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The BRAF-inhibitor PLX4720 inhibits CXCL8 secretion in BRAFV600E mutated and normal thyroid cells: a further anti-cancer effect of BRAF-inhibitors

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CXCL8 is a chemokine secreted by normal and thyroid cancer cells with proven tumor-promoting effects. The presence of BRAFV600E mutation is associated with a more aggressive clinical behavior and increased ability to secrete CXCL8 by papillary-thyroid-cancer cells. Aim of this study was to test the effect of the BRAF-inhibitor (PLX4720) on the basal and TNF- α -induced CXCL8 secretions in BRAFV600E mutated (BCPAP, 8305C, 8505C), in RET/PTC rearranged (TPC-1) thyroid-cancer-cell-lines and in normal-human-thyrocytes (NHT). Cells were incubated with increasing concentrations of PLX4720 alone or in combination with TNF- α for 24-hours. CXCL8 concentrations were measured in the cell supernatants. PLX4720 dose-dependently inhibited the basal and the TNF- α -induced CXCL8 secretions in BCPAP (F: 14.3, $p < 0.0001$ for basal and F: 12.29 $p < 0.0001$ for TNF- α), 8305C (F: 407.9 $p < 0.0001$ for basal and F: 5.76 $p < 0.0001$ for TNF- α) and 8505C (F: 55.24 $p < 0.0001$ for basal and F: 42.85 $p < 0.0001$ for TNF- α). No effect was found in TPC-1 (F: 1.8, $p = 0.134$ for basal; F: 1.6, $p = 0.178$ for TNF- α). In NHT an inhibitory effect was found only at the highest concentration of PLX4720 (F: 13.13 $p < 0.001$ for basal and F: 2.5 $p < 0.01$ for TNF- α). Cell migration assays showed that PLX4720 reduced both basal and CXCL8-induced cell migration in BCPAP, 8305C, 8505C and NHT but not in TPC-1 cells. These results constitute the first demonstration that PLX4720 is able to inhibit the secretion of CXCL8 in BRAFV600E mutated thyroid cancer cells indicating that, at least some, of the anti-tumor activities of PLX4720 could be exerted through a lowering of CXCL8 in the thyroid-cancer-microenvironment.

BRAF mutation leading to V600E amino acid substitution is identified in the majority of melanomas^{1,2}. BRAFV600E mutation is also the most frequent genetic lesion in papillary thyroid cancer (PTC), being present in nearly 40% of cases³⁻⁶. The presence of BRAFV600E mutation was reported to be associated with a more aggressive clinical behavior of papillary thyroid cancer (PTC)^{3,6-9}. The V600E mutation strongly enhances BRAF kinase activity by inserting a negatively charged residue adjacent to the phosphorylation site at T598 and mimicking phosphorylation at Thr598 and Ser601 residues^{1,10,11}, with the final result of increasing ERK1/2 phosphorylation¹.

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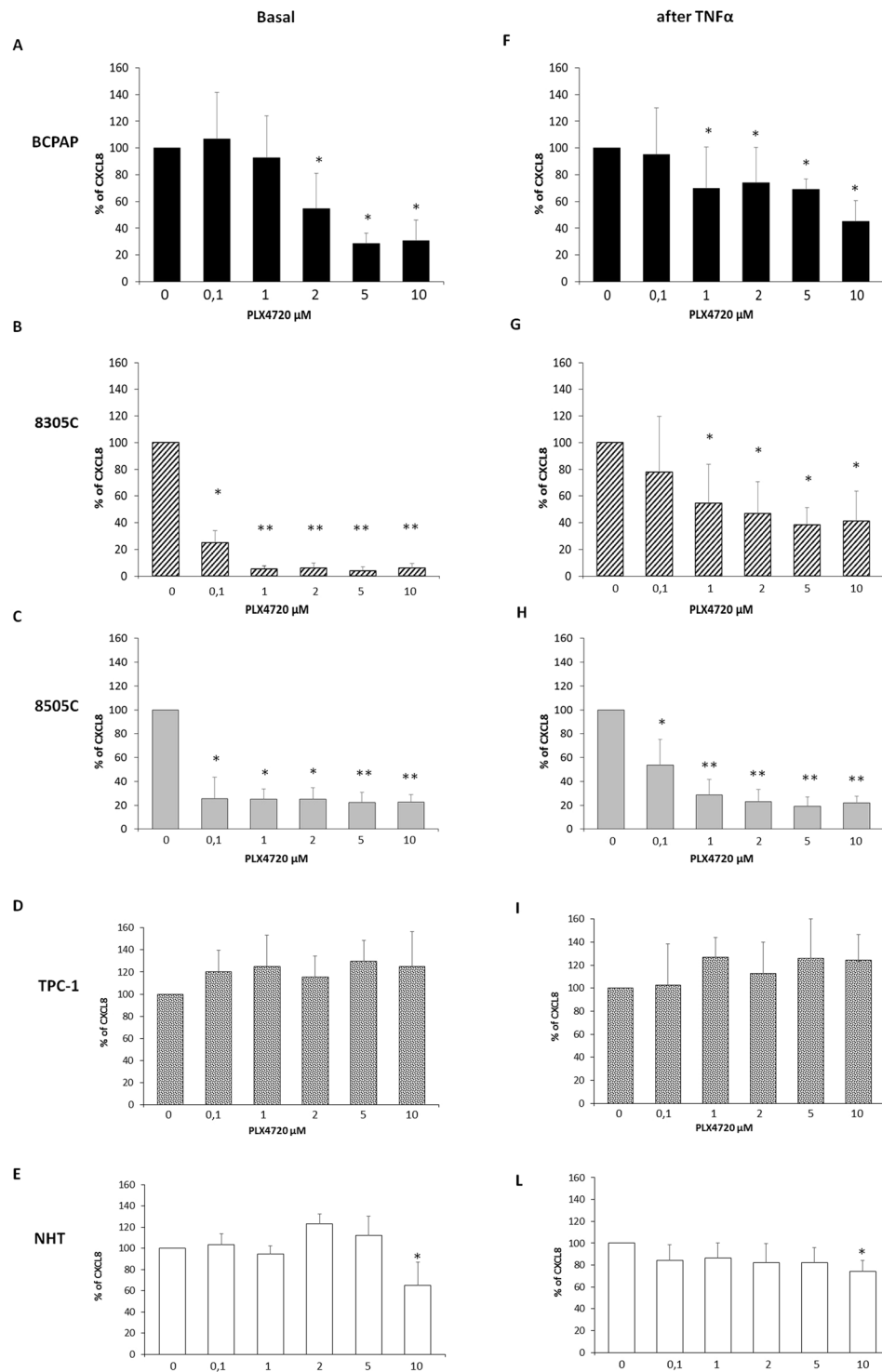


Figure 1. Panel A PLX4720 inhibit the basal CXCL8 secretion in BCPAP (ANOVA F: 14.3; $p < 0.0001$), the inhibitory effect was significant starting by 2 μ M concentration (Post Hoc analysis by Bonferroni $*p < 0.001$ vs. basal). Panel B PLX4720 inhibited the basal CXCL8 secretion in 8305C (ANOVA F: 407.9; $p < 0.0001$) (ANOVA F: 407.9; $p < 0.0001$), the inhibitory effect was significant starting from 0.1 μ M (Post Hoc analysis by Bonferroni $*p < 0.001$ vs. basal, $**p < 0.001$ vs. 0.1 μ M). Panel C PLX4720 inhibited the basal CXCL8 secretion in 8505C (ANOVA F: 55.24; $p < 0.0001$), the inhibitory effect started from 0.1 μ M (Post Hoc analysis by Bonferroni $*p < 0.001$ vs. basal, $p < 0.001$ vs. 0.1 μ M). Panel D Basal secretion of CXCL8 was not inhibited by PLX4720 in TPC-1 cell lines at any concentrations (ANOVA F: 1.8, $p = 1.34$). Panel E Basal secretion of CXCL8 was inhibited by PLX4720 in NHT (ANOVA F: 13.13; $p < 0.001$) being significant only at the higher concentration of 10 μ M (Post Hoc analysis by Bonferroni $*p < 0.01$ vs. basal). Panel F PLX4720 inhibit the TNF- α -stimulated CXCL8 secretion in BCPAP (ANOVA F: 12.29 $p < 0.0001$), the inhibitory effect was significant starting by 1 10 μ M (3.13; t anPost Hoc analysis by Bonferroni $*p < 0.001$ vs. TNF- α). Panel G PLX4720 inhibit the TNF-

α -stimulated CXCL8 secretion in 8305C (ANOVA F: 5.76 $p < 0.0001$), the inhibitory effect started from 1 rom *Post Hoc* analysis by Bonferroni * $p < 0.001$ vs. TNF- α). Panel H PLX4720 inhibit the TNF- α -stimulated CXCL8 secretion in 8505C (ANOVA F: 42.85 $p < 0.0001$), the inhibitory effect started from 0.1 rom *Post Hoc* analysis by Bonferroni * $p < 0.001$ vs. TNF- α , ** $p < 0.001$ vs. 0.1 ni * $p < 0.001$ vs. Panel I TNF- α -stimulated CXCL8 secretion was not inhibited by PLX4720 in TPC-1 cell lines at any concentrations (ANOVA F: 1.6, $p = 1.78$). Panel E TNF- α -stimulated CXCL8 secretion was inhibited by PLX4720 in NHT (ANOVA F: 2.5; $p < 0.01$) being significant only at the higher concentration of 10 μM (*Post Hoc* analysis by Bonferroni * $p < 0.001$ vs. TNF- α alone).

The BRAFV600E oncoprotein confers constitutive activation to the MAPK pathway, which accelerates tumor growth in experimental models of thyroid cancer¹².

Thyroid cancer microenvironment is composed of a mixture of immune cells and soluble mediators, which are present within and surrounding primary thyroid tumors^{13–15}. Among soluble mediators, specific chemokines not only attract different types of immune cell into the tumor site, but also produce pro-tumorigenic actions, including pro-angiogenetic, cyto-proliferative and pro-metastatic effects^{13–15}. Thus, the composition of thyroid cancer microenvironment, and in particular the levels of CXCL8 are currently regarded as a key factor for driving tumor progression¹⁶. In this scenario, no previous study investigated the role of BRAFV600E oncogene in influencing the production of CXCL8 within thyroid cancer microenvironment. Our group recently demonstrated that the BCPAP thyroid cancer cell line, harboring the BRAFV600E mutation, does secrete high amounts of CXCL8 both in basal culture conditions and after incubation with TNF α ¹⁷. CXCL8 has well-known pro-tumorigenic effects in several human cancers including the thyroid ones^{16,18} and its targeting was shown to hamper tumor progression^{18,19}. Indeed, *in vitro* and *in vivo* studies demonstrated that CXCL8 plays a crucial role in promoting epithelial-mesenchymal transition (EMT) and migration/metastatization of thyroid cancer cells^{20,21}. Supporting these actions, the administration of recombinant CXCL8 in xenografted mice with PTC significantly increased mortality^{19,22,23} while targeting of CXCL8 with an anti-CXCL8 monoclonal antibody significantly prolonged survival¹⁹. Previous attempts to inhibit CXCL8 secretion in normal and neoplastic thyroid cells were only partially effective owing to the presence of multiple intracellular pathways leading to CXCL8 secretion. Thus, lowering CXCL8 concentrations in thyroid cancer microenvironment requires specific strategies depending on the specific oncogenic background of neoplastic cells^{22,24,25}.

Pharmacological compounds with BRAF kinase blocking activity were shown to inhibit the secretion of CXCL8 in melanoma cell lines harboring the BRAFV600E mutation², but their effect in thyroid cancer cells remains to be investigated. Among them, the Plexxikon compound PLX4720 (7-azaindole derivative) is an orally administrable selective inhibitor of BRAFV600E with proven *in vitro* and *in vivo* therapeutic efficacy in melanoma models^{26–28}. As for the thyroid, PLX4720 was shown to inhibit the proliferation of BRAFV600E mutated thyroid cancer cell lines *in vitro*²⁹ and to reduce tumor growth and distant metastases in a human thyroid cancer model in mice³⁰. Whether this BRAF-inhibitor also exerts a CXCL8-lowering effect, in thyroid cancer cells remains to be investigated. Aim of the present study was to test the effect of PLX4720 on the secretion of CXCL8 in normal human thyroid (NHT) cells, in the BCPAP, 8305C and 8505C thyroid cancer cell line harboring the BRAF V600e mutation and in the TPC-1 thyroid cancer cell line bearing the RET/PTC re-arrangement. The potential effect of PLX4720 on the basal and CXCL8-induced thyroid cell migration was also investigated.

Results

Effects of treatment with PLX4720 in terms of inhibition of CXCL8 secretion in thyroid cancer cell lines and NHT.

CXCL8 concentrations were assayed in the supernatants of, BCPAP, 8305C and 8505C in basal and TNF- α -stimulated condition. As previously reported, TNF- α elicited a significant increase in the concentrations of CXCL8 in the supernatants of BCPAP^{17,31} but also in 8305C and 8505C cells (data not shown). The treatment with PLX4720 significantly and in a dose-dependent manner, inhibited the secretion of CXCL8 in: BCPAP (ANOVA F: 14.3; $p < 0.0001$) (Fig. 1A), in 8305C (ANOVA F: 407.9; $p < 0.0001$) (Fig. 1B) and in 8505C (ANOVA F: 55.24; $p < 0.0001$) (Fig. 1C). *Post Hoc* analysis performed by Bonferroni demonstrated some differences in the strength of inhibition induced by PLX4720 in different cell types. Indeed, significant inhibition of the basal secretion of CXCL8 started from a 2 μM concentration of PLX4720 in BCPAP ($p < 0.001$ vs. basal) (Fig. 1A) and from a 0.1 μM concentration of PLX4720 in 8305C (Fig. 1B) and 8505C (Fig. 1C); ($p < 0.001$ vs. basal for both 8305C and 8505C).

PLX4720 also produced a significant and dose-dependent inhibition of the TNF α -induced CXCL8 secretion in BCPAP cells (ANOVA F: 12.29 $p < 0.0001$) (Fig. 1F), in 8305C cells (ANOVA F: 5.76 $p < 0.0001$) (Fig. 1G) and in 8505C cells (ANOVA F: 42.85 $p < 0.0001$) (Fig. 1H). Also in this case, some differences among cell types were observed. Indeed, *Post Hoc* analysis by Bonferroni showed that the CXCL8 inhibiting effect of PLX4720 was significant starting from 1 μM in BCPAP (Fig. 1F) and in 8305C (Fig. 1G) ($p < 0.001$ vs. TNF- α alone for both cells) while in 8505C, a significant inhibition of the TNF- α -stimulated CXCL8 secretion started from a 0.1 μM concentration of PLX4720 ($p < 0.001$ vs. TNF- α alone) (Fig. 1H).

At difference with the above findings, PLX4720 did not produce any effect in terms of CXCL8 inhibition on TPC-1 cell lines, both in basal (ANOVA F: 1.8; $p = 0.134$) and in TNF- α -stimulated conditions (ANOVA F: 1.6; $p = 0.178$) (Fig. 1D–I). However, treatment with PLX4720 inhibited both the basal (ANOVA F: 13.13; $p < 0.001$) and the TNF- α -stimulated CXCL8 secretion (ANOVA F: 2.5; $p < 0.01$) in NHT (Fig. 1E–L). *Post Hoc* analysis by Bonferroni evidenced a significant inhibition of the basal and the TNF- α -stimulated CXCL8 secretion only at the highest (10 μM) PLX4720 concentration ($p < 0.01$ vs. basal; $p < 0.001$ vs. TNF- α alone) (Fig. 1E–L).

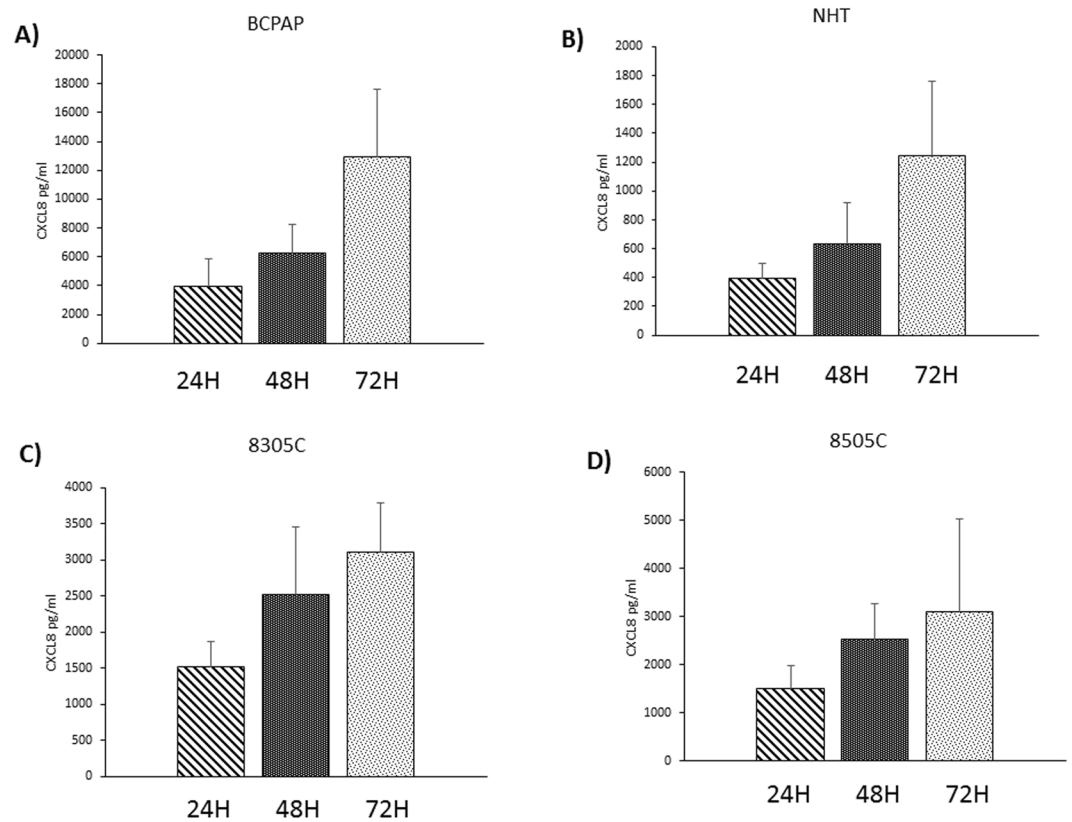


Figure 2. CXCL8 concentrations progressively increased throughout the 72-h time course in untreated thyroid cells. Panel A BCPAP cells (ANOVA: $F = 13.39$; $p < 0.0001$). Panel B NHT cells (ANOVA: $F = 2.7$; $p < 0.002$). Panel C 8305C cells (ANOVA $F = 6.76$; $p < 0.009$). Panel D 8505C cells (ANOVA: $F = 11.09$; $p < 0.001$).

Overall, the inhibition of CXCL8 secretion was paralleled by the inhibition of ERK phosphorylation, as assessed by Western blot, in all cell types with the exception of TPC-1 cells in which no CXCL8 inhibiting effect and no reduction of ERK phosphorylation was found (Supplemental Fig. S1).

Time course of PLX4720 inhibition of CXCL8 secretion. To assess whether the CXCL8-inhibiting effect of PLX4720 could vary over time, a time course of CXCL8 secretion following treatment with PLX4720 was performed. For this set of experiments those cells in which an inhibitory effect on CXCL8 secretion was exerted by PLX4720 (NHT, BCPAP, 8505C and 8305C) were treated with PLX4720 (0, 0.1, 1, 2, 5, 10 μM) for 24, 48 and 72 hours. A first finding was that, throughout the time course the basal levels of CXCL8 progressively increased in all cell types (although at a different magnitude) (ANOVAs: $F = 13.39$; $p < 0.0001$ for BCPAP; $F = 6.76$; $p < 0.009$ for 8305C; $F = 11.09$; $p < 0.001$ for 8505C; $F = 2.7$; $p < 0.002$ for NHT) (Fig. 2A–D). The percentages of inhibition of the CXCL8 secretion were compared for each concentration of PLX4720 at 24-h, 48-h and 72-h in each cell type. Separated ANOVAs for each cell type were performed to assess possible differences in percentages of inhibition of CXCL8 secretion by PLX4720 at each concentration throughout the time course. As shown in Fig. 3(A–D), no significant change in the inhibitory power of a given concentration of PLX4720 as assessed at 24 h, 48 h and 72 h could be found in any cell type. These findings would suggest that the CXCL8-inhibitory effect of PLX4720 is maintained (at a similar strength of inhibition) up to 72 hours.

Migration Assays. To assess whether the PLX4720-mediated inhibition of CXCL8 secretion would produce any biological consequence, cell migration assays were performed.

The transwell migration assay showed that rh-CXCL8 induced migration in all cell types. In details, the following results were found: $150 \pm 20\%$ migrated cells in BCPAP (ANOVA $F = 27.9$ $p < 0.0001$; *Post Hoc* CXCL8 vs. basal $p < 0.0001$), $150 \pm 0.1\%$ migrated cells in 8305C (ANOVA $F = 161.7$ $p < 0.0001$; *Post Hoc* CXCL8 vs. basal $p < 0.0001$), $120 \pm 30\%$ migrated cells in 8505C (ANOVA $F = 21.6$ $p < 0.0001$; *Post Hoc* CXCL8 vs. basal $p < 0.05$), $130 \pm 20\%$ migrated cells in NHT, (ANOVA $F = 25.3$ $p < 0.0001$; *Post Hoc* CXCL8 vs. basal $p < 0.05$), $140 \pm 10\%$ migrated cells in TPC-1, (ANOVA $F = 4.4$ $p < 0.01$; *Post Hoc* CXCL8 vs. basal $p < 0.05$) Fig. 4(A–E). The incubation with PLX4720 inhibited basal cell migration of BCPAP (mean: $56 \pm 10\%$ migrated cells; *Post Hoc* PLX4720 vs basal $p < 0.005$), 8305C (mean: $57 \pm 8\%$ migrated cells; *Post Hoc* PLX4720 vs basal $p < 0.0001$), 8505C (mean: $37 \pm 17\%$ migrated cells; *Post Hoc* PLX4720 vs basal $p < 0.005$) and NHT (mean: $62 \pm 18\%$ migrated cells; *Post Hoc* PLX4720 vs basal $p < 0.005$) Fig. 4(A–D). On the other hand, PLX4720 did not inhibit the basal migration of TPC-1 (mean: $108 \pm 24\%$ migration; *Post Hoc* PLX4720 vs basal NS) Fig. 4E. Finally, co-treatment with PLX4720 and rh-CXCL8 had an inhibitory effect on the CXCL8-induced cell migration of BCPAP (mean: $85 \pm 20\%$

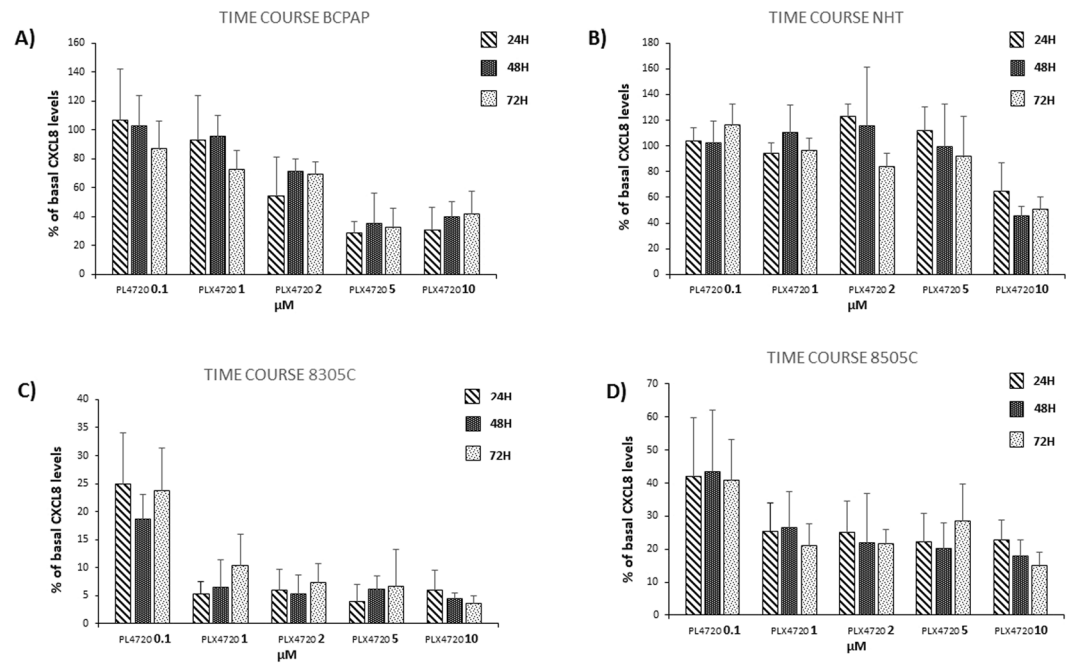


Figure 3. Time course of PLX4720 inhibitory effect on CXCL8 secretion. No significant change in the inhibitory power of PLX4720 at a given concentration was found throughout the 72-h time course. Panel A BCPAP (ANOVAs: PLX4720 0.1 μM F: 0.6; NS; 1 μM F: 1.7; NS; 2 μM F: 1.7; $p = 0.2$; 5 μM F: 2.7; $p = 0.7$; 10 μM F: 1.0, NS). Panel B NHT (ANOVAs: PLX4720 0.1 μM F: 1.7, NS; 1 μM F: 2.8; NS; 2 μM F: 3; NS; 5 μM F: 1; NS; 10 μM F: 2.3, NS). Panel C 8305C (ANOVAs: PLX4720 0.1 μM F: 1.6, NS; 1 μM F: 2.6; NS; 2 μM F: 0.5; NS; 5 μM F: 0.4; NS; 10 μM F: 1.3, NS). Panel D 8505C (ANOVAs: PLX4720 0.1 μM F: 0.02, NS; 1 μM F: 0.2; NS; 2 μM F: 0.1; NS; 5 μM F: 1.2; NS; 10 μM F: 2.9, NS).

migrated cells; *Post Hoc* CXCL8+ PLX4720 vs CXCL8 $p < 0.001$), 8305C (mean: $24 \pm 10\%$ migration; *Post Hoc* CXCL8+ PLX4720 vs CXCL8 $p < 0.0001$) 8505C (mean: $58 \pm 20\%$ migration; *Post Hoc* CXCL8+ PLX4720 vs CXCL8 $p < 0.0001$) and NHT (mean: $60 \pm 10\%$ migration; *Post Hoc* CXCL8+ PLX4720 vs CXCL8 $p < 0.0001$) but not in TPC-1 cells (mean: $120 \pm 30\%$ migration; *Post Hoc* CXCL8+ PLX4720 vs CXCL8 NS). Figure 4(A–E).

As a further confirmation of these findings, wound healing assay was performed (Supplemental Fig. S2).

Discussion

The results of the present study show that the selective BRAF-inhibitor, PLX4720, inhibits the basal and the TNF α -stimulated secretion of CXCL8 in BRAFV600E mutated thyroid cancer cell lines (BCPAP, 8305C and 8505C). The inhibitory effect of PLX4720 was observed also in primary cultures of NHT, even if only at the maximal concentration. On the other hand, PLX4720 did not produce any significant effect in terms of CXCL8 inhibition in TPC-1 cells harboring RET-PTC rearrangement.

It should be highlighted that, the strength of the inhibitory effect of PLX4720 (both in terms of minimal concentration able to elicit a significant inhibitory effect and in terms of percentages of inhibition) was different in relation to the specific cell type. Indeed, the CXCL8-inhibiting effect of PLX4720 was stronger in 8305C and 8505C tumor cell lines as compared to BCPAP which would fit with the notion that not all tumors may show the same effects from a single targeted agent³². The finding that the inhibitory effect of PLX4720 on CXCL8 secretion was not limited to BRAFV600E mutated cells, but also occurred in NHT could be regarded as unexpected and deserves to be discussed.

The fact that the concentrations of CXCL8 in the supernatants of NHT cells treated with PLX4720 progressively increased up to 72 hours of culture and that the percentage of inhibition versus untreated NHT cells remained constant would not support the hypothesis of a cytotoxic effect exerted by PLX4720. The issue of a potential cytotoxic effect of PLX4720 was addressed by previous studies on both normal and thyroid cancer cells demonstrating lack of any cytotoxic effect^{29,33}. In particular, Nucera *et al.* demonstrated that PLX4720 treatment (1 μM or 10 μM which corresponds to the concentrations used in the present study) did not lead to apoptosis as assessed by flow cytometric analysis³³. Taken together these data indicate that the inhibition of CXCL8 in NHT cells is not due to a cytotoxic effect of PLX4720. A more likely explanation stems from the study by Tsai *et al.*, who demonstrated that PLX4720 does not inhibit wildtype BRAF kinase activity in several tumor cell lines, but only at higher concentrations than those effective on cells harboring a mutated BRAFV600E kinase²⁷. These data provide an explanation for the high concentration of PLX4720 required to produce a significant inhibition of the CXCL8 in NHT cells. The inhibitory effect of PLX4720 on the secretion of CXCL8 by NHT cells should not be underestimated because these cells are by far the most abundant ones in a neoplastic thyroid, being their contribution to the global secreted amounts of CXCL8 in the tumor microenvironment not negligible²⁵.

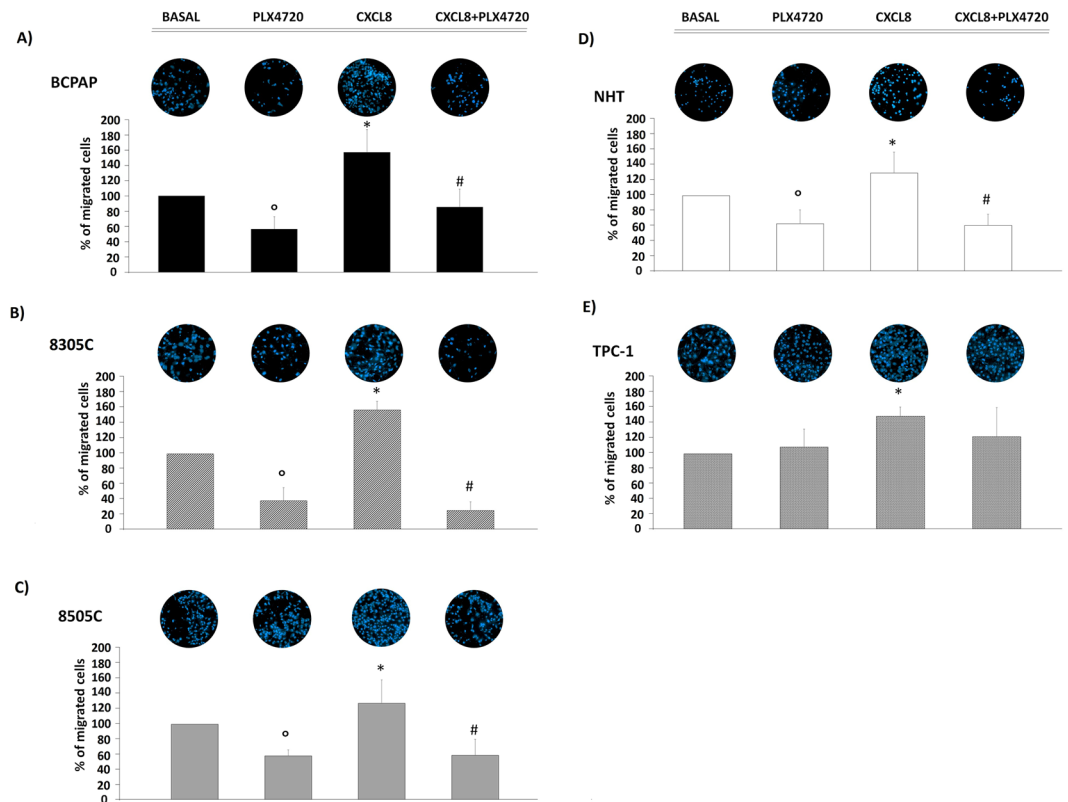


Figure 4. Effect of PLX4720 on the basal and rh-CXCL8-induced migration in BCPAP, 8305C, 8505C, NHT and TPC-1 thyroid cancer cells. Representative images and the respective histograms after 16 hours of migration within the trans-well migration chamber system in BCPAP, 8305C, 8505C, NHT and TPC-1 (Panels A–E respectively). Panel (A) in BCPAP the treatment with PLX4720 10 μ M reduced the basal migration (ANOVA $F = 27.9$, $p < 0.0001$; *Post Hoc* $^{\circ}p < 0.005$ vs. basal). The incubation of BCPAP with rh-CXCL8 produced a significant increase of BCPAP migration (*Post Hoc* $*p < 0.0001$ vs. basal). The co-incubation with rh-CXCL8 and PLX4720 10 μ M significantly inhibited the BCPAP migration induced by rh-CXCL8 (*Post Hoc* $^{\#}p < 0.001$ vs. CXCL8). Panel (B) in 8305C, the treatment with PLX4720 10 μ M reduced the basal migration of 8305C (ANOVA $F = 161.7$, $p < 0.0001$; *Post Hoc* $^{\circ}p < 0.005$ vs. basal). The incubation of 8305C with rh-CXCL8 produced a significant increase of 8305C migration (*Post Hoc* $*p < 0.0001$ vs. basal). The co-incubation with rh-CXCL8 and PLX4720 10 μ M significantly inhibited the 8305C migration induced by rh-CXCL8 (*Post Hoc* $^{\#}p < 0.0001$ vs. CXCL8). Panel (C) in 8505C the treatment with PLX4720 10 μ M reduced the basal migration of 8505C (ANOVA $F = 21.6$, $p < 0.0001$; *Post Hoc* $^{\circ}p < 0.005$ vs. basal). The incubation of 8505C with rh-CXCL8 produced a significant increase of 8505C migration (*Post Hoc* $*p < 0.0001$ vs. basal). The co-incubation with rh-CXCL8 and PLX4720 10 μ M significantly inhibited the 8505C migration induced by rh-CXCL8 (*Post Hoc* $^{\#}p < 0.0001$ vs. CXCL8). Panel (D) in NHT, the treatment with PLX4720 10 μ M reduced the basal migration of NHT (ANOVA $F = 25.3$, $p < 0.0001$; *Post Hoc* $^{\circ}p < 0.005$ vs. basal). The incubation of NHT with rh-CXCL8 produced a significant increase of NHT migration (*Post Hoc* $*p < 0.05$ vs. basal). The co-incubation with rh-CXCL8 and PLX4720 10 μ M significantly inhibited the NHT migration induced by rh-CXCL8 (*Post Hoc* $^{\#}p < 0.0001$ vs. CXCL8). Panel (E) in TPC-1 the treatment with PLX4720 10 μ M did not reduce the basal migration of TPC-1 (ANOVA $F = 4.4$, $p < 0.01$, *Post Hoc* NS vs. basal). The incubation of TPC-1 with rh-CXCL8 produced a significant increase of TPC-1 migration (*Post Hoc* $*p < 0.05$ vs. basal). The co-incubation with rh-CXCL8 and PLX4720 10 μ M did not reduce the TPC-1 migration induced by rh-CXCL8 (*Post Hoc* $^{\#}$ NS vs. CXCL8). Bar graphs show the corresponding analysis of migrated cells on the lower side of the transwell filter. Basal migration was conventionally estimated as 100%.

A further aspect deserving to be commented is that the inhibitory power of PLX4720 (expressed as % of inhibition) for a given cell type and a given concentration of PLX4720 was maintained at a similar degree over a 72-h time course indicating lack of any rebound effect of the CXCL8 inhibiting effect. The existence of a rebound effect was previously demonstrated by Montero-Conde *C et al.*, showing that the effects of BRAF inhibition in BRAF mutant thyroid cancer cells, were hampered (after 72 hours) by the release of negative feedback mechanisms³⁴.

PLX4720 was previously demonstrated to exert several anticancer effects in both xenografts models of melanoma and thyroid cancer. In animal models of melanoma, treatment with PLX4720 leads to a reduction of tumor growth and induces a higher expression of apoptotic genes and cell cycle arrests in neoplastic cells^{26,27}.

As far as thyroid tumors are concerned, treatment with PLX4720 lead to a reduction of proliferation, migration and invasion of thyroid cancer cells^{29,33}.

In this regards, the results of the present study suggest that at least some of the anti-cancer effect of PLX4720 might be exerted also through a reduction of CXCL8 within the tumor microenvironment. Indeed, even if CXCL8 inhibition might not be the only mechanism, the results of migration assays experiments clearly indicated that PLX4720 was able to reduce the migration of thyroid cells. In particular, this effect was evident only in those cells in which the treatment by PLX4720 also produced an inhibition of the CXCL8 secretion (BCPAP, 8305C, 8505C, NHT) but not in those cells in which PLX4720 did not inhibit CXCL8 secretion (TPC-1). These results highlight that the ability of PLX4720 in inhibiting CXCL8 secretion would in turn, result in potentially relevant biological consequences.

The results of the present study constitute the first demonstration of a CXCL8-inhibiting effect exerted by PLX4720 in thyroid cancer cells and should be regarded as potentially relevant. Indeed, CXCL8 is a chemokine with extensively described pro-tumorigenic effects which include influencing of tumor cell growth, angiogenesis, invasiveness and EMT^{16,20,35,36}. Furthermore targeting/lowering of CXCL8 levels is known to produce beneficial effects in thyroid cancer. Interestingly, a previous *in vivo* study showed that patients with melanoma harboring the BRAFV600E mutation experience a significant decrease in the circulating levels of CXCL8 while treated with BRAF-inhibitors².

In this view, the present study was specifically aimed at evaluating whether PLX4720 (already proven to have several anti cancer effects) was also able to inhibit CXCL8 secretion which could be regarded “*per se*” as beneficial against cancer. In conclusion, the here reported results, would suggest a further anti-cancer effect of PLX4720 which is manifested by its ability to reduce CXCL8 secretion. Although *in vivo* studies will be required to assess the therapeutic benefits deriving from the inhibition of CXCL8 secretion in thyroid tumor microenvironment, this could represent an alternative therapeutic strategy, at least for those minority of patients who are refractory to conventional therapies.

Materials and Methods

Primary cultures of NHT. Surgical specimens of normal human thyroid were obtained from the contralateral disease-free lobe of patients who underwent thyroidectomy for a solitary non-functioning nodule (n = 3). The study was approved by the Institutional Board of ICS-Maugeri. Before surgery, written informed consent to the study was obtained from all patients. All the experiments were performed in accordance with the relevant guidelines and regulations. Surgical specimens were minced and then incubated with collagenase type II (Sigma, Saint Louis, MO, USA) 5 mg/ml, in 5 ml of Coon's F12 medium, for 4 h at 37 °C as previously described³⁷. Then, 10 ml of Coon's F12 medium were added, following which, cells were filtered, spun at 1000 × g for 10 min, washed with Coon's F12 medium, spun again, and finally re-suspended in complete medium containing 5% newborn calf serum and a mixture of six hormones including insulin (5 µg/ml), hydrocortisone (50 µg/ml), transferrin (5 µg/ml), somatostatin (10 ng/ml), gly-his-lysine (10 ng/ml) and bovine TSH (1 mU/ml).

Thyroid tumor cell lines BCPAP, TPC-1, 8305C and 8505C. Human thyroid cancer cell lines, BCPAP harboring the BRAF V600E mutation and TPC-1 bearing the RET/PTC rearrangement, were a gift of Prof. M. Santoro (Medical School, University “Federico II” of Naples, Naples, Italy). These cell lines had been previously tested and authenticated by DNA analysis. Cancer cells were propagated in Dulbecco's Modified Eagle Medium (DMEM) (Sigma, Saint Louis, MO, USA) supplemented with 10% fetal bovine serum (Sigma, Saint Louis, MO, USA), 2 mM L-glutamine and 100 U/ml penicillin/streptomycin (Sigma, Saint Louis, MO, USA) as previously described³⁷. Human thyroid cancer cell lines, 8305C and 8505C harboring the BRAF V600E mutation were previously tested and authenticated by DNA analysis. 8305C cells were propagated in Minimum Essential Medium (MEM) (Sigma, Saint Louis, MO, USA) supplemented with 10% fetal bovine serum (Sigma, Saint Louis, MO, USA), 2 mM L-glutamine and 100 U/ml penicillin/streptomycin (Sigma, Saint Louis, MO, USA). 8505C cells were propagated in RPMI medium (Sigma, Saint Louis, MO, USA) supplemented with 10% fetal bovine serum (Sigma, Saint Louis, MO, USA), 2 mM L-glutamine and 100 U/ml penicillin/streptomycin (Sigma, Saint Louis, MO, USA). Cells were incubated with the chosen stimuli in serum-free medium.

Basal and TNF- α -induced CXCL8 secretion in BCPAP, 8305C, 8505C, TPC-1 thyroid cancer cell lines and in primary cultures of NHT in the presence or absence of increasing concentrations of PLX4720. For the CXCL8 secretion assays, 3000 cells were seeded into 96-well plates in complete medium. After adherence to the plastic surface, BCPAP, TPC-1, 8305C, 8505C and NHT cells were incubated for 24 h in serum-free medium (basal condition) with increasing concentrations of PLX4720 (0, 0.1, 1, 2, 5, 10 µM) (Sigma Aldrich). Incubation with PLX4720 was done in the absence (basal secretion) and in the presence (stimulated condition) of TNF- α 10 ng/ml. All experiments were performed in triplicates.

Time course of PLX4720 inhibition of CXCL8 secretion in BCPAP, 8305C, 8505C, TPC-1 thyroid cancer cell lines and in primary cultures of NHT. For the CXCL8 secretion assays, 3000 cells were seeded into 96-well plates in complete medium. After adherence to the plastic surface, BCPAP, 8305C, 8505C and NHT cells were incubated for 24, 48 and 72 hours with or without PLX4720 0, 0.1, 1, 2, 5, 10 µM. All experiments were performed in triplicates.

ELISA for CXCL8. CXCL8 was measured in cell supernatants of BCPAP, TPC-1, 8305C, 8505C and NHT cells using commercially available kits (R&D Systems, Minneapolis, MN). The mean minimum detectable concentration of CXCL8 was 3.5 pg/ml. The intra- and inter-assay coefficients of variation were 3.4% and 6.8%, respectively. Samples were assayed in duplicates.

Western blot for pERK. Cells were seeded in 6 well plates at a concentration of 500000 cells/well and treated for 24 hours with increasing concentrations of PLX4720. Subsequently cells were washed three times with PBS and incubated 10 minutes in buffer RIPA supplemented with protease inhibitors on ice. Cells were then scraped and centrifuged. The amounts of proteins in the supernatants were quantified using the BCA kit (Thermo Scientific). Western blots were conducted by standard procedure on 15 µg protein. 1:1000 anti-phospho ERK1/2 (αP-ERK1/2) or anti-ERK1/2 antibodies (αERK1/2; Cell Signaling Technology, Inc., Danvers, MA, USA) were used. Bands were quantified by densitometry with the Quantity One program version 4.6.1. (BioRad). (Supplemental materials).

Migration. The cell migration assay was performed with the trans-well migration chamber system (Merck Millipore, Milan, Italy), as previously described³⁷. Briefly, NHT, TPC-1, BCPAP, 8305C and 8505C were cultured for 24 hours with fresh medium alone or supplemented with 10 µM of PLX4720, 100 ng/ml of CXCL8 (R&D Systems, Minneapolis) or their combination. After treatment, 20×10^3 cells/well were seeded in the upper chambers of the 96-well plate with polycarbonate inserts having 0.3Cm²/well membrane area and 8 µm pore size. In each condition the lower chambers were filled with 150 µl of the corresponding medium. Cells were left to migrate for 16 hours at 37 °C and 5% CO₂. At the end of the incubation, samples were analyzed as previously described³⁸. Briefly, cell inserts were washed three times with PBS and migrated cells on the underside of the membrane were fixed with 4% paraformaldehyde for 20 minutes. Cell nuclei were then stained with Hoechst 33342 (1:1000) (Life Technologies, Monza, Italy). Finally, the membranes were cut out with a scalpel, and mounted onto glass slides with DACO reagent (Life Technologies, Monza, Italy). Three replicates were evaluated for each condition. Images were acquired using an Olympus BX51 microscope (Olympus, Deutschland GmbH, Hamburg, Germany). The number of migrated cells was counted analyzing 12 random fields of the membranes per condition. Data are expressed as % of mean numbers of migrated cells ± standard deviation (SD).

Wound-healing assay. For wound-healing assay cells were seeded in a 24-well plate. When cells reached nearly the 90% of cell confluence, a scratch wound was created in the monolayer using a sterile 200 µl pipette tip. Cells were then treated with fresh medium alone, or supplemented with 10 µM of PLX4720. Phase contrast images were captured between 0 and 24 hours using an Olympus IX53 microscope (Olympus, Deutschland GmbH, Hamburg, Germany). Data are expressed as the percentages of the remaining gap area after 24 hours relative to the initial gap area (0 hours). The area was measured using the LCMicro software (Olympus Soft Imaging Solutions GmbH) (Supplemental materials).

Statistical analysis. Statistical analysis was performed using the SPSS software (SPSS, Inc., Evanston, IL). Mean group values were compared by using one-way ANOVA for normally distributed variables. *Post Hoc* analysis was performed according to the Bonferroni's correction for multiple comparisons. Values are reported as mean ± SD unless otherwise noted. A *p* value < 0.05 was considered statistically significant.

Data Availability

There are no restrictions on the availability of materials or information.

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Author Contributions

F.C., O.A., F.L. and M.R. conceived and designed the experiments. F.C., V.C., M.M., L.C. and M.D. conducted the experiments. F.C. and V.C. and F.L. performed the study. F.C. and S.T. carried out the data analysis. L.P. provided thyroid cancer cells 8305C and 8505C and contributed to the interpretation of the results obtained. F.C. and M.R. wrote this manuscript. L.C. revised results, data analysis and the text. All authors have read and approved the final manuscript.

Additional Information

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