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Therapeutic potential of curcumin in regenerative dentistry

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Introduction: Natural compounds have emerged as promising candidates in drug development due to their potent immunomodulatory anti-inflammatory, antibacterial, analgesic, and healing properties. They have shown significant therapeutic potential in clinical applications, such as mouth rinses, toothpastes, and localized delivery systems. The use of natural alternatives can contribute to tackling antimicrobial resistance. Among natural compounds, curcumin has gained particular attention, demonstrating robust anti-cancer, antibiotic, and anti-inflammatory activities in numerous *in vivo* studies, while exhibiting a favorable safety profile for the treatment of various diseases. In this study, the remedial effects of curcumin and its metabolite, tetrahydrocurcumin, on dental pulp were explored. In addition, these results were compared with our previous findings on the effects of these natural compounds on periodontal ligament and gingival epithelial cells, further broadening our understanding of their therapeutic potential in oral disease such as caries and periodontitis.

Methods: RNA sequencing was used to investigate the differentially expressed genes in dental pulp cells following treatments with curcumin and tetrahydrocurcumin.

Results: We show that treatment of dental pulp cells with 1 μ M of curcumin or tetrahydrocurcumin is sufficient to promote Wnt signaling pathway in dental pulp cells. Curcumin treatment promotes the upregulation of cellular metabolism and enhances cellular response to stress. Our enrichment analysis shows that treatment with tetrahydrocurcumin modulates the extracellular matrix and angiogenesis.

Conclusions: The findings of this study highlight the cytoprotective and regenerative properties of curcumin and tetrahydrocurcumin. These properties could be leveraged as a therapeutic approach to promote tissue regeneration in oral diseases.

KEYWORDS

dental pulp, curcumin, tetrahydrocurcumin, tissue regeneration, natural therapeutic

Introduction

Natural compounds play a vital role in modern medical practices and health strategies due to their potential in modulating inflammation, oxidative stress, and cellular signaling pathways (1). Modern extraction techniques have improved the stability and bioavailability of many herbal compounds, making them even more effective for clinical applications (2). They can be used in treating chronic diseases such as diabetes, neurodegenerative disorders, cardiovascular conditions, and cancer (3–5). Incorporation of herbal medicine into clinical practices allows for the investigation of synergistic interactions between traditional remedies and modern pharmaceuticals (6–8).

Recent advances in regenerative dentistry are revolutionizing oral healthcare. Mesenchymal stem cells (MSCs) derived from dental pulp, periodontal ligament, or

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bone marrow possess remarkable potential for differentiating into various cell types, such as osteoblasts, odontoblasts, and fibroblasts (9-11). These properties enable their application in regenerating damaged oral tissues. In addition, the use of hydrogels and nanomaterials to support cell growth and differentiation enhances the precision and success of tissue regeneration. However, regulatory considerations, cost, and long-term stability remain challenging factors in incorporating these advances in the clinical setting (12-17). Natural availability, affordability, and the relatively low toxicity of some natural compounds render them a potential alternative in regenerative approaches. Multiple independent studies have demonstrated wound healing and anti-inflammatory properties of these compounds in vivo (18-21). As an example, citrus extract exhibits antibacterial, anti-inflammatory, and tissueregenerative characteristics (22). Bioactive compounds, such as polyphenols, terpenoids, alkaloids, saponins, and peptides, have been shown to be efficient in the management of periodontal disease (23). Curcumin, a polyphenolic compound derived from the rhizome of Curcuma longa, has gained scientific interest for its diverse pharmacological properties, including anti-inflammatory, antioxidant, and regenerative effects. Curcumin exerts its regenerative effects primarily by modulating signaling pathways crucial for cell proliferation, differentiation, and survival. It can regulate signaling pathways, such as Wnt/β-catenin and PI3K/Akt/ mTOR, and enhance stem cell viability and differentiation in neural repair (4, 5). In skin regeneration, curcumin has been shown to promote fibroblast migration and angiogenesis, which are key processes for wound healing (4). In bone regeneration, curcumin enhances osteoblast differentiation and mineralization by modulating RANKL/OPG signaling and reducing oxidative stressinduced damage (4, 6). Despite anti-inflammatory and proregenerative properties of curcumin, poor bioavailability and rapid metabolism and elimination of this compound have led to its limited clinical application. Tetrahydrocurcumin (THC) is one of the major metabolites of curcumin. The chemical structure of curcumin and THC are very similar, with THC lacking the double bonds that exist in the central seven-carbon chain of curcumin. THC is reported to have superior bioavailability than curcumin and exhibit a different molecular mechanism and biological activity. THC exhibits cytoprotective and anti-oxidative properties and has been proposed to be more potent than curcumin in the treatment of various conditions (24-26). In this study, the impact of curcumin and THC treatment on dental pulp was investigated. Using RNA sequencing, differentially expressed genes in human dental pulp cells were investigated to find an effective natural alternative in regenerative approaches in oral cavities.

Methods

Cell culture

Primary human dental pulp cells used in these assays were a kind donation from Dr. Ana Angelova at King's College London. Original cell harvest experiments were undertaken with the understanding and written consent of each individual and in full accordance with the World Medical Association Declaration of Helsinki (version 2002). The study was approved and followed the guidelines set by the Ethical Committee for human studies at King's College Hospital, King's College University of London. These cells were expanded in Full DMEM media with 10% fetal bovine serum (FBS).

Viability

Curcumin and tetrahydrocurcumin compounds were provided by Colgate Palmolive. For viability, 20,000 cells/cm² dental pulp cells were plated in triplicates in 96-well plates and incubated at 37°C, 5% CO₂ for 24 h. Curcumin and tetrahydrocurcmin were used at concentrations 100 µM, 50 µM, 10 µM, 1 µM and 0.5 µM to treat dental pulp cells. As the stock of these compounds was prepared in dimethylsulfoxide (DMSO), DMSO was used as the vehicle-only control in all the experiments in addition to the media-only control. Cell viability was assessed using an MTS Assay Kit (Cell Proliferation, Colorimetric, ab197010). A colorimetric plate reader (Thermo Multiskan Ascent 354 microplate reader) was used to read the absorbance at 490 nm. Normal distribution of results was tested using the Shapiro-Wilk test. Statistical significance against 100% viability in media was reported using a one-way ANOVA and Dunnett's multiple comparisons in GraphPad Prism 8.3.0. The adjusted p-value is reported in graphs according to New England Journal of Medicine guidelines: p < 0.001 (***), p < 0.002 (**), p < 0.033(*), and p > 0.12 (ns).

Quantitative polymerase chain reaction (qPCR)

Dental pulp cells were plated at 0.05×10^6 cells in triplicates in 24-well plates and incubated for 24 h (37°C, 5% CO₂/95% air, 100% humidity) using a standard culture medium. Media only and DMSO (vehicle only) were used as the negative and vehicle controls, respectively. 6-Bromoindirubin-3'-oxime (BIO), a known Wnt pathway activator inhibitor, was used as a positive control at 50 nM (Sigma, St. Louis, MO, USA). After 24 h of treatment with 1 µM of curcumin, tetrahydrocurcumin, and controls, cells were lysed with Trizol for RNA extraction. RNA was reverse transcribed using random primers (M-MLV Reverse Transcriptase kit; Promega, Madison, WI, USA) according to the manufacturer's instructions. Gene expression was then assayed by real-time qPCR using SYBR Green (Roche, Basel, Switzerland) on a Rotor-Gene Q cycler (Qiagen, Hilden, Germany) system. Beta-actin was used as a housekeeping gene (Forward-GGCTGTATTCCCCTCCATCG, Reverse-CCAGTTGGTAACAA TGCCTGT) and Axin2 as the read-out for Wnt pathway activity (Forward-TGACTCTCCTTCCAGATCCCA, Reverse-TGCCCAC ACTAGGCTGACA). Reactions were performed in triplicate and relative changes to housekeeping gene expression were calculated using the $2-\Delta\Delta CT$ method, where CT is the threshold cycle. Groups were then analyzed with one-way ANOVA followed up with multiple comparison tests in GraphPad Prism 8. The adjusted *p*-value is reported in graphs according to the *New England Journal* of *Medicine* guidelines: p < 0.001 (***), p < 0.002 (**), p < 0.033 (*), and p > 0.12 (ns).

Bulk sequencing

Dental pulp cells were seeded at a density of 0.05×10^6 in triplicate and treated with freshly made 1 µM curcumin and tetrahydrocurcumin for 24 h. RNA was isolated using Trizol and Qiagen Mini Kit (Qiagen, Germany). After assessment of quality and quantity, RNA was sent for bulk sequencing. Partek RNA sequencing pipeline was used. All algorithms used to analyze the data were rerun with default settings, unless otherwise indicated. The quality of the sequencing reads was examined using FastQC (v0.11.4) (https://www.bioinformatics. babraham.ac.uk/projects/fastqc/). Raw sequencing reads (100-nt, paired-end) were trimmed using Trim Galore! (v1.001.001) (https://www.bioinformatics.babraham.ac.uk/projects/trim_galore/). Traces of ribosomal RNA and mitochondrial RNA were removed using Bowtie2 (v2.2.5) (27). Reads were aligned to the human reference genome GRCh38 using STAR (v2.7.3a) aligner with multi-sample setting (28). Mapping and alignment quality were examined using FastQC and duplicate reads were removed using the Mark Duplicates function of the Picard tools (v2.17.11) (http://broadinstitute.github.io/picard/). Reads were annotated using the Partek E/M with Ensembl Transcripts release 104. Samples were then visualized and explored using unsupervised methods and clustered based on principal component analysis (PCA), UMAP, tSNE, and hierarchical clustering. Differentially expressed genes (DEGs) were analyzed between media controls versus curcumin or tetrahydrocurcumin using DESeq 2 (v3.5) (29). DEGs with FDR value ≤ 0.05 and fold change ≥ 1.5 were filtered out for further analysis. For gene ontologies, Enrichr biological function and bioplanet pathways were used. For functional enrichment analysis, G profiler was used with a significance threshold set at Bonferroni correction and the term size set at 50. For the analysis of shared upregulated genes in pulp, periodontal ligament (PDL), and gingival epithelial cells, we looked at the DEG from our previous study (30) and compared it to the DEG from the dental pulp cells.

Results

Curcumin and tetrahydrocurcumin promote Wnt signaling pathway in dental pulp cells

To evaluate the impact of curcumin and tetrahydrocurcumin on pulp, the effect of these compounds on dental pulp cell viability was assessed at concentrations of 100 μ M, 50 μ M, 10 μ M, 1 μ M and 0.5 μ M (Figure 1). At 1 and 0.5 μ M, both compounds resulted in more than 90% viability of pulp cells. Therefore, 1 μ M was selected as the optimum concentration for further downstream analysis. Wnt signaling pathway is crucial in the development and regeneration of many tissues in the oral cavity. *Axin2* is a downstream target of this pathway, and its expression is commonly used as a read-out of the level of Wnt signaling pathway activation (31, 32). Therefore, *Axin2* expression level analysis showed that 1 μ M of curcumin and THC can promote Wnt signaling pathway in dental pulp cells.

Curcumin and tetrahydrocurcumin promote matrix organization and multiple signaling pathways in dental pulp cells

After determining the suitable concentration of curcumin and tetrahydrocurcumin for the treatment of pulp cells, 1 μ M of these compounds was used to investigate the DEGs. Media and DMSO treatments were used as controls (see Supplementary Material). At an FDR value ≤ 0.05 and fold change ≥ 1.5 , treatment with curcumin as well as tetrahydrocurcumin resulted in more than 100 DEGs (Figure 2). Curcumin treatment resulted in the upregulation of cytoplasmic translation, gene expression, macromolecule biosynthesis, and oxidative phosphorylation. Treatment with tetrahydrocurcumin resulted in the upregulation of collagen fibril organization, regulation of apoptosis, platelet aggregation, and supramolecular fiber organization and upregulation of many signaling pathways, such as TGFB, Integrin, FAS, BDNF, and IL4 regulation of apoptosis.

Shared upregulated genes in pulp, PDL, and gingival cells after treatment with curcumin and tetrahydrocurcumin

We previously showed the impact of curcumin and tetrahydrocurcumin on PDL and gingival cells (30). Therefore, we asked if there are any shared upregulated genes between human dental pulp cells, PDL, and human gingival epithelial cells (HGEP) upon treatment with curcumin and tetrahydrocurcumin. To do this, all upregulated genes in these three cell types at an FDR value ≤ 0.05 and fold change ≥ 1.5 (Figure 3) were analyzed. Treatment with curcumin results in the highest number of upregulated genes in pulp cells. There were 53 shared upregulated genes between pulp and PDL. Gene ontology (GO) enrichment analysis of these shared genes using g: profiler demonstrates enrichment in response to toxic substance and oxidative stress, tissue development, and cell proliferation. UCHL1 and Tripartite Motif Containing 16 Like (Pseudogene) (TRIM16l) were upregulated in all cell types upon treatment with curcumin. Treatment with tetrahydrocurcumin results in the highest number of upregulated genes in pulp cells and 27 shared upregulated genes between pulp and PDL. GO enrichment analysis of these shared genes using g: profiler demonstrates enrichment in angiogenesis, coagulation, and muscle contraction. There were no shared upregulated genes between pulp, PDL, and gingival epithelial cells upon treatment with tetrahydrocurcumin.



Discussion

We previously demonstrated anti-inflammatory and regenerative potential of curcumin and tetrahydrocurcumin on human PDL and gingival epithelial cells (30). In this study, we investigated the regenerative potential of curcumin and tetrahydrocurcumin on dental pulp cells. Our results show that 1 μ M of curcumin and tetrahydrocurcumin promote Wnt signaling pathway in dental pulp cells without adverse effects on viability, similar to their effect on PDL and gingival epithelial cells. Wnt signaling pathway is associated with response to tissue damage (33–38), suggesting a regenerative potential of curcumin and its metabolite on dental pulp cells.

RNA sequencing demonstrates that treatment of dental pulp cells with curcumin results in upregulation of peptide biosynthetic process, macromolecule biosynthesis, mitochondrial ATP synthesis, cytoplasmic translation, and oxidative phosphorylation, suggesting a role in cellular metabolism and cytoprotection. Effect of curcumin on mitochondrial biogenesis has been shown previously. Recent studies have highlighted a multifaceted role for curcumin in cellular metabolism. This is through modulation of several metabolic pathways: scavenging reactive oxygen species (ROS), increasing antioxidant enzyme activity, regulating energy metabolism, and influencing the AMP-activated protein kinase (AMPK) pathway (39, 40). When compared to its impact on PDL and gingival epithelial cells, treatment with curcumin resulted in a higher number of upregulated genes in pulp cells. Shared upregulated genes between pulp and PDL were enriched in response to toxic substance and oxidative stress, tissue development, and cell proliferation. This suggests that curcumin has a role in cytoprotection and modulation of the cellular response to stress, such as inflammation and oxidative damage in oral tissues. Interestingly, TRIM16l and UCHL1 were upregulated in dental pulp, PDL, and gingival epithelial types upon treatment with curcumin. TRIM16 plays significant roles in cellular processes such as protein homeostasis, autophagy, and regulation of cellular responses to stress. This is particularly important for its involvement in the management of protein aggregates under oxidative or proteotoxic stress, especially in conditions like neurodegeneration and cancer (41, 42). UCHL1, a mitochondrial 10-formyltetrahydrofolate dehydrogenase, is an important mediator of many cellular functions and bone and lipid metabolism (43-47). It has recently been shown that UCLH1 can mediate MAPK signaling pathway in orthodontic tooth movement (48). Ubiquitination is a posttranslational modification that affects many cellular pathways, such



Differential gene expressions after treatment of pulp cells with curcumin and tetrahydrocurcumin. Volcano plots of differentially expressed genes after treatment of pulp cells with curcumin (A). Gene ontology (GO) of selected biological function (B) and pathways (C) in curcumin treatment at FDR value ≤ 0.05 and fold change ≥ 1.5 . Volcano plots of differentially expressed genes after treatment of pulp cells with curcumin (D). GO of selected biological function (E) and pathway (F) in treatment with tetrahydrocurcumin (F) at FDR value ≤ 0.05 and fold change ≥ 1.5 .

as immune response, angiogenesis, cell proliferation, apoptosis, and DNA repair. Deubiquitinating enzymes reversely modify proteins (49). Ubiquitination and deubiquitination play key roles in many diseases such as cancer (50). Interestingly, UCLH1 plays an important role in deubiquitination and TRIM16 can promote ubiquitination of proteins (47, 51). Upregulation of UCHL1 in pulp, PDL, and gingival epithelial cells upon curcumin treatment demonstrates that this natural compound has an important role in maintaining cellular proteostasis, particularly under stress conditions, making it a potential natural preventative and therapeutic candidate for oral diseases. Treatment of dental pulp cells with tetrahydrocurcumin resulted in collagen fibril organization, platelet aggregation, regulation of apoptotic signaling pathway, and intracellular sequestering of iron ion. Signaling pathways such as TGFB, Integrin, FAS, BDNF, and IL4 regulation of apoptosis are also upregulated upon treatment with tetrahydrocurcumin. Integrins are expressed in dental pulp and play a role in cell attachment on extracellular matrix proteins (52, 53). Integrin a5 promotes odontogenetic differentiation of dental pulp stem cells due to extracellular matrix deposition (54). Integrins also have a role in dental pulp stem cell senescence through



Shared and unique upregulated genes in pulp, periodontal ligament (PDL), and gingival epithelial cells. Venn diagram showing shared and unique upregulated genes after treatment of pulp, PDL, and human ginigval epithelial cells (HGEP) cells with curcumin (A) and tetrahydrocurcumin (B). Gene ontology (GO) enrichment analysis by g: profiler of the shared upregulated genes in pulp and PDL after treatment with curcumin demonstrates biological processes (C) and reactome (D). GO enrichment analysis by g: profiler of the shared upregulated genes in pulp and PDL cells after treatment with tetrahydrocurcumin demonstrates biological processes (E) and reactome (F). Graphical illustration guide (G).

mTOR signaling pathway (55). FAS pathway has been shown to modulate apoptosis and regulate immunomodulation in dental pulp stem cells (56, 57). BDNF has a role in inflammation mediate odontoblasts differentiation and can (58).Upregulation of these pathways in THC treatment suggests a role tissue remodeling and regeneration in for tetrahydrocurcumin in dental pulp cells. Furthermore, shared upregulated genes between pulp and PDL after tetrahydrocurcumin treatment were enriched in angiogenesis and coagulation, suggesting a regenerative potential of this natural compound on dental pulp cells.

Regenerative potential of curcumin has been shown in multiple independent studies (59-64). Curcumin plays an antiinflammatory role in dental pulp stem cells and can regulate proliferation and senescence of dental follicle cells. Curcuminloaded biomaterials have been developed to enhance tissue regeneration and reduce local inflammation (65-67). However, curcumin's poor bioavailability has led to innovative approaches to improve absorption and enhance its effectiveness in vivo (68). Tetrahydrocurcumin, with its improved bioactivity, has shown promise in preclinical models of ischemic injury, enhancing angiogenesis while reducing fibrosis and oxidative damage (4, 6). Similarly, anti-inflammatory properties of tetrahydrocurcumin have been shown in independent studies. This natural compound can reduce the release of TNF- α and IL-6 and influence other inflammatory pathways, such as NF-kB, JAK/STAT, and MAPK, which are activated in chronic diseases and metabolic disorders (4, 5). Tetrahydrocurcumin has been shown to be neuroprotective and pro angiogenesis and exert vascular protection in the brain (69-72). Interestingly, THC has been shown to suppress angiogenesis in murine adipose tissue and in human osteosarcoma (73, 74). This suggests that the impact of tetrahydrocurcumin on angiogenesis may be tissue dependent and can be harnessed to promote tissue regeneration or suppress tissue invasion and metastasis. Notably, THC has also been shown to be effective in the treatment of cancer (75). The results of this study show that curcumin and tetrahydrocurcumin promote Wnt signaling pathway in dental pulp, suggesting a role in tissue development and regeneration. In our study, curcumin exhibits cytoprotective properties and can regulate cellular metabolism and response to stress in dental pulp. The wider benefits of curcumin in cellular response to stress in oral tissue can be proposed by the upregulation of UCHL1 in dental pulp, PDL, and gingival epithelial cells, suggesting that curcumin can serve as a natural prophylactic and therapeutic candidate for pulp and PDL. Tetrahydrocurcumin affects different signaling pathways in dental pulp. It can promote regulation of the extracellular matrix, angiogenesis, and apoptotic signaling, suggesting a role in homeostasis and tissue regeneration. Enrichment in blood vessel development and morphogenesis in pulp and PDL after tetrahydrocurcumin treatment suggests this natural compound can serve as a candidate for oral wound healing.

Conclusion

The results of this study suggest that curcumin and tetrahydrocurcumin exhibit cytoprotective and regenerative potential on oral tissue. Investigation of differentially expressed genes provides a comprehensive analysis of affected biological processes and detection of any adverse effect these compounds may have. To overcome the limitations of bulk RNA sequencing and fully recognize the therapeutic potential of these compounds in treating oral diseases, further studies and *in vivo* validation are recommended.

Data availability statement

RNASeq datasets generated and/or analyzed during the current study are available in the NCBI repository with accession number PRJNA1192542.

Ethics statement

Original cell harvest experiments were undertaken with the understanding and written consent of each participant and in full accordance with the World Medical Association Declaration of Helsinki (version 2002). The study was approved and followed the guidelines set by the Ethical Committee for human studies at King's College Hospital, King's College University of London. The studies were conducted in accordance with the local legislation and institutional requirements. The human samples used in this study were acquired from primarily isolated as part of your previous study for which ethical approval was obtained. Written informed consent for participation was not required from the participants or the participants' legal guardians/next of kin in accordance with the national legislation and institutional requirements.

Author contributions

AB: Conceptualization, Data Curation, Formal analysis, Investigation, Methodology, Validation, Visualization, Writing – original draft, Writing – review & editing. PS: Funding Acquisition, Project Administration, Supervision, Writing – review & editing.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Generative AI statement

The author(s) declare that no Generative AI was used in the creation of this manuscript.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fdmed.2025. 1537478/full#supplementary-material

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