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Quercetin potentiates UVB-induced c-Fos expression: Implications for its use as a chemopreventive agent

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Abstract

Quercetin (Qu) is currently being investigated as a chemopreventive agent for a number of cancers including non-melanoma skin cancer induced by ultraviolet (UV) light. We previously reported that Qu degradation has important consequences on signaling and cell biology. In the current study, we report that Qu induces c-Fos mRNA and protein expression through activation of p38 and CREB, and Qu potentiates UVB-induced c-Fos expression. Inclusion of ascorbic acid (AA) in cell culture medium stabilizes Qu and completely prevents both Qu- and UVB- induced p38 and CREB activation, leading to a blockade of c-fos gene expression through reduced CREB/CRE binding. AA stabilizes c-Fos mRNA, increasing steady state levels even when c-fos gene expression is suppressed, but this has no effect on c-Fos protein levels in either mock- or UVB-irradiated cells. We report that Qu blocks mTOR signaling and inhibits c-Fos protein expression directly through this mechanism, since co-treatment with Qu and AA resulted in complete suppression of UVB-induced c-Fos protein expression even in the presence of significantly increased mRNA levels. We further confirmed that this was not due to increased protein turnover since inhibition of proteasome activity with MG-132 did not raise c-Fos protein levels in Qu+AA treated cells. Together, these data indicate that while Qu has been reported to have some beneficial properties as a chemopreventive agent, it is also capable of inducing c-fos expression, a cellular event important for the promotion phase of tumor development, if it is not stabilized.

Keywords

quercetin; c-Fos; ultraviolet light; p38; mTOR

Introduction

Currently, there is a major focus on the use of natural products as chemopreventive agents for non-melanoma skin cancer (NMSC) and other types of cancer because of their anti-oxidant and anti-inflammatory properties in addition to their potential to inhibit numerous signaling

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pathways involved in cell proliferation, transformation, migration, and survival (1–3). Flavonoids represent one of the most actively studied classes of molecules for their potential to prevent cancer. However, the cellular effects of these compounds are often very complicated due to the general lack of specificity for any particular intracellular target, and the end effect on cell fate is often the combination of multiple cellular activities. In the current study, we investigated some of the cellular effects of quercetin (Qu; 3,3',4',5,7-pentahydroxyflavone) on ultraviolet B (UVB)-induced signaling in human keratinocytes. Our original interest in Qu was with regard to the compound's known phosphatidylinositol 3-kinase (PI3-K) inhibitory activity, and we initially reported that a concentration of 50 μ M was required for consistent suppression of UVB-induced PI3-K activation in HaCaT human keratinocytes due to the cell-independent rapid degradation of Qu in aqueous cell culture medium (4). At this concentration, we demonstrated that Qu actually displays pro-oxidant activity, an effect that could be inhibited by including 1 mM ascorbic acid (AA) in the growth medium to stabilize Qu. However, this reduced the pro-apoptotic effect of Qu, leading to the interpretation that the generation of reactive species during Qu degradation could be a beneficial action through increased killing of initiated cells. On the contrary, our current findings suggest that there are significant cellular consequences to Qu degradation that could lead to elevated promotion of the initiated cells that survive Qu treatment, which raises questions regarding the chemopreventive efficacy in UV-induced skin cancer.

Chronic exposure to ultraviolet light is the primary cause of NMSC, the most common type of cancer in the U.S. (5). UVB (280–320 nm) comprises between 1% and 10% of the total UV light that reaches the Earth's surface. UVB is known to act as a complete carcinogen, and our laboratory and others have demonstrated that UVB-induced activation of the transcription factor activator protein-1 (AP-1) plays a functional role in UVB-induced tumor promotion (6). AP-1 stimulates cell proliferation through regulation of cell cycle protein expression [reviewed in (7)] and also controls cellular transformation since inhibition of AP-1 has been shown to block the effect of tumor promoting agents (6,8–12).

The AP-1 transcription factor complex is comprised of dimers of Jun and Fos protein family members. The exact constituents depend on the stimulus or the physiological condition, and the different dimer pairs can activate different sets of genes [reviewed in (13)]. In HaCaT keratinocytes, the UVB-activated AP-1 complex is composed of c-Fos and JunD (14). While JunD expression is unaffected by UVB irradiation, c-Fos levels have been shown to increase with UVB treatment in a manner that correlates with AP-1 activation, suggesting that alterations in c-Fos protein levels are a major driving factor behind the level of AP-1 activity. In addition, c-Fos expression has been shown to be required for malignant progression of skin tumors (15). Increased cellular oxidative stress can lead to p38 activation and ultimately c-fos gene expression by stimulating cyclic AMP response element binding (CREB) protein binding to the c-fos promoter (16–18). In the current study we worked to determine the effect of Qu treatment on c-Fos protein levels in UVB-irradiated HaCaT keratinocytes. We report that Qu actually increased c-Fos protein levels and potentiated the UVB response. This effect could be prevented completely by stabilizing Qu with AA, since treatment with Qu+AA completely inhibited the UVB-induced increase in c-fos promoter activity and also inhibited protein synthesis by suppressing mTOR signaling. These findings demonstrate an important cellular effect of Qu resulting from its pro-oxidant activity. Although we previously reported that Qu treatment may be beneficial by killing initiated cells (4), failure to stabilize Qu could have detrimental effects that limit any chemopreventive efficacy of the compound. Even stabilization of Qu with AA was shown to be a transient event, which might not prevent the tumor promoting event of elevated c-Fos expression that occurs as Qu degrades. Importantly, these findings highlight some of the complexities of working with natural products and suggest that some of these compounds may require a significant amount of characterization to validate them as chemopreventive agents and avoid deleterious side effects.

Materials and Methods

Materials

Quercetin dihydrate and ascorbic acid were purchased from Sigma-Aldrich (St. Louis, MO.). Phospho-specific antibodies for p38 (T180/Y182), CREB (S133), Akt (S473), mTOR (S2448), p70S6K (T389), and 4EBP (S65) were from Cell Signaling Technology (Danvers, MA.). c-Fos antibodies were purchased from Santa Cruz Biotechnology (Santa Cruz, CA). Real-time PCR probes (c-Fos: Hs00170630_m1 and GAPDH: Hs99999905_m1) and TaqMan reagents were purchased from Applied Biosystems (Foster City, CA). Actinomycin D and MG-132 were from EMD Biosciences/Calbiochem (San Diego, CA).

Cells

The human keratinocyte cell line, HaCaT, was established from cells obtained from adult sun damaged skin and have been described previously (19–21). HaCaT cells contain UV-signature mutations and express mutant dysfunctional p53 and a defective NF- κ B signaling pathway, which are common findings in UV-initiated keratinocytes in human skin. However, these cells maintain p38, PI3-K and AP-1 signaling pathway functionality compared to normal human keratinocytes and were thus chosen as an appropriate cell line to study UVB-induced c-Fos expression and upstream signaling events. FL-30 cells were developed from HaCaT cells and stably express firefly luciferase driven by a full length c-fos promoter element (14). The cells were cultured in Dulbecco's modified Eagle's Medium (DMEM) with 10% fetal bovine serum and 100 units/ml penicillin/streptomycin at 37°C and in 5% CO₂. The cells were cultured to 80–90% confluence and maintained in serum-free DMEM (SFM) for 24 h prior to UVB exposure.

UVB irradiation of HaCaT cells

HaCaT cells were pretreated with quercetin and/or ascorbate for 1 hr prior to UVB irradiation. The concentration of 50 μ M quercetin was chosen based on our previously published study (4) and the fact that this concentration is feasible for chemoprevention studies using topical Qu. After incubation, HaCaTs were washed once in PBS and irradiated with a dose of 250 J/m² using a bank of two SF20 UVB lamps (National Biological Corp.) providing a peak emission of 313 nm. Control cells were treated in the same manner and mock irradiated. Following irradiation, HaCaT cells were again washed with PBS and returned to DMEM containing appropriate drug treatments. In experiments using MG-132, there was no initial pretreatment and MG-132 was only used following exposure to UVB.

Western blotting

Cells were lysed in RIPA buffer and protein concentration was determined as previously described (4). A total of 40 μ g protein was resolved by SDS-PAGE and transferred to a PVDF membrane. Membranes were blocked in Tris buffered saline containing 0.1% Tween-20 (TBST) and either 5% nonfat dry milk or 5% BSA (for phospho-specific antibodies). After washing in TBST, membranes were incubated with HRP-conjugated secondary antibodies and then washed extensively in TBST. Antigen-antibody complexes were detected using Amersham ECL Detection Reagent (GE Healthcare, Buckinghamshire, UK). Densitometric analysis was performed using NIH ImageJ software and density values were normalized to the untreated control, which was assigned a value of 1.0.

c-fos-Luciferase assay

FL-30 cells were treated and UVB-irradiated in triplicate for each independent experiment. Six hours after exposure to UVB, cells were lysed and a total of 20 μ g protein per sample replicate were assayed for luciferase activity according to the manufacturer's instructions for

the Luciferase Assay System (Promega, Madison, WI) using a TD 20/20 luminometer (Turner Designs, Sunnyvale, CA). The experiment triplicates were averaged and the means from each independent experiment were subsequently averaged and analyzed by Student's *t* test for statistical significance.

RNA extraction and real-time PCR analysis

For the measurement of steady state RNA levels, cells were treated as indicated above. In RNA stability assays, cells were pretreated for 30 min with 5 µg/ml actinomycin D prior to treatment with Qu and AA for the indicated times. Cellular RNA was isolated from treated HaCaT cells by phenol-chloroform extraction using the Ambion ToTally RNA Kit (Applied Biosystems, Foster City, CA) according to manufacturer's instructions. Extracted RNA was treated with DNase using the Ambion DNA-free Kit and purified using the RNeasy MinElute Cleanup Kit from Qiagen (Valencia, CA). Purified RNA was then reverse transcribed using the High Capacity cDNA Reverse Transcription Kit from Applied Biosystems. Real-time PCR was performed using an ABI PRISM 7700 Sequence Detection System and TaqMan probes specific for human *c-fos* and GAPDH. Relative expression was calculated to be $2^{-\Delta\Delta C_t}$, where ΔC_t is equal to the difference in cycle number (C_t) between *c-Fos* and GAPDH for each sample and $\Delta\Delta C_t$ is equal to the difference between ΔC_t for each treatment and the untreated control. Each individual assay was performed in triplicate for each gene product and a total of 3 independent experiments were performed.

Electromobility shift assay

After treatment and UVB-irradiation, nuclear extracts were isolated from HaCaT cells as described previously (22). The following sequences derived from the human *c-fos* promoter containing the CRE were used to generate double stranded oligos for EMSA analysis: CRE1: 5'-GAGCCCGTGACGTTTACACT-3', and CRE2: 5'-TGAGTGTAACGTCACGG-3'. They were annealed and 5'-overhangs were labeled with [³²P]dCTP (Perkin Elmer, Waltham, MA). For gel shift assays, 5 µg of nuclear protein extracts were incubated at room temperature for 20 min with a mixture containing 10 mM Hepes (pH 7.9), 0.1 mM EDTA, 50 mM KCl, 2.5 mM DTT, 10% glycerol, 0.5% Triton X-100, and 0.05 mg/ml polyIdC. Labeled probe was added and the reactions were incubated at room temperature for another 30 min. Reactions were then fractionated on a non-denaturing 6% polyacrylamide gel in 0.25 X Tris borate EDTA. The gels were dried and visualized with an MD Storm phosphorimager (Molecular Dynamics, Sunnyvale, CA).

Results

Quercetin induces p38 and CREB phosphorylation leading to *c-fos* promoter activity unless stabilized by ascorbic acid

We previously demonstrated that quercetin (Qu) is unstable in DMEM, but can be stabilized for several hours by inclusion of 1 mM ascorbic acid (AA) in the medium (4). Western analysis of protein from HaCaT keratinocytes indicated that 250 J/m² UVB induced p38 phosphorylation (Figure 1A). Treatment with 50 µM Qu increased p38 phosphorylation in both mock irradiated and UVB-irradiated HaCaT cells compared to the controls. When 1 mM AA was included in the growth medium, Qu treated cells displayed no increase in phospho-p38 levels, and the response to UVB was completely lost. CREB, a substrate downstream of p38 signaling, displayed a nearly identical response to UVB and Qu as p38, where there were increases in CREB phosphorylation with both UVB and Qu alone and in combination and a significant reduction in the UVB response when cells were treated in the presence of AA (Figure 1B). Total p38 and CREB levels were unaffected by Qu, AA or UVB treatment.

Transcriptional activation of c-fos is primarily driven by binding of CREB to the cyclic AMP response element (CRE) in the promoter region (17). We examined the effects of Qu on c-fos promoter activity using FL-30 cells, a HaCaT cell line that has been stably transfected with a firefly luciferase reporter construct where expression of luciferase is driven by a full length human c-fos promoter. Irradiation of FL-30 cells with 250 J/m² UVB resulted in a 5.1-fold increase in luciferase expression over control, while Qu treatment alone induced a 2.7-fold increase in luciferase expression (Figure 2A). Qu treatment of UVB-irradiated HaCaT cells significantly reduced luciferase expression compared to UVB alone, although luciferase levels were still significantly increased over the untreated control (2.1-fold over control). In this assay system, luciferase expression is dependent on both gene induction and protein translation, and the potential for Qu to act on both of these pathways will be addressed below.

The effects of Qu on the c-fos promoter were further analyzed by performing an EMSA using probe containing the CRE from the c-fos promoter. We determined that UVB irradiation, treatment with Qu alone, and combining Qu treatment with UVB irradiation all increased CRE binding and that CRE binding was highest in nuclear extracts from UVB-irradiated HaCaT cells (Figure 2B). The level of CRE binding that resulted from these treatments followed a similar trend to that seen in the c-fos-luciferase assay in Figure 2A. Similar to p38 and CREB phosphorylation, inclusion of AA in the medium during the time of treatment abrogated the effect of Qu on CRE binding. Taken together, these data indicate that Qu induces c-fos promoter activity to levels greater than that seen in untreated cells, but partially blocks UVB-induction of the c-fos promoter. Furthermore, these results suggest that co-treatment with Qu and AA (Qu+AA) suppresses both basal and UVB-induced promoter activity.

Quercetin increases c-fos mRNA, but stabilization with ascorbic acid prevents UVB-induced c-Fos mRNA expression

Total cellular RNA was isolated from Qu- and UVB-treated HaCaT cells 2 hr after exposure to UVB, and reverse transcribed cDNA was analyzed by real-time PCR with a probe specific for human c-fos. UVB-irradiated cells displayed an 8.4-fold increase in c-fos message compared to mock irradiated controls (Figure 3A). Qu treatment also increased the level of c-fos mRNA by 50-fold over control, a level much greater than that induced by UVB and potentiated the UVB response (126-fold over control). Interestingly, AA supplementation also increased steady state mRNA levels in both mock- and UVB-irradiated HaCaT cells, a finding that differed from the effect of AA on c-fos promoter activity. Although this increase was not statistically significant compared to control levels, treatment with both Qu+AA did in fact result in a moderate increase.

Since this increase in steady state c-fos mRNA caused by supplementation of DMEM with AA could not have been the result of increased transcriptional activity, we next investigated the possibility that AA was increasing the stability of c-fos mRNA. HaCaT cells were treated with actinomycin D to inhibit mRNA synthesis and subsequently treated with Qu, AA, or both. Through real-time PCR analysis, we determined that AA supplementation alone resulted in a marked reduction in the rate of c-fos mRNA decay compared to control (Figure 3B). Qu treatment had no significant effect on mRNA stabilization alone, but actually reduced the effect of AA.

Quercetin potentiates UVB-induced c-Fos protein expression in HaCaT keratinocytes, and ascorbate-stabilized quercetin inhibits increases in c-Fos protein levels

Our laboratory has previously reported that UVB irradiation increases the levels of c-Fos protein (14,17,18). We analyzed c-Fos protein levels by Western analysis and confirmed that 250 J/m² UVB did increase expression over the level expressed in mock irradiated HaCaT cells (Figure 3C). Treatment with Qu alone also increased c-Fos protein to a level comparable to

that induced by UVB, and treatment of cells with both Qu and UVB resulted in level of c-Fos that was substantially increased over either Qu or UVB treatment alone. These data corroborate our previous finding for the Qu and UVB effects on mRNA expression. Contrary to the effect of AA on steady state mRNA levels, there was no effect of AA treatment alone on c-Fos protein expression in either mock- or UVB-irradiated cells compared to their respective controls. Interestingly, treatment with Qu+AA showed only a slight increase in c-Fos protein expression in mock-irradiated cells, and a complete suppression of c-Fos protein expression even when cells were irradiated with UVB, suggesting that the stabilization of c-fos mRNA did not translate to increased protein.

Quercetin blocks mTOR signaling and inhibits c-Fos protein translation

One explanation for the apparent disconnect between mRNA and protein levels is that Qu independently affects protein synthesis and mRNA expression. To address this possibility, we investigated the effects of Qu on mTOR signaling since this pathway is known to regulate *de novo* protein synthesis and since we and others have reported that Qu has inhibitory activity against PI3-K, a kinase upstream of Akt and mTOR activation (4,23,24). We determined that mTOR phosphorylation at Ser2448, an activating phosphorylation, was slightly increased with UVB (Figure 4A), as was phosphorylation of the downstream mTOR effectors 4EBP (S65; Figure 4B) and p70S6K (T389; Figure 4C). Qu suppressed phosphorylation of each of these proteins in both mock and UVB-irradiated cells, and when the cells were treated with Qu in the presence of AA, the level of suppression was increased, a finding that is consistent with our previous report that AA stabilizes Qu in aqueous medium (4).

Lastly, we treated HaCaT cells with Qu+AA in the presence of the proteasome inhibitor MG-132 in order to prevent c-Fos turnover and more specifically address the effects of stabilized Qu on c-Fos protein synthesis (Figure 4D). HaCaT cells treated with MG-132 displayed significantly increased c-Fos protein levels due to inhibition of protein degradation, which resulted in a reduction of the effect of Qu on UVB-induced c-Fos protein levels. However, cells treated with Qu+AA still displayed low levels of c-Fos protein in the presence of MG-132 resulting from a sustained reduction of c-Fos translation.

Discussion

Non-melanoma skin cancer (NMSC) is the most commonly diagnosed of all human malignancies and is the cause of 2,000 deaths annually and millions of dollars in health care costs. Clearly, new treatment strategies are needed to prevent the development of precancerous actinic keratoses and squamous cell tumors caused by prolonged exposure to ultraviolet light. Quercetin is currently being evaluated as a potential chemopreventive agent in multiple types of cancer in part because it has established PI3-K and MAPK inhibitory activities [for review, see (25)]. Our initial interest in Qu was specifically due to this activity, since it has been shown that the PI3-K and MAPK signaling pathways are activated in response to UVB irradiation and both PI3-K and p38 are upstream of c-Fos expression and AP-1 activation (16,26,27). Natural products and small molecule inhibitors of these pathways would protect against promotion of initiated cells by reducing AP-1 activation and could be potentially useful in cancer prevention.

The findings from the current study are summarized in Figure 5. We determined that Qu potentiated the UVB-induced c-fos gene expression, a direct result of increased p38 and CREB phosphorylation. The inhibitory effect of Qu on PI3-K was not sufficient to prevent the elevation of c-Fos protein levels that directly resulted from increased gene expression. These effects of Qu could be prevented by stabilizing the compound with AA, which resulted in a complete inhibition of UVB-induced c-fos mRNA and protein expression and all upstream signaling. These findings exemplify the difficulties that can arise when working with natural products.

We previously reported that Qu could potentially act as a chemopreventive agent due to its ability to kill initiated cells through PI3-K inhibition and pro-oxidant effects (4). However, considering the results of the current study, Qu degradation could potentially have the consequence of increasing AP-1 activity and negating the chemopreventive effect. Dietary Qu has been shown to be effective in preventing cancer in animal models in colon (28–33), breast (34), and lung tissues (35), as well as squamous cell carcinoma (SCC) of the tongue (36). In NMSC models, the effects of Qu are unclear. In one study, dietary Qu had no effect on UV-induced SCC (37), while topically-applied Qu decreased papilloma formation in a chemically-induced skin cancer mouse model (38). We have tested topical Qu in a UVB-induced skin cancer model and found no significant reduction of SCC incidence, tumor multiplicity, or tumor burden compared to vehicle-treated controls (data not shown). There are several possible explanations for the differences in these reported results. First, orally administered Qu may not adequately distribute to the skin to elicit any preventative effect. Second, it is possible that Qu is more effective against chemically-induced skin cancer, although Kato et al. (38) used a topical dose that was 5 times more concentrated than the dose we used (5% vs. 1%, respectively). Interestingly, a recent study evaluating myricetin, a flavonol with great structural similarity to Qu, demonstrated significant efficacy in preventing UVB-induced skin tumor formation when much lower amounts were used (0.001–0.003% myricetin/treatment vs. 1% quercetin/treatment in our study) (39). There have been no reports describing myricetin stability, but we have evidence that myricetin does not induce c-fos expression in cultured HaCaT cells when used at the same concentration as Qu in the current study (data not shown). Lastly, the formulation used to deliver Qu topically may affect the long-term stability of Qu in the skin, and as demonstrated in the current study, Qu degradation can affect cell signaling and the cellular response to UVB. Overall, these studies suggest that the effectiveness of Qu is dependent on the specific type of cancer and the method of administration. Furthermore, our findings suggest that any aqueous-based topical formulation of Qu developed for human use would have to contain a Qu stabilizing agent that significantly slowed Qu degradation in order to prevent the detrimental effects of Qu degradation. If Qu is to be pursued as a topical chemopreventive agent for non-melanoma skin cancer, further studies are needed to determine if there are optimal conditions under which Qu can be effective.

Our findings, combined with the large number of reported cellular effects of various natural products, highlight the complexities and the careful considerations required to properly evaluate these compounds for their chemopreventive potential, particularly with regard to the proper dosing required to achieve desired experimental or clinical outcomes. Some of these studies highlight an important conflict in the reports of *in vitro* and *in vivo* effects of Qu. For example, pioneering studies by Nagao et al. (40) identified numerous flavanoids, including quercetin, with mutagenic potential. However, oral Qu has been demonstrated to display no carcinogenic effect in rats (41), despite the known relationship between mutagenicity and carcinogenicity. In addition, other studies using different *in vitro* models demonstrate conflicting findings to our own. Ying, et al. (42) recently published a study demonstrating that 10 μ M Qu blocks ICAM expression in part through inhibition of c-fos induction. As previously stated, we had determined that this dose was ineffective at preventing UVB-induced PI3-K activity, our primary goal, but raising the concentration to an effective level caused increased sensitization to UVB-induced c-fos expression. This property may be detrimental to the chemopreventive effect of Qu, but might be useful for other purposes, like increasing the sensitivity of tumor cells to agents like cisplatin as reported by Sharma, et al. (43). These facts all indicate that Qu is capable of eliciting different effects that need to be well characterized to fully understand how the compound might be useful in preventing or treating cancer. Quercetin is an excellent example of how natural products need to be thoroughly examined before drawing premature conclusions regarding their preventive or therapeutic efficacy.

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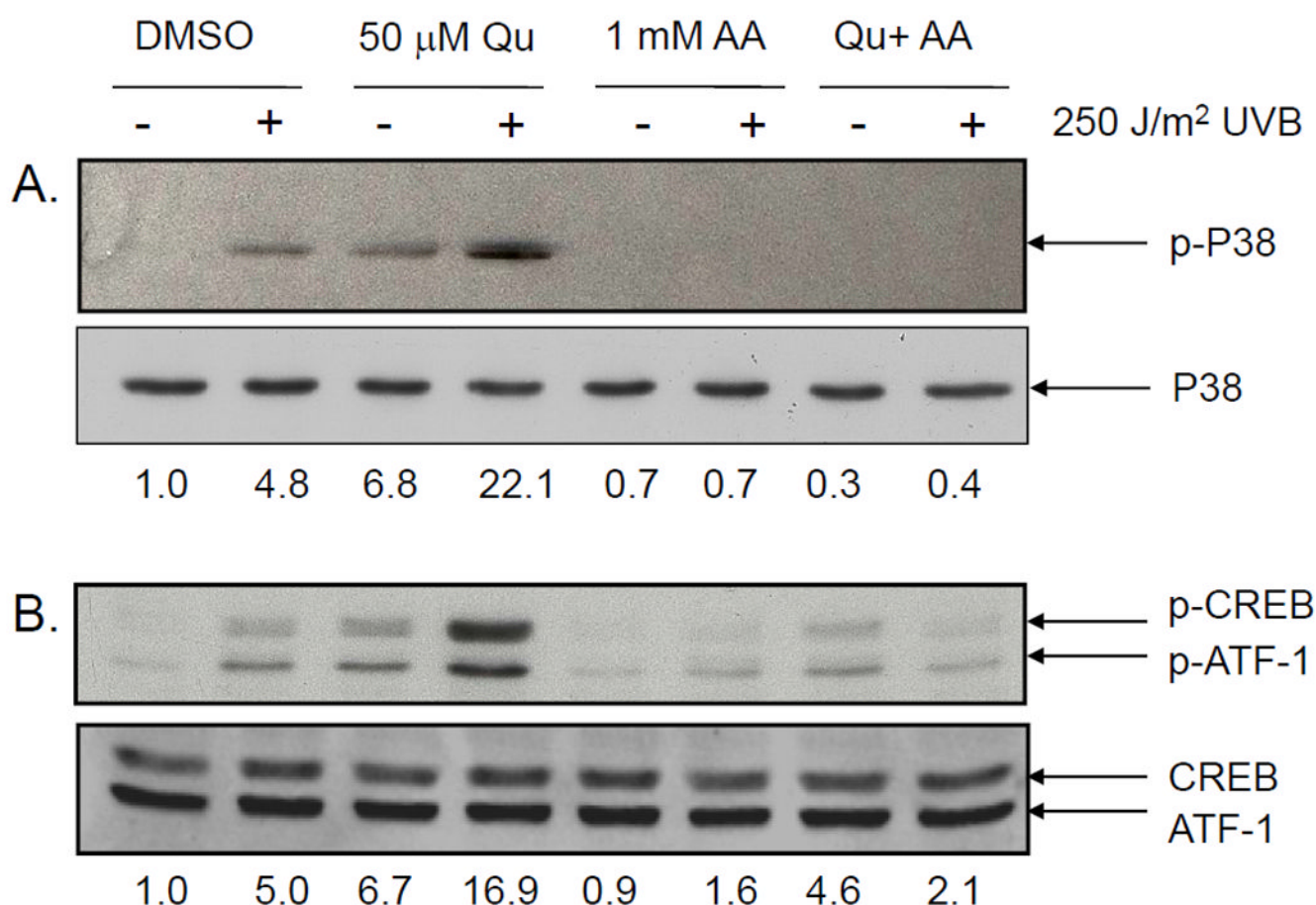


Figure 1. Quercetin induces p38 and CREB phosphorylation unless stabilized by ascorbic acid
 HaCaT cells were treated with 50 μ M Qu for 1 hr prior to exposure to 250 J/m² UVB irradiation and for 2 hr post UVB. Cells were either treated in normal DMEM or in DMEM supplemented with 1 mM ascorbic acid. (A) p38 phosphorylation was induced by UVB and Qu alone, and the response was potentiated in cells treated with both UVB and Qu. Treatment with 1 mM ascorbic acid completely inhibited the UVB- and Qu-induced increase in p38 phosphorylation. (B) CREB phosphorylation in UVB-irradiated and Qu-treated HaCaT cells was similar to that seen with p38 phosphorylation. Both Qu and UVB treatment individually increased CREB phosphorylation and the combined treatment potentiated the response. Again, AA significantly reduced CREB phosphorylation in response to both UVB and Qu and prevented the potentiated response seen in cells treated with both Qu and UVB. The antibodies used for both p-CREB and CREB recognize p-ATF-1 and ATF-1, respectively, as well. Relative band densities are represented numerically immediately below each set of blots.

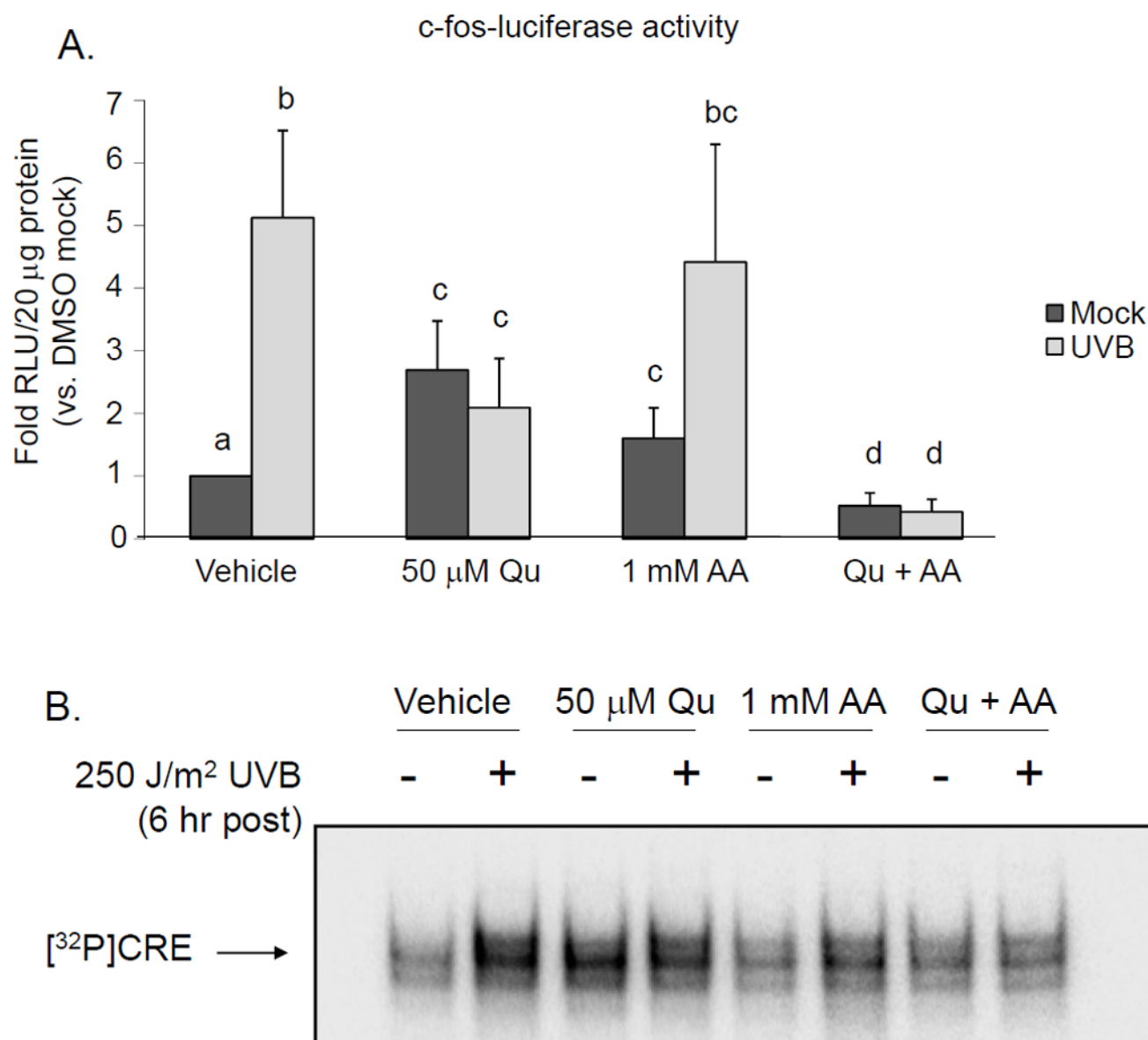


Figure 2. Quercetin stimulates c-fos promoter activity and CRE binding

Cells were treated with 50 µM Qu +/- 1 mM AA for 1 hr prior to irradiation with 250 J/m² UVB and following irradiation for 6 hr. (A) UVB induced a 5.1-fold increase in c-fos-luciferase activity over untreated controls. Treatment with Qu induced a 2.7-fold increase, but also inhibited UVB-induced c-fos-luciferase activity by 74%. AA treatment significantly increased the basal level of c-fos-luciferase activity by 1.6-fold, but did not affect the response to UVB. However, supplementation with AA and treatment with Qu suppressed basal c-fos-luciferase activity and completely inhibited the response to UVB. Data represent the mean from 4 independent experiments and statistically significant differences between groups ($p < 0.05$) are indicated by labels containing different letters. Any two treatment conditions with a label containing the same letter are not significantly different. (B) CRE binding was assessed using nuclear lysates from Qu-treated HaCaT cells and a double stranded oligo containing the CREB binding sequence. CRE binding was increased in lysates extracted from UVB-irradiated cells and from Qu-treated cells. Qu apparently did not affect CRE binding induced by UVB, as it

was not different from Qu alone. AA had little to no effect on CRE binding, but Qu+AA completely suppressed basal and UVB-induced CRE binding. These data are representative of 3 independently performed experiments.

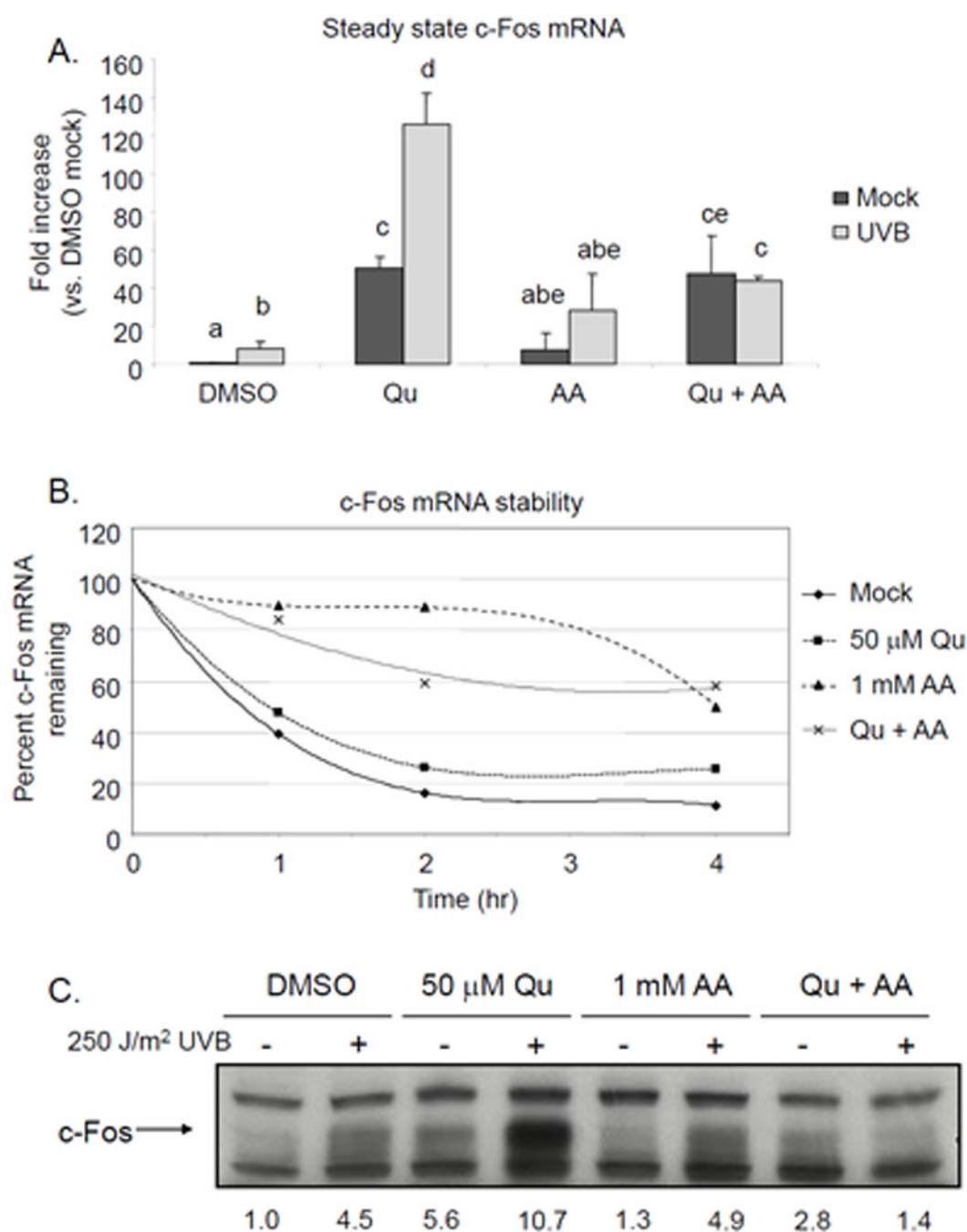


Figure 3. Quercetin treatment increases c-fos mRNA expression and potentiates UV-Binduced c-Fos protein production

(A) HaCaT cells were pretreated with 50 μ M Qu +/- 1 mM AA 1 hr prior to irradiation with 250 J/m² UVB and for 2 hr post-UVB, at which point RNA was extracted and purified. Real time RT-PCR was performed using probes specific for c-fos and GAPDH. The fold increase of steady state c-fos mRNA was determined as indicated in *Materials and Methods*. UVB induced an 8.4-fold increase in c-fos mRNA and Qu alone induced a 50-fold increase. Qu treatment in UVB-irradiated cells resulted in a potentiated response, where the induction of c-fos mRNA was equal to 126-fold over control cells. AA treatment alone raised steady state c-fos mRNA levels by 7.8-fold and the UVB-induced increase was 28-fold. AA combined with

Qu treatment raised steady state c-fos levels 47-fold over control and the UVB response in these cells was completely inhibited, with c-fos levels remaining at 43-fold over control. These data are means from 3 independently performed experiments and statistically significant differences between groups ($p < 0.05$) are indicated by labels containing different letters. Any two treatment conditions with a label containing the same letter are not significantly different. (B) HaCaT cells treated with Qu, AA or both were concurrently treated with actinomycin D to prevent the synthesis of new mRNA. At the indicated time points, mRNA was extracted and real time RT-PCR was performed and data were analyzed as indicated above. Data were normalized to 0 hr, set to 100% for each treatment condition. AA stabilized c-fos mRNA, which likely contributed to the increase in steady state c-fos mRNA detected in the AA treated samples in (A). Qu did not stabilize c-fos mRNA and reduced the stabilizing effect of AA. These data are a representative sample from 2 independently performed experiments. (C) HaCaT cells were pretreated with 50 μ M Qu \pm 1 mM AA for 1 hr prior to irradiation with 250 J/m² UVB and following irradiation for 6 hr, at which point protein lysates were extracted and Western analysis was performed using specific antibodies for c-Fos. Treatment with either Qu or UVB irradiation alone induced c-Fos protein expression over control, but cells treated with both Qu and UVB irradiation displayed a potentiated increase in c-Fos protein. Supplementation with AA had no effect on c-Fos protein levels in mock and UVB-irradiated HaCaT cells. However, Qu treatment in AA-supplemented cells completely blocked the UVB-induced increase in c-Fos protein levels. Relative band densities are represented numerically immediately below the blot. Data are representative of 3 independently performed experiments.

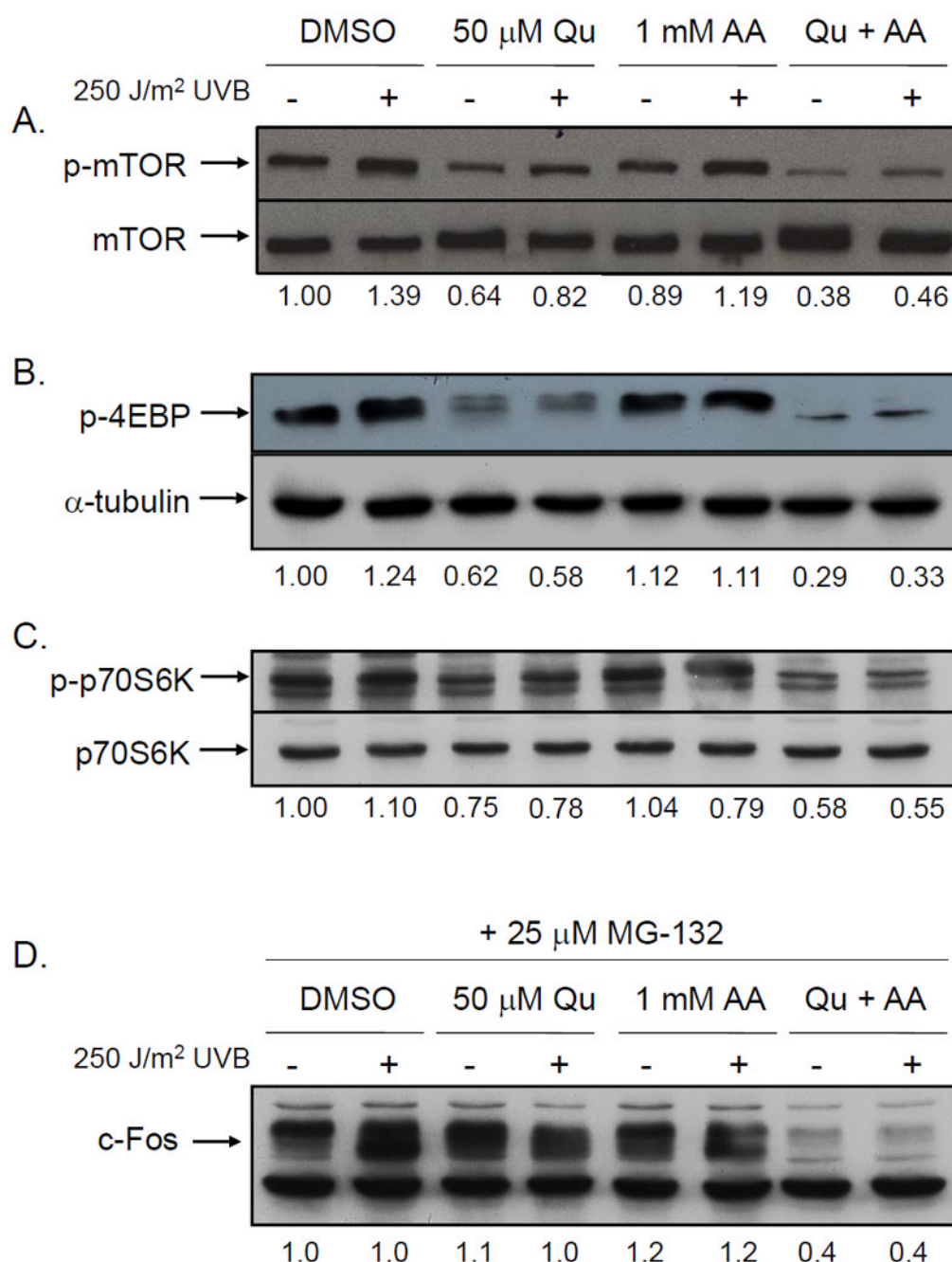


Figure 4. Quercetin inhibits mTOR signaling, but only blocks c-Fos protein synthesis when stabilized by AA

HaCaT cells were pretreated with 50 μ M Qu \pm 1 mM AA for 1 hr prior to irradiation with 250 J/m² UVB and following irradiation for 2 hr. Protein lysates were extracted and Western analysis was performed using specific antibodies for (A) phospho-mTOR, (B) phospho-4EBP, and (C) phospho-p70S6K. UVB irradiation increased phosphorylation of each of these signaling intermediates, whereas treatment with Qu suppressed basal and UVB-induced phosphorylation. AA supplementation of DMEM alone had no effect on either mock or UVB-induced phosphorylation. However, inhibition of phosphorylation by Qu was enhanced in the presence of AA for mTOR, 4EBP, and p70S6K. Total mTOR and α -tubulin levels were

unchanged across all treatment conditions. Relative band densities are represented numerically immediately below each set of blots. Data are representative of 3 independently performed experiments. (D) HaCaT cells were pretreated with 50 μ M Qu \pm 1 mM AA for 1 hr prior to irradiation with 250 J/m² UVB and following irradiation for 6 hr. Immediately following exposure to UVB, cells were treated with 25 μ M MG-132 to inhibit proteasome activity. Following the treatment period, cell lysates were prepared and Western analysis was performed to determine the effect of treatment on c-Fos protein levels. Treatment with MG-132 increased c-Fos levels in all treatment conditions compared to cells with no MG-132 treatment (Figure 4) except for in cells treated with Qu+AA, in which there was no increase. Relative band densities are represented numerically immediately below the blot. Data are representative of 3 independently performed experiments.

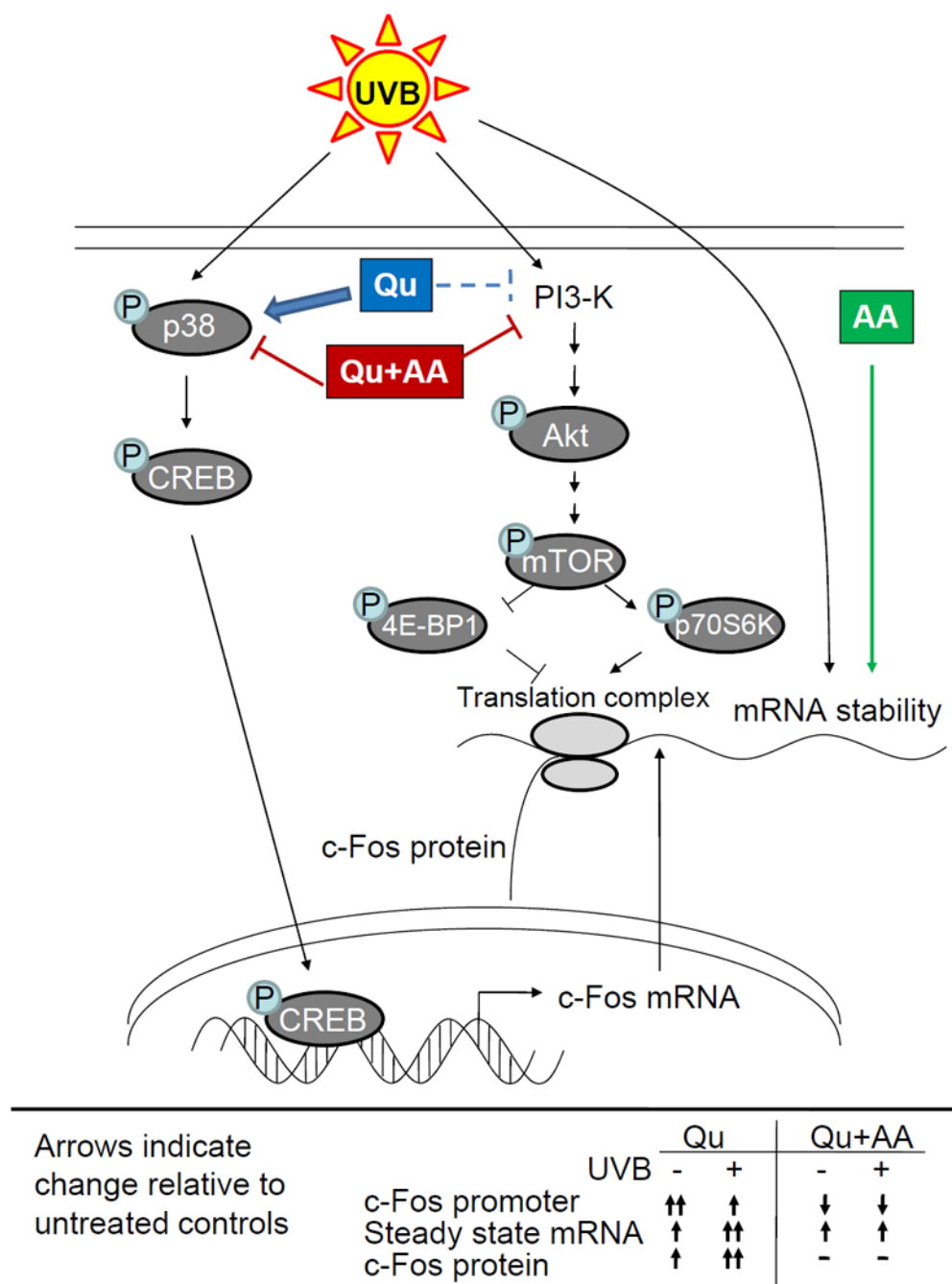


Figure 5. Proposed mechanism of quercetin-mediated induction and inhibition of c-Fos mRNA and protein expression

Quercetin induces c-fos gene expression through upstream activation of p38 and CREB and potentiates the response to UVB, leading to an increase in steady state c-fos mRNA levels. Co-treatment with Qu and ascorbic acid blocks p38 and CREB activation and reduces c-fos promoter activity (confirmed by the observation that CREB/CRE binding is reduced). c-fos mRNA levels are actually increased after treatment with Qu+AA as a result of increased c-fos mRNA stability, but the response to UVB is completely inhibited. PI3-K, also activated by UVB and upstream of mTOR, is known to be inhibited by Qu and with greater efficacy by Qu+AA. Simultaneous inhibition of mTOR signaling by Qu+AA prevents c-Fos protein synthesis

even in the presence of increased mRNA levels. With Qu alone, inhibition of this pathway is less efficacious and more transient as a result of Qu degradation, leading to increased c-Fos protein driven by significantly increased levels of c-fos mRNA.