



Article Combination of Lycopene and Curcumin Synergistically Alleviates Testosterone-Propionate-Induced Benign Prostatic Hyperplasia in Sprague Dawley Rats via Modulating Inflammation and Proliferation

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Abstract: Background: Benign prostatic hyperplasia (BPH) is a progressive urological disease occurring in middle-aged and elderly men, which can be characterized by the non-malignant overgrowth of stromal and epithelial cells in the transition zone of the prostate. Previous studies have demonstrated that lycopene can inhibit proliferation, while curcumin can strongly inhibit inflammation. This study aims to determine the inhibitory effect of the combination of lycopene and curcumin on BPH. Method: To induce BPH models in vitro and in vivo, the BPH-1 cell line and Sprague Dawley (SD) rats were used, respectively. Rats were divided into six groups and treated daily with a vehicle, lycopene (12.5 mg/kg), curcumin (2.4 mg/kg), a combination of lycopene and curcumin (12.5 mg/kg + 2.4 mg/kg) or finasteride (5 mg/kg). Histologic sections were examined via hematoxylin and eosin (H&E) staining and immunohistochemistry. Hormone and inflammatory indicators were detected via ELISA. Network pharmacology analysis was used to fully predict the therapeutic mechanism of the combination of lycopene and curcumin on BPH. Results: Combination treatment significantly attenuated prostate hyperplasia, alleviated BPH pathological features and decreased the expression of Ki-67 in rats. The upregulation of the expression of testosterone, dihydrotestosterone (DHT), 5α -reductase, estradiol (E2) and prostate-specific antigen (PSA) in BPH rats was significantly blocked by the combination treatment. The expression levels of inflammatory factors including interleukin (IL)-1 β , IL-6 and tumor necrosis factor (TNF)- α were strongly inhibited by the combination treatment. From the network pharmacology analysis, it was found that the main targets for inhibiting BPH are AKT1, TNF, EGFR, STAT3 and PTGS2, which are enriched in pathways in cancer. Conclusion: The lycopene and curcumin combination is a potential and more effective agent to prevent or treat BPH.

Keywords: BPH; lycopene; curcumin; synergistic effect; inflammation

1. Introduction

Benign prostatic hyperplasia (BPH), the most common chronic disease among aging men [1], can potentially lead to lower urinary tract symptoms (LUTSs) [2–4]. Epidemiological studies show the incidence of BPH increases with age, which reaches 40–50% at the age of 50–60 and up to 80–90% after the age of 80 [5,6]. Histologically, BPH is characterized by the hyperproliferation of the glandular epithelium and supporting stromal cells of



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Copyright: © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). the prostate gland [7]. At the biochemical level, BPH is considered to be caused by the imbalance between androgen and estrogen and the over-expression of growth factors in stromal and epithelial cells [8].

Although the underlying mechanisms of BPH development are not fully understood, various pathogenic mechanisms have been proposed, including oxidative stress, inflammation, the imbalance of proliferation and apoptosis and hormonal imbalances [9]. Indeed, the androgenic pathway is thought to be predominant. Dihydrotestosterone (DHT), a powerful prostatic and rogen, is converted from testosterone by 5α -reductase [10]. With a high affinity, DHT binds to the androgen receptor, which initiates the transcription of genes that encode differentiation and growth-promoting factors. This process subsequently stimulates prostatic proliferation [11]. Furthermore, inflammation is a prevalent observation in the prostate, evident on both histological and biochemical levels, thereby promoting the advancement of BPH [12]. The inflammation observed in BPH may result in tissue damage, and the cytokines generated by inflammatory cells may stimulate the production of growth factors and angiogenesis in the prostate, as a form of wound-healing response [13]. The major pathological feature of BPH is the hyperplasia of prostatic epithelial and stromal cells, caused by an imbalance between cell growth and apoptosis [14]. Apoptosis is programmed cell death which occurs on a regular basis to keep a homeostatic balance between cell proliferation and death rates [15].

The management of BPH typically entails the utilization of Alpha-1 adrenergic receptor blockers, 5α -reductase enzyme inhibitors (e.g., finasteride (FN)) and surgical intervention, or a combination of these therapeutic modalities [16]. Research has shown that different types of drugs have different side effects. Alpha-1 blockers can be associated with orthostatic hypotension, and 5α -reductase inhibitors are associated with sexual dysfunction [17,18].

Interestingly, natural plant extracts are emerging as an innovative modality to treat BPH and show significant improvement or even curative effects, and they have less side effects than common conventional medical treatments [19]. Lycopene (LY) is a natural pigment with anti-oxidant and anti-proliferative activities. Epidemiological studies have shown that high intake of LY is related to the reduced risk of prostatic diseases like BPH and prostatic cancer [20–22]. Zou et al. has found that 30 mg/kg Maca and 7.5 mg/kg LY can effectively inhibit the progression of BPH. However, the synergistic effect of these two drugs was not researched in this study [23]. Curcumin (CUR), derived from the rhizome of *Curcuma longa* L., is a medicinal herb with a long history of use in the Indian system of medicine for the treatment of various health conditions over many centuries [24]. CUR possesses a wide range of therapeutic effects such as anti-inflammatory, anti-oxidant, anticancer and anti-microbial [25–27]. There is a research suggesting that 50 mg/kg curcumin via daily oral administration showed an inhibitory effect on BPH in rats [28].

However, the utilization of a solitary plant extract for the treatment of BPH necessitates a higher dosage, which may impose a greater strain on the liver and kidney functions [29]. Furthermore, monotherapy typically targets a singular aspect, and prolonged usage may result in the development of drug resistance [30]. The implementation of combined therapy presents a compelling strategy for managing chronic ailments. As assessed using the International Prostate Symptom Score, combination therapy has demonstrated an enhanced likelihood of better recovery and response rates, while also mitigating drug resistance in patients with long-standing chronic conditions [30]. Therefore, this study aims to evaluate the potential synergistic effects of co-administering LY and CUR in mitigating the development of BPH in rats, while also gaining insight into the multi-target mechanism underlying this combined therapeutic approach.

2. Results

2.1. Effects of LY and CUR on the Viability of BPH-1 Cells In Vitro

The effects of LY, CUR and a combination of LY and CUR on the viability of BPH-1 cells were detected via a CCK-8 assay. As shown in Figure 1, LY (Figure 1A) and CUR (Figure 1B) inhibited the cell proliferation in a dose-dependent manner, and when the



inhibition rate of cell proliferation reached about 50%, the concentrations of LY and CUR were 800 μ g/mL and 10 μ g/mL, respectively.

Figure 1. Effects of LY and CUR on the viability of BPH-1 cells. (**A**) The dose-dependent effect of LY on cell viability; (**B**) the dose-dependent effect of CUR on cell viability; (**C**) the effect of LY/CUR combinations on cell viability. All values are means \pm SD (n = 6). Within each panel, the values with the superscription symbol are significantly different from that of the control; * p < 0.05, ** p < 0.01 and *** p < 0.001.

Since CUR showed more sensitive dose-dependent activity than LY, the lower doses of LY (200 µg/mL) and CUR (5, and 10 µg/mL) were selected for evaluating the combination effect, and the nature of the combination was determined by calculating the CI. As shown in Figure 1C, the LY and CUR combination had more potent activities than the corresponding individual treatment. Compared with the control, LY (200 µg/mL), CUR (5 µg/mL), CUR (10 µg/mL), the LY (200 µg/mL)/CUR (5 µg/mL) combination and the LY (200 µg/mL)/CUR (10 µg/mL) combination inhibited the BPH-1 cell viability by 37.19% (p < 0.001), 30.19% (p < 0.001), 43.23% (p < 0.001), 49.13% (p < 0.001) and 64.81% (p < 0.001), respectively. The CI for the LY (200 µg/mL)/CUR (10 µg/mL) combination treatment was 0.9 < 1. The results indicated that the LY and CUR combination had a synergistic effect on inhibiting the proliferation of BPH-1 cells.

2.2. Effects of LY and CUR Combination Treatments on BPH Development in Rats

Rats were induced for BPH via the subcutaneous injection of TP and treated with FN or experimental drugs. After 8 weeks of induction with TP, both the prostatic weight (Figure 2A) and prostate index (prostate weight (mg)/body weight (g)) (Figure 2B) in the BPH group were significantly higher than those in the control group, indicating that the BPH model was successfully established. Of note, the TP-induced increases in prostate

weight and prostate index were significantly attenuated by the LY and CUR treatments. The prostate weights of rats in the LY, CUR and COM groups were significantly reduced by 23.08% (p < 0.05), 26.78% (p < 0.05) and 47.35% (p < 0.01), compared with those in the BPH group (Figure 2A). In parallel, the prostate indices in the rats in the LY, CUR and COM groups were significantly reduced by 26.2% (p < 0.05), 22.18% (p < 0.05) and 46.26% (p < 0.01), compared with those in the BPH group (Figure 2B). The LY/CUR combination further increased the individual treatment effect, and the CI for the combination treatment was 0.229 < 1, suggesting that the LY/CUR combination may provide more a potent treatment regimen than individual treatments in inhibiting BPH development.



Figure 2. Effects of experimental treatments on BPH development in rats. (**A**) Prostatic weight; (**B**) prostate index; (**C**) quantified results of epithelium thickness of prostate (ETP); (**D**) time-dependent body weight change; (**E**) body weight of rats at week 8; (**F**) representative H&E staining images. All values are mean \pm SD (n = 6). Within each panel (**A**–**C**), the values with the superscription symbols are significantly different from those of the corresponding control; * p < 0.05, ** p < 0.01 and *** p < 0.001, when compared with the BPH group; ### p < 0.001, compared with the CON group; one-way ANOVA followed by Tukey's comparison test.

The epithelium thicknesses of the prostate (ETPs) were determined in the H&E-stained prostate tissues. The ETP in the BPH group was significantly increased by 241%, compared with that in the CON group (Figure 2C,F). Treatments with LY, CUR and the LY/CUR combination significantly reduced the ETP by 62.51% (p < 0.001), 58.45% (p < 0.001) and 65.16% (p < 0.001), respectively, compared with the BPH group.

To sum up, these data indicate that the combination of LY and CUR can inhibit the development of BPH and is more effective than LY and CUR alone.

2.3. Effects of LY and CUR Combination Treatments on Serum Levels of Hormones in Rats

As shown in Figure 3A–D, the serum levels of DHT, 5 α -reductase, T and E2 were indicated as being significantly higher in the BPH group by 1.37-, 1.17-, 1.23- and 1.27-fold compared with the CON group, suggesting TP administration had resulted in the upregulation of the serum level of DHT, 5 α -reductase, T and E2 and the consequent benign prostate hyperplasia. Of note, treatments with LY, CUR and the LY/CUR combination significantly reduced the expression of DHT by 10.17%, 19.97% (p < 0.05) and 29.06% (p < 0.01), respectively, compared with the BPH group. The expression levels of 5 α -reductase in the LY, CUR and COM group were significantly decreased by 29.87% (p < 0.05), 27.48% (p < 0.05) and 38.3% (p < 0.01), respectively, compared with the BPH group. Compared with the BPH group, the regulation of testosterone via LY and CUR was not significant, but the expression of T in the COM group was reduced by 20.35% (p < 0.05). The expression of E2 and PSA in the COM group was decreased by 24.58% (p < 0.05) and 22.77% (p < 0.05), respectively, compared with the BPH group.



Figure 3. Effects of experimental treatments on serum levels of hormones. (**A**) Dihydrotestosterone (DHT), (**B**) 5 α -reductase, (**C**) testosterone (T), (**D**) estradiol (E2) and (**E**) prostate-specific antigen (PSA) in rats. All values are mean \pm SD (n = 6). # p < 0.05 vs. CON group, * p < 0.05, ** p < 0.01 vs. BPH group, one-way ANOVA followed by Tukey's comparison test.

2.4. Effects of LY and CUR Combination Treatments on Inflammatory Responses in Rats

As shown in Figure 4A–C, BPH induction stimulated the secretion of TNF- α , IL-1 β and IL-6 into blood and significantly increased the serum levels by 1.24, 1.4 and 1.22 in comparison with the CON group. However, the TNF- α , IL-1 β and IL-6 levels in the COM group were significantly lower than those in the BPH group by 29.13% (p < 0.01), 51.85% (p < 0.001) and 45.6% (p < 0.001), respectively.



Figure 4. Assessment of treatment effects on serum levels of inflammatory cytokines TNF- α (**A**), IL-1 β (**B**) and IL-6 (**C**). All values are mean \pm SD (n = 6). # p < 0.05 vs. CON group, * p < 0.05, ** p < 0.01, *** p < 0.001 vs. BPH group, one-way ANOVA followed by Tukey's comparison test.

2.5. Effects of LY and CUR Combination Treatments on Prostate Cell Proliferation

To explore the inhibitory effect of drugs on proliferation, cell proliferation marker Ki-67 was detected. As shown in Figure 5, the expression level of Ki-67 in the BPH group was much higher than that in the CON group, as the gray value of the BPH group was twice that of the CON group. After treatment, the expression level of Ki-67 in the LY, CUR and COM group decreased by 17.27% (p < 0.05), 22.47% (p < 0.05) and 29.13% (p < 0.01), respectively, compared with the BPH group.



Figure 5. Immunohistochemistry (IHC) of prostatic tissues. (**A**) Animal prostates were collected and stained by cell proliferation marker Ki-67, in rats treated by (**a**) control group (CON), (**b**) BPH model group (BPH), (**c**) BPH+FN group (FN), (**d**) BPH+LY group (LY), (**e**) BPH+CUR group (CUR) or (**f**) BPH + LY + CUR group (COM), respectively. The arrows point to brown areas representing the expression of Ki-67. (**B**) Quantitative analysis of gray value was carried out by ImageJ. All values are mean \pm SD (n = 3). ### p < 0.001 vs. CON group, * p < 0.05, ** p < 0.01 vs. BPH group, one-way ANOVA followed by Tukey's comparison test.

2.6. Network Pharmacology Analysis

The process of gathering and merging data pertaining to BPH-related targets from Targetnet, UniProt and Genecards yielded a total of 1282 targets that met the relevance threshold of greater than 6. A comparative analysis was conducted between these targets and the predicted targets of LY and CUR, resulting in the identification of 52 common targets that were deemed as potential candidates associated with the anti-BPH activity of LY (No. 1–12) and CUR (No. 13–52) (Table 1, Figure 6). Notably, there were no overlapping targets between LY and CUR, indicating that these compounds may impede the progression of BPH through the modulation of distinct targets.

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10Dumspectinity prospinatus creeptor alphaCDCATACDCATA11Peroxisome proliferator-activated receptor alphaRPARAP1027612Retinoic acid receptor alphaRARAP1027613Epidemal growth factor receptor erbB1EGRRP0053314Serine / threonine-protein kinase AKTAKT1P3174915Serine / threonine-protein kinase B-rafBRAFP1505616Apoptosis regulator Bcl-2BCL2P1041517Signal transducer and activator of transcription 3STAT3P4076318Cyclooxygenase-2PTCS2P335420Tyrosime-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine / threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPI,P0518629Serine / threonine-protein kinase Aurora-AAURKAO1496531Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NF2212O1623633Matrix metalloproteinase 13MMP13P4545234Tyro	10	Dual-specificity phosphatase Cdc25A	CDC25A	P30304
11111111121202/10912Retinoic acid receptor alphaRARAP1027613Epidermal growth factor receptor erbB1EGFRP0053314Serine/threonine-protein kinase AKTAKT1P3174915Serine/threonine-protein kinase B-rafBRAFP1505616Apoptosis regulator Bd-2BCL2P1041517Signal transducer and activator of transcription 3STAT3P4076319TNF-alphaTNFP0137520Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase Atarora-AAURKAO1496531Kibosomal protein S6 kinase 1RP3443P344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase AktriJAK1 <td>10</td> <td>Parovisome proliferator activated receptor alpha</td> <td>DDA DA</td> <td>007869</td>	10	Parovisome proliferator activated receptor alpha	DDA DA	007869
12Extension admit explore appliedINAC1102/013Epidemial growth factor receptor erbB1EGFRP0053314Serine/threonine-protein kinase B-rafBRAFP1505616Apoptosis regulator Bcl-2BCL2P1041517Signal transducer and activator of transcription 3STAT3P4076318Cyclooxygenase-2PTGS2P3535420Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P4044924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496531Ribosomal protein S6 kinase 1RP56KB1P2343334Tyrosine-protein kinase JAK1JAK1P244335Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anthydrase IXCA9Q1679037GUoxygenase-1PTGS1P2321938Tol-like receptor (LLR7/TLR9)TLR9Q9N245234Type-1 angioteniase 13MMP13P245235Arachido	11	Patinois acid recentor alpha		D10276
13Epidemial grown lactor receptor eropsPORTPORTS14Serine/threonine-protein kinase AKTAKTIP3174915Serine/threonine-protein kinase B-rafBRAFP1505616Apoptosis regulator Bcl-2BCL2P1041517Signal transducer and activator of transcription 3STAT3P4076318Cyclooxygenase-2PTGS2P3535419TNF-alphaTNFP0137520Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4944131Ribosomal protein S6 kinase 1RP56KB1P234332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase I3MMP13P4545234Tyrosine-protein kinase IAK1JAK1P2344835Arachidonate 5-lipoxygenaseALOX5P09917 <td>12</td> <td>Enidownal arouth factor receptor alpha</td> <td></td> <td>P00522</td>	12	Enidownal arouth factor receptor alpha		P00522
14Serine/threonine-protein kinase ArAArAArAF11F174915Serine/threonine-protein kinase FrafBRAFP1055616Apoptosis regulator Bcl-2BCL2P1041517Signal transducer and activator of transcription 3STAT3P4076318Cyclooxygenase-2PTGS2P3535420Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAFIP0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 50 ki anse 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P1091736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1SPHK1 <td< td=""><td>13</td><td>Soring /throoping protoin kinges AVT</td><td></td><td>F00355 D21740</td></td<>	13	Soring /throoping protoin kinges AVT		F00355 D21740
13Serine / utreomine-protein kinase b-ratDKAP1300016Apoptosis regulator Bcl-2BCL2P1011517Signal transducer and activator of transcription 3STAT3P4076318Cyclooxygenase-2PTGS2P3535419TNF-alphaTNFP0137520Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine /threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B33P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496533Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonica shydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor RBCXCR2P2502541Macrophage activator inhibitor-1SERIPINE1P0512142Matrix metalloproteinase 3MPG51P2345835Arachidonate 5-lipoxygenase-1 <td>14</td> <td>Serine/ inteonine-protein kinase AK1</td> <td></td> <td>F31/49 D1505(</td>	14	Serine/ inteonine-protein kinase AK1		F31/49 D1505(
16Apoptosis regulator bc/2bc/2F1041517Signal transducer and activator of transcription 3STAT3P1076318 $C_yclooxygenase-2$ PTCS2P3335419TNF-alphaTNFP10137520Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EF300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 65 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1SFHR1P0733334Tyrosine-protein kinase 1SFHR1P0733335Arachidonate 5-lipoxy	15	Serine/threenine-protein kinase b-rai	DKAF	P15056
17Signal transducer and activator of transcription 3SIA13P40/6318CC(cloxygenase-2PTGS2P3353419TNF-alphaTNFP0137520Tryrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496531Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2O163633Matrix metalloproteinase 13MMP13P4345234Trrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Clycoxygenase-1PTCS1P2321938Toll-like receptor TBCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Ype-1 angiotensin II receptor (by homology)ACTR11P3055644 <td>16</td> <td>Apoptosis regulator BCI-2</td> <td>BCL2</td> <td>P10415</td>	16	Apoptosis regulator BCI-2	BCL2	P10415
18Cyclooxygenase-2PTCS2P3333419TNF-alphaTNFP0137520Tyrosine-protein kinase JAK2JAK2O6067421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2344335Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512141Macrophage colony stimulating factor receptorCSF1RP0733342Shphingosine	17	Signal transducer and activator of transcription 3	SIAI3	P40763
19 $1NF$ $P01375$ 20Tyrosine-protein kinase JAK2JAK2 $O60674$ 21Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300 $O09472$ 23Serine / threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein S6 kinase 1RPS6KB1P2344332Nuclear factor erythroid 2-related factor 2NFE12Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachionate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679039Plasminogen activator inhibitor-1SERPINE1P0512141Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NNA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17A	18	Cyclooxygenase-2	PIGS2	P35354
20I prosine-protein kinase JAK2JAK2O606/421Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Trosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q9NR9638Toll-like receptor RLY/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342NDAM17ADAM17P7853644HERGKCNH2Q1280945ADAM17ADAM	19	TNF-alpha	INF	P01375
21Matrix metalloproteinase 9MMP9P1478022Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFEL2Q1623633Matrix metalloproteinase JAK1JAK1P2345834Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q9NR9639Plasminogen activator inhibitor-1SERPINE11P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSFIRP033342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensiae 3MMP3P0855644HERGKCNH2Q1280945ADAM17ADAM17P748748Type-1 angiotensiae 3MMP3<	20	Tyrosine-protein kinase JAK2	JAK2	O60674
22Histone acetyltransferase p300EP300Q0947223Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 66 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945	21	Matrix metalloproteinase 9	MMP9	P14780
23Serine/threonine-protein kinase RAFRAF1P0404924Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)ACTR1P3055644HERGKCNH12Q1280945ADAM17ADAM17P7853646Matrix metalloprote	22	Histone acetyltransferase p300	EP300	Q09472
24Matrix metalloproteinase 14MMP14P5028125Estradiol 17-beta-dehydrogenase 3H5D17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 66 kinase 1RP56K811P23443332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase 1AK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P385644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4<	23	Serine/threonine-protein kinase RAF	RAF1	P04049
25Estradiol 17-beta-dehydrogenase 3HSD17B3P3705826DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 56 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE11P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP033342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP14	24	Matrix metalloproteinase 14	MMP14	P50281
26DNA topoisomerase II alphaTOP2AP1138827Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein S6 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aninopeptidase NANPEPP15144 <t< td=""><td>25</td><td>Estradiol 17-beta-dehydrogenase 3</td><td>HSD17B3</td><td>P37058</td></t<>	25	Estradiol 17-beta-dehydrogenase 3	HSD17B3	P37058
27Matrix metalloproteinase 7MMP7P0923728Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein S6 kinase 1RP56KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (ILR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE11P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensing II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P78536MMP348Tyrosinase IVDPP4P2748748Tyrosinase IVANPEPP151445011-beta-hydroxysteroid dehydrogenase [NAD+1]HPGDP15428	26	DNA topoisomerase II alpha	TOP2A	P11388
28Alkaline phosphatase, tissue-nonspecific isozymeALPLP0518629Serine/threonine-protein kinase Aurora-AAURKAO1496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein S6 kinase 1RPS6KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2845551Carbonic anhydrase IICA2P00918 <td>27</td> <td>Matrix metalloproteinase 7</td> <td>MMP7</td> <td>P09237</td>	27	Matrix metalloproteinase 7	MMP7	P09237
29Serinê/threonine-protein kinase Aurora-ÁAURKAOl496530Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein 56 kinase 1RPS6KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2845551Carbonic anhydrase IICA2P009185215-hydroxysteroid dehydrogenase [NAD+1]HPGDP15428	28	Alkaline phosphatase, tissue-nonspecific isozyme	ALPL	P05186
30Glycogen synthase kinase-3 betaGSK3BP4984131Ribosomal protein S6 kinase 1RPS6KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE11P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+1]HPGDP15428	29	Serine/threonine-protein kinase Aurora-Á	AURKA	O14965
31Kibosomal protein S6 kinase 1RPS6KB1P2344332Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P385644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1CA2P0091851Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+1]HPGDP15428	30	Glycogen synthase kinase-3 beta	GSK3B	P49841
32Nuclear factor erythroid 2-related factor 2NFE2L2Q1623633Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)ACTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+1]HPGDP15428	31	Ribosomal protein S6 kinase 1	RPS6KB1	P23443
33Matrix metalloproteinase 13MMP13P4545234Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+1]HPGDP15428	32	Nuclear factor erythroid 2-related factor 2	NFE2L2	Q16236
34Tyrosine-protein kinase JAK1JAK1P2345835Arachidonate 5-lipoxygenaseALOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP167949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxystaglandin dehydrogenase [NAD+1]HPGDP15428	33	Matrix metalloproteinase 13	MMP13	P45452
35Árachidonate 5-lipoxygenaseÁLOX5P0991736Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE11P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyporstaglandin dehydrogenase [NAD+]HPGDP15428	34	Tyrosine-protein kinase JAK1	JAK1	P23458
36Carbonic anhydrase IXCA9Q1679037Cyclooxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P284551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	35	Arachidonate 5-lipoxygenase	ÁLOX5	P09917
37Cycloxygenase-1PTGS1P2321938Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP167949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	36	Carbonic anhydrase IX	CA9	O16790
38Toll-like receptor (TLR7/TLR9)TLR9Q9NR9639Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1647949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	37	Cvclooxygenase-1	PTGS1	P23219
39Plasminogen activator inhibitor-1SERPINE1P0512140Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1614949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	38	Toll-like receptor (TLR7/TLR9)	TLR9	O9NR96
40Interleukin-8 receptor BCXCR2P2502541Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1614949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	39	Plasminogen activator inhibitor-1	SERPINE1	P05121
41Macrophage colony stimulating factor receptorCSF1RP0733342Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP167949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	40	Interleukin-8 receptor B	CXCR2	P25025
42Sphingosine kinase 1SPHK1Q9NYA143Type-1 angiotensin II receptor (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP167949Aminopeptidase NANPEPP151145011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	41	Macrophage colony stimulating factor receptor	CSF1R	P07333
43Type-1 angiotensin II recentro (by homology)AGTR1P3055644HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2845551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+1]HPGDP15428	42	Sphingosine kinase 1	SPHK1	O9NYA1
1019 Fundation (19 fundation)19 Fundation10000044HERGKCNH2Q1280945ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	43	Type-1 angiotensin II receptor (by homology)	AGTR1	P30556
11111112120045ADAM17ADAM17P7853646Matrix metalloproteinase 3MMP3P0825447Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	44	HFRG	KCNH2	012809
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47Dipeptidul neutroproteinase of Dipeptidyl peptidase IVDPP4P2748748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	46	Matrix metalloproteinase 3	MMP3	P08254
1112/40748TyrosinaseTYRP1467949Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	47	Dipentidyl pentidase IV	DPP4	P27487
49Aminopeptidase NANPEPP151445011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	48	Tyroginage	TVR	P14679
5011-beta-hydroxysteroid dehydrogenase 1HSD11B1P2884551Carbonic anhydrase IICA2P009185215-hydroxyprostaglandin dehydrogenase [NAD+]HPGDP15428	40	Aminopentidase N	ANPEP	P15144
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52 15-hydroxyprostaglandin dehydrogenase [NAD+] HPGD P15428	51	Carbonic anhydraes II		P00018
	52	15-hvdroxyprostaglandin dehvdrogenase [NAD+]	HPGD	P15428

Table 1. Common targets of BPH-LY and CUR interconnection.



Figure 6. The predicted molecular candidate targets between the BPH- and LY-associated targets (**A**), or between the BPH- and CUR-associated targets (**B**).

In order to investigate the potential mechanisms underlying the synergistic effects of lycopene and curcumin against BPH, a total of 52 common targets were inputted into the STRING database to construct an original protein–protein interaction (PPI) network. The resulting network, consisting of 51 nodes and 339 edges, was generated from the tsv file (as shown in Figure 7). The analysis revealed five key targets, namely AKT1 (degree = 42), TNF (degree = 37), EGFR (degree = 33), STAT3 (degree = 32) and PTGS2 (degree = 30).



Figure 7. Protein–protein interaction network. The PPI network was constructed using Cytoscape and analyzed using NetworkAnalyzer. Different colors represent the degree. Node size is proportional to the degree of interaction.

A KEGG pathway enrichment analysis was conducted on the 52 common targets (Figure 8) with a significance level of p < 0.01. The results revealed the top three significantly enriched KEGG pathways, which were pathways in cancer (hsa05200), EGFR tyrosine kinase inhibitor resistance (hsa01521) and endocrine resistance (hsa01522).



Figure 8. KEGG enrichment analysis of the anti-BPH targets of lycopene and curcumin.

3. Materials and Methods

3.1. Chemicals and Reagents

Testosterone propionate (TP) was purchased from Ningbo Second Hormone Factory (Ningbo, Zhejiang, China), and FN was purchased from Zhejiang CONBA Pharmaceutical Co., Ltd. (Hangzhou, Zhejiang, China). LY was purchased from Xinjiang Keyu Technology Co., Ltd. (Urumchi, Xinjiang, China). CUR was purchased from Yuanye Biotechnology Co., Ltd. (Shanghai, China). BPH-1 human benign prostatic hyperplasia cell line cells were purchased from Beina Biology (Beijing, China). ELISA kits for determining rat di-hydrotestosterone (DHT), 5α -reductase, testosterone (T), estradiol (E2), prostate-specific antigen (PSA), interleukin (IL)-1 β , IL-6 and tumor necrosis factor (TNF)- α were purchased from Jiangsu Meibiao Biotechnology Co., Ltd. (Yancheng, Jiangsu, China). Protein extraction solution was purchased from Solarbio Science & Technology Co., Ltd. (Beijing, China).

3.2. Cell Culture

Cell culture reagents were purchased from Gibco (Life Technologies, Gaithersburg, MD, USA) unless otherwise stated. The BPH-1 cells were cultured in DMEM medium supplemented with 10% (v/v) heat-inactivated FBS and 100 units/mL penicillin–streptomycin at 37 °C in a humidified atmosphere of 5% CO₂.

3.3. Cell Viability Assay

Cell viability was assessed by using the Cell Counting Kit 8 (CCK-8) assay (Dojindo, Kumamoto, Japan). Briefly, BPH-1 cells (5×10^3 /well) were seeded in 96-well tissue culture plates and grown for 24 h. Cells in the treated groups were then treated with LY (50, 100, 200, 400 and 800 µg/mL), CUR (1.25, 2.5, 5, 10 and 20 µg/mL) and the LY/CUR combinations (200/5 and 200/10 µg/mL) for 24 h, being dissolved in the vehicle (DMEM medium). Cells in the control group were treated with the vehicle alone. The cell viability was measured via CCK-8 analysis following the instructions provided by the vendor. The experiments were triplicated and independently repeated at least twice.

3.4. Animal Study

Male Sprague Dawley rats (4 months old; 250–300 g), purchased from Guangdong Medical Laboratory Animal Center (Guangzhou, China), were housed with free access to food and water at the animal facility of South China University of Technology with a controlled environment of a 12 h light/dark cycle, temperature ($22 \pm 2 °C$), and humidity ($55 \pm 9\%$). After one week of acclimatization, the rats were castrated and recovered for one week before BPH induction via the subcutaneous injection of TP (5 mg/kg BW, dissolved in 200 µL of olive oil) daily for 8 weeks (Figure 1). Rats in the control group were injected subcutaneously with olive oil alone. The animal study was reviewed and approved by the Animal Ethics Committee of South China University of Technology.

The castrated rats were randomly assigned into one of the following six experimental groups (n = 6/group) and received the corresponding treatment daily starting 4 weeks after the initiation of BPH induction (Table 2): (i) control group (CON): orally administered 1 mL of olive oil; (ii) BPH model group (BPH): administered orally with 1 mL of olive oil; (iii) BPH + FN group (FN): orally administered FN (5 mg/kg BW); (iv) BPH + LY group (LY): orally administered LY (12.5 mg/kg BW); (v) BPH + CUR group (CUR): orally administered CUR (2.4 mg/kg BW); (vi) BPH + LY + CUR group (COM): orally administered LY and CUR (12.5 + 2.4 mg/kg BW). After 28 consecutive days of treatments, the experiment was concluded. Subsequently, the rats were euthanized, and expeditiously, blood samples and prostates were extracted for subsequent analyses. A diagram of the experimental protocol is shown in Figure 9.

Group	LY mg/(kg.d) BW	CUR mg/(kg.d) BW	FN mg/(kg.d) BW	Oilve Oil (mL)
(i) CON	0	0	0	1
(ii) BPH	0	0	0	1
(iii) FN	0	0	5	1
(iv) LY	12.5	0	0	1
(v) CUR	0	2.4	0	1
(vi) COM	12.5	2.4	0	1

Table 2. Experimental groups.

3.5. Histopathology and Immunohistochemistry (IHC)

The prostate tissues underwent fixation in 4% paraformaldehyde at 4 °C for 24 h, followed by dehydration and paraffin embedding and sectioning at a thickness of 5 μ m. Hematoxylin and eosin (H&E) staining was utilized for histopathology, while immunohistochemistry (IHC) involved boiling the tissue slices in critic acid (pH 6.0) for 30 min for

antigen retrieval, soaking them in 3% H_2O_2 for 10 min and blocking them with 2% normal goat serum (diluted 1:10 in PBS) for 30 min at 20 °C. The slices were then incubated with primary antibody overnight and secondary antibody for 30 min. Fields were selected using a systematic random sampling scheme [31]. Images were acquired using a light microscope (Nikon Eclipse E100, Tokyo, Japan), and the density was quantified by using ImageJ 1.50i.



Figure 9. Experimental flow chart.

3.6. Enzyme-Linked Immunosorbent Assay (ELISA)

The serum levels of DHT, T, E2, 5α -reductase, PSA, IL-1 β , IL-6 and TNF- α were measured via ELISA following the protocols provided by the manufacturer.

3.7. Network Pharmacology

The gene targets of LY and CUR were identified through a comprehensive approach that involved the utilization of the Drugbank database, the Swisstarget Prediction database and the published literature. The target organism selected for this study was Homo sapiens. The BPH-associated gene targets were obtained from the Targetnet database, the UniProt database and the Genecards database. To establish a gene library of anti-BPH targets for LY and CUR, a comparative and analytical approach was employed to identify common targets between BPH-associated targets and the predicted targets of LY and CUR.

The collected BPH target library and the LY or CUR target library were analyzed using Venny (https://bioinfogp.cnb.csic.es/tools/venny/) (accessed on 12 December 2022) to obtain the LY- and CUR-BPH target intersection. The relationship between the targets was established using the STRING database (https://string-db.org/) (accessed on 12 December 2022), and the network characteristics of the LY- and CUR-target association network were analyzed using Cytoscape 3.7.2. Finally, the cross-targets were analyzed, and a pathway enrichment diagram was generated using the KEGG database.

3.8. Statistical Analysis

Quantitative data were reported as means \pm SD and subjected to statistical analysis. Statistical significance was assessed using one-way analysis of variance (ANOVA) followed by Tukey's comparison test, with GraphPad Software (version 6.02 for windows, San Diego, CA, USA) employed for this purpose. A *p*-value of less than 0.05 was deemed statistically significant. The figures display *p*-values for comparisons between treatment groups and the BPH group, with *, *p* < 0.05; **, *p* < 0.01; ***, *p* < 0.001. *p*-values for comparisons between the BPH group and the CON group are also presented on the figures, with #, *p* < 0.05; ##, *p* < 0.01; ###, *p* < 0.001.

The combination index (CI) was calculated to determine the nature of the LY and CUR combinations based on the methods described with appropriate modifications [32,33]. In brief, the expected value of the combination effect between treatment 1 and treatment 2 was calculated as [(observed treatment 1 value)/(control value)] × [(observed treatment 2 value)]/(control value)] × (control value); and the CI was calculated as the ratio of (observed value of the combination effect)/(expected value of the combination effect). The CI values of <1, >1 and =1 indicate a synergistic, an antagonistic and an additive effect, respectively.

4. Discussion

In this work, we demonstrated (1) that the combination effectively attenuated BPH development in rats through the quantitative analysis of histopathology and the prostate index, (2) that the combination treatment decreased the production of both dihydrotestosterone (DHT) and 5α -reductase, (3) that inflammatory responses can be greatly reduced via combination therapy in the BPH model of rats and (4) that AKT1, TNF, EGFR, STAT3 and PTGS2 are main targets which are associated with the anti-BPH activity of lycopene and curcumin.

Although BPH is prevalent among elderly men worldwide, its pathogenesis remains unclear, impeding the advancement of efficacious and secure treatments. The potential mechanisms underlying pathogenesis include systemic/local hormonal and vascular changes that are age-related, as well as the imbalance of cell apoptosis and proliferation and persistent inflammation [34]. Increasing evidence suggests that inflammation may have a key role in BPH development and progression. Inflammation is frequently accompanied with the infiltration of inflammatory cells in prostates, and the infiltrated cells produce cytokines, which may stimulate local growth factor production and angiogenesis [35].

Indeed, four weeks of the oral combination of LY and CUR administration showed promising effects in inhibiting BPH in rats. In particular, combined treatment nearly normalized the prostatic weight and index compared with the BPH group with few side effects. In addition, the inhibition in the progression of BPH in the COM group was more effective than in the LY and CUR group alone. Histologically, the combination treatment was able to significantly improve the prostatic structural organization and reduce fibrotic tissue formation.

In this research, we found that the combination of LY and CUR can effectively reduce the production of DHT, 5α -reductase, E2 and PSA (Figure 10). Disordered hormone production is a widely recognized risk factor for BPH due to the crucial role played by hormones such as estrogen, testosterone and DHT in regulating cellular proliferation and apoptosis in the prostate gland. DHT exhibits a 3-fold higher affinity for androgen receptors (ARs) than testosterone, which is converted to DHT via 5α -reductase activity. The serum concentration of DHT has been found to be positively correlated with the incidence and progression of BPH [36,37]. E2, a metabolite of T, is synthesized by the enzyme aromatase and is expressed in the urogenital tract along with fat [38]. Studies have demonstrated a positive correlation between elevated serum estrogen levels or an increased estrogen/androgen ratio and the development of BPH [39].



Figure 10. Molecular mechanism diagram.

BPH is significantly associated with chronic inflammation during the aging process. A study demonstrated that the histopathological analysis of 3942 BPH cases revealed a prevalence of 43.1% of inflammatory features, primarily characterized by chronic and mild inflammation [40]. The presence of T lymphocytes and macrophages within the prostate gland leads to the upregulation of cell growth factors and inflammatory cytokines in cases of BPH [41]. Inflammatory factors like interleukin (IL)-6, IL-1 α and IL-1 β may be secreted by the infiltrated lymphocytes, which can stimulate both epithelial and stromal cell proliferation [42,43]. In this research, the inflammatory factors including TNF- α , IL-6 and IL-1 β were both significantly reduced by the combined treatment.

The network pharmacology analysis identified 52 matched targets among the predicted targets of LY, CUR and the osteoporosis-associated targets, 12 targets for LY and 40 targets for CUR, which proves that LY and CUR inhibit the progression of BPH via different targets, and their combined treatment covers more comprehensive targets, and thus can improve the therapeutic efficiency. In the PPI network, AKT1 and TNF are the top-ranked genes. AKT1 plays a crucial role in cellular proliferation and viability, exhibiting expression in a diverse range of tissues [44]. TNF primarily facilitates the adhesion and migration of inflammatory cells through upregulating the expression of cell adhesion factors, thereby exerting a crucial influence on the onset and progression of prostatic inflammation. According to Mostafa et al., the passive expansion of peripheral blood vessels facilitates the acquisition of adequate oxygen and nutrients by tissues, thereby promoting prostate cell proliferation and ultimately leading to the development of BPH [45]. Further pathway analysis suggested that the LY and CUR combination's inhibition of BPH may be in part through regulating pathways in cancer.

In conclusion, we have demonstrated that combined (LY and CUR) treatment can synergistically inhibit the progression of BPH by reducing the expression of inflammatory factors and cell proliferation marker Ki-67 and regulating the hormone level. Additionally, the network pharmacology analysis suggests that the combination of LY and CUR can cover more comprehensive targets and thus improve the therapeutic efficiency. Moreover, it is also important to improve the targeting and bioavailability of drugs in the future due to the poor water solubility and bioavailability of curcumin. Nanotechnology like lipid-based nanoparticles, exosomes and carrier-free nanodrugs would be the focus of our future research [46–49]. Additionally, translational trials are further required to determine if the combination of LY and CUR is indeed beneficial and more effective for BPH patients.

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Abbreviations

BPH	benign prostatic hyperplasia
TP	testosterone propionate
LY	lycopene
CUR	curcumin
FN	finasteride
COM	combination
CON	control
H&E	hematoxylin and eosin
DHT	dihydrotestosterone
IL	interleukin
TNF-α	tumor necrosis factor-alpha
IHC	immunohistochemistry
ELISA	Enzyme-linked Immunosorbent Assay
E2	estradiol
PSA	prostate-specific antigen
CCK-8	Cell Counting Kit 8
Т	testosterone

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