Chrysin, a natural flavonoid enhances steroidogenesis and steroidogenic acute regulatory protein gene expression in mouse Leydig cells

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Abstract

During the aging process of males, testosterone biosynthesis declines in testicular Leydig cells resulting in decreases in various physiological functions. To explore the possibility of delaying the decline using food supplements, we have studied steroidogenic effects of a natural flavonoid, chrysin, in mouse Leydig cells. Chrysin dramatically increased cyclic AMP (cAMP)-induced steroidogenesis in MA-10 mouse Leydig tumor cells. This result was confirmed using Leydig cells isolated from mouse testes. The steroidogenic effect of chrysin is not associated with an increase in expression of the P450 side-chain cleavage enzyme, required for the conversion of cholesterol to pregnenolone. In addition, when 22(R)hydroxylcholesterol was used as a substrate, chrysin induced a non-significant increase in steroid hormone, suggesting that the majority of the observed increase in steroidogenesis was due to the increased supply of substrate cholesterol. These observations were corroborated by showing that chrysin induced a marked increase in the expression of steroidogenic acute regulatory (StAR) protein, the factor that controls mitochondrial cholesterol transfer. Also, chrysin significantly increased StAR promoter activity and StAR mRNA level. Further studies indicated that this compound depressed expression of DAX-1, a repressor in StAR gene transcription. In the absence of cAMP, chrysin did not increase steroidogenesis. However, when a sub-threshold level of cAMP was used, StAR protein and steroid hormone were increased by chrysin to the levels seen with maximal stimulation of cAMP. These results suggest that while chrysin itself is unable to induce StAR gene expression and steroidogenesis, it appears to function by increasing the sensitivity of Leydig cells to cAMP stimulation.

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Introduction

It is well documented that blood testosterone concentration declines progressively during the course of male aging, a phenomenon that is associated with decreased bone density, muscle mass and strength, sexual function, and other physiological functions (Vermeulen 2000, Hogervorst et al. 2001, Matsumoto 2002, Kaufman & Vermeulen 2005). Several longitudinal studies on age-related changes in blood testosterone indicated that the incidence of hypogonadism in males increased with age (Harman et al. 2001, Feldman et al. 2002). Therefore, delaying the decline in blood testosterone is clinically significant for improving the health of aging males.

Testosterone is principally synthesized in testicular Leydig cells under the regulation of luteinizing hormone (LH) and then released into blood (Miller 1988). In the first step of testosterone biosynthesis, the substrate cholesterol is metabolized by the steroidogenic enzyme, P450 side-chain cleavage enzyme (P450scc), and converted to pregnenolone that is further converted to testosterone through the action of several steroidogenic enzymes (Payne & Youngblood 1995, Payne & Hales 2004). To initiate steroidogenesis, cholesterol must be transferred from the outer to inner membrane of mitochondria, since the P450scc enzyme is located on the matrix side of the inner mitochondrial membrane and the cholesterol content of the inner membrane is limited (Churchill & Kimura 1979, Privalle et al. 1983). However, the aqueous space between the outer and inner mitochondrial membranes acts as a barrier to cholesterol transfer because of its hydrophobic nature. As a result, diffusion of cholesterol through this water barrier is very slow and is not able to supply sufficient substrate for adequate testosterone production (Schumacher et al. 1985, Phillips et al. 1987). Therefore, the mitochondrial cholesterol transfer becomes the rate-limiting step in testosterone biosynthesis. We have
previously reported that the steroidogenic acute regulatory (StAR) protein plays a critical function at this step by facilitating the cholesterol transfer (Clark et al. 1994, Lin et al. 1995, Wang et al. 1998). It was well demonstrated that LH stimulation of Leydig cells induces StAR gene expression, and the level of StAR protein in Leydig cells significantly affects testosterone production (Stocco 2001). However, StAR protein level also declines during Leydig cell aging, and mitochondrial cholesterol transfer in aged rat Leydig cells is defective (Leers-Sucheta et al. 1999, Luo et al. 2001, Culty et al. 2002). These studies indicated that the age-related decrease in StAR gene expression is involved in the decrease in testosterone biosynthesis in aging Leydig cells. Our previous studies suggested that it is possible to delay the decline in testosterone biosynthesis by increasing the StAR gene expression (Wang et al. 2003a, 2005). Based on the observations in these studies, we were able to reverse the age-related declines in StAR protein expression and blood testosterone concentration by feeding the aging rats with a drug to inhibit the activity of cyclooxygenase-2 (COX2), an enzyme whose activity is inhibitory to StAR gene expression (Wang et al. 2005). However, aging is a long process and the long-term application of pharmacological levels of COX2 inhibitors is limited by their potential side effects on health. Therefore, we have attempted to identify natural compounds in food and food supplements that have fewer side effects and effectively increase StAR gene expression and testosterone biosynthesis in Leydig cells. A natural flavonoid, chrysin, was previously used to increase blood testosterone (Brown biosynthesis in Leydig cells. A natural flavonoid, chrysin, was effectively increase StAR gene expression and testosterone in food and food supplements that have fewer side effects and Therefore, we have attempted to identify natural compounds inhibitors is limited by their potential side effects on health.

Steroid hormone production

MA-10 cells were cultured for 30 min in serum-free Waymouth’s medium in six-well plates with or without chrysin (as described in the Figure Legends) and then 0.05 mM dbcAMP was added to the culture for 6 h. The medium was collected at the end of each experiment and stored at −80 °C. Progesterone concentrations in the medium were determined by RIA (Resko et al. 1974).

Mouse Leydig cell isolation and cell culture

B6–129S7 mice aged 3 months (Jackson Laboratory, Bar Harbor, ME, USA) were euthanized in a chamber filled with carbon dioxide followed by cervical dislocation. Testes were collected from the mice. Leydig cells were isolated from the testes using density gradient centrifugation as described previously (Hutson et al. 1996). The cells were cultured in 12-well plates with DMEM/F-12 medium containing 0.1% BSA at 32°C and 5% CO2. After 2 h of culture, the cells were incubated with chrysin for 30 min in DMEM/F-12 medium without BSA and then 0.05 mM dbcAMP was added to the culture for 4 h. The culture medium and Leydig cells were collected and stored at −80°C. Testosterone concentrations in the medium were determined by RIA. All procedures were approved by the Texas Tech University Health Sciences Center Animal Care and Use Committee.

Western blot analysis

StAR protein, P450scc, and DAX-1 protein in Leydig cells were detected by western blot analysis as described previously (Townson et al. 1996). Western blot analysis experiments were performed at least three times and the results of one representative experiment are shown for each figure.

Transfection

MA-10 cells were cultured in 12-well plates (0·2 × 10^6 cells per well) overnight. The cells in each well were transfected with 0·5 μg DNA of the StAR promoter/luciferase plasmid PGL2/StAR expressing firefly luciferase driven by the −966 bp sequence of the StAR promoter (Caron et al. 1997). Transfections also included 6·0 ng of the pRL-SV40 vector DNA (a plasmid that constitutively expresses Renilla luciferase
under the control of the SV40 promoter, Promega). Transfections were performed using FuGENE6 Transfection Reagent (Roche) following the manufacturer’s instructions. After 48 h of culture, the cells were utilized for further experiments.

Luciferase assays

Following experiments, the cells were washed three times with ice-cold PBS and lysed with Passive Lysis Buffer (Promega). The supernatants were utilized for luciferase assays using a Dual Luciferase Reporter Assay System following the manufacturer’s instructions (Promega). The Relative Light Units (determined by dividing the reading from the PGL2/StAR promoter by the reading from Renilla luciferase) were measured using a TD-20/20 luminometer (Turner Designers, Sunnyvale, CA, USA).

RT-PCR

In experiments designed to determine StAR or DAX-1 mRNA expression, cells were washed three times with cold PBS and used for total RNA purification using TRIzol reagent in accordance with the manufacturer’s instructions (Invitrogen). The first-strand cDNA was synthesized from total RNA using the Reverse Transcription System (Promega). PCR for StAR was performed as previously described (Rao et al. 2003). PCR for DAX-1 was performed with the forward primer 5’-TCCTGTACCACGCT-TATGTG-3’ and the reverse primer 5’-TCGAAGTG-CAGGTGATCTTG-3’, for 32 cycles at 94°C for 1.5 min, 55°C for 1.5 min, 72°C for 2.5 min, and then followed by incubation at 72°C for 15 min. β-Actin was used as an internal marker in PCR for DAX-1.

Protein kinase A activity assays

Protein kinase A (PKA) activity was assayed as described previously (Wang et al. 2003a), using the SignaTec cAMP-dependent protein kinase assay system (Promega) following the manufacturer’s instruction. PKA activity was expressed as picomoles of 32P incorporated per min per μg protein.

Statistical analysis

Each experiment was repeated at least three times. Statistical analysis of the data was performed with ANOVA using the GraphPad Prism 4 system (GraphPad Software, San Diego, CA, USA). The data are shown as the mean ± S.E.M.

Results

Chrysin-enhanced steroid hormone production

To study the effect of chrysin on Leydig cell steroidogenesis, MA-10 mouse Leydig cells were incubated with increasing concentrations of chrysin and 0.05 mM dbcAMP. Incubation of MA-10 cells with chrysin induced a concentration-dependent increase in steroid hormone production. When the chrysin concentrations in the culture were increased from 0 to 10 μM, progesterone production was increased from 8.4 to 42.2 ng/ml (Fig. 1). To verify the results, the Leydig cells isolated from mice were cultured in the medium containing 10 μM chrysin and 0.05 mM dbcAMP. Incubation of the mouse Leydig cells with chrysin significantly enhanced steroidogenesis, with testosterone production being increased 2.4-fold over that of the cells stimulated with 0.05 mM dbcAMP alone (Fig. 2).

P450scc expression was analyzed by western blot analysis to determine whether the increase in steroid hormone is due to an increase in P450scc expression. As shown in the Fig. 3, chrysin did not affect P450scc expression. Further, the hydrophilic substrate, 22(R)-hydroxysterol, was used to examine the effects of chrysin on the activities of steroidogenic enzymes. It was shown that chrysin induced a slight, but non-significant increase in progesterone production (Fig. 3).

Chrysin-enhanced StAR protein expression

Western blot analysis was performed to determine whether StAR protein levels in the chrysin-treated Leydig cells were altered. Incubation of MA-10 cells with the increasing concentrations of chrysin resulted in an increase in StAR protein expression in a concentration-dependent manner, paralleling the increase in steroid hormone production (Fig. 1). Also, incubation of the mouse Leydig cells with 10 μM chrysin enhanced cAMP-induced StAR protein expression concomitantly with the increase in testosterone production (Fig. 2).

![Figure 1](https://www.endocrinology-journals.org/assets/fig1.png)

**Figure 1** Concentration-dependent increases in StAR protein expression and steroidogenesis in MA-10 mouse Leydig cells treated with chrysin. MA-10 cells were cultured with increasing concentrations of chrysin for 30 min and then 0.05 mM dbcAMP was added to the culture for 6 h. (A) The cells were collected and 20 μg cell lysate protein was used for western blot analysis of StAR protein. (B) The culture medium was collected and progesterone production was assessed by RIA. *P<0.05, **P<0.01 compared with the cells treated with 0.05 mM dbcAMP alone.
Chrysin-enhanced StAR gene transcription

Luciferase assay of StAR promoter activity and RT-PCR analysis of StAR mRNA levels were performed to determine the effect of chrysin on StAR gene transcription. Similar to its effects on StAR protein expression and steroidogenesis, the increasing levels of chrysin significantly increased cAMP-induced StAR promoter activity and StAR mRNA levels in a concentration-dependent manner. In MA-10 cell cultures, 10 μM chrysin induced a 12-fold increase in StAR promoter activity (Fig. 4).

Synergistic interaction between chrysin and cAMP in steroidogenesis

MA-10 cells were incubated with increasing concentrations of dbcAMP in the medium containing 10 μM chrysin to study the interaction between chrysin and cAMP. The results indicated that chrysin did not increase StAR protein expression and steroid hormone production in the absence of dbcAMP. However, in the presence of sub-threshold level of dbcAMP (0.05 mM), chrysin dramatically increased StAR protein expression and steroidogenesis (Fig. 5). Similarly, 0.05 or 0.1 mM dbcAMP alone is unable to induce significant increases in StAR protein and steroid hormone. However, in the presence of 10 μM chrysin, low level of cAMP is able to induce maximal levels of StAR protein expression and steroidogenesis, with 0.1 mM dbcAMP increasing progesterone production to tenfold over that seen in the paired group treated with dbcAMP alone.

Figure 2 Chrysin-enhanced StAR protein expression and testosterone production in the Leydig cells isolated from mouse testes. The Leydig cells were isolated from the testes of 3 month-old mice. The cells were incubated with chrysin and H89 for 30 min and then 0.05 mM dbcAMP was added to the culture for 4 h. (A) The cells were collected and 5 μg cell lysate protein was used for western blot analysis of StAR protein. (B) The culture medium was collected and testosterone production was assessed by RIA. *P<0.05, **P<0.01 compared with control.

Figure 3 P450scc expression and steroid hormone production in MA-10 mouse Leydig cells using 22(R)hydroxycholesterol as a substrate. MA-10 cells were cultured with 25 μM 22(R)hydroxycholesterol. The cells were incubated with increasing concentrations of chrysin for 30 min and then 0.05 mM dbcAMP was added to the culture for 6 h. (A) The cells were collected and 20 μg cell lysate protein was used for western blot analysis of P450scc protein. (B) The culture medium was collected and progesterone production was assessed by RIA. There is no significant difference in steroid production among the groups (P>0.05).

Figure 4 Concentration-dependent increase in StAR gene transcription in MA-10 mouse Leydig cells treated with chrysin. MA-10 cells were cultured with increasing concentrations of chrysin for 30 min and then 0.05 mM dbcAMP was added to the culture for 6 h. (A) The cells were collected for total RNA isolation and StAR mRNA was analyzed by RT-PCR using the rRNA L19 as an internal marker. (B) MA-10 cells were transfected with a StAR promoter/luciferase plasmid (PGL2/StAR) and a pRLSV40 vector, a plasmid that constitutively expresses Renilla luciferase. The cell lysate was used for luciferase assays using a Dual Luciferase Reporter Assay System. *P<0.05, **P<0.01 compared with the cells treated with 0.05 mM dbcAMP alone.

Figure 5 Synergistic interaction between chrysin and cAMP in steroidogenesis

MA-10 cells were incubated with increasing concentrations of dbcAMP in the medium containing 10 μM chrysin to study the interaction between chrysin and cAMP. The results indicated that chrysin did not increase StAR protein expression and steroid hormone production in the absence of dbcAMP. However, in the presence of sub-threshold level of dbcAMP (0.05 mM), chrysin dramatically increased StAR protein expression and steroidogenesis (Fig. 5). Similarly, 0.05 or 0.1 mM dbcAMP alone is unable to induce significant increases in StAR protein and steroid hormone. However, in the presence of 10 μM chrysin, low level of cAMP is able to induce maximal levels of StAR protein expression and steroidogenesis, with 0.1 mM dbcAMP increasing progesterone production to tenfold over that seen in the paired group treated with dbcAMP alone.
DAX-1 expression in chrysin-treated cells

To further understand how chrysin enhanced StAR gene transcription, DAX-1 protein was detected by western blot. The results shown in Fig. 6 indicated that expression of DAX-1 protein was maintained at a high level in MA-10 cells. While stimulation of the cells with 0.05 mM dbcAMP did not reduce DAX-1 protein, co-incubation with 10 μM chrysin markedly reduced DAX-1 protein in MA-10 cells. When DAX-1 protein was reduced, StAR protein and steroidogenesis were significantly increased. Similar results were obtained in RT-PCR analysis of DAX-1 mRNA. The results in Fig. 7 indicated that while chrysin alone only slightly reduced DAX-1 expression, the co-action of chrysin and 0.05 mM dbcAMP dramatically reduced DAX-1 protein and mRNA levels.

The essential role of PKA activity in chrysin-enhanced steroidogenesis

PKA activity in chrysin-treated Leydig cells was inhibited using an inhibitor H89 to determine the role of PKA activity in the increased StAR protein expression. As shown in Fig. 6, while 10 μM chrysin dramatically enhanced cAMP-induced StAR protein and steroid hormone production, the inhibition of PKA activity resulted in a marked reduction in StAR protein expression. Progesterone production by MA-10 cells was reduced to 5% by the PKA inhibitor. Testosterone production by the isolated Leydig cells was also reduced to 2-5% (Fig. 2).

To verify the role of PKA in the observed results, PKA activities in MA-10 cells were assayed. Results from the assays indicated that while PKA activity in the cells was significantly increased by 0.05 mM dbcAMP, it was not affected by 10 μM chrysin. When 25 μM H89 was added to the culture, dbcAMP-increased PKA activity was significantly reduced (Fig. 7).

The essential role of lipoxygenase activity in chrysin-enhanced steroidogenesis

The above results and previous studies (Wang et al. 2000, 2002) encouraged us to determine the role of arachidonic acid (AA)-mediated signaling in chrysin-induced steroidogenesis. MA-10 cells were cultured for 30 min in serum-free Waymouth’s medium containing chrysin and H89. Then 0-05 mM dbcAMP was added to the culture for 6 h. (A) The cells were collected and 20 μg cell lysate protein was used for western blot analysis of DAX-1 protein. (B) The culture medium was collected and progesterone production was assessed by RIA. ***P<0.001 compared with the cells treated with 0-05 mM dbcAMP alone.
Discussion

Chrysin is one of the natural flavonoids present in plants (Rapta et al. 1995, Williams et al. 1997). It is present in honey and propolis at high levels (Gambelunghe et al. 2003). In addition to its activities in anti-cancer (Cardenas et al. 2006), anti-oxidation (Lapidot et al. 2002), and anti-inflammation (Cho et al. 2004), chrysin has also been reported to be an inhibitor of aromatase (Kellis & Vickery 1984, Ibrahim & Abul-Hajj 1990), an enzyme that converts testosterone to estrogen. Based on these studies, chrysin has been considered to be a testosterone-increasing agent by blocking the further metabolism of testosterone by aromatase (Dhawan et al. 2002). We have been interested in its testosterone-increasing activity and examined it with MA-10 mouse Leydig cells. The results of the present study suggest that chrysin increases testosterone biosynthesis mainly by enhancing cAMP-induced StAR gene expression in Leydig cells.

The steroidogenic effect of chrysin was indicated by the concentration-dependent increase in steroid hormone production in MA-10 mouse Leydig cells incubated with increasing concentrations of chrysin in culture medium containing 0·05 mM dbcAMP. This effect was confirmed in primary cultures utilizing Leydig cells isolated from mouse testes, with testosterone production being dramatically enhanced when the cells were treated with chrysin. While chrysin enhanced dbcAMP-stimulated steroidogenesis, it did not affect P450scc expression. Also, when water-soluble 22(R)hydroxylcholesterol was used as the substrate, steroid hormone production in each group reached the maximal level, and there is no significant difference in steroidogenesis among the groups. In these groups, chrysin only induced a slight, non-significant increase in steroid hormone production. These observations indicated that the majority of the increase in steroid hormone did not result from increases in the activities of steroidogenic enzymes, but rather, from the increased supply of the substrate cholesterol to the inner mitochondrial membrane. These observations were corroborated by the results from western blot analysis of StAR protein in MA-10 Leydig cells and also in primary cultures of Leydig cells isolated from mouse testes. In these cells, chrysin induced a marked increase in StAR protein, the factor that facilitates mitochondrial cholesterol transfer to the P450scc enzyme. Also, the increase in StAR protein occurred concomitantly with the increase in steroid hormone production. Moreover, reduction in StAR protein expression by inhibiting PKA activity abolished the chrysin-enhanced steroidogenesis in Leydig cells.

Normally, low levels of dbcAMP, such as 0·05 mM, are unable to induce significant increases in StAR protein and steroidogenesis in MA-10 Leydig cells. However, in the presence of chrysin, 0·05 mM of dbcAMP significantly
increased StAR protein and steroid hormone production. The effectiveness of dbcAMP stimulation was dramatically enhanced in the chrysin-treated cells, with StAR protein expression and steroidogenesis reaching their maximal levels at much lower concentrations of dbcAMP than those in the cells without chrysin. On the other hand, although chrysin strongly enhanced cAMP-stimulated steroidogenesis, in the absence of cAMP chrysin did not induce a significant increase in StAR protein or steroid hormone. Also, inhibition of PKA activity reduced chrysin-induced StAR protein and steroidogenesis. These results suggested that this flavonoid is unable to stimulate StAR protein expression by itself, but rather interacted with cAMP and reduced the threshold of cAMP-stimulated StAR protein expression, resulting in increase in the sensitivity of Leydig cells to cAMP stimulation.

How chrysin interacted with cAMP to increase sensitivity of Leydig cells to cAMP stimulation is unknown. It was demonstrated that in cAMP-stimulated steroidogenesis, both cAMP–PKA phosphorylation and AA-mediated signaling pathways are required for StAR gene expression (Wang et al. 2000). These two signaling pathways synergistically interact with each other, with an increase of signal in either pathway dramatically enhancing sensitivity of Leydig cells to cAMP stimulation (Wang et al. 2002, 2003). Based on these studies, we have examined the roles of these two signaling pathways in chrysin–enhanced steroidogenesis. The results from PKA activity assays indicated that while PKA activity is critical for the observed increase in StAR protein expression, chrysin did not enhance the cAMP–PKA phosphorylation signaling. However, when AA metabolism by 5-lipoxygenase was blocked, the increases in StAR protein and steroid hormone in the chrysin-treated cells were reversed, suggesting that AA–mediated signaling is involved in the chrysin–increased StAR protein expression. Since 5-lipoxygenase–generated AA metabolites significantly enhanced steroidogenic sensitivity of Leydig cells to cAMP stimulation (Wang et al. 2003), it is interesting to determine if chrysin enhances this AA–mediated signaling.

While the manner in which chrysin interacted with cAMP is awaiting further elucidation, the evidence from the present study indicates that this interaction enhanced StAR gene expression at the level of transcription. This could be readily seen from the luciferase assays of StAR promoter activity, which indicated that incubation of cAMP-treated MA–10 cells with chrysin induced a concentration–dependent increase in StAR promoter activity. Similar results were obtained in RT-PCR analysis of StAR mRNA levels. To further understand how this natural flavonoid enhanced StAR gene transcription, the transcription factors regulating StAR promoter activity were examined. It was found that when MA–10 Leydig cells were incubated with chrysin in a medium containing 0.05 mM dbcAMP, DAX–1 protein and DAX–1 mRNA levels in the cells were markedly reduced. DAX–1 protein is a powerful transcriptional repressor that binds to a hairpin structure in StAR promoter DNA and inhibits StAR gene transcription (Zazopoulos et al. 1997). The results suggest that interaction between chrysin and cAMP enhance StAR gene transcription by depressing DAX–1 gene expression. The results in Figs 6 and 7 further indicated that DAX–1 is constitutively expressed in MA–10 Leydig cells, generating a tonic inhibition of StAR gene transcription (Zazopoulos et al. 1997). Thus, the reduction of DAX–1–generated tonic inhibition resulted in the increase in sensitivity of Leydig cells to cAMP stimulation.

In addition to the transcriptional regulation, StAR gene expression is also regulated at the level of post–transcription. Regarding this, the studies in the recent years have described a cAMP–regulated turnover and stability of StAR mRNA (Zhao et al. 2005), in which cAMP selectively induced an unstable form of StAR mRNA to facilitate rapid increase and decrease in StAR expression. Also, the effect of cAMP on the post–transcriptional regulation of StAR expression involves A–kinase anchoring proteins (AKAPs). AKAPs recruit StAR mRNA and PKA to the mitochondria and facilitate StAR translation and phosphorylation (Dyson et al. 2008, Papadopoulous et al. 2007). However, whether chrysin enhanced the cAMP–regulated post–transcriptional regulation of StAR gene expression needs to be clarified by experimental evidence.

Taken together, the present study suggests a potential for the application of chrysin to delay the age–related decline in StAR gene expression and testosterone biosynthesis in Leydig cells. This is supported by studies using aged male rats, in which supplementation with chrysin for 30 days significantly improved the reproductive functions of the animals (Dhawan et al. 2007). The effectiveness of chrysin might be attenuated by its low bioavailability in some food supplements (Gambelunghe et al. 2003). Recent studies have attempted to improve the effectiveness of chrysin by modification of its molecular structure (Dao et al. 2004). Several chrysin derivatives with improved activities were reported. Interestingly, some of these derivatives are able to bind to the COX2 molecule and selectively inhibits COX2 activity (Cho et al. 2004). We have previously reported that inhibition of COX2 activity dramatically enhanced StAR gene expression and steroidogenesis in aging Leydig cells (Wang et al. 2005). Thus, these derivatives with COX2–inhibiting activity may significantly improve the steroid synthesis–promoting activity of chrysin.

Obviously, more studies, especially long–term clinical studies, are needed to test the efficacy of chrysin or its derivatives on delaying the age–related decline in blood testosterone. Since the decline in testosterone is associated with decreases in various physiological functions, further studies on the steroidogenic effect of chrysin may be beneficial for aging males, especially for those suffering from hypogonadism.

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References


Zhao D, Duan H, Kim YC & Jefcoate CR 2005 Rodent StAR mRNA is substantially regulated by control of mRNA stability through sites in the 3′-untranslated region and through coupling to ongoing transcription. *Journal of Steroid Biochemistry and Molecular Biology* **96** 155–173.

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