

Inhibition of HIV-1 by curcumin A, a novel curcumin analog

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Abstract: Despite the remarkable success of combination antiretroviral therapy at curtailing HIV progression, emergence of drug-resistant viruses, chronic low-grade inflammation, and adverse effects of combination antiretroviral therapy treatments, including metabolic disorders collectively present the impetus for development of newer and safer antiretroviral drugs. Curcumin, a phytochemical compound, was previously reported to have some in vitro anti-HIV and anti-inflammatory activities, but poor bioavailability has limited its clinical utility. To circumvent the bioavailability problem, we derivatized curcumin to sustain retro-aldol decomposition at physiological pH. The lead compound derived, curcumin A, showed increased stability, especially in murine serum where it was stable for up to 25 hours, as compared to curcumin that only had a half-life of 10 hours. Both curcumin and curcumin A showed similar inhibition of one round of HIV-1 infection in cultured lymphoblastoid (also called CEM) T cells ($IC_{50}=0.7 \mu\text{M}$). But in primary peripheral blood mononuclear cells, curcumin A inhibited HIV-1 more potently ($IC_{50}=2 \mu\text{M}$) compared to curcumin ($IC_{50}=12 \mu\text{M}$). Analysis of specific steps of HIV-1 replication showed that curcumin A inhibited HIV-1 reverse transcription, but had no effect on HIV-1 long terminal repeat basal or Tat-induced transcription, or NF- κ B-driven transcription at low concentrations that affected reverse transcription. Finally, we showed curcumin A induced expression of *HO-1* and decreased cell cycle progression of T cells. Our findings thus indicate that altering the core structure of curcumin could yield more stable compounds with potent antiretroviral and anti-inflammatory activities.

Keywords: curcumin A, HIV-1, reverse transcription, heme oxygenase-1

Introduction

Curcumin, a natural phytochemical derived from turmeric has been reported to have a number of biological effects including inhibition of HIV-1 infection.^{1,2} Curcumin was shown to inhibit HIV-1 transcription,³ particularly through Tat-activated HIV-1 transcription.³⁻⁵ It was also shown to prevent the release of HIV-1 from chronically infected cells,⁶ inhibit in vitro HIV-1 protease⁷ and HIV-1 integrase,^{8,9} and demonstrated in silico binding to HIV-1 reverse transcriptase,¹⁰ protease,¹¹ and integrase.¹² However, these in vitro effects are yet to be confirmed with live virus in cultured cells. Additionally, curcumin was shown to protect neurons¹³ and to have a positive effect on spatial memory in HIV-1-associated dementia in a rat model.¹⁴

Curcumin affects cell cycle¹⁵ and also induces *HO-1*.² *HO-1* plays a critical role in the regulation of inflammation and immune response.¹⁶⁻¹⁸ Induction of *HO-1* inhibits HIV-1,¹⁹ hepatitis C virus,²⁰ and Ebola virus.²¹ We recently showed that *HO-1* induction led to the production of macrophage inflammatory cytokines and decreased surface expression of CCR-5 in lipopolysaccharide (LPS)-treated monocyte-derived macrophages providing an additional molecular mechanism of HIV-1 inhibition by LPS in macrophages.²² Thus, it was of interest to analyze the effect of curcuminoids on *HO-1* expression.

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Despite many reported pharmacological properties of curcumin^{23–25} and its low toxicity in clinical trials,^{2,26} it demonstrated poor absorption, rapid metabolism, rapid systemic elimination,^{27–29} and overall poor bioavailability.^{28–31} We sought to improve the bioavailability of curcumin by developing structural analogs of curcumin by taking a previously reported approach in which β -diketone moiety of curcumin was removed,²⁸ and to test these analogs for HIV-1 inhibition and *HO-1* induction.

We synthesized a novel curcuminoid, curcumin A and obtained a pure compound. We analyzed stability of curcumin A in parallel to curcumin in mouse serum, media, fetal bovine serum-free media, and complete media. We also analyzed stability of curcumin and curcumin A in phosphate-buffered saline (PBS) and in PBS supplemented with 10% bovine serum albumin (BSA). We then tested curcumin A and the parental curcumin for HIV-1 inhibition in cultured T cells and primary peripheral blood mononuclear cells (PBMCs) infected with vesicular stomatitis virus glycoprotein-pseudotyped HIV-1. We also determined toxicity of the compounds in T cells and PBMCs. To establish a mechanism of the curcumin-mediated inhibition of HIV-1, we analyzed the effect of curcumin on early and later stages of HIV-1 infection using single round HIV-1 infection with vesicular stomatitis virus glycoprotein-pseudotyped HIV-1. Finally, we tested *HO-1* induction by curcumin and curcumin A and their effect on cell cycle progression. Our study demonstrates that curcumin A is a stable curcuminoid that has strong antiretroviral activity by targeting HIV-1 reverse transcription (RT) and inducing *HO-1*. Thus, altering the core structure of curcumin could yield more stable compounds with potent antiretroviral and anti-inflammatory activities.

Materials and methods

Cells were obtained from ATCC, (American Type Culture Collection, Manassas, VA, USA). PBMCs were purchased from Astarte Biologics (Redmond, WA, USA).

Plasmids

HIV-1 proviral vector pNL4-3.Luc.R-E- (courtesy of Prof Nathaniel Landau, NYU School of Medicine, New York, NY, USA) was obtained from the NIH AIDS Research and Reference Reagent Program. HIV-1 LTR luciferase expression vectors were kindly provided by Dr Manuel López-Cabrera (Unidad de Biología Molecular, Madrid, Spain):³² HIV-1 LTR (–105 to +77) followed by the luciferase reporter gene, (HIV LTR 2 \times NF- κ B 3 \times SP1), HIV-1 LTR, NF- κ B and SP1 sites followed by luciferase reporter gene.

Synthesis of curcumin A

The chemical synthesis of the acyclic analog of curcumin (curcumin A) was accomplished via a slight modification of the previously reported procedure.²⁸ In brief, acetic acid (50 mL) was saturated with hydrogen chloride for a period of 1 hour at 0°C. A mixture of vanillin (3.04 g, 20 mmol, 2.0 equiv) and acetone (580 mg, 10 mmol, 1.0 equiv) was slowly added to acetic acid. The reaction mixture was slowly brought to room temperature and stirred for 48 hours. The crude reaction mixture was poured into ice-cold water (200 mL) and the precipitated solid was isolated via vacuum filtration. It was dried and purified using silica gel flash column chromatography. Gradient elution with 30%–60% ethyl acetate in hexanes resulted in the isolation of the desired product (1.93 g, 59% yield) as yellowish orange solid.

Solubility of curcumin A

To explore the solubility of curcumin A, an excess amount of curcumin A was added to 1 mL of polyethylene glycol 400 in a 5 mL stopper vial, vortex mixed, equilibrated, and analyzed by high-performance liquid chromatography, using a previously established technique.³³ The solubility was 86.17 \pm 4.34 mg/mL, suggesting that curcumin A has good solubility in a pharmaceutically approved excipient.

Single round HIV-1 replication assay

CEM-T cells or PBMCs were infected with VSVG-pseudotyped pNL4-3.Luc.R-E-virus (HIV-1 Luc) prepared as previously described.³⁴ PBMCs were stimulated by treating with 2.5 μ g/mL phytohemagglutinin (PHA) for 24 hours and then activated for another 24 hours with 10 units/mL IL-2 before the infection with HIV-1 Luc. PBMCs were infected with HIV-1 Luc, cultured at 0.5 \times 10⁶ cells/mL in 6-well plates at 37°C and 5% CO₂ for 24 hours and then treated with indicated concentration of compounds. The cells were collected after 48 hours in culture, washed with PBS, and resuspended in 100 μ L of PBS. Then, 100 μ L of reconstituted luciferase buffer (LucLite Kit; PerkinElmer Inc., Waltham, MA, USA) was added to each sample and, after 10 minutes incubation, the lysates were transferred into white plates (PerkinElmer Inc.) and luminescence measured using Labsystems Luminoscan RT equipment (PerkinElmer Inc.).

Calcein-AM uptake cell viability assays

PBMCs were cultured as described above in 96-well plates at 37°C. To measure cytotoxicity with calcein, cells were supplemented with 0.2 μ M calcein-AM (Thermo Fisher Scientific, Waltham, MA, USA) for 30 minutes at 37°C.

Fluorescence was measured using the luminescence spectrometer implementing an excitation wavelength of 495 nm and emission filters at 515 nm. To measure cellular viability with trypan blue, the cells were supplemented with 0.2% trypan blue, transferred to a plastic disposable counting chamber, and counted on a Cellometer Automatic Cell Counter (Nexcelcom Bioscience, Lawrence, MA, USA).

Cell cycle analysis of CEM-T cells treated with curcumin A

Approximately one million cells were fixed in 70% ethanol at -20°C for 2 hours and stained with propidium iodide (10 mg/mL) containing RNase A (1 mg/mL) for 30 minutes. The data were acquired in BD FACS Calibur (BD Biosciences, San Jose, CA, USA), and analyzed using FlowJo software. Unpaired Student's *t*-test was used to determine statistical significance.

Analysis of *HO-1* messenger (m)RNA expression

CEM-T cells or PBMCs were treated with curcumin or curcumin A (2 μM) for 24 hours. Total RNA was extracted from cultured cells using TRIzol reagent according to the manufacturer's protocol (Thermo Fisher Scientific). Total RNA (100 ng) was reverse-transcribed to complementary (c)DNA using SuperscriptTM RT-PCR (reverse transcription polymerase chain reaction) kit (Thermo Fisher Scientific); hexamers and oligo-dT were used as primers. For RT-PCR analysis, cDNA was amplified using Roche LightCycler 480 (Hoffman-La Roche Ltd., Basel, Switzerland) and SYBR Green1 Master mix (Hoffman-La Roche Ltd.). PCR was carried with denaturation at 95°C for 10 seconds, annealing at 60°C for 10 seconds, and extension at 72°C for 10 seconds for 45 cycles. For quantification of mRNA levels for *HO-1*, 18S ribosomal (r)RNA was used as a house keeping normalization standard. The following primer sequences were used for *HO-1*, forward-CTTCTTCACTTCCCAACA; reverse-GCTCTGGTCCTTGGTGTCAT, amplicon size 193; for 18S rRNA, forward-GCTGTTGCTACATCGACCTTT; reverse-CTCCA-GGTTTTGCAACCAGT, amplicon size 168. Mean crossing point values for *HO-1* and 18S rRNA were determined and relative changes in gene expression were analyzed using an algorithm known as $\Delta\Delta\text{Ct}$ method. Unpaired Student's *t*-test was used to determine statistical significance.

Quantitative PCR for viral DNA

CEM-T cells or stimulated PBMCs were infected with HIV-1 Luc and treated with 2 μM curcumin, curcumin A or 1 μM azidothymidine (AZT) and further cultured for 48 hours. Total DNA was

extracted from 4×10^6 cells using lysis buffer (10 mM Tris-HCl pH 8, 10 mM ethylenediaminetetraacetic acid, 5 mM NaCl, 200 $\mu\text{g}/\text{mL}$ proteinase K). The cells were lysed for 20–30 minutes at room temperature and proteinase K was inactivated by heating to 95°C for 5 minutes. For the RT-PCR analysis, 100 ng DNA was amplified using Roche Light Cyclers 480 and SYBR Green1 Master mix. PCR was carried out with initial preincubation for 5 minutes at 45°C and then for 3 minutes at 95°C followed by 45 cycles of denaturation at 95°C for 15 seconds, annealing and extension at 60°C for 45 seconds, and final extension at 72°C for 10 seconds. Quantification of early-LTR and late-LTR was carried out using β -globin DNA as a normalization standard. Primer sequences for early-LTR, forward-GGCTAACTAGG-GAACCCACTG, reverse-CTGCTAGAGATTTTCCACACT-GAC; for late-LTR forward-TGTGTGCCCGTCTGTTGTGT, reverse-GAGTCCTGCGTCGAGAGATC; and for globin, forward-CAACCTCAAACAGACACCATGG, reverse-TC-CACGTTACCTTGCCC.³⁵ Mean crossing point values for early-LTR, late-LTR, and β -globin were determined and $\Delta\Delta\text{Ct}$ method was used to calculate relative expression levels. Unpaired Student's *t*-test was used to test statistical significance.

Analysis of curcumin A stability

Serum incubation

Curcumin or curcumin A was dissolved in dimethylsulfoxide (DMSO) (10 mM), and then diluted with PBS to 100 μM , which was mixed with an equal volume of mouse serum. The final concentration of curcumin or curcumin A was 50 μM , and the amount of organic solvent in the mixture was less than 1% (v/v). The mixture was incubated at 37°C and samples were collected at 0 hours, 4 hours, 8 hours, 12 hours, 16 hours, 20 hours, and 24 hours. The samples (60 μL) were mixed with 240 μL of cold acetone, vortexed for 2 minutes, kept at -20°C for 30 minutes, and then proteins were removed by centrifugation at $13,000 \times g$ for 5 minutes. The supernatant was transferred to a clean test tube and evaporated to dryness using a Speedvac concentrator. The pellet was reconstituted in 50 μL of acetonitrile, and a 10 μL aliquot was injected for analysis.

Media or PBS incubation

Curcumin or curcumin A was dissolved in DMSO (10 mM), and then diluted with Roswell Park Memorial Institute (RPMI) 1640 media, PBS or PBS supplemented with 10% BSA. The solutions were incubated at 37°C and samples were collected at different time points within 24 hours incubation. The sample (100 μL) was mixed with 400 μL of cold acetone, vortexed for 2 minutes, kept at -20°C for 30 minutes, and then proteins were removed by centrifugation at $13,000 \times g$ for 5 minutes.

The supernatant was transferred to a clean test tube and evaporated to dryness using a Speedvac concentrator. The pellet was reconstituted in 50 μ L of acetonitrile, and a 10 μ L aliquot was injected onto the nano C18 column for the liquid chromatography-mass spectrometry (LC-MS) for analysis.

Nano-LC-MS condition

The samples were loaded onto a nano-C18 column attached to Shimadzu nano-LC coupled in-line to LTQ Orbitrap XL tandem mass spectrometer (Thermo Fisher Scientific). The injection volume was 10 μ L. The mobile phase consisted of a 0.1% formic acid aqueous solution (A) and a 0.1% formic acid acetonitrile solution (B). The gradient elution program was as follows: 0–6.02 minutes, 1% B; 6.02–6.11 minutes, 1%–2% B; 6.11–20 minutes, 2%–80% B; 20–25 minutes, 80% B; 25–30 minutes, 80%–85% B; 30–31 minutes, 80%–2% B; 31–40 minutes, 2% B (v/v). The flow rate was set to 600 nL/min. The compounds were ionized by electrospray ionization and detected by Orbitrap

at 30,000 mass resolution (full scan, m/z 150–2,000). The spray voltage, capillary temperature, and capillary voltage were set to 2.0 kV, 200°C, and 39.5 V, respectively.

Results and discussion

Synthesis of curcumin A

Presence of β -diketone moiety in curcumin makes it prone to decomposition at physiological pH (see structure in Figure 1A). To exclude the possibility that curcumin contained contaminants, we analyzed its composition by LC-MS analysis and did not find demethoxycurcumin or bisdemethoxycurcumin, typical contaminants (Figure 1A). To improve curcumin stability, we synthesized curcumin A (as described in Materials and methods) in which β -diketone moiety was removed (Figure 1B). Removal of the β -diketone moiety from curcumin yielded yellowish orange solid of molecular weight 327.1232 Mp 100–102°C.

Curcumin A structure was verified by nuclear magnetic resonance analysis ($^1\text{H-NMR}$) (CDCl_3 , 400 MHz, ppm) δ 7.66

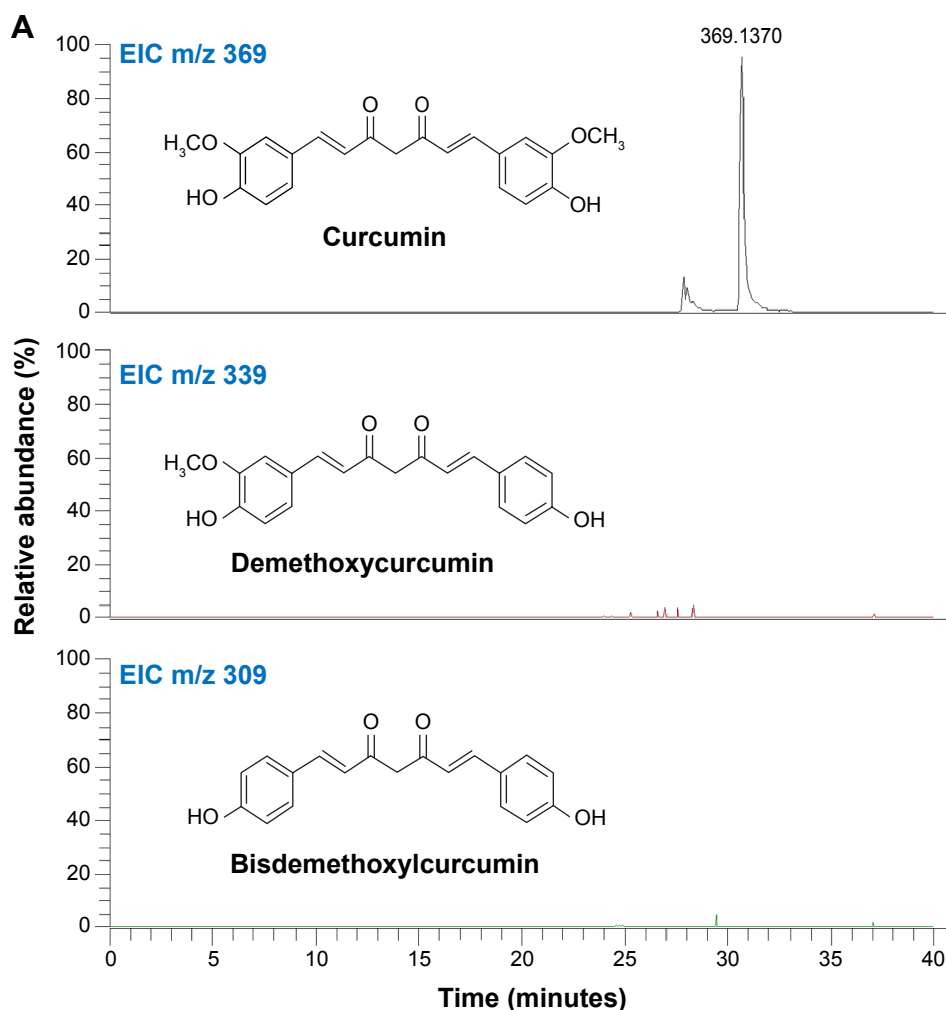


Figure 1 (Continued)

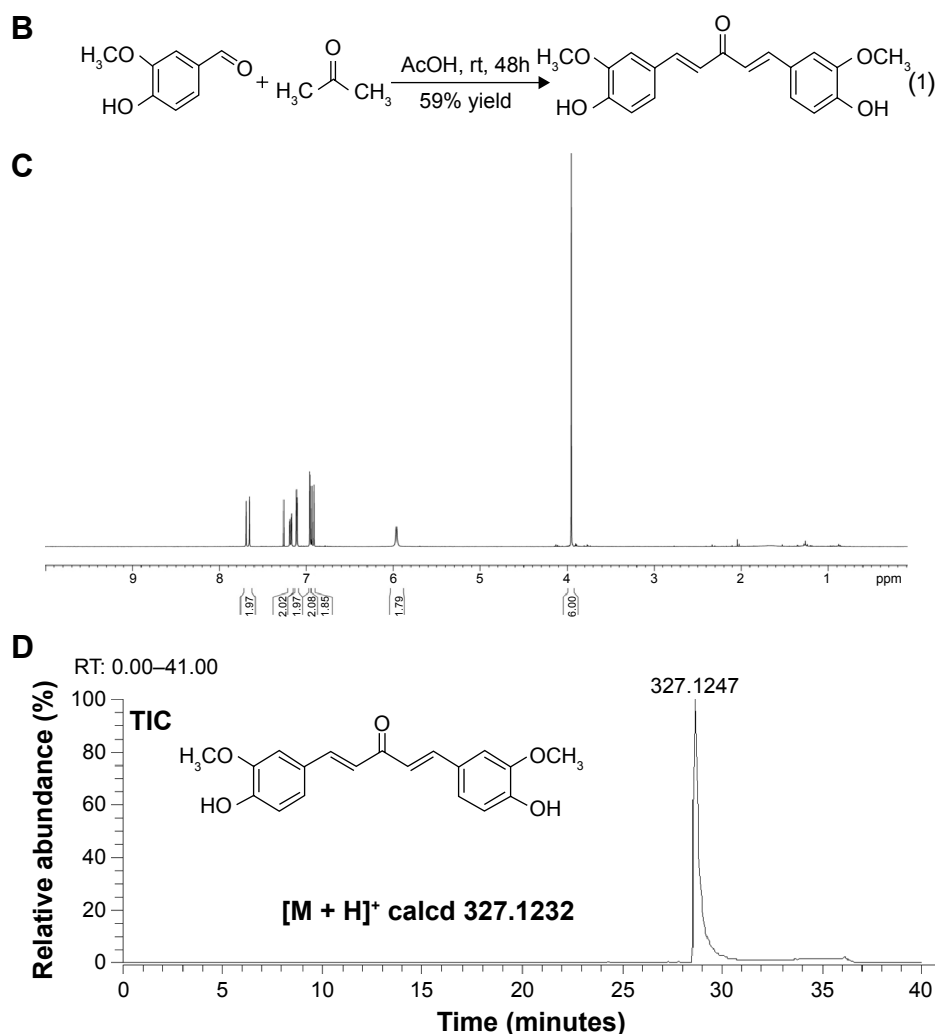


Figure 1 Synthesis and structures of curcumin A.

Notes: (A) The purity analysis of curcumin by nano-LC-FTMS. (B) General step for the synthesis of curcumin A. (C) ¹H-NMR spectrum of purified curcumin A in CDCl₃. The spectrum was recorded on 400 MHz NMR instrument. (D) Nano-LC-FTMS analysis of curcumin A.

Abbreviations: TIC, total ion current; EIC, extract ion chromatogram; rt, room temperature; h, hours; RT, reverse transcription; LC-FTMS, liquid chromatography - Fourier transform mass spectrometry; calcd, ; ¹H-NMR, .

(d, *J* = 15.6 Hz, 2H), 7.18 (dd, *J* = 8.0, 1.6 Hz, 2H), 7.11 (d, *J* = 2 Hz, 2H), 6.96 (d, *J* = 1.6 Hz, 2H), 6.91 (d, *J* = 9.6 Hz, 2H), 5.96 (s, 2H), 3.95 (s, 6H) (Figure 1C). Other spectroscopic data (carbon-13 nuclear magnetic resonance spectroscopy, infrared spectroscopy, and high-resolution mass spectrometry) for curcumin A were consistent with the previously reported literature and its predicted chemical structure (not shown). Curcumin A purity was further analyzed by LC-MS high resolution mass spectrometry which shows a single peak with the expected molecular weight (Figure 1D). Thus we obtained pure curcumin A with the expected structure which was analyzed for its stability and inhibition of HIV-1.

Curcumin A has an improved stability

To analyze stability of curcumin A and compare it to curcumin, the compounds were incubated with mouse

plasma, in complete or minimal cell culture media, and in PBS. The compounds were extracted from the incubation mixtures and resolved by nano-LC-MS mass spectrometry as described in Materials and methods. While curcumin had a half-life of 10 hours in mouse serum, curcumin A was stable for up to 24 hours of incubation (Figure 2A). In serum-free media, both curcumin and curcumin A were degraded, but curcumin A had a longer half-life (about 3 hours) compared to curcumin (about 1 hour) (Figure 2B). In complete media, curcumin and curcumin A were equally and relatively stable (Figure 2C). Incubation in PBS again showed increased stability of curcumin A (Figure 2D). To test whether BSA has a stabilizing effect on curcumin, curcumin and curcumin A were incubated in PBS supplemented with 10% BSA (Figure 2E). No stabilization and even a slight increase in degradation of curcumin was

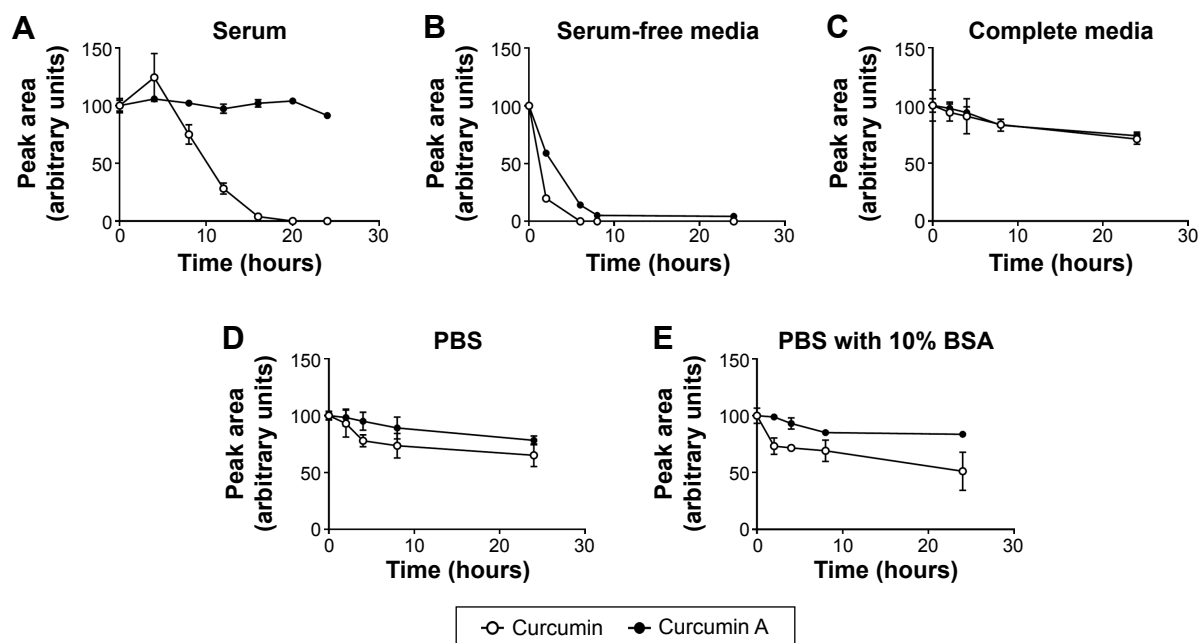


Figure 2 Stability of curcumin A in serum, serum-free media, complete media, PBS, and PBS supplemented with 10% BSA.

Notes: Degradation of curcumin and curcumin A at different time points spanning 24 hours incubation, in presence of mouse serum (A), RPMI cell culture medium (B), RPMI supplemented with 10% fetal bovine serum (C), in PBS (D), and in PBS supplemented with 10% BSA (E).

Abbreviations: PBS, phosphate-buffered saline; BSA, bovine serum albumin; RPMI, Roswell Park Memorial Institute.

observed suggesting that BSA has no stabilizing effect on curcumin (Figure 2E). Taken together, curcumin A has improved stability and longer half-life, in serum, serum-free media, and PBS.

Curcumin A inhibits one round of HIV-1 infection

The effects of curcumin and curcumin A on one round of HIV-1 infection were first analyzed in cultured CEM-T cells infected with VSVG-pseudotyped HIV-1 pNL4-3 virus expressing luciferase in place of *nef* (HIV-1 Luc).³⁴ The cells were treated with the compounds for 24 hours and luciferase activity was measured as an indicator of HIV-1 replication. One round HIV-1 infection was equally inhibited by curcumin (the half maximal inhibitory concentration [IC_{50}]=0.7 μ M) and curcumin A (IC_{50} =0.8 μ M) (Figure 3A). Viability of CEM-T cells was analyzed using calcein-AM assay which measured fluorescence of the calcein converted intracellularly from the non-fluorescent calcein-AM taken up by the cells. Curcumin and curcumin A reduced viability of CEM-T cells when added to the cells for 24 hours, with curcumin being slightly more toxic (IC_{50} =1.26 μ M) than curcumin A (IC_{50} =2.4 μ M) (Figure 3B). We next analyzed the effect of curcumin and curcumin A on single round HIV-1 infection in primary PBMCs. PBMCs were activated by treatment with PHA and IL-2 (see Materials and methods

for details), infected with HIV-1 Luc for 24 hours and then treated with curcumin or curcumin A for another 24 hours. Both curcumin and curcumin A markedly inhibited HIV-1 infection with curcumin A being more potent (IC_{50} =2 μ M) than curcumin (IC_{50} =12 μ M) (Figure 3C). Both curcumin and curcumin A reduced viability of PBMCs as analyzed using trypan blue, with IC_{50} s of 35 μ M and 22 μ M, respectively (Figure 3D). Thus, curcumin A inhibits one round HIV-1 infection comparable to curcumin in CEM-T cells and displays more potent activity in primary PBMCs. Also, curcumin A demonstrated a better therapeutic window in both cultured CEM-T cells and PBMCs.

Curcumin and curcumin A inhibit HIV-1 RT

Previously, curcumin was shown to inhibit HIV-1 transcription by having an effect on chromatin remodeling.³⁶ To elucidate if curcumin and curcumin A inhibited HIV-1 transcription in the context of single round HIV-1 infection, we analyzed the expression of HIV-1 encoded *gag* and *env* mRNAs in CEM-T cells infected with HIV-1 Luc virus. Expression of these mRNAs was significantly reduced in infected cells treated with curcumin or curcumin A (Figure 4A). To determine whether the inhibition was due to the effect of curcumin or curcumin A on HIV-1 transcription, 293T cells were transfected with a combination of a reporter expressing luciferase under the control of HIV-1 LTR and a Tat expressing vector or

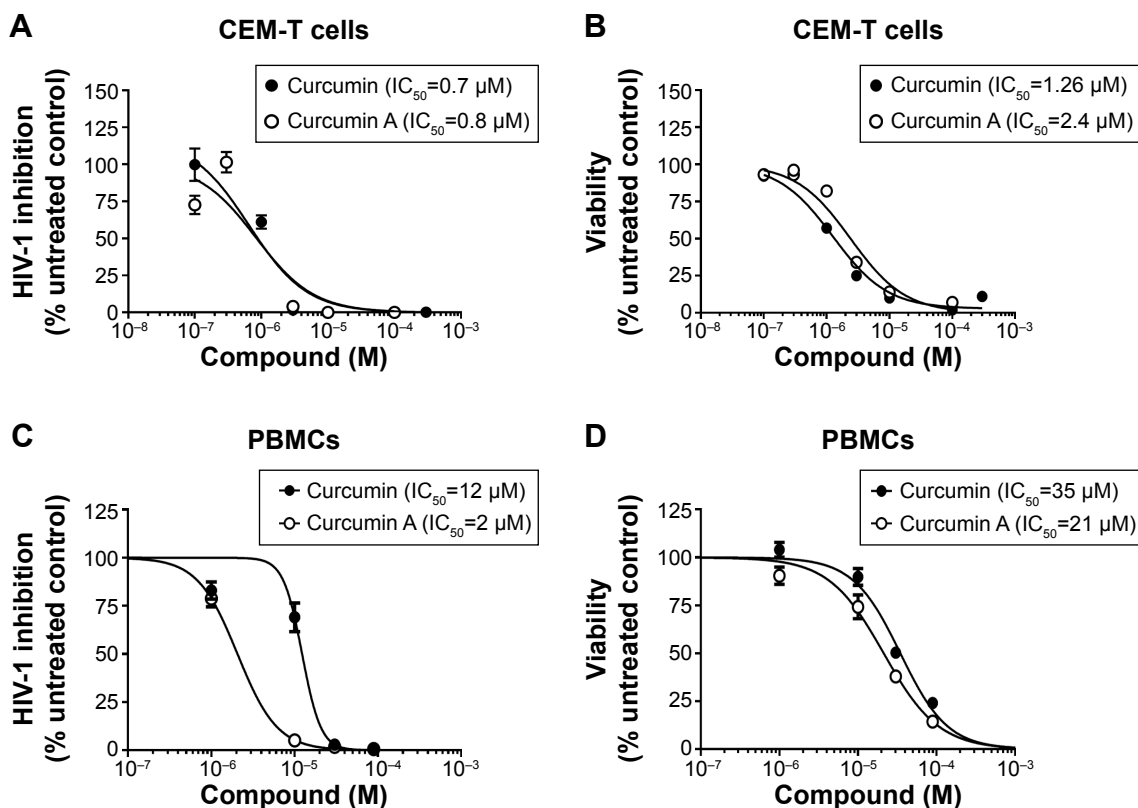


Figure 3 Effect of curcumin and curcumin A on one round of HIV-1-Luc replication and cellular viability in CEM-T cells and PBMCs.

Notes: CEM-T cells (A) or peripheral blood mononuclear cells (PBMCs) activated with PHA and IL-2 (C) were infected with VSVG-pseudotyped pNL4-3.Luc.R-E- (HIV-1 Luc) virus and treated with the indicated concentrations of curcumin or curcumin A for 48 hours at 37°C; then the cells were lysed and luciferase activity was measured. (B) CEM-T cells were treated with the indicated concentrations of curcumin or curcumin A for 48 hours at 37°C. CEM-T cells were treated with 0.4 μ M calcein-AM for 30 minutes and calcein fluorescence was measured at 485 nm excitation and 515 nm emission on the luminescence spectrometer equipped with the robotic arm (PerkinElmer LS 50B; PerkinElmer Inc., Waltham, MA, USA). (D) Activated PBMCs were treated with the indicated concentrations of curcumin or curcumin A for 48 hours at 37°C and viability of cells was measured by trypan blue exclusion method. IC_{50} s shown on panels (A–D) were determined with GraphPad Prism 6 Software (GraphPad Software, Inc., La Jolla, CA, USA).

Abbreviations: PHA, phytohemagglutinin; VSVG, vesicular stomatitis virus glycoprotein G; IC_{50} , the half maximal inhibitory concentration; M, molar concentration.

proviral pNL4-3 vector expressing luciferase in place of *nef*. Neither curcumin nor curcumin A at 2 μ M concentrations had an effect on Tat-induced HIV-1 transcription or transcription from proviral pNL4-3 plasmid (Figure 4B). We also analyzed the effect of curcumin and curcumin A on basal HIV-1 LTR transcription and NF- κ B driven transcription in 293T cells transfected with HIV-1 LTR luciferase vector or HIV LTR 2 \times NF- κ B 3 \times SP1 vector (described in Materials and methods). There was no effect of curcumin or curcumin A on basal HIV-1 transcription or NF- κ B driven transcription (Figure 4C) at 2 μ M concentration. We reasoned that curcumin and curcumin A might affect an earlier stage of HIV-1 infection and thus indirectly reduce the subsequent HIV-1 transcription step. We analyzed early and late RT by quantifying HIV-1 DNA for early and late LTR.³⁵ Both curcumin and curcumin A inhibited early LTR similarly or better than the established HIV-1 inhibitor, AZT (Figure 4D). In contrast, only curcumin A inhibited late LTR formation, similar to AZT, whereas

curcumin had no significant effect (Figure 4D). Thus, curcumin inhibited one round HIV-1 infection with an inhibitory effect on HIV-1 RT, while curcumin A showed more potent inhibition of both early and late HIV-1 RT.

Curcumin and curcumin A induce expression of *HO-1*

Previously, curcumin was shown to induce *HO-1*.² Activation of *HO-1* by heme was also shown to inhibit HIV-1.¹⁹ Thus we examined the effects of curcumin and curcumin A on *HO-1* in primary PBMCs and also in cultured CEM-T cells. The cells were treated with 1 μ M curcumin A or curcumin and *HO-1* expression was analyzed by mRNA quantification with quantitative PCR using 18S RNA for normalization. Both curcumin A and curcumin induced *HO-1* in CEM-T cells and PBMCs (Figure 5A). Thus, the HIV-1 inhibitory effects of curcumin and curcumin A could partly be due to the activation of *HO-1*.

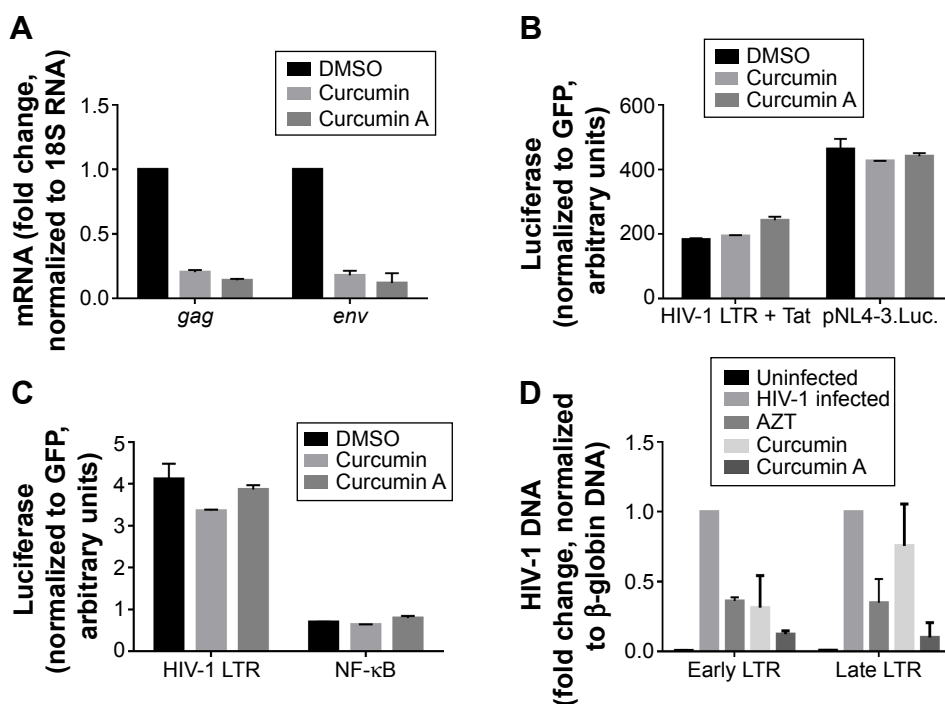


Figure 4 Effect of curcumin and curcumin A on HIV-1 messenger (m)RNA expression, HIV-1 transcription and HIV-1 reverse transcription.

Notes: (A) Effect of curcumin A on HIV-1 mRNA expression. CEM-T cells infected with HIV-1 Luc were treated with DMSO, 1 μ M curcumin or 1 μ M curcumin A for 48 hours as indicated. RNA was extracted, reverse transcribed, and analyzed with primers for HIV-1 *gag* and *env* genes by real-time PCR on Roche 4800 (Hoffman-La Roche Ltd., Basel, Switzerland) using 18S RNA as a reference. (B, C). Effect of curcumin and curcumin A on Tat-induced and basal HIV-1 transcription. In panel (B) 293T cells were transiently transfected with a vector contacting HIV LTR followed by the luciferase reporter (HIV LTR) and Tat expression vector or pNL4-3.Luc. In panel (C) 293T cells were transfected with HIV-1 LTR expression vector or with HIV LTR with the inactivated SPI sites (NF- κ B). For normalization, the cells were also co-transfected with GFP expressing vector. At 24 hours post-transfection the cells were treated with 1 μ M curcumin or curcumin A for 24 hours. Then the cells were lysed and luciferase activity was measured. GFP fluorescence was measured in parallel and used for normalization. (D) Effect of curcumin and curcumin A on HIV-1 reverse transcription. CEM-T cells were infected with HIV-1 Luc and then treated with DMSO, AZT, 1 μ M curcumin or curcumin A as indicated for 6 hours. DNA was extracted and analyzed by real-time PCR on Roche 4800 using primers for early and late LTR and β -globin gene as a reference.

Abbreviations: DMSO, dimethyl sulfoxide; mRNA, messenger RNA; PCR, polymerase chain reaction; AZT, azidothymidine.

Treatment with curcumin A affects cell cycle progression

Curcumin and its analogs were previously shown to have an effect on cell cycle progression and to cause apoptosis of cancer cells.³⁷ Thus we examined the effects of curcumin A and curcumin on the cell cycle progression of CEM-T cells. While curcumin had no effect on the cell cycle progression of CEM-T cells, the cells treated with curcumin A showed increased S phase accumulation (1.4 times) and decreased G2/M phase accumulation (three times) (Figure 5B). Thus this reduction of the cell cycle progression could be contributory to the effect of curcumin A.

Conclusion

We show here that curcumin A, which we derivatized from curcumin, has improved stability in serum and serum-free media over the parent compound, and also retained the ability of the parent compound to inhibit HIV-1 infection in CEM-T cells and PBMCs. Phase 1 clinical trials of curcumin have shown that efficacy of curcumin is limited by its poor bioavailability,

resultant from its poor absorption, rapid metabolism, and rapid systemic elimination.²⁷ This underscores the relevance of development of derivatives which retain the rich biological activities of curcumin, but also withstand the rapid degradation that has plagued the desired pharmacodynamics effect of the parent compound: curcumin A, the derivative we obtained in pure form, is stable in serum up to 24 hours as opposed to curcumin which is degraded by 50% within 10 hours. We also found that at 2 μ M concentration curcumin A and curcumin inhibited HIV-1 RT while having no effect on HIV-1 transcription. Further analysis of the mechanism of HIV-1 inhibition pointed to *HO-1* induction, in line with earlier reports of HIV-1 inhibition by *HO-1*,¹⁹ and alteration of cell cycle kinetics as putative mechanisms. The intended route of future administration of curcumin A is intranasal. Recently, curcumin was formulated as a nanoemulsion and successfully tested for intranasal delivery.³³ We have also developed micro- and nano-emulsions of curcumin A and are now optimizing these proprietary formulations for brain delivery through intranasal administration. In conclusion, the improved stability of

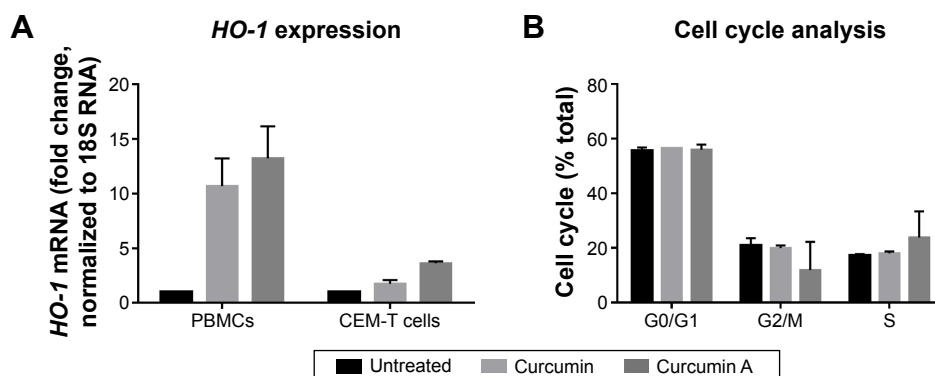


Figure 5 Effect of curcumin and curcumin A on *HO-1* expression and cell cycle progression.

Notes: (A) Expression of *HO-1*. CEM-T cells were treated with 1 μ M curcumin or curcumin A. DMSO was used as vehicle control. After 24 hours treatment, RNA was extracted, reverse transcribed, and analyzed by real-time PCR for *HO-1* using 18S RNA as a housekeeping control gene. (B) Effect of curcumin and curcumin A on the cell cycle progression. CEM-T cells were treated with 1 μ M curcumin or curcumin A for 24 hours and then fixed with 70% ethanol, stained with propidium iodide, and analyzed by fluorescence assisted cell sorting (FACS). Data were analyzed using BD FACS Calibur software (BD Biosciences, San Jose, CA, USA). Cells accumulated in G0/G1, G2/M and S phases of cell cycle are shown. Cell cycle phases: G0, resting phase; G1, gap 1 phase; G2, gap 2 phase; M, mitosis phase, and S, synthesis phase. All results are shown as a mean of three independent measurements \pm standard deviation.

Abbreviations: DMSO, dimethyl sulfoxide; mRNA, messenger RNA; PCR, polymerase chain reaction; PBMCs, peripheral blood mononuclear cells.

curcumin A over curcumin, and the retention of the in vitro anti-HIV-1 property of curcumin hold promise for the future potential clinical benefit of this derivative in antiretroviral therapy.

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Author contributions

AAK and EN developed curcumin and curcuminoids, including curcumin A. AAK, EN, and SN conceived the study. NK and CM analyzed antiviral and cellular effects of curcumin A. XL and TA conducted MS and stability analysis. AI purified HIV-1 virus and helped to conduct antiviral analysis. MH helped in preparing the manuscript. AAK, SN, and EN analyzed the data and wrote the manuscript. All authors contributed toward data analysis, drafting, and revising the manuscript.

Disclosure

The authors report no conflicts of interest in this work.

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