting (Fig. 6A and B) (38). GRK2 overexpression induced a significant increase in the degree of A_{2B}AR desensitization, as shown in Fig. 6C. Furthermore, the high levels of GRK2 protein were able to overcome almost completely the inhibitory effect of TNF- α on A_{2B}AR functionality (Fig. 6D). The receptor reduced functional responses to its selective agonist in a time-dependent manner even in the presence of the cytokine. These data support our hypothesis that TNF- α impaired Gs protein-coupled receptor desensitization by affecting, at least in part, the activity of intracellular GRK2 proteins, as demonstrated for the A_{2B}AR.

DISCUSSION

TNF- α has been widely reported to exert opposite and conflicting effects on MSCs (6–9), primarily depending on cytokine concentration, time of exposure, and stage of cell differentiation (7, 9). In this study, a low concentration of TNF- α was demonstrated to enhance and accelerate the A_{2B}AR-mediated differentiation of MSCs toward an

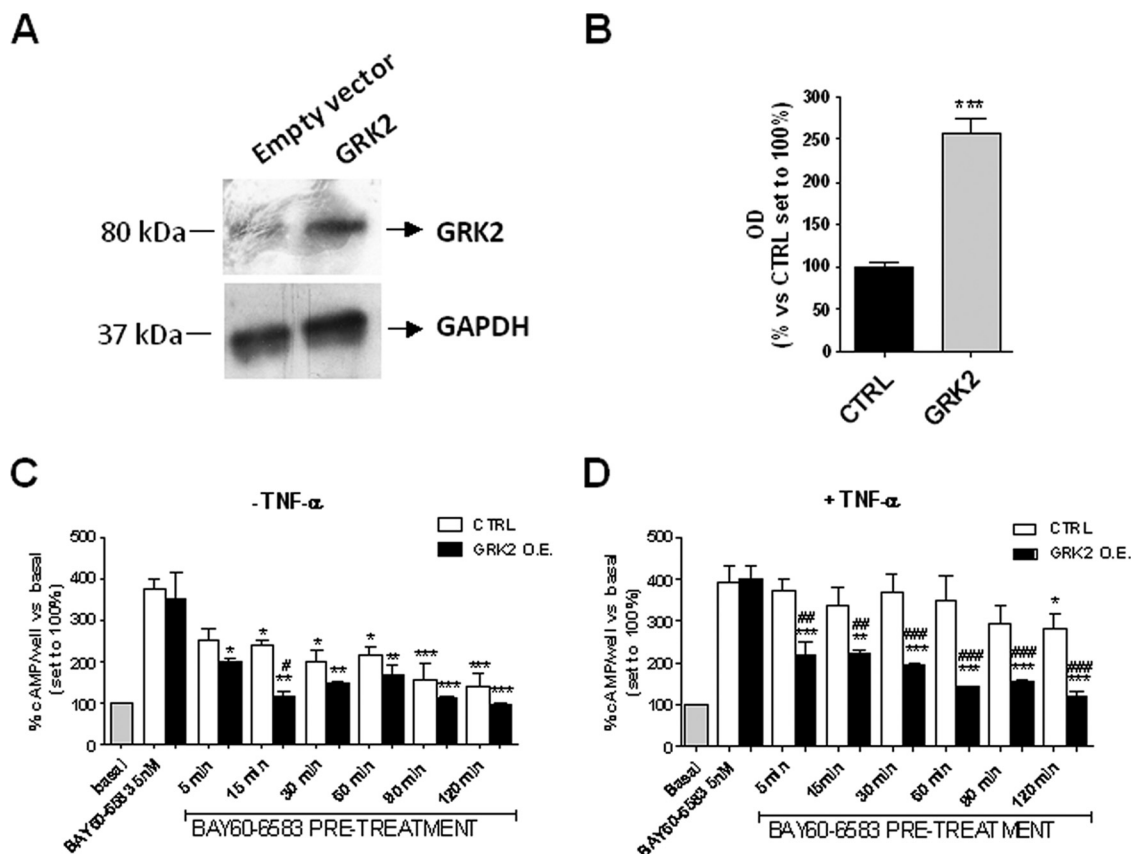


FIG 6 Effect of GRK2 overexpression on A_{2B}AR functional response during MSC differentiation. (A and B) MSCs were cultured in osteogenic medium and then transfected with a GRK2 plasmid as reported in Materials and Methods. At the end of the treatment period, the cells were lysed and the GRK2 protein levels were evaluated using Western blot analysis. GAPDH was used as the loading control. (A) Representative Western blots. (B) Densitometric analysis of the immunoreactive bands performed using the ImageJ program. The data were expressed as percent optical density (OD) with respect to the level for untreated cells (empty vector), which was set to 100% (mean values ± SEM; n = 3). ***, P < 0.001 versus the control. (C and D) MSCs were cultured in osteogenic medium in the absence (C) or presence (D) of TNF-α and then transfected with a GRK2 plasmid as reported in Materials and Methods. On the 5th day of differentiation, cells were incubated with 5 nM BAY60-6583 for different times (5 to 120 min) in the absence or presence of 1 ng/ml TNF-α. After extensive washing, cells were treated for 15 min with 5 nM BAY60-6583. Intracellular cAMP levels were evaluated as reported in Materials and Methods. The data were expressed as cAMP percentages with respect to the level for untreated cells (basal), which was set to 100% (mean values ± SEM; n = 3). *, P < 0.05; **, P < 0.01; ***, P < 0.001 versus respective BAY60-65383 treatments; #, P < 0.05; ##, P < 0.01; ###, P < 0.001 versus TNF-α.

osteoblast phenotype according to the consolidated evidence that low, submaximal concentrations (1 to 10 ng/ml) of TNF-α, particularly in the early phase of tissue injury, have an anabolic effect (42, 43). The mechanisms involved in the cytokine-elicited action on osteogenesis remain to be elucidated. Dissecting such molecular aspects is a crucial goal for a therapeutic intervention aimed at controlling bone remodeling, under both physiological and pathological conditions. Here, the interplay between TNF-α and the A_{2B}AR was investigated in MSCs, examining in particular GPCR desensitization, a process primarily mediated by intracellular GRK proteins (32, 44, 45). The induction of osteogenesis mediated by low TNF-α concentrations was primarily related to a reduction of GRK2 expression, leading to an enhancement of A_{2B}AR functionality.

Several pharmacological strategies targeting GPCRs for promoting osteogenesis (10, 14, 15, 18) have been limited by the loss of receptor functionality that reduces the response to agonists over time. Notably, the occurrence of desensitization processes has been linked to the switch from anabolic to catabolic events in bone remodeling, suggesting that manipulating the GPCR desensitization machinery is a useful strategy for the treatment of bone diseases (12, 13, 46).

TNF-α has been demonstrated to play an important role in bone healing by affecting MSC behavior (7). Consistent with recent literature (42, 43), here a low TNF-α concen-

tration was confirmed to stimulate MSC differentiation to osteoblasts within 15 days of cell treatment. The presence of the cytokine in osteogenic medium significantly increased Runx2 and ALP expression levels, thereby favoring MSC mineralization. Of note, this effect was not associated with significant changes in cell proliferation. TNF- α not only affects the differentiation processes, it also positively or negatively influences the rate of proliferation (47, 48). In our experimental model of induced MSC differentiation, TNF- α caused a complete switch to a differentiation process blocking the proliferative machinery.

Among the different GPCRs involved in bone remodeling and osteoblastogenesis, the purinergic receptor A_{2B}AR has arisen recently (15–19). Of note, this receptor responds to micromolar concentrations of adenosine, which is released under pathological conditions, such as stress or inflammation (49).

A functional interplay between A_{2B}ARs and TNF- α has been shown in different cell models, including glioma (23), intestinal epithelial (21), and vascular smooth-muscle (20) cells. In particular, the cytokine has been shown to enhance the adenosine-mediated responses under inflammatory conditions by regulating A_{2B}AR responses at different levels, including an upregulation of protein expression and/or an increase of receptor functional responsiveness by inhibiting receptor phosphorylation (21, 23).

To date, no evidence is available on the regulation of A_{2B}ARs by TNF- α in MSCs or the role of this dialogue in controlling the bone remodelling process. Here, TNF- α was shown to induce a significant increase in the A_{2B}AR-mediated osteogenic effects. The cytokine treatment increased the receptor mRNA and protein levels only in undifferentiated cells. Thus, the enhancement of the GPCR responses elicited by TNF- α was primarily ascribed to a direct effect on GRK2, which is the most important kinase involved in GPCR desensitization in response to agonist stimulation. Spurney et al. have demonstrated that GRK2 and β -arrestin are temporally regulated during osteoblast differentiation in a pattern that would tend to enhance GPCR responsiveness and favor MSC differentiation (13). When an injury occurs, the release of cytokines and other soluble factors in the cellular microenvironment may regulate the activity of these kinases and in turn affect the GPCR final biological outcome (50, 51). In particular, the increase of Gs-coupled receptor responsiveness and the consequent accumulation of higher levels of cAMP are crucial events in pushing osteoblast precursors to a differentiated phenotype. In this respect, Sinha and collaborators have recently demonstrated that cAMP signaling downstream of Gs-coupled receptors plays a critical role in determining the commitment of osteoblast precursors to bone with respect to adipocyte (52).

TNF- α has been shown to prevent Gs-coupled receptor desensitization by regulating GRK2 association with plasma membranes, thereby inhibiting receptor phosphorylation and desensitization in different cell lines (22). These data may support a beneficial role of low cytokine concentrations in potentiating GPCR functional responses. In MSCs, we demonstrated that TNF- α caused a significant downregulation of GRK2 protein level without affecting its gene expression. Kinase regulation was shown to partially involve the Mdm2-ubiquitin-dependent proteasome pathway, consistent with previous data (34, 36, 37, 40). Because TNF- α -mediated GRK2 downregulation was not completely counteracted by the use of a proteasome inhibitor, we speculate that additional mechanisms are involved in such an outcome. In this respect, GRK2 turnover has been linked to its phosphorylation by c-Src (53) or by mitogen-activated protein kinase (54); these effects, together with Mdm2 recruitment, can be favored by the β -arrestin machinery (35), which has been demonstrated to play different scaffold functions to coordinate both Mdm2-dependent and -independent processes implicated in GRK2 stability (35, 40, 53, 54).

The decrease in GRK2 expression was accompanied by a reduction of GRK2-A_{2B}AR association in response to agonist stimulation. Globally, these events caused an impairment of receptor desensitization, producing a time-prolonged accumulation of intracellular cAMP upon A_{2B}AR stimulation (Fig. 7). In this scenario, it is likely that proinflammatory soluble factors regulate the commitment of MSC toward osteoblasts

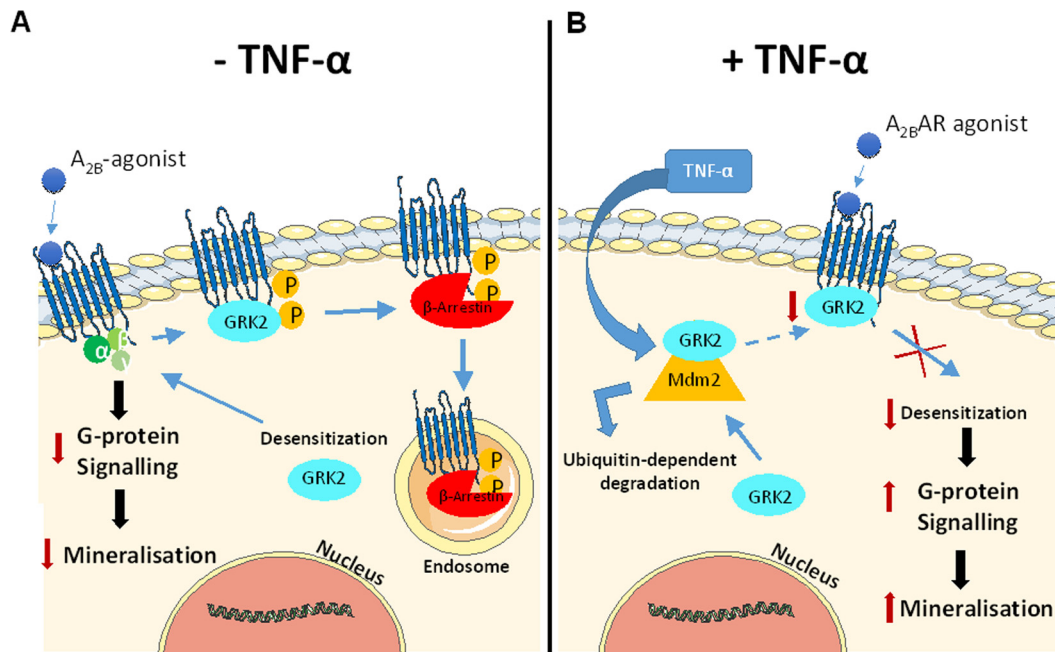


FIG 7 Cartoon illustrating the modulatory effect of TNF- α on A_{2B}AR responses in MSCs. (A) Agonist-mediated A_{2B}AR stimulation induces GRK2 recruitment, with consequent receptor phosphorylation and reduction of the functional responsiveness. (B) TNF- α reduces GRK2 levels and its association with A_{2B}AR by recruitment of the Mdm2/ubiquitin proteasome pathway. The GPCR desensitization process was impaired in the presence of the cytokine, thus enhancing the prodifferentiating effects elicited by A_{2B}AR stimulation.

by modulating the activity of desensitizing proteins and consequently increasing Gs-coupled receptor activity.

The unequivocal involvement of GRK2 in TNF- α -mediated regulation of A_{2B}AR was demonstrated by using a specific GRK2 inhibitor and by overexpressing GRK2 in MSCs. The chemical inhibition of GRK2 by the synthetic peptide KRX29 enhanced the osteogenic effects evoked by the A_{2B}AR agonist, similarly to the TNF- α -evoked effect. Otherwise, the overexpression of the GRK2 isoform in MSCs almost completely overcame the inhibitory effect of TNF- α on A_{2B}AR desensitization, thus confirming GRK2 as a target for the cytokine. Because GRK2 overexpression did not show a complete reversal of A_{2B}AR desensitization kinetics, additional factors, such as TNF- α -induced A_{2B}AR upregulation or the involvement of other GRK subtypes, can be speculated.

These data indicate that the release of cytokines in the inflammatory environment dictates MSC differentiation and represents a useful target to enhance bone formation, favoring the response evoked by different Gs-coupled receptors endowed with anabolic properties on bone.

MATERIALS AND METHODS

Materials. The chemicals 2-[6-amino-3,5-dicyano-4-[4-(cyclopropylmethoxy)phenyl]pyridin-2-ylsulfanyl]acetamide (BAY60-6583) and *N*-(4-acetylphenyl)-2-[4-(2,3,6,7-tetrahydro-2,6-dioxo-1,3-dipropyl-1H-purin-8-yl)phenoxy]-acetamide (MRS1706) were purchased from Tocris Bioscience (Bristol, United Kingdom). An RNeasy minikit was obtained from Qiagen. The Script cDNA synthesis kit was furnished by Bio-Rad s.r.l. Fluocycle II SYBR was from Euroclone (Milan, Italy). TNF- α was purchased from Sigma-Aldrich.

Cell cultures. Human bone marrow MSCs and cell culture medium were purchased from Lonza (Milan, Italy). The cell line was monitored for DNA profiling by HLA-DP beta and confirmed as human by PCR. The cells were used until passage 5 for fewer than 6 months after resuscitation.

Cells were subcultured in normal growth medium (MSCGM; Lonza) and plated (5×10^3 cells/cm²). The medium was changed to remove nonadherent cells every 3 to 4 days, and the cells were used at passages 0 to 3. For osteoblast differentiation studies, cells were seeded (9×10^3 cells/cm²) and cultured as previously described (15).

Cell viability assay. MSCs were seeded in 96-well microplates (5×10^3 cells/well) and cultured in proliferation or osteogenic medium for the indicated days in the absence (control) or in the presence of TNF- α (0.1 ng/ml to 10 ng/ml). To evaluate cell viability, a 3-(4,5-dimethylthiazol-2-yl)-5-(3-

TABLE 1 Human primers used for real time RT-PCR

Protein	Primer nucleotide sequences ^a	Product size (bp)
Runx2	FOR, 5'-GGCCTGGTGTAAATGGT-3'; REV, 5'-AGGCTGTTTGACGCCATAGT-3'	178
ALP	FOR, 5'-CTGCAAGGACATCGCCTATC-3'; REV, 5'-CATCAGTTCGTCTTGGGGTA-3'	101
A _{2B} AR	FOR, 5'-TCCATCTTCAGCCTTCTGGC-3'; REV, 5'-AAAGGCAAGGACCCAGAGGA-3'	128
β -Actin	FOR, 5'-GCACTCTCCAGCCTTCTCC-3'; REV, 5'-GAGCCGCCGATCCACACG-3'	254

^aFOR, forward; REV, reverse.

carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium (MTS) assay was used by following the manufacturer's instructions (Promega, Milan, Italy).

Real-time RT-PCR analysis. MSCs were cultured in normal growth medium and were treated with osteogenic medium in the presence or absence of TNF- α (1 ng/ml) and BAY60-6583 (5 nM), alone or in combination. The treatments were repeated every 3 days, and the expression levels of osteogenic markers (Runx2, osterix, ALP, and osteocalcin) were quantified after 0, 5, and 15 days of treatment. Moreover, MSCs were cultured in the presence or absence of TNF- α (1 ng/ml) in normal growth medium for 48 h or in osteogenic medium for 5 days. Gene expression then was assessed by real-time reverse transcription-PCR (RT-PCR) as previously described (55), and the primers used for the analysis are indicated in Table 1.

Mineralization assay. MSCs were seeded (9×10^3 cells/cm²) and then treated in the absence (control) or in the presence of TNF- α (1 ng/ml), BAY60-6583 (5 nM), MRS1706 (1 μ M), and KRX29 (1 μ M), either alone or in combination. Treatments were repeated every 3 days, and the mineralization was quantified after 15 or 21 days of treatment. The rate of mineralization was quantified using alizarin red staining (19).

Western blot analysis. MSCs were differentiated under osteogenic conditions for 48 h or 5 days in the absence (control) or presence of TNF- α (1 ng/ml). When indicated, MSCs were pretreated with 10 μ M proteasome/calpain inhibitor MG-132 (37) for 3 h and then incubated in osteogenic medium for 48 h in the absence (control) or presence of TNF- α (1 ng/ml).

Thereafter, cells were lysed (and 30 μ g of protein added with Laemmli solution) and loaded for SDS-PAGE (7.5%). Proteins were electrotransferred into polyvinylidene difluoride (PVDF) membranes and incubated overnight at 4°C using the primary antibodies anti-A_{2B}AR (sc-28996; 1:150; Santa Cruz Biotechnology), anti-GRK2 (sc-562; 1:200; Santa Cruz Biotechnology), and anti-glyceraldehyde-3-phosphate dehydrogenase (anti-GAPDH) (G9545; 1:5,000; Sigma-Aldrich) and by the appropriate peroxidase-conjugated secondary antibodies. Signals were detected using a chemiluminescent substrate (ECL; PerkinElmer, Waltham, MA). ImageJ software was used to perform the densitometric analysis of the immunoreactive bands (version 1.41; NIH, Bethesda, MD).

Measurement of cAMP levels during BAY60-6583 desensitization induction in MSCs. MSCs (2×10^4 cells/well) were plated in 24-well plates, and after 24 h, osteogenic differentiation was induced in the presence or absence of TNF- α (1 ng/ml) for 0, 5, or 15 days. At each time of differentiation, cells were stimulated with BAY60-6583 (5 nM) for 15 min and cAMP levels were quantified. In the desensitization assays, MSCs (differentiated for 0, 5, or 15 days in the absence or presence of TNF- α) were pretreated with BAY60-6583 (5 nM) for different times (5 to 120 min) and then washed and stimulated with BAY60-6583 (5 nM) for 15 min in the presence of 1 U/ml of adenosine deaminase (ADA) and the phosphodiesterase inhibitor Ro 20-1724 (20 μ M) (56). At the end of treatments, intracellular cAMP levels were quantified using a competitive protein binding method (57).

In order to investigate the effect of GRK2 inhibition on A_{2B}AR functional responses during MSC differentiation, the cells were cultured in osteogenic medium for 0, 5, or 15 days. At each time point, the cells were challenged with 5 nM BAY60-6583 for different times (5 to 120 min) in the absence or presence of the GRK2 inhibitor KRX29 (1 μ M).

GRK2 association with A_{2B}AR or Mdm2. To test the A_{2B}AR-mediated recruitment/association with GRK2, as well as Mdm2 involvement in GRK2 regulation, a quantitative immune-enzymatic assay on crude MSC lysates was used (38, 58, 59). MSCs were differentiated for 0, 5, or 15 days in the absence or presence of TNF- α (1 ng/ml) and then stimulated for 5 min with 5 nM BAY60-6583. Conversely, for Mdm2-GRK2 association, MSCs were incubated in osteogenic medium for 48 h or 5 days in the absence or presence of TNF- α (1 ng/ml). At the end of treatments, cells were suspended in lysis buffer containing 1% protease inhibitor cocktail (Sigma-Aldrich, Milan, Italy). Cell lysates (30 μ g/well) were incubated for 60 min into wells precoated with a rabbit anti-A_{2B}AR or anti-Mdm2 antibody; after washes, each well was incubated for 15 min with 1% bovine serum albumin (BSA) to block nonspecific sites and then for 2 h at room temperature with a mouse primary anti-GRK2 antibody. Wells then were washed and incubated for 1 h with an anti-mouse horseradish peroxidase (HRP)-conjugated antibody and washed again (38). The colorimetric substrate kit allowed a colorimetric quantification of the receptor-GRK2 association.

GRK2 overexpression. GRK2 overexpression was obtained by transfecting MSCs using the polyethyleneimine method (38). MSCs were cultured in osteogenic medium for 48 h in the absence or presence of TNF- α . The cells were incubated with 1 μ g of either GRK2 plasmid (OriGene, MD) or the corresponding empty vector (OriGene, MD) for 30 h. Transfected MSCs were rapidly seeded in 24-well plates and subjected to the cAMP assay as described above. GRK2 overexpression was confirmed by Western blot analysis.

Statistical analysis. A nonlinear multipurpose curve-fitting program, GraphPad Prism (version 5.00), was used for data analysis and graphic presentation. Data are reported as the means \pm standard errors

of the means (SEM) from 3 or 4 different experiments. Statistical analyses were performed using a one-way analysis of variance (ANOVA) study followed by the Bonferroni test for repeated measurements. Differences were considered statistically significant for P values of <0.05 .

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We have no conflicts of interest to declare.

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