Ca\textsuperscript{2+} signalling plays a role in celastrol-mediated suppression of synovial fibroblasts of rheumatoid arthritis patients and experimental arthritis in rats

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Background and Purpose: Celastrol exhibits anti-arthritic effects in rheumatoid arthritis (RA), but the role of celastrol-mediated Ca\textsuperscript{2+} mobilization in treatment of RA remains undefined. Here, we describe a regulatory role for celastrol-induced Ca\textsuperscript{2+} signalling in synovial fibroblasts of RA patients and adjuvant-induced arthritis (AIA) in rats.

Experimental Approach: We used computational docking, Ca\textsuperscript{2+} dynamics and functional assays to study the sarcoplasmic/endoplasmic reticulum Ca\textsuperscript{2+} ATPase pump (SERCA). In rheumatoid arthritis synovial fibroblasts (RASFs)/rheumatoid arthritis fibroblast-like synoviocytes (RAFLS), mechanisms of Ca\textsuperscript{2+}-mediated autophagy were analysed by histological, immunohistochemical and flow cytometric techniques. Anti-arthritic effects of celastrol, autophagy induction, and growth rate of synovial fibroblasts in AIA rats were monitored by microCT and immunofluorescence staining. mRNA from joint tissues of AIA rats was isolated for transcriptional analysis of inflammatory genes, using siRNA methods to study calmodulin, calpains, and calcineurin.

Key Results: Celastrol inhibited SERCA to induce autophagy-dependent cytotoxicity in RASFs/RAFLS via Ca\textsuperscript{2+}/calmodulin-dependent kinase kinase-β–AMP-activated protein kinase–mTOR pathway and repressed arthritis symptoms in AIA rats. BAPTA/AM hampered the in vitro and in vivo effectiveness of celastrol. Inflammatory- and autoimmunity-associated genes down-regulated by celastrol in
Conclusion and Implications: Celastrol triggered Ca\textsuperscript{2+} signalling to induce autophagic cell death in RASFs/RAFLS and ameliorated arthritis in AIA rats mediated by calcium-dependent/-binding proteins facilitating the exploitation of anti-arthritic drugs based on manipulation of Ca\textsuperscript{2+} signalling.

What is already known
- Celastrol exhibits anti-arthritic effects by targeting Tregs, Th17 cells, and inhibiting NF-κB signalling.

What this study adds
- Celastrol inhibits SERCA to induce autophagy-dependent cytotoxicity in RASFs/RAFLS and represses arthritis in vivo.
- Block of calcium signalling attenuates the \textit{in vitro} and \textit{in vivo} effectiveness of celastrol.

What is the clinical significance
- Manipulation of Ca\textsuperscript{2+} signalling may represent an alternative approach to the treatment of RA.

1 | **INTRODUCTION**

Rheumatoid arthritis (RA) is a systemic autoimmune condition exhibiting polyarthritis and multiple organ disorders (Tsujimura, Saito, Nawata, Nakayamada, & Tanaka, 2008). A variety of cell populations such as lymphocytes, macrophages, and synovial fibroblasts (SLFs) are critically involved in the pathogenesis of RA. Rheumatoid arthritis fibroblast-like synoviocytes (RAFLS) and their primary form, rheumatoid arthritis synovial fibroblasts (RASFs) derived from patient SLFs, are considered key cellular participants in arthritic joints. RASFs in diseased articular tissues over-proliferate and they provide a barrier that hinders the delivery of anti-arthritic agents to the inflamed sites (Bartok & Firestein, 2010). The invasive capacity of RASFs also worsens disease progression by aggravating RA joint erosion via mechanical destruction of normal synovial tissues, neighbouring or distal to the inflamed site (Tolboom et al., 2005). These factors, together with the multidrug- and apoptosis-resistant properties of RASFs (Fang et al., 2013; Igarashi, Hirano, Yahagi, Saika, & Ishihara, 2014; Kim et al., 2011; Maillefert et al., 1996), lead to refractoriness in RA and support RASFs as a promising pharmacological target for the treatment of RA.

Ca\textsuperscript{2+} plays an important role in autoimmunity and inherited immunological dysregulation. In systemic lupus erythematosus, the pro-inflammatory cytokine expression of autoreactive T- and B-lymphocytes upon autoantigens stimulation is manipulated by Ca\textsuperscript{2+} signalling (Tsokos, 2011). During RA progression, autoreactive T-lymphocytes respond to autoantigens through a molecular mechanism similar to that in systemic lupus erythematosus, suggesting the involvement of Ca\textsuperscript{2+} signalling in RA (Sakaguchi et al., 2003). Our recent PCR screen for in vitro treatments in RASFs (Dias et al., 2018) suggested a possible link between Ca\textsuperscript{2+}-modulated gene regulation and different pathogenic factors of RA, in which calcium-binding proteins, such as calmodulin and calpains, are prerequisites for the effective downstream signalling of RA. This cellular Ca\textsuperscript{2+} signalling is closely related to the rigorous maintenance of Ca\textsuperscript{2+} homeostasis, exerted by various Ca\textsuperscript{2+} pumps (Berridge, 2012; Clapham, 2007). The sarcoplasmic/endoplasmic reticulum (SR/ER) Ca\textsuperscript{2+}-ATPase (SERCA) is the most extensively studied Ca\textsuperscript{2+} transporter located in the SR/ER membrane (Stammers et al., 2015). Recently, SERCA inhibition has been identified as a novel therapeutic strategy for tumourigenesis by triggering cytotoxicity in cancer cells (Denmeade & Isaacs, 2005; Michelangel & East, 2011). Suppression of SERCA can severely disrupt Ca\textsuperscript{2+} homeostasis to induce the ER stress response, thereby causing mitochondrial damage through Ca\textsuperscript{2+} overloading and caspase activation, leading to apoptosis and autophagic cell death (Wong et al., 2013). Moreover, SERCA inhibitors can induce autophagic cell death in both apoptosis- and multidrug-resistant cancer cells (Janssen et al., 2009). For instance, the SERCA inhibitor thapsigargin induces autophagy-related cell death in RASFs via ER stress activation (Kato, Ospelt, Gay, Gay, & Klein, 2014). By manipulating Ca\textsuperscript{2+} signalling to induce autophagic cell death, such findings have suggested the potential use of SERCA inhibitors in RASFs that are resistant to apoptosis.

\textit{Tripterygium wilfordii Hook F} (TwHF), a Chinese medicinal plant, is an effective medication for RA (Tao & Lipsky, 2000; Jiang et al., 2015; Ma, Zhou, Fan, & Sun, 2016). A recent clinical investigation revealed that TwHF-based remedies produced better therapeutic outcomes than disease-modifying antirheumatic drugs alone in controlling symptoms of active RA (Lv et al., 2015). Another randomized trial also demonstrated that TwHF is more effective than the first-line RA drug sulfasalazine (Goldbach-Mansky et al., 2009). Additionally, RA patients who failed to respond to conventional therapy were sensitive and showed good tolerance, to multiple therapeutic doses of TwHF (Tao, Younger, Fan, Wang, & Lipsky, 2002). Celastrol, a triterpene isolated from TwHF, suppressed autoimmune arthritis and reduce bone damage by modulating osteo-immune crosstalk (Nanjundaiah et al., 2012). Moreover, celastrol inhibited IL-17A- and LPS-stimulated
migration of RAFLS through the suppression of NF-κB-mediated MMP-9 expression (Li et al., 2012; Li et al., 2013). Celastrol also suppresses arthritis in adjuvant-induced arthritis (AIA) rats by altering the balance between pathogenic and regulatory T cells in the inflamed joints (Astry et al., 2015). Such beneficial effects have been related to the repression of Th17 cell polarization, associated with the IL-17 signalling pathway (Cascao et al., 2012; Han et al., 2015). Although celastrol exhibits anti-inflammatory effects by suppressing cell proliferation, invasion, and bone resorption in the treatment of RA (Cascao, Fonseca, & Moita, 2017), mechanistic studies regarding the proliferation, invasion, and bone resorption in the treatment of RA—celastrol exhibits anti-inflammatory effects by suppressing cell proliferation, invasion, and bone resorption in the treatment of RA (Cascao, Fonseca, & Moita, 2017), mechanistic studies regarding the Ca2+-mobilizing effect of celastrol on RA treatment are scarce. Given that celastrol is able to down-regulate a panel of inflammatory- and autoimmune-associate genes by mobilizing Ca2+ in RASFs (Dias et al., 2018), the current study aimed to examine whether celastrol is able to inhibit SERCA and activate Ca2+ signalling pathways to mitigate RA, in the well-established experimental AIA model in rats (Astry et al., 2015), which may provide a new strategy for treating RA via activation of Ca2+ signalling.

2 | METHODS

2.1 | Cell culture

All cells were obtained from the American Type Culture Collection (Rockville, MD, USA) unless otherwise stated. Immortalized wild-type and Bax-Bak double-knockout mouse embryonic fibroblasts (MEFs) were kindly provided by Professor Shigeomi Shimizu (Tokyo Medical and Dental University, Medical Research Institute, Japan). RASFs were freshly isolated from RA patients in Guangdong General Hospital, Guangdong Academy of Medical Sciences, Guangzhou (China), with research ethics committee approval number: GDREC 2015391H. RAFLS (MH7A) are the immortalized cell line purchased from ATCC. All media were supplemented with 10% FBS and the antibiotics penicillin (50 U·ml⁻¹) and streptomycin (50 μg·ml⁻¹; Invitrogen, Paisley, Scotland, UK). All cell cultures were incubated at 37°C in a 5% humidified CO₂ incubator.

2.2 | Isolation of RASFs from RA patients

This study involving human tissue samples was approved by the Research Ethics Committee (Guangdong General Hospital, Guangdong Academy of Medical Sciences) with approval number: GDREC 2015391H. RASFs were isolated from synovium from RA patients who had undergone knee surgery for synovectomy. The diagnosis of RA was made according to American Rheumatism Association revised criteria 1987 for classification of RA (Arnett et al., 1988). Synovial strips were cut into pieces and cultured in DMEM containing 20% FBS in 25-cm² culture flask. Medium was changed every 3 days, and the synovial tissues were collected from the cultured medium after 2 weeks. RASFs were trypsinized and diluted with DMEM containing 20% FBS before subculturing into other flasks. Cells were maintained at 37°C in a humidified incubator supplied with 5% of CO₂. The purification of RASFs was evaluated by staining for CD90. Cultured RASFs from passages 5–7 were employed for experimental studies.

2.3 | Endogenous autophagy detection

In brief, celastrol-treated RASFs or RAFLS cells grown on coverslips were fixed with 4% paraformaldehyde (Sigma) for 20 min and then rinsed with PBS. The cells were then permeabilized in methanol for 2 min and anti-LC3 was used in the analysis. The coverslips were then mounted with FluorSave™ mounting media (Calbiochem, San Diego, CA, USA). Fluorescence localization and quantitation of LC3 autophagosomes were visualized using the API Delta Vision Live-cell Imaging System (Applied Precision Inc., GE Healthcare Company, Washington, USA). The percentage of cells with punctuate immunofluorescence LC3 staining was calculated by the number of immunofluorescence-positive cells (≥10 dots per cell) over the total number of cells in the same field. A minimum of 1,000 cells from randomly selected fields were scored.

2.4 | Cytotoxicity assays

Cytotoxicity was assessed using the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT: 5.0 mg·ml⁻¹) assay. Briefly, 4 × 10⁵ BAX-Bak WT or DKO MEFs were seeded per well in 96-well plates. After overnight culture, the cells were then exposed to test compounds (0.039–100 μmol·L⁻¹) for 72 hr. Cells treated with DMSO were used as control. Subsequently, MTT (10 μL) was added to each well for 4 hr followed by the addition of 100-μL solubilization buffer (10% SDS in 0.01 mol·L⁻¹ HCI) and overnight incubation. A₅₇₀ nm was measured on the next day. The percentage of cell viability was calculated by the formula: Cell viability (%) = A₅₇₀ treated/A₅₇₀ control × 100. Data were obtained in triplicate from five independent experiments.

2.5 | Annexin V detection by flow cytometry analysis

Apoptosis was detected by Annexin V staining kit (BD Biosciences, San Jose, CA, USA). In brief, RASFs or RAFLS treated with the indicated doses of celastrol for 24 hr were harvested and detected by flow cytometry using FITC-Annexin V and Propidium Iodide staining, according to the manufacturer’s instructions. The population of apoptotic cells were quantitatively determined by flow cytometer (BD FACSaria III, San Jose, CA, USA). Data acquisition and analysis were conducted with CellQuest (BD Biosciences) in triplicate from five independent experiments.

2.6 | Measurement of cytoplasmic calcium dynamics

Intracellular cytosolic Ca²⁺ dynamics was determined using the FLIPR Calcium 6 Assay Kit (Molecular Devices) according to the manufacturer’s instructions. In brief, RASFs were plated in black wall/clear bottom 96-multiwell plates (Costar, Tewksbury, MA, USA) at a density...
of 10,000 cells per well and incubated overnight before treatment. Next day, the RASFs were treated with calcium 6 reagent for 2 hr at 37°C and 5% CO₂. Indicated concentrations of celastrol or thapsigargin were then added to the wells and immediately subjected to data acquisition on the SpectraMax Paradigm Multi-Mode Microplate Reader (Molecular Devices) at room temperature using a 1-s reading interval throughout five independent experiments.

2.7 | Measurement of intracellular free calcium

Intracellular free calcium level were determined by Fluo-3, AM dye. Briefly, RAFLS cells were washed twice with MEM medium after treatment with indicated concentrations of celastrol for 4 hr. Then, cell suspensions were incubated with 5-µM Fluo-3, AM at 37°C for 30 min. The cells were washed twice with HBSS and then subjected to FACS analysis. At least 10,000 events were analyzed. Data were obtained in triplicate from five independent experiments.

2.8 | Computational docking

The co-crystal structure of SERCA complexed with thapsigargin was retrieved from the Protein Data Bank (PDB). To prepare the protein for docking, Autodock Tools·1.5.6rc3 was used to remove crystallographic water molecules and to repair missing atoms. The output files after preparation was in PDBQT format. The ligands to be docked were drawn using ChemDraw software and converted into 3D structures using Open Babel. All 3D structures were saved in PDBQT format. The docking grid box was created and defined by the thapsigargin binding site on SERCA. Celastrol was docked on the selected sites of SERCA using Autodock 4. At each grid point, the energy of a particular ligand configuration was predicted. Docking parameters were set to 250 runs and 2,500,000 energy evaluations for each cycle using the Lamarckian Algorithm. The binding energies and the number of conformations in each cluster were attained from the docking log files (dlg).

2.9 | Measurement of SERCA activity

Ca^{2+} ATPase (SERCA1A) was extracted and purified from hind leg muscle of female rabbits. The effect of celastrol on Ca^{2+} ATPase activity was then determined using the enzyme-coupled method utilizing LDH and pyruvate kinase as previously described (Michelangeli, Colyer, East, & Lee, 1990). The SERCA inhibitory raw data were fitted to the allosteric concentration-effect equation using Figure P (Biosoft, Cambridge, UK):

\[
\text{Activity} = \frac{\text{minimum activity} + (\text{maximum activity-minimum activity})}{(1 + ([I]/IC_{50})^3)}.
\]

Data were obtained in triplicate from five independent experiments.

2.10 | Protein extraction and western blotting

The antibody-based procedures used in this study comply with the recommendations made by the British Journal of Pharmacology. After drug treatment, adherent and floating cells were lysed with RIPA. Protein concentrations were determined using the Bio-Rad protein assay (Bio-Rad Laboratories, Inc., Hercules, CA, USA). After electrophoretic separation, gels were blotted and stained with primary antibodies. The binding of the antibody was visualized by peroxidase-coupled secondary antibody using the ECL Western Blotting Detection Reagents (Invitrogen, Paisley, Scotland, UK). Band intensities were quantified by using the software ImageJ (NIH, Bethesda, MD, USA). Data were obtained from five independent experiments.

2.11 | LC-MS/MS measurement of ATP metabolites

Celastrol-treated RAFLS were harvested in 12 ml of ice-cold PBS. The cell pellet was then treated with 150 µl of 15% trichloroacetic acid (TCA) containing 7.5 µl of 20.0-µM [13C, 15N]ATP as internal standard and placed on ice for 10 min. After centrifugation at 12,100 x g for 15 min, the acidic supernatant was separated and neutralized twice with 80 µl mixture of trioctylamine and 1.12-trichlorotrifluoroethane (a volume ratio of 45 to 55), the samples were then ready for LC-MS/MS analysis. A Thermo Fisher TSQ LC-MS/MS system consisted of an Accela Autosampler, an Accela pump, and a Quantum Access triple quadrupole mass spectrometer. Data acquisition was performed using the Xcalibur software version 2.0.7, and data processing was carried out using the Thermo LCquan 2.5.6 data analysis program. The chromatographic separation was performed using X Terra-MS C18 column (150 mm x 2.1 mm i.d., 3.5 µm, Waters, Milford, MA). The two eluents were as follows: (A) 5-mM HA-0.5% DEA in water, pH 10 was adjusted with acetic acid; and (B) 50% acetonitrile in water. The mobile phase consisted of linear gradients of A and B: 0–15 min, 100–80% A (v/v); 15–35 min, 80–70% A; 35–45 min, 70–45% A; 45–46 min, 45–0% A; 46–50 min, 0–0% A; and 51–70 min, 100–100% A. The liquid flow rate was set at 0.3 ml-min^{-1}, and the column temperature was maintained at 35°C. Data were obtained from five independent experiments.

2.12 | Dual luciferase assay

Briefly, RAFLS were co-transfected with Renilla and one of the luciferase reporter plasmid of ATF6, CHOP, GRP-78, and GRP94 respectively. The next day, cells were treated with indicated concentrations of celastrol and thapsigargin for 24 hr. Cells then harvested with 100 µl of passive lysis buffer before luminescence measurements. According to the protocol of the dual luciferase reporter assay system (Promega Corp., E1910), 100 µl of the firefly luciferase reagent (LARII) was first added to each test sample, followed by addition of 100 µl of the Renilla luciferase reagent and firefly quenching (Stop & Glo). The data were represented as the ratio of firefly to Renilla luciferase activity (Fluc/Rluc) from five independent experiments.
2.13 AIA in rats and treatment

All animal care and experimental procedures were approved by the Animal Ethical Committee of Department of Health and Supervision, Macao Special Administrative Region of China, and carried out in accordance to the “Institutional Animal Care and User Committee guidelines” of the Macau University of Science and Technology. Animal studies are reported in compliance with the ARRIVE guidelines (Kilkenny, Browne, Cuthill, Emerson, & Altman, 2010; McGrath & Lilley, 2015) and with the recommendations made by the British Journal of Pharmacology. Drug preparation, treatment, and animal data collection were conducted blindly and independently by three investigators.

The AIA model was established using 6-week-old male Sprague Dawley rats (RRID:MGI:5651135) weighing 130 ± 20 g (four per cage) which were purchased from Guangdong Medical Lab Animal Center. The animals were housed in a temperature-controlled room with 12-hr light/dark cycles and given ad libitum access to food and water.

Arthritis was induced in rats by inoculation with complete Freund’s adjuvant. Non-viable desiccated Mycobacterium tuberculosis (BD, USA) was emulsified in mineral oil (Sigma, USA) to yield 5 mg·ml⁻¹ of M. tuberculosis. The rats were injected intradermally at the base of the tail with 100-μl of this emulsion. The first signs of inflammation were observed on Day 9 after adjuvant injection. Arthritic scores were evaluated and recorded every 3 days. Each paw was evaluated and scored individually on a scale from 0 to 4. The scoring criteria was as follows: 0, no evidence of erythema and swelling; 1, erythema and mild swelling confined to the tarsals or ankle joint; 2, erythema and mild swelling extending from the ankle to the tarsals; 3, erythema and moderate swelling extending from the ankle to the metatarsal joints; and 4, erythema and severe swelling encompass the ankle, foot and digits, or ankyllosis of the limb.

Forty-eight male rats were randomly divided into five experimental groups as follows: (a) healthy control group (n = 8) without treatments; (b) vehicle-treated control group (n = 8), AIA rats receiving the same vehicle and administration route of the drug in each experiment; (c) positive control group (n = 8), AIA rats receiving (by gavage) MTX (7.6 mg·kg⁻¹·week⁻¹ in a volume of 10 ml·kg⁻¹ body weight). (d) AIA rats receiving 1 mg·kg⁻¹ of celastrol (Cel; n = 8); (e) AIA rats receiving 1 mg·kg⁻¹ of celastrol + 3.5 mg·kg⁻¹ BAPTA/AM (BM; n = 8). AIA rats were treated i.p. with celastrol daily in the first 19 days and then treated every 2 days until Day 36, whereas BM was administered daily. Celastrol (China cdmust, A0106) and BAPTA/AM (Santa Cruz, USA) were dissolved in the vehicle (PEG400: H₂O: Ethanol = 6:3:1) and administered i.p. in a volume of 1 ml·kg⁻¹ body weight. MTX was dissolved in a vehicle of propylene glycol: Tween-80: normal saline, 50:5:45.

At the end of the treatment period, the rats were killed (cervical dislocation during pentobarbital anaesthesia) and the left hind paw was amputated and fixed in 4% PFA, then scanned using in vivo microCT scanner (SkyScan 1176, Bruker, Belgium). The following scanning parameters were used to obtain high-quality images of the joint: 35-μm resolution, 85 kV, 385 μA, 65-ms exposure time, 0.7° rotation step in 360°, and a 1-mm Al filter. The images were reconstructed using NRecon software (Bruker-micro CT, Belgium).

MicroCT score was obtained from five disease-related indices of the microCT analysis for calcaneus (bone mineral density, bone volume fraction, cortical mineral density, trabecular number, and total porosity). MicroCT score was calculated using the formula as follows: (Acquired value – minimum value)/(maximum value – minimum value) or 1 – (Acquired value – minimum value)/(maximum value – minimum value). The final microCT score is equally averaged from these five indices of disease.

2.14 Immunofluorescent staining of rat synovial tissues

Synovial tissues from all treatment groups were fixed and embedded in paraffin for microtome sectioning and immunofluorescence staining. After deparaffinization, the tissue sections (5μm thick) were subjected to antigen retrieval (EnVision™ FLEX Target Retrieval Solution, High pH (50×)). Anti-LC3B or anti-vimentin were used in the analysis. The coverslips were then mounted with FluorSave™ mounting media (Calbiochem, San Diego, CA, USA) for fluorescence imaging. The expression of LC3 and vimentin was captured by API DeltaVision Live-cell Imaging System (Applied Precision Inc., GE Healthcare Co., Washington, USA).

2.15 siRNA transfection, RNA extraction, and cDNA synthesis

RASFs were transfected with a mixture of siRNAs using Lipofectamine® 3000 (Invitrogen) according to the manufacturer’s protocol. To maximize the knockdown efficiency for the calcium binding protein (calmodulin, calpains, and calcineurin), the siRNA of these gene isoforms were mixed together before the transfection: siRNA CALM1 + CALM2 + CALM3 for calmodulin, siRNA CAPN1 + CAPN2 for calpains, and siRNA PPP3CA + PPP3CB + PPP3CC for calcineurin. All siRNA were purchased from Qiagen (Cambridge, USA).

RNA was extracted from the (AIA) animal joint tissues or immortalized RASFs using FavorPrep™ Blood/Cultured Cell Total RNA Purification Mini Kit (Favorgen Biotech Corp.). RNA concentration was determined using the NanoDrop 2000c Spectrophotometer (Thermo Scientific) and 1 μg of total RNA was used to synthesize the corresponding cDNA using the Transcriptor Universal cDNA Master mix (Roche, USA).

2.16 Real-time quantitative PCR

Quantitative PCR was performed using cDNA prepared from 1 μg of RNA of treated immortalized RASFs, with PowerUp™ SYBR® Green Master Mix (Applied Biosystems) in the Viia™ 7 Real Time PCR System (Applied Biosystems), together with primers synthesized (Tech Dragon Ltd., Hong Kong) using our own designed templates. Primer
sequences were designed employing ThermoFisher Scientific's online OligoPerfect™ Designer software and then verified with NCBI's Primer-BLAST software to confirm specific recognition of the target genes. Gene expression levels were normalized to GAPDH, (control), and analysed using the $2^{-\Delta\Delta CT}$ method. Five independent experiments with six replicates per group were analysed for each primer. Primer sequences are designed as below:

**ADGRE5**
- Forward: 5′-CTTTCCCGATTCTTGCAGCA-3′
- Reverse: 5′-TCCATCAGTTCTCATCACCA-3′

**BMP1**
- Forward: 5′-CAACACGGTCGAGTATG-3′
- Reverse: 5′-CAGTGTGGATTCATCCACA-3′

**CALM1**
- Forward: 5′-TCAGCTGACCAGAAGAACAGA-3′
- Reverse: 5′-TCAGCTGACCAGAAGAACAGA-3′

**CALM2**
- Forward: 5′-ATGCTGATGGTAATGGCACA-3′
- Reverse: 5′-AAAGCCATCAGACAACAGA-3′

**CALM3**
- Forward: 5′-ATTGCTAGATGATGTAATG-3′
- Reverse: 5′-TTCTGGCCCTTCTCAGGATT-3′

**CAPN1**
- Forward: 5′-AACCAGCTGAGAAGAACAGA-3′
- Reverse: 5′-AAAAGCACGAGACCAACTC-3′

**CAPN2**
- Forward: 5′-AGGGGCTTCTTCTCAACTC-3′
- Reverse: 5′-AAGCTCAGATAGCTCAGC-3′

**CAST**
- Forward: 5′-GAAGTCGATGAGGCAAAAGC-3′
- Reverse: 5′-ATCACCCAAGCCAATGAGTC-3′

**CD40**
- Forward: 5′-GAGATCAATTTCCTGCCAGCA-3′
- Reverse: 5′-GAACTGCTTCTCCAACGTTC-3′

**CMTM1**
- Forward: 5′-ATCACCCAAGCCAATGAGTC-3′
- Reverse: 5′-GCAAGTTGGAAGGGTAC-3′

**ERBB2**
- Forward: 5′-AACCAGCTGCTCACCAC-3′
- Reverse: 5′-CAGAACTCCTCCACCAGCAG-3′

**FGF10**
- Forward: 5′-GCATGTGCGGAGCTACAATC-3′
- Reverse: 5′-GCTGACCTTCCGTTCTTCTC-3′

**GLMN**
- Forward: 5′-GGGACAGACAGACGACGACT-3′
- Reverse: 5′-AAGGCACTCAGACAACCAGA-3′

**HRH1**
- Forward: 5′-ACCCCAAGAGGATGAAAT-3′
- Reverse: 5′-GCCTGACATGTGCAACAATATC-3′

**IFNAR1**
- Forward: 5′-AAGCTCAATTGGTCTCCTCC-3′
- Reverse: 5′-AAGCTCAGATAGCTCAGC-3′

**IK**
- Forward: 5′-TTCTGGCCCTTCTCAGGATT-3′
- Reverse: 5′-CCAAGGACCTGCTACAAA-3′

**IL1R1**
- Forward: 5′-ATGATGTTCGTCCTTCTGC-3′
- Reverse: 5′-TGAATCTCTGGAGGCTTGTTC-3′

**IL4R**
- Forward: 5′-TTCTGGCCCTTCTCAGGATT-3′
- Reverse: 5′-CCAAGGACCTGCTACAAA-3′

**LEPR**
- Forward: 5′-AATGACTTTCACAGCCAATC-3′
- Reverse: 5′-GGTCTTCCACACAGAATC-3′

**NFATC3**
- Forward: 5′-TTCTGGCCCTTCTCAGGATT-3′
- Reverse: 5′-CCAAGGACCTGCTACAAA-3′

**NFKB1**
- Forward: 5′-GCCCTGATATGGCCACCA-3′
- Reverse: 5′-GCCCTGATATGGCCACCA-3′

**NFRKB**
- Forward: 5′-TCGAGCTGTTCCTTCCAGT-3′
- Reverse: 5′-GGAAGAGGGCAGCTAATTCC-3′

**NFX1**
- Forward: 5′-CCAAGTTTATGGCGCAGTTT-3′
- Reverse: 5′-TGGAAGGGCAGCTAATTCC-3′

**PPP3CA**
- Forward: 5′-GGGACTTCCAAAGACGGCTTC-3′
- Reverse: 5′-GCCCTGATATGGCCACCA-3′
2.17 | Data and statistical analysis

The data and statistical analysis comply with the recommendations of the British Journal of Pharmacology on experimental design and analysis in pharmacology (Curtis et al., 2018). Data are expressed as means ± SEM. All the histological samples were scored blindly and independently by two investigators. Each experiment was conducted independently at least five times. Statistical analysis was carried out with GraphPad Prism 7 (RRID:SCR_002798). Data were analysed with one-way ANOVA. Dunnett’s or Tukey’s post hoc test was carried out only if P < .05 was accepted. Values of P < .05 were taken to indicate statistically significant differences between means.

2.18 | Materials

All chemicals and reagents were purchased from Sigma, unless otherwise stated. The following reagents from other suppliers were used: thapsigargin (Calbiochem, 586005, USA), compound C (Calbiochem, 171260, USA), BAPTA/AM (Santa Cruz, USA), baflomycin A1 (Calbiochem, 196000, USA), STO-609 (Calbiochem, 570250, USA), celastrol (cdmust, A0106, China), methotrexate (MTX; LC labs, MA, USA), RIPA (CST, 9806, USA), Flu-3, AM (Life Technologies, F14218, USA), EnVision™ FLEX Target Retrieval Solution, High pH (50x; DAKO, DM828, USA), antibodies light-chain 3B (LC3B) rabbit mAb (Cell signaling, 2775/3868, USA, RRID:AB_915950), phospho-p70S6 kinase (Thr^389) rabbit mAb (CST, 9205, USA), p70S6 kinase rabbit mAb (CST, 9202, USA), phospho-AMP-activated protein kinase (AMPK; Thr^172) rabbit mAb (CST, 2531, USA), AMPK rabbit mAb (CST, 2532, USA), vimentin mouse mAb (Abcam, 8978, USA), anti-β-of actin mouse monoclonal IgG1 (Santa Cruz, sc-47778, USA), rabbit anti-mouse IgG (H + L) secondary antibody TRITC (Invitrogen, PA1-28565, USA), and GOXMO TRITC HIGH XADS (Invitrogen, A16083, USA).

2.19 | Nomenclature of targets and ligands

Key protein targets and ligands in this article are hyperlinked to corresponding entries in http://www.guidetopharmacology.org, the common portal for data from the IUPHAR/BPS Guide to PHARMACOLOGY (Harding et al., 2018), and are permanently archived in the S186 of Concise Guide to PHARMACOLOGY 2017/18 (Alexander, Christopoulos et al., 2017; Alexander, Fabbro et al., 2017a,b; Alexander, Kelly et al., 2017; Alexander, Striessnig et al., 2017).

3 | RESULTS

3.1 | Celastrol targets SERCA for Ca^2+ mobilization in RASFs/RAFLS

Flow cytometry analysis of fluo-3 AM dye-stained RAFLS showed that celastrol released Ca^{2+} in a time- and concentration-dependent manner (Figure 1a). Celastrol is a triterpenoid compound and another triterpenoid, saikosaponin-d, caused the accumulation of cytosolic Ca^{2+} via inhibition of SERCA (Wong et al., 2013). Thus celastrol may also target SERCA for Ca^{2+} release and we therefore studied the changes in calcium dynamics, induced by celastrol in the presence of a known SERCA inhibitor, thapsigargin. As shown in Figure 1b, FLIPR calcium 6 assay demonstrated that either thapsigargin or celastrol alone changed the calcium dynamics in RASFs, but celastrol was much less potent than thapsigargin. Pre-incubation of RASFs with thapsigargin abolished the effects of celastrol on Ca^{2+} mobilization (Figure 1b), suggesting that celastrol targeted the SERCA-dependent Ca^{2+} stores which had been depleted by thapsigargin treatment, as a negative control, pretreatment with a different kinase inhibitor, compound C, which inhibits AMPK, with no reported interaction with SERCA, did not affect the changes in calcium dynamics induced by celastrol.

Computational docking results further revealed that celastrol and thapsigargin preferentially bound to the same pharmacophore on SERCA. The lowest binding energies for thapsigargin and celastrol were -12.37 and -10.27 kcal·mol^{-1} respectively (Figure 1c). Using purified sarcoplasmic reticulum (SR) membranes from rabbit skeletal muscle, that expressed the SERCA1A isoform (Wong et al., 2013),...
Celastrol inhibited SERCA1A in a concentration-dependent manner (Figure 1d), and the data fit to an allosteric concentration-effect equation. RT-PCR analysis further identified the SERCA 2 isoform as the most abundant type of SERCA in RASFs (Figure 1e), suggesting that celastrol was also effective in the inhibition of the SERCA2 isoform for calcium mobilization, as indicated in Figure 1b.
3.2 | Celastrol induces autophagic flux to activate autophagic cell death in RASFs and RAFLS

Mobilization of cytosolic Ca\(^{2+}\) induces autophagy and autophagic cell death in cancer cells (Wong et al., 2013). We therefore tested the effects of celastrol on autophagy and cell death in RASFs/RAFLS and found that autophagy and cell death in RASFs/RAFLS was induced by celastrol, in a concentration- and time-dependent manner (Figures 2a and S1A and B). The celastrol-induced endogenous autophagic LC3-II puncta formation in RAFLS (Figure 2b) and LC3-II protein conversion in RASFs (Figure 2c) were similarly blocked by the PI3K inhibitor, wortmannin, which is an autophagy inhibitor (Klionsky et al., 2016). Of note, celastrol significantly increased the rate of LC3-II conversion in the presence of the lysosomal protease inhibitor bafilomycin A1, compared with the administration of the inhibitors alone (Figure 2d). These results are therefore consistent with the premise that celastrol-increased Ca\(^{2+}\) mobilization induces autophagy and the resultant cell death of SLFs through the enhancement of autophagic flux.

3.3 | Celastrol activates CaMKKβ-AMPK-mTOR signalling pathway to induce Ca\(^{2+}\)-dependent autophagy in RASFs/RAFLS

We have previously shown that autophagy can be activated by calcium mobilization via the Ca\(^{2+}\)/calmodulin-dependent kinase kinase-β (CaMKKβ)-AMPK-mTOR pathway (Wong et al., 2013). Here, we showed that celastrol activated AMPK phosphorylation and reduced the phosphorylated form of p70S6K, a downstream target of mTOR (Figure 3a). In addition, the AMPK inhibitor compound C (non-toxic dose: 5 μM) significantly inhibited celastrol-induced endogenous LC3-II puncta formation and LC3-II protein conversion in both RAFLS and RASFs (Figure 3b,c). As autophagy can be activated by an upstream kinase of AMPK, the CaMKKβ (Wong et al., 2013), treatment with the CaMKKβ inhibitor, STO-609 (non-toxic dose: 25 μM), abolished celastrol-induced endogenous LC3-II puncta formation and LC3-II protein conversion in RAFLS and RASFs (Figure 3d,e), suggesting that celastrol increased the level of cytosolic Ca\(^{2+}\) which induced autophagy induction through the CaMKKβ-AMPK-mTOR pathway.

3.4 | Calcium chelator BAPTA/AM abolishes celastrol-induced autophagic cell death in RASFs/RAFLS and apoptosis-resistant fibroblasts

The synovial environment in RA patients favours the survival of FLS and prevents their removal by apoptosis and this resistance of FLS to apoptosis has been considered as a major therapeutic challenge (Kim et al., 2011). Thus, induction of autophagic cell death might serve as an alternative way of eliminating apoptosis-resistant RASFs/RAFLS in RA (Bartok & Firestein, 2010; Turner & Filer, 2015). Hence, we further validated whether block of Ca\(^{2+}\) release affects celastrol-induced autophagic cell death in RASFs/RAFLS. Of note, the calcium chelator BAPTA/AM suppressed celastrol-induced endogenous autophagic LC3-II puncta formation in RAFLS (Figure 4a) and inhibited LC3-II protein conversion in RASFs (Figure 4b). Concomitantly, annexin V-Pi staining showed that both BAPTA/AM and the autophagy inhibitor, wortmannin, reduced celastrol-mediated autophagic cell death (Figure 4c). These findings suggest that mobilization of Ca\(^{2+}\) is necessary for celastrol to induce autophagy and autophagic cell death in SLFs. Furthermore, we mimicked the apoptosis-resistant phenotype of RASFs using apoptosis-resistant fibroblasts, for example, Bax-Bak double-knockout (DKO) MEFs. Interestingly, while the resistant factor of DKO fibroblasts was 1.77 to >14 times more resistant to cell death by the anti-inflammatory agents, leflunomide, tacrolimus, baricitinib, sulfasalazine, and hydroxychloroquine, colchicine, celastrol, and MTX showed little difference in cell death sensitivity in both Bax-Bak wild-type and DKO fibroblasts (Figure 5a). As expected, celastrol was able to induce autophagic puncta formation and autophagic cell death in these apoptosis-resistant fibroblasts, whereas co-administration of BAPTA/AM substantially blocked the celastrol-induced autophagic effects and cell death (Figure 5b,c). All these findings supported the view that celastrol eliminated SLFs and apoptosis-resistant fibroblasts through calcium-induced autophagic cell death.

3.5 | Celastrol-mediated Ca\(^{2+}\) release induces energy loss and ER stress activation in RAFLS

Abnormal cytosolic Ca\(^{2+}\) levels can disrupt calcium homeostasis and activate the ER stress response in cells, which further induces the release and overload of Ca\(^{2+}\) in mitochondria, leading to cytochrome c release and mitochondrial membrane potential loss (Denmeade & Isaacs, 2005). By LC-MS/MS analysis, we found that the amounts of ATP, ADP, and AMP metabolites in the mitochondria was decreased in the celastrol-treated RAFLS (Figure 5d), suggesting mitochondrial damage in RAFLS which can be caused by the perturbation of calcium homeostasis (Todd, Lee, & Glimcher, 2008). As disruption of calcium homeostasis can also lead to activation of the ER stress response as well as apoptosis and autophagic cell death (Todd, Lee, & Glimcher, 2008), we therefore investigated whether celastrol could activate the ER stress response using the dual-luciferase reporter assays. Both celastrol and thapsigargin, a positive regulator for the elevation of cytosolic Ca\(^{2+}\), markedly increased the expression of ER stress markers ATF6, CHOP, GRP78, and GRP94 in RAFLS (Figure 5e), indicating that celastrol may similarly induce apoptosis and autophagic cell death via Ca\(^{2+}\) release and ER stress activation in RAFLS.

3.6 | Celastrol suppresses AIA in rats via mobilization of Ca\(^{2+}\)

Although celastrol has exhibited anti-arthritic effect in vivo (Li et al., 2013; Venkatesha, Yu, Rajaiah, Tong, & Moudgil, 2011) through inhibition of cytokines, chemokines, and inflammatory mediators (Cascao, Fonseca, & Moita, 2017), the role of celastrol-mediated Ca\(^{2+}\) in RA therapy remains unclear. Therefore, the possible anti-arthritic effect
FIGURE 2 Effect of celastrol-induced autophagy in RA synovial fibroblasts. (a) The correlation between celastrol-mediated autophagy and cytotoxicity. The concentration-dependent cytotoxicity of celastrol (Cel) was measured by annexin V-propidium iodide flow cytometry in parallel with dose-dependent autophagic puncta formation in RAFLS treated with DMSO or 0- to 2-μM celastrol for 24 hr. Y-axis: PI; X-axis: annexin-V FITC. (b) Effect of wortmannin (Wort) in celastrol-induced autophagy in RAFLS. RAFLS were treated with DMSO or 2-μM celastrol for 24 hr in the presence or absence of 1-μM wortmannin. The cells were counted only with the increased puncta pattern of LC3 fluorescence (≥10 dots/cell) in immunofluorescence-positive cells over the total number of cells in the same field. Fluorescence images were captured at 60× magnification; scale bar, 15 μm. Bar charts demonstrate the quantitation of the increase in autophagic cells. (c) Effect of Wort in celastrol-mediated LC3-II conversion. RASFs were treated with DMSO or 2-μM celastrol for 24 hr in the presence or absence of 1-μM wortmannin. (d) Effect of celastrol in autophagic flux. RASFs were treated with DMSO or 2-μM celastrol in the presence or absence of lysosomal protease inhibitors, Bafilomycin A1 (BafA1) 50 nM for the indicated times. Western blotting with antibodies against LC3 conversion (LC3-I, 18 kDa; LC3-II, 16 kDa) and β-actin. The data are expressed as fold matched control values relative to the DMSO-treated control. The data shown are the mean ± SEM of five independent experiments. *P < .05, significantly different from DMSO-treated control or celastrol-treated group.
FIGURE 3  Effect of celastrol in CaMKKβ–AMPK–mTOR signalling cascade. (a) Activation of the AMPK–mTOR signalling pathways by celastrol (Cel). RASFs were treated with DMSO or 2 μM celastrol for the indicated times, and rapamycin (Rap, 400 nM) was used as a positive control. Western blot analysis with antibodies against p-AMPK, total AMPK, p-p70S6K, total p70S6K, and β-actin. (b) Effect of compound C (CC) on celastrol-induced autophagic puncta formation in RAFLS. RAFLS were treated with DMSO or 1 μM celastrol for 24 hr with or without 5 μM compound C before fluorescence microscopic analysis. Bar chart represents the quantitation of autophagic cells with an increased percentage of LC3 puncta. (c) Effect of compound C (CC) on celastrol-mediated LC3-II conversion. RASFs were treated with DMSO or 2 μM celastrol for 24 hr in the presence or absence of 5 μM compound C. (d) Effect of CaMKKβ inhibitor in celastrol-induced autophagy in RAFLS. RAFLS were treated with DMSO or 1 μM celastrol for 24 hr with or without 25 μM STO-609 (CaMKKβ inhibitor) before fluorescence microscopy scoring. Bar chart represents the quantitation of autophagic cells with increased percentage of LC3 puncta. (e) Effect of CaMKKβ inhibitor, STO-609 in celastrol-mediated LC3-II conversion. RASFs were treated with DMSO or 2 μM celastrol for 24 hr in the presence or absence of 25 μM STO-609. Western blotting with antibodies against LC3 conversion (LC3-I, 18 kDa; LC3-II, 16 kDa) and β-actin. The data are expressed as fold matched control values relative to the DMSO-treated control. The data are expressed as the mean values ± SEM of five independent experiments. *P < .05, significantly different from DMSO-treated control group.
FIGURE 4  Effect of Ca2+ chelator BAPTA/AM in celastrol-mediated autophagy and cell death in RAFLS. (a) Autophagic effect of celastrol (Cel) in RAFLS in the presence of Ca2+ chelator BAPTA/AM. RAFLS were treated with DMSO or 1-μM celastrol for 24 hr in the presence or absence of 10-μM BAPTA/AM (BM). Bar charts representing the quantitation of cells with increased autophagic puncta are shown. Fluorescence images were captured at 60× magnification; scale bar, 15 μm. (b) Effect of BAPTA/AM in celastrol-mediated LC3-II conversion in RASFS. Western blotting with antibodies against LC3 conversion (LC3-I, 18 kDa; LC3-II, 16 kDa) and β-actin. Bar chart represents the quantitation of LC3-II conversion. (c) Recovery effect of BAPTA/AM and Wort in celastrol-mediated autophagic cell death in RAFLS. RAFLS treated with celastrol for 24 hr were measured using flow cytometry after annexin V-PI staining. Y-axis: PI; X-axis: annexin-V FITC. The data are expressed as the mean values ± SEM of five independent experiments. *P < .05, significantly different from DMSO-treated control or celastrol-treated group.
FIGURE 5 Activation of ER stress and Ca²⁺ signalling in celastrol-mediated autophagic cell death in the apoptosis-resistant fibroblasts. (a) Comparison of drug-resistance effects on various anti-inflammatory agents caused by celastrol in apoptosis-defective Bax-Bak DKO fibroblasts. MEFs were incubated with compounds for 72 hr, and the MTT cytotoxicity assay was performed to determine their cytotoxicity. The IC₅₀ values shown on the chart are the means ±SEM from five independent experiments. (b) Effect of BAPTA/AM (BM) in celastrol (Cel)‐induced autophagy in Bax-Bak DKO apoptosis-defective fibroblasts. Bar charts show the quantitation of the increase in autophagic cells. Fluorescence images were captured at 60× magnification; scale bar, 15 μm. (c) Effect of BAPTA/AM in celastrol‐induced autophagic cell death in the apoptosis-defective fibroblasts. The percentage of cell death in celastrol-treated Bax-Bak DKO MEFs with or without BAPTA/AM was measured using flow cytometry after annexin V staining. (d) Effect of celastrol in mitochondrial ATP generation in RAFLS. RAFLS treated with DMSO or 1-μM celastrol for 24 hr were subjected to mitochondrial extraction. Mitochondrial ATP, ADP, and AMP were measured by HPLC-MS (N = 6). (e) Activation of the ATF6, CHOP, GRP78, and GRP94 signalling in celastrol-treated RAFLS. ATF6-, CHOP-, GRP78-, and GRP94-containing dual-luciferase reporter plasmid-transfected RAFLS were treated with DMSO or 1-μM celastrol for 24 hr, and thapsigargin (TG, 1 μM) was used as a positive control. Cell lysates were harvested for dual-luciferase reporter assay. The data are expressed as the mean values ± SEM of five independent experiments. *P < .05, significantly different from DMSO-treated control or celastrol-treated group.
of celastrol via Ca\(^{2+}\) mobilization was examined in the AIA rat model. As shown in Figure 6a,b, celastrol significantly reduced the arthritic score and hind paw volume without affecting other organs or body weight in comparison with vehicle-treated AIA rats (Figure S2). Interestingly, addition of the calcium chelator BAPTA/AM partly decreased the anti-arthritic effect of celastrol (Figure 6c). MicroCT analysis was used to evaluate the severe swollen joints and bone destruction in the vehicle-treated AIA rats by direct comparisons of bone mineral density, bone volume fraction, cortical mineral density, trabecular number, and total porosity to other treatment groups. As shown in Figures 6d and S2C and D, the overall microCT and radiological scores indicated that inflammation and bone destruction were significantly better with the treatment of celastrol or MTX. MicroCT analysis of bone volume fraction, trabecular number, and total porosity indicated that the AIA rats treated with the combination of celastrol and BAPTA/AM displayed severe bone destruction compared to the rats treated with celastrol alone. As shown in Figures 6c–d, S2B–D, and Video S1, the mean microCT score dropped significantly from 0.54 to 0.33, while the mean radiological score rose from 1.33 to 1.83. The induction of autophagy in the synovial joints of the rat model after celastrol treatment was then examined by fluorescent immunohistochemical analysis. The antibodies LC3B (red) was used to detect the formation of autophagosomes, while vimentin (green) was used as a marker to indicate the proliferation and epithelial–mesenchymal transition of fibroblasts which are critical to the pathogenesis of RA. As shown in Figure 6e, the administration of celastrol (1 mg·kg\(^{-1}\)) alone clearly triggered the induction of autophagy when compared with untreated and healthy controls. In addition, the expression of vimentin was significantly suppressed in celastrol-treated rats compared with AIA rats. The suppression of vimentin was greater after celastrol than that in AIA rats treated with 7.6 mg·kg\(^{-1}\) of MTX, suggesting that the therapeutic efficacy of celastrol had been achieved at 1 mg·kg\(^{-1}\). The celastrol-induced effects on autophagy activation and vimentin suppression were mitigated by co-treatment of celastrol and BAPTA/AM, which decreases intracellular Ca\(^{2+}\) levels. Accordingly, the anti-arthritic effect of celastrol was partly attributable to the elevated Ca\(^{2+}\) and its downstream activation of the autophagic pathway.

3.7 The calcium-dependent/-binding proteins calmodulin, calpains, and calcineurin play critical roles in celastrol-mediated anti-arthritis effect in AIA rats

Celastrol mobilized cytosolic Ca\(^{2+}\) in patient-derived RASFs and regulated 23 out of 370 genes associated with inflammation and autoimmunity (Dias et al., 2018). Accordingly, expression of these Ca\(^{2+}\)-induced genes was validated using the joint tissues from our AIA model. Real-time PCR analysis indicated that except for CD40, GLMN, and TRAP1, the expression of the other 20 genes was almost significantly down-regulated in our AIA model treated with celastrol, whereas co-treatment with the Ca\(^{2+}\) chelator, BAPTA/AM abolished the down-regulation of genes mediated by celastrol (Figure 7). These results suggest the possible therapeutic role of Ca\(^{2+}\) signalling in RA condition. Ca\(^{2+}\) is a second messenger that is required for the signal transduction of many cellular processes (Clapham, 2007), and calmodulin, calcineurin, and calpains are well-known downstream modulators of Ca\(^{2+}\)-dependent enzymes, such as protein kinases, phosphatases, and proteases (Villalobo A, 2018). Therefore, the role of celastrol-mediated Ca\(^{2+}\) signalling in the regulation of inflammatory and autoimmunity-associated gene expression was verified by knockdown of these three Ca\(^{2+}\)-binding/-dependent proteins. To ensure high efficiency in transcription and knockdown of genes, we used the RAFLS cells for validation of these 23 genes of interest by specific siRNA knockdown of calmodulin, calcineurin, and calpains individually, prior to celastrol treatment. In Figure S3, effective knockdown of the target genes was achieved by transfecting the cocktail of siRNAs specific to different gene isoforms of calmodulin (CALM1 + CALM2 + CALM3), calpains (CAPN1 + CAPN2), and calcineurin (PPP3CA + PPP3CB + PPP3CC). Under the same gene knockdown condition, 21 out of 23 genes (fold change values <0.75) were down-regulated compared to RAFLS transfected with scrambled control siRNA followed by celastrol (Ctrl siRNA + Cel), while one gene, CD40, was up-regulated (fold change value >1.5) when compared to cells transfected with control siRNA (Ctrl siRNA alone; Figure 8). Importantly, the individual knockdown of calmodulin, calcineurin, or calpains restored the gene expression pattern of celastrol-treated RAFLS to different extents (Genes siRNA + Cel). These findings suggest that the three Ca\(^{2+}\)-dependent regulators, calmodulin, calcineurin, and calpains, are the upstream regulators of these 22 inflammatory genes in response to treatment with celastrol.

As shown in the proposed mechanisms depicted in Figure 9, 22 out of 23 genes were modulated entirely or partly, by at least two of the investigated Ca\(^{2+}\)-dependent/-binding proteins, and calcineurin was the most frequently involved protein. For example, the expression of CD40, HRH1, LEPR, NFKB1, STAT3, TOLIP, and TRAP1 was modulated by calmodulin and calcineurin; IFNAR1 and TLR6 were modulated by calpains and calcineurin; the remaining genes, including BMP1, CAST, ERBB2, FGF10, GLMN, IK, NFATC3, NFKB1, and NFX1, were suppressed by all three Ca\(^{2+}\)-dependent/-binding proteins. Furthermore, four genes were modulated by only one of the Ca\(^{2+}\)-dependent/-binding proteins: CMTM1 was suppressed only by calmodulin, ADGRE5 and IL4R were suppressed only by calpains, and IL1R1 was suppressed only by calcineurin. These data suggest that several calmodulin- and calpain-independent pathways are activated in the transcription of selected inflammatory and autoimmunity-associated genes after treatment with celastrol.

4 DISCUSSION

Celastrol, the major bioactive component of the Chinese medicinal plant Tripterygium wilfordii Hook F, has been used as a conventional treatment for RA in China and other Asian countries, demonstrating significant anti-arthritic effects in both experimental arthritis models and RA patients (Tao & Lipsky, 2000; Tao, Younger, Fan, Wang, & Lipsky, 2002; Goldbach-Mansky et al., 2009; Venkatesha, Yu, Rajaiah, Tong,
FIGURE 6  The anti-arthritic effect of celastrol on adjuvant-induced arthritis (AIA) in rats via calcium mobilization. (a) The arthritic scores and hind paw swelling of celastrol (Cel)-treated AIA rats. (b) Representative images of hind paw swelling from AIA rats after treatment. (c) Representative microCT images of hind joints of AIA rats after treatment. Yellow arrows indicate bone erosion. Healthy control group and four groups of rats were treated with vehicle, MTX (7.6 mg·kg⁻¹), celastrol (1 mg·kg⁻¹), or celastrol (1 mg·kg⁻¹) combined with BAPTA/AM (BM; 3.5 mg·kg⁻¹) after arthritis induction for 36 days. Hind paw volumes (ml) and arthritic scores were determined every 3 days. The data are expressed as the mean values ± SEM (n = 8). (d) The microCT scores of celastrol-treated AIA rats. MicroCT scores were obtained from five disease-related indexes of microCT analyses of the calcaneus - bone mineral density, bone volume fraction, cortical mineral density, trabecular number, and total porosity. The data are expressed as the mean values ± SEM from n = 8 rats per group. *P < .05, significantly different from vehicle-treated AIA group. \( \Delta P < .05 \) for microCT score comparison between celastrol treatment group and celastrol plus BAPTA/AM treatment group. (e) The autophagic effect and proliferative rate of synovial fibroblasts in synovium of AIA rat treated by celastrol. Synovium tissues isolated from five rats of each treatment group were section and immune-stained with antibodies against autophagic marker LC3B and synovial fibroblasts marker vimentin prior to secondary antibody (GOXMO TRITC HIGH XADS) treatment. The fluorescence images shown are the representative images from five independent animals. Bar charts show the ImageJ (RRID:SCR_003070) quantitation of the fluorescence signal from LC3B and vimentin. *P < .05, significantly different from vehicle-treated AIA group. For celastrol-treated AIA group \( \Delta P < .05 \), significantly different from Cel + BM-treated AIA group.
FIGURE 7  Effect of BAPTA/AM in the expression of celastrol-regulated inflammatory and autoimmunity genes in the joint tissues of AIA rats. After the completion of drug treatment in the AIA rat model, the joint tissues harvested from the AIA vehicle control group (AIA Ctrl), celastrol 1 mg·kg⁻¹ group (Cel), and celastrol 1 mg·kg⁻¹ plus BAPTA/AM group (BM + Cel) were prepared for RNA isolation, reverse transcription into cDNA, and real-time qPCR analysis of 23 genes of interest related to inflammation and autoimmunity (Dias et al., 2018). Among these 23 genes identified in the human RASF studies, 20 of their counterparts in rats have been identified as being down-regulated by celastrol-induced increase of calcium concentration in a similar pattern. After RT-qPCR, gene expression was normalized to GAPDH, relative to AIA vehicle control, and analysed using the $2^{-\Delta\Delta CT}$ method. The data are presented as the mean ± SEM, *$P < .05$, significantly different from AIA or Cel + BM-treated group.
FIGURE 8 Effect of the Ca^{2+}-dependent modulators in celastrol-mediated inflammatory and autoimmunity gene expression. RAFLS were treated with celastrol (Cel; 1 μM) with or without transfection of a mixture of siRNA for calmodulin genes (CALM1 + CALM2 + CALM3), calpain isoforms catalytic subunit genes (CAPN1 + CAPN2), or calcineurin isoforms catalytic subunit genes (PPP3CA + PPP3CB + PPP3CC) and compared to cells transfected with control siRNA. After RT-qPCR analysis of 22 genes associated with inflammation and autoimmunity, gene expressions were normalized to GAPDH, relative to control siRNA, and analysed using the $2^{-\Delta\Delta Ct}$ method. The data are the mean ± SEM from five independent experiments. *P < .05, significantly different from untreated or Cel + siRNA group.
FIGURE 9  Schematic diagram showing the proposed signalling pathways for inflammatory and autoimmunity-associated gene expression via Ca²⁺ and Ca²⁺-dependent/-binding proteins. Upper part, colour-coded representation of the pathways by which the genes of interest can be affected by the calcium signalling pathway proteins in response to the cytosolic Ca²⁺ rise induced by celastrol treatment; lower part, network map linking the calcium signalling pathway proteins to their effects on the calcium modulated genes and five RA pathogenic factors (anti-apoptosis, proliferation, inflammation and immunity, angiogenesis, migration, and invasion). Genes and pathways are colour-coded according to the upper part. RA pathogenesis factor colours are only linked to their coloured asterisks ( *) and are independent of the gene and pathway colour-coding.
Ca²⁺ release by manipulating Ca²⁺ homeostasis of RASFs instead of other engineered articular cartilage (Natoli et al., 2010). Although the molecular targets responsible for the celastrol-mediated antiarthritic effect have been identified (Cascao et al., 2012; Li et al., 2012; Li et al., 2013; Astry et al., 2015; Jiang et al., 2015; Lv et al., 2015), the mechanistic role of Ca²⁺ signalling underpinning such therapeutic effects remains unclear. Emerging evidence suggests that Ca²⁺ flux may be involved in the pathogenesis of RA. For example, significant differences in ER Ca²⁺ concentrations were found in synovial fluid T cells from patients with chronic inflammatory arthritis after T cell receptor stimulation, when compared to normal T cells (Carruthers, Arrol, Bacon, & Young, 2000). The activation of neutrophils isolated from RA patients was correlated to the Ca²⁺ entry regulated by CD147-induced modifications of lipid rafts, which modulates the downstream activation of the cation channel TRPM7 (Wang et al., 2014). Apart from cellular activation, Ca²⁺ flux is associated with synovial accumulation of autoantigen in RA patients by triggering the activation of peptidylarginine deiminase and subsequent citrullination of vimentin in macrophages (Vossenaar et al., 2004). In addition, conformational changes resulting in altered ATPase activity of SERCA1 have been reported in AIA rats which are correlated with the systemic inflammatory status of the animal model (Strosova et al., 2011). Such observations reinforce the idea of targeting Ca²⁺ mobilization in cytoplasm or the balance of Ca²⁺ homeostasis as an alternative therapeutic approach for improving RA.

Reports from several studies have demonstrated the possibility of manipulating the activity of Ca²⁺ channels with biological agents or medicinal compounds for the treatment of RA. Wang et al. have shown that verapamil, a Ca²⁺ channel inhibitor, antagonized TNF-α-mediated inflammation in collagen-induced arthritis mice via the reduction of Ca²⁺ (Wang et al., 2016). The blockade of store-operated Ca²⁺ release-activated channels with the use of neutralizing antibodies can also effectively suppress the activity of T cells and B cells derived from RA patients (Liu et al., 2017). Intriguingly, ionomycin-mediated Ca²⁺ modulation enhanced the tensile properties of developing engineered articular cartilage (Naitöli et al., 2010).

In this study, we examined the potential of inhibiting RA progression by manipulating Ca²⁺ homeostasis of RASFs instead of other myeloid cells, as this cell type has been shown to be extensively involved in RA pathogenesis ranging from pro-inflammatory mediator synthesis to mechanical damage of joint tissues. AIA rats were treated with celastrol given intraperitoneally, because earlier pharmacokinetic analysis demonstrated low plasma levels of celastrol after oral administration. The maximal plasma concentration (Cmax) in a rat model with oral celastrol (1,000 μg·kg⁻¹) only reached 13.75 ± 7.94 μg·L⁻¹, compared with 38.83 ± 12.83 μg·L⁻¹ after intravenous injection (100 μg·kg⁻¹; Zhang et al., 2012). Interestingly, the absorption of celastrol was enhanced by co-treatment with other bioactive components of Tripterygium wilfordii Hook F (Zhang et al., 2012), providing insight to the improvement of drug delivery of celastrol. We have demonstrated that the pharmacological mechanism of celastrol in the arthritic condition involves the activation of Ca²⁺ signalling through inhibition of SERCA. Addition of the calcium chelator BAPTA/AM, which is a highly selective agent for Ca²⁺, leads to a partial inhibition of the therapeutic effect of celastrol in our experimental model of AIA rats. Furthermore, celastrol induced ER stress, thereby inducing apoptotic and autophagic cell death in apoptosis-resistant fibroblasts, RASFs and RAFLS, a finding that may offer an alternative way to eliminate apoptosis-resistant SLFs of RA patients. In fact, celastrol-induced cytotoxicity via the up-regulation of ER stress and glycogen synthase kinase-3β signalling pathways (Feng et al., 2013), which is closely related to the phosphorylation status of the enzyme and may, in part, be regulated by heat shock protein 90 (Zhang et al., 2008). In addition, celastrol can also target the proteasome to activate the pro-apoptotic unfolded protein response pathway and autophagy in cancer cells (Fribley et al., 2015; Wang et al., 2012). Therefore, the multi-target nature of celastrol has aroused safety concerns. In fact, the actions of celastrol appeared to be more selectively expressed in the inflammatory RAFLS and cancer cells. It is possible that the ER stress of these cells is already at a high level, such that further stimulation after celastrol exposure, may exceed a critical level leading to induction of ER stress-associated cytotoxicity. On the contrary, in normal cells exhibiting basal (low) levels of ER stress, the administration of celastrol may only up-regulate the ER stress to the cytoprotective threshold of the pathway (rather than cytotoxic levels), which could partly explain the higher tolerance towards celastrol treatment.

Downstream signalling pathways induced by the elevation of cytosolic Ca²⁺ levels are mediated by secondary Ca²⁺ sensors and calmodulin is a universal Ca²⁺-binding protein which regulates the activity of a variety of enzymes, such as protein kinases and phosphatases, as well as other signalling proteins, including membrane receptors, channels, and structural proteins (Villalobo A., 2018). In chronic inflammatory arthritis, abnormal activation of calcineurin (a Ca²⁺-calmodulin activated phosphatase) in synoviocytes was observed which could contribute to the corresponding pathogenesis (Yoo et al., 2006). Calpains, a set of Ca²⁺-activated proteases, have also been shown to participate in cartilage destruction in collagen-induced arthritic knee joints of mice (Szomor, Shimizu, Fujimori, Yamamoto, & Yamamuro, 1995). Therefore, we propose that celastrol-mediated Ca²⁺ mobilization modulates a panel of inflammatory- and autoimmunity-associated genes through the Ca²⁺-dependent/-binding proteins. Knockdown of calmodulin, calpains, and calcineurin revealed novel preliminary Ca²⁺ signalling pathways that link pharmacological targets of celastrol to their downstream gene expression based on the proposed Ca²⁺-modulated gene network (Dias et al., 2018). The current findings not only explain the role of the reported genes but also provide a new mechanistic insight into the regulation of inflammation- and autoimmunity-associated gene expression via Ca²⁺ signalling pathways. Therefore, the ameliorating effect of celastrol in the AIA model of RA is partly attributable to Ca²⁺-mediated autophagic cell death in apoptosis-resistant RAFLS. Celastrol may alleviate the symptoms of RA patients by down-regulation of the inflammatory response genes in RASFs via activation of Ca²⁺ signalling pathways. Notably, celastrol exhibits its therapeutic effect via targeting other immune-component cells such as regulatory T cells and Th17 cells (Astry et al., 2015; Cascao et al., 2012; Han et al., 2015) and/or other molecular pathways such as IL-
17A and NF-κB signalling (Li et al., 2012; Li et al., 2013) as mentioned previously independent of Ca²⁺ signalling, which is supported by the observation that BAPTA/AM alone was unable to completely block the therapeutic effect of celastrol.

SERCA is a novel therapeutic target for cancer treatment, especially in multidrug-resistant tumours (Wong et al., 2013; Xu, Shao, & Zeng, 2013). The current study indicates the potential therapeutic role of SERCA in refractory RA through eliminating apoptosis- and multidrug-resistant SLFs via calcium-induced autophagic cell death. With the essential role of SERCA in maintaining normal cellular homeostasis, specific SERCA inhibitors such as thapsigargin are likely to result in significant toxicity to normal SLFs. This limitation has led to an increasing interest in the development of prodrug strategies to reduce the off-target-organ side effects and systemic toxicities (Yuan, Quan, et al., 2012). For instance, methacyclamide copolymer-based dexamethasone prodrug, hydrogen peroxide-sensitive prodrugs of MTX, and aminopterin have been synthesized for the treatment of RA. While maintaining a comparable therapeutic efficacy, these prodrugs have resulted in a safer toxicity profile and an increased therapeutic window, compared to their parent drugs (Peiro Cadahia et al., 2018; Yuan, Nelson, et al., 2012). Therefore, it is worthwhile to further investigate whether the toxicity of celastrol can be minimized using similar prodrug strategies (Wang, Liu, Wang, He, & Chen, 2011) to provide a safer agent for treating RA patients who are less responsive to current anti-arthritis agents.

5 | CONCLUSION

This study has provided new insights into how celastrol regulates a panel of inflammatory and autoimmunity-associated genes via activation of Ca²⁺-binding-dependent proteins. Through interlinking the possible genes regulating apoptosis, proliferation, inflammation, immu-
nity, angiogenesis, and migration/invasion process of AIA models after treatment with celastrol, the current investigation has revealed the detailed molecular mechanisms of celastrol in the effective treatment of RA via calcium signalling.

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CONFLICT OF INTEREST

The authors declare no conflicts of interest.

AUTHOR CONTRIBUTIONS

V.K.W.W., C.L.Q., and S.W.X. designed and conducted the autopho-
yg, animal experiments, data analysis, and also drafted the manu-
scrip. B.Y.K.L., W.Z., and H.W. carried out the AIA model experiment. F.M. conducted the SERCA activity assay and proof-
reading for the whole manuscript. I.R.D.S.R.D. and S.W.F.M. performed the PCR array and real-time PCR experiments. H.P. performed the microCT data analysis. S.H. and T.E. conducted the computational docking. Y.H., N.Z., and Y.Q.Q. performed all flow cytometry experiments. T.W.C., X.C., and L.Y. performed the western blot experiments. W.J.Y. conducted the luciferase report assay. W.Z. and Z.L. conducted the ATP metabolite experiments. Y.X. and R.L. isolated the RASFs from RA patients. Q.J. revised the manuscript and gave valuable suggestions. L.L. conceived the idea, supervised all research, and revised the manuscript.

DECLARATION OF TRANSPARENCY AND SCIENTIFIC RIGOUR

This Declaration acknowledges that this paper adheres to the princi-
iples for transparent reporting and scientific rigour of preclinical research as stated in the BJP guidelines for Design & Analysis, Immunoblotting and Immunohistochemistry, and Animal Experimentation, and as recommended by funding agencies, publishers and other organisations engaged with supporting research.

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**SUPPORTING INFORMATION**

Additional supporting information may be found online in the Supporting Information section at the end of the article.

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