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In vitro responses of human dermal fibroblasts to mechanical strain: A systematic review and meta-analysis

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In vitro research in the field of mechanotransducive regulation of dermal fibroblasts is characterized by highly variable methodology and contradictory results. The primary objective of this systematic review was to establish how in vitro mechanical stretch affects human dermal fibroblast function, by means of a quantitative synthesis of all available evidence. The secondary objectives were to examine the effects of covariates related to donor age, fibroblast origin, experimental treatments, and mechanical stimulation parameters on dermal fibroblast responsiveness to mechanical strain. Summary outcomes for fibroblast proliferation and collagen production were combined using a fixed-effects meta-analytical model. Subgroup analysis and meta-regression were used to investigate the effects of different conditions on the summary outcomes. Mechanical strain was found to not affect fibroblast proliferation in neonatal fibroblasts, while adult fibroblasts proliferation was significantly increased. Collagen production was significantly increased in response to mechanical stimulation, with Vitamin C stimulation as the most important covariate. Stretching frequency emerged as positively associated with fibroblast proliferation and negatively associated with collagen production. We conclude from this study that distinct differences exist in the effects of mechanical stretching between dermal fibroblasts from neonatal and adult donors, which will help to further elucidate the pathophysiological mechanism behind tension-induced scarring.

KEYWORDS

skin, biomechanics, keloid, hypertrophic scar, mechanotransduction

Abbreviation: ECM, extracellular matrix; ROUT, robust regression and outlier removal.

1 Introduction

Extensive cutaneous wound healing unavoidably leads to scarring. Excessive scarring in the form of a hypertrophic scar or keloid can occur if the process of wound healing is disordered (Ogawa, 2017). These raised and thickened scars involve a pronounced and prolonged activation of myofibroblasts during the remodeling stage of wound healing (Hinz, 2016), resulting in excessive production and reduced degradation of extracellular matrix (ECM); thus, contributing to a build-up of ECM (Hinz, 2016).

Excessive scarring seems to occur disproportionately at anatomical sites with higher resting tension or high cyclical tension (Barnes et al., 2018). Inplane mechanical tension in general is considered to have harmful effects on cutaneous wound healing, and on the quality of the subsequent scar. A myriad of therapies exist aimed at mitigating the negative effects of tension by diminishing in plane stresses, such as optimized suturing techniques (Marsidi et al., 2020) or devices that disperse tension to the surrounding tissue (Al-Mubarak and Al-Haddab, 2013).

The concept of mechanotransduction involves reciprocal interactions between cells and their mechanical environment (Hinz, 2016), and the subsequent translation into biochemical signals (Martino et al., 2018). The discovery of mechanically sensitive ion channels was the first step to unravelling the potential molecular basis behind scar exacerbation caused by tension (Guharay and Sachs, 1984). This field covers clinical research (Krishnan et al., 2016), animal models (Son and Hinz, 2021), in vitro studies (Shirakami et al., 2020), and in silico studies (Tsuge et al., 2020). In vitro studies play an important role in elucidating the role of mechanobiology in scar formation on a cellular and biomolecular level, however effects described in these articles vary widely and are often contradictory (Wong et al., 2011) These studies involve the seeding of primary cells on extensible membranes, and the subsequent analysis of outcomes while controlling for relevant deformation parameters. Increased attention has been paid recently to improving the translational value of animal studies by means of systematic reviews and meta-analyses (Hooijmans et al., 2014), given the abundance of narrative reviews on this topic of which none evaluate the methodological quality of individual studies. Although the framework for systematic reviewing of clinical trials does not easily lend itself to application in in vitro research (Hooijmans et al., 2014), sparse examples of its successful adaptation exist (Mikolajewicz et al., 2018; McKee et al., 2019).

In this work, we aimed to critically assess the effect of *in vitro* mechanical stretching on human dermal fibroblasts using a systematic review approach. The primary objective of this systematic review and meta-analysis was to characterize the effect of *in vitro* mechanical stretch on the functional parameters of human dermal fibroblasts, namely proliferation and collagen production. The secondary objective was to examine the effects of covariates related to donor age, fibroblast origin, experimental treatments, and parameters of mechanical stimulation on the dermal fibroblast responsiveness to mechanical strain. Finally, we evaluated the consistency of reporting of *in vitro* studies in order to improve their translational potential.

2 Results

2.1 Study characteristics

The systematic literature search (Figure 1) resulted in a total of 11,230 articles. Two additional articles were found by screening

references of relevant articles. Title and abstract screening of the remaining articles resulted in the inclusion of 48 relevant full-text papers. Most of these articles were written in the context of cutaneous scar research. Two papers on bioengineered tendon tissue, and four papers on cardiovascular research were included as they involved experiments on dermal fibroblasts. All individual experiments that met the inclusion criteria were included, resulting in more datapoints than papers.

2.2 Methodological quality

The research described in the 48 papers was of heterogeneous design. Donor location varied widely, foreskin being the most commonly used donor tissue being (n = 16) (Figure 2C) and nineteen articles not specifying donor location. The stretching device used in the experiments was described in all articles; 17 papers reported the use of a commercial device from Flexcell, others (n = 20) used custom-built devices (Figure 2D). Methodological quality was scored using a predefined set of items assessing methodological quality (Figure 2A, B, Supplementary Table S5). Items related to the materials used were generally well reported, with only two articles omitting culturing medium and conditioning altogether, and six articles missing a detailed description of the stretching device. Demographical information on cell donors was absent in the vast majority of articles (n = 35). Similarly, the number of cell donors was severely underreported (n = 31). Details on the extensible membrane and its coating or treatment were provided by all authors. Eight papers were missing one or two details on either the load duration, frequency, and magnitude. Conversely, the number of repetitions of experiments was poorly reported in nineteen articles. Four articles made no mention of sample size at all, and a further fifteen articles were vague or included a range of repetitions per analysis used. Most papers reported the use of an unstretched control sample, with only seven articles using an inappropriate control group or making no mention of an unstretched control. Further scrutiny resulted in a number of methodological flaws in articles regarding unstretched control groups, including the use of t = 0 measurements as a control group, and selective omission of control group data. Nineteen articles included an explicit statement on the identical exposure of both the experimental and control group. In nine cases the exposure was demonstrably different from the intervention group.

2.3 Data extraction and processing

Two outcome measures were chosen based on reporting frequency and relevance to the molecular mechanism of mechanotransduction: fibroblast proliferation and collagen production. This resulted in the inclusion of 20 studies for quantitative synthesis, 19 for fibroblast proliferation (Supplementary Table S1) and 10 for collagen synthesis (Supplementary Table S2). Two frequently reported outcomes measures, matrix reorganization or reorientation and sample contraction, were reported too heterogeneously and no scale-free overall effect size could be determined. Reporting of cell proliferation varied, with some giving detailed growth curves and some only differences between the cell counts at a single time point. Therefore, to combine the data from cell proliferation datasets we estimated the difference between the growth constants in experimental (k_s) and control (k_c) conditions under the assumption of exponential



growth as $k_s k_c = \frac{ln Y_s/Y_c}{t}$ (where Y_s and Y_c are cell numbers or densities reported at time *t*). The performance of this assumption was tested by fitting the time-series dataset from Nishimura et al. (Nishimura et al., 2007) (Figure 3A) to an exponential growth curve (Figure 3B) to estimate growth constants. The correlation between estimated growth constants and the calculated ones was 0.913 for means, and 0.757 for variance (Pearson's R, Figures 3C, D). Variance was consistently larger in the calculated method, indicating a more conservative estimate which is appropriate given the complex error propagation. For collagen production, the normalized mean difference between experimental and control conditions in each dataset was calculated.

2.4 Analysis of heterogeneity

Heterogeneity was analyzed in both datasets. Single-study exclusion analysis of the fibroblast proliferation dataset did not affect the overall effect size of the outcome (Figure 3E). Cumulative study exclusion analysis determined that 44% of studies needed to be excluded to obtain a homogenous dataset and their removal would result in a decreased effect size (Figure 3F). In the collagen production dataset, single study exclusion analysis identified three studies that contributed to heterogeneity, and their exclusion affected the overall effect size of the outcome (Figure 3G). Cumulative study exclusion analysis demonstrated that 80% of the studies needed to be excluded to attain heterogeneity (Figure 3H). The removal of these studies resulted in a decreased effect size. Funnel plot analysis of the fibroblast proliferation dataset (Figure 3I) demonstrated the bi-modal distribution of the study level effect data points with one peak around the fixed-effects estimate and another one at the higher values. Funnel plot of the collagen production dataset (Figure 3J) demonstrated that the study level effects were distributed around the fixed-effects estimate.

2.5 Meta-analysis

2.5.1 Fibroblast proliferation

Since the distribution of study level effects in fibroblast proliferation showed an apparent division between two groups (Figure 3), we examined if two separate populations in this dataset



Demographics include age and sex. (C) Frequency of reporting of specific donor-site locations. (D) Frequency of use for different stretching devices in selected papers.



FIGURE 3

Data processing and heterogeneity analysis. (A–D) Overview of processing and validation of cell proliferation data that starts from original graph (A), reproduced with permission from Nishimura et al. (Nishimura et al., 2007)), which was fited with exponential growth curves for control Y and experimental group Z (B). The difference between growth constants obtained from fitting was compared to the k_s - k_c calculated from a single time point as correlation for means (C) and standard deviation (D). (E–H) Heterogeneity analysis was performed for the cell proliferation (E, F) and collagen production (G, H) using single (E, G) and cumulative (F, H) study exclusion. (I, J) Publication bias and effect size distributions were assessed using funnel plots for cell proliferation (I) and collagen production (J) datasets. Study level data is indicated by black circles. The outliers in the collagen dataset are indicated with red circles.

corresponded to fibroblast donor age. The reporting of donor age allowed to separate data from neonatal and adult donors, while the studies in which donor age was not properly reported were combined together (Figure 4). Neonatal fibroblasts showed an apparent insensitivity to stretch-induced proliferation (Ks-Kc: 0.00[CI:-0.01: 0.01]) while adult fibroblasts exhibited a clear increase in proliferation



neonates (red), adults (green) and studies with unspecified donor age (blue).

after mechanical stretch (Ks-Kc: 0.14[CI:0.12:0.16]). In the group of non-reported age, fibroblast proliferation showed the same insensitivity to stretch-induced proliferation as neonatal fibroblasts (Ks-Kc: 0.01[CI:-0.01:0.03]). While neonatal cell responded more to uniaxial than to biaxial stretch, adult cells showed the opposite response, with an extensive response to biaxial stretch (Figures 4A, B, D). Proliferation of neonatal cells was not affected by Vitamin C, however a very small subset of cells of non-reported age showed a positive effect of Vitamin C on stretch-induced proliferation. Unfortunately, none of the articles on adult cells described adding Vitamin C to the culture medium. While all the neonatal fibroblast originated from foreskin, adult fibroblasts were obtained from different locations including foreskin (Figure 4C). Although adult fibroblasts from different locations responded similarly to the stretch, notable outliers were breast skin and foreskin (however this was only one dataset), which demonstrated a remarkable low stretch-induced proliferation compared to other donor site locations.

2.5.2 Collagen production

The collagen production dataset contained a small number of studies that demonstrated effect sizes that were orders of magnitude larger than the rest of the dataset (Figure 5). The ROUT method was thus used to identify five outliers in the collagen dataset that were subsequently omitted from further analysis. Overall, mechanical stretching increased collagen production (NMR: 0.44[CI:0.44:0.45]). Consistent with its known function, addition of Vitamin C almost doubled the stimulatory effect of mechanical stretching. Foreskin fibroblast derived from neonatal donors also produced more collagen in response to mechanical stretching. Collagen production was evenly stimulated by uniaxial and biaxial stretching, and an even greater effect was seen in circumferentially stretched fibroblasts.

2.6 Meta-regression

The effects of the continuous covariates strain magnitude and frequency on stretch-induced fibroblast proliferation and collagen production was investigated using a fixed-effects meta-regression model. Stretch-induced proliferation (Figures 6A, B, D) and stretch-induced collagen production (Figures 6C, E) positively correlated with strain magnitude in both neonatal and adult cells. However, in all three cases, the slope of the association was minimal, and goodness of fit was poor. Inspection of the plots suggests a non-linear association between increasing magnitude and proliferation,



FIGURE 5

Effect of mechanical stretching on collagen prodution by fibroblasts. (A) The forest plot for the normalized mean differences between collagen production by mechanically stretched and control human dermal fibroblasts. Studies are stratified according to age of fibroblast donors. Black circles/lines indicate study level effect sizes with 95% CI. Diamonds/shaded bands indicate overall effect sizes with 95% CI. Red asterisks and arrows indicate outliers that were not included in the overall effect size calculation. (B) Subgroup analysis for the use of vitamin C stimulation during culture, skin origin of fibroblasts and different stretching mechanisms. Black circles/lines indicate effect size and 95% CI within subgroups, with the number of included datasets indicated in the parentheses, shaded bands indicate the 95% CI for the overall effect.



FIGURE 6

Effect of the parameters of mechanical stretching on the fibroblast proliferation and collagen production. The relationship between (A, C, D) stretching magnitude (%) and (B, E) frequency (Hz), and fibroblast proliferation in cultures of neonatal (A, B) and adult (D) origin, or collagen production (C, E) were assessed using a fixed-effects linear meta-regression model. Meta-regression is shown as a black line, with the slope (β_{inter}) and goodness of fit (\mathbb{R}^2) indicated. 95% CI and prediction intervals are indicated by dark and light shaded red bands, respectively.

with an initial increase and subsequent downward slope toward 20% strain. An increased strain frequency was strongly and positively associated with an increase in stretch-induced proliferation in neonatal cells (Figure 6B), and strongly and negatively associated with stretch induced collagen production (Figure 6E).

3 Discussion

This study constitutes the first systematic review and metaanalysis of *in vitro* studies assessing mechanobiology of human fibroblasts. The primary goal was to characterize the effect of *in vitro* mechanical stretch on human dermal fibroblasts. Two parameters, fibroblast proliferation and collagen deposition, were studied in a sufficient number of studies to perform a meta-analysis.

In total, 71 and 42 discrete experimental estimates on cell proliferation and collagen production were extracted, respectively. We found, for the first time, that only fibroblasts from adult donors increased both their proliferation and collagen production in response to mechanical stretch, while neonatal fibroblasts mainly responded by increasing their collagen production.

Meta-analysis on this data demonstrated the absence of stretch-induced proliferation in neonatal fibroblasts, while this effect was certainly present for adult fibroblasts. Scarless wound healing in the human fetus is an established fact (Larson et al., 2010). Our results seem to suggest that age-related difference in dermal fibroblast's reactivity to mechanical tension may exist for a longer duration than previously thought.

In contrast, dermal fibroblasts from donors of all ages showed an increase in production of collagen type I in response to mechanical stimulation, although the degree of increase was variable, partially due to the differences in the use of vitamin C during culture. Unfortunately, other types of collagens were not reported abundantly enough to perform a quantitative synthesis, making an analysis of other types of collagen relevant to scar formation impossible (Oliveira et al., 2009).

We have found that stretch magnitude and the frequency of stretching differentially affected fibroblast proliferation and collagen secretion. Although there was no significant correlation between the stretch magnitude and fibroblast function, detailed data examination suggested that the relationship may be non-linear. In contrast, stretch frequency significantly correlated with both fibroblast proliferation (positive correlation) and collagen production (negative correlation). These data may explain the dichotomous effects of observed in clinical studies (Schouten et al., 2011; Wong et al., 2011). Thus, our studies propose a new direction of research in identifying an optimal mechanical stretching protocol to reduce scarring.

This study has several limitations. First, the quality of the 48 identified articles was critically appraised using a modified scoring system based on previous work by Golbach et al. (2016), and several important flaws were identified. Reporting of the demographic characteristics of fibroblast donors was particularly poor and seldom contained information on donor sex, age, and fibroblast site location. Moreover, the number of donors used per experiment were reported only in a minority of articles. Improved reporting on the number of donors will increase the generalizability of results. Similarly, increased reporting on donor characteristics will allow for more in-depth assessment of their role in scar formation. Second, we have identified deficiencies in reporting of experimental conditions. A minority of articles unequivocally stated that the conditions in the control experiments were identical to the experimental group. Cell passage numbers were not consistently reported, despite their known impact on proliferation. Moreover, although the majority of articles described clear samples sizes, a vague description or range of sample sizes was still a problem in many papers. Finally, 27 out of 48 articles did not include a conflict-of-interest statement. These findings emphasize the need for better reporting of basic science studies and the requirements of a specific quality assessment tools for the synthesis studies. Even though the articles included for quantitative syntheses had relatively high scores for reporting quality (Supplementary Tables S1, S2), reported conclusions are subject to the deficiencies in reporting identified above.

Furthermore, to control the scope, this study solely investigated the effect of mechanical strain given its predominant role, even though other types of mechanical deformation such as shear and compression may play important roles. We were also limited in the number of analyzed outcomes. More specifically, we were not able to include matrix reorganization and fibroblast differentiation due to lack of consistent reporting of the outcomes. Similarly, even though fibroblast subpopulations (e.g. reticular *versus* papillary fibroblasts) are thought to behave differentially in scar formation (Woodley, 2017), we could not assess a potential difference due to a lack of data.

To conclude, this study decisively demonstrated the stimulatory effect of mechanical stretching on human dermal fibroblast proliferation and collagen production and identified donor age and the frequency of mechanical stretching as important determinants of the outcome. Our study uniquely suggests that reducing proliferation while simultaneously increasing collagen production by adult cells may be beneficial for mimicking the mechanical stretching response in neonatal cells. Our analysis also suggests that potential strategies to achieve this may include treatment with ascorbic acid. These results provide concrete recommendations for future experimental research to further elucidate the physiological mechanism behind stretchinduced scarring, for the development of novel treatment strategies, and in optimization of tissue engineering protocols.

4 Methods

4.1 Software

Articles retrieved through database search were imported into EndNote X9 (Clarivate Analytics, Philadelphia, United States) for duplicate removal. The web application Rayyan (Ouzzani et al., 2016) was used for title and abstract screening. META-LAB (Mikolajewicz and Komarova, 2019), a plug-in developed in MATLAB R2016b was used for data extraction, meta-analysis, and meta-regression. Data fitting and statistics were carried out in GraphPad Prism9 (SanDiego, CA, United States) and Microsoft Excel was used for further data processing. The open-source graphics editing software Inkscape was used to edit plots and figures.

4.2 Search strategy

A systematic search of available literature was performed in the following databases: PubMed, Embase.com, Clarivate Analytics/Web

of Science Core Collection on 1 June 2020. The systematic search was updated on 26 May 2021. The full search strategy can be found in Supplementary Table S3. Additional articles were identified by checking references. The PRISMA statement was followed for reporting in this systematic review (Moher et al., 2009), with the exception of protocol registration as basic science reviews are not accepted in protocol registries.

4.3 Selection of literature

Title and abstract screening were performed individually by two authors, LvH and YM. Disagreement on individual cases was resolved through discussion by the two authors. A third author (MLG) was available in case of unresolvable differences. Studies were included if the experimental protocol involved exposing human dermal or scar fibroblasts to mechanical stretch. Exclusion criteria were as follows: language other than English, animal studies, clinical studies, and mechanical stimulation that was based on other mechanisms (e.g., pressure). Reviews, letters, editorials, and conference abstracts were also excluded.

4.4 Assessment of methodological quality

Studies were ranked for quality using an assessment tool based on a tool previously described by (Golbach et al., 2016). The assessment included twelve items on the experimental setup, intervention, and reporting characteristics (Supplementary Table S4). Items were scored numerically based on the "completeness" of reporting in the relevant studies, in which only the quality of the experiments that fall within the scope of this review were assessed. Total aggregate quality scores were calculated at the study level. Assessment of methodological quality was done separately by two authors (LvH and YdJ), and mutual agreement was sought on divergent scores by discussion between these two authors.

4.5 Data extraction

Studies were included for quantitative synthesis when data on cell proliferation or collagen deposition were provided for both intervention and control groups. Data was extracted concerning experimental design, methodology, outcome measures, and results. The following results were exported into an Excel spreadsheet: sample size, means, and standard errors/deviations. When a range of sample sizes was provided, the lowest value was used. When sample size was not explicitly stated, n = 3 was assumed. In the case of unreported measure of variance, it was assumed to be a standard error. Graphical data was extracted using the META-LAB toolbox for basic research (Mikolajewicz and Komarova, 2019). When data was insufficient or unclear, the authors were contacted to elucidate the data.

4.6 Outcome measures and effect sizes

Two outcome measures were sufficiently reported for quantitative analysis: 1) cell proliferation and 2) collagen type I production. When

collagen subtypes were reported, type I collagen data was extracted. When subtypes were not specified, overall collagen volumes were extracted.

4.6.1 Study-level outcome for cell proliferation

To account for differences in the duration of intervention and methodology, cell proliferation datasets were treated as follows. We assumed that cell growth occurs exponentially as $Y_i = Y_0 \times e^{k_i \times t}$, where Y_i is cell number at time t in experimental (Y_s) and control (Y_c) groups, Y_o is the number of cells at baseline, which we assumed to be identical for the two groups, and k_i are the growth constants. The effect of experimental conditions was then assessed as the difference between growth constants in experimental and control groups: $k_s - k_c \frac{\ln Y_e/Y_c}{t}$. The variance was calculated using arithmetic error propagation technique as $SD_i = \sqrt{\left(\frac{SD_{Ye}}{Ye}\right)^2 + \left(\frac{SD_{Ye}}{Ye}\right)^2}$. The longest datasets from Nishimura (Nishimura et al., 2007) were used to validate this normalization strategy. Data was fitted to an exponential growth curve using non-linear regression. Growth constants attained through these methods were subtracted and compared to the experimental method by calculating the correlation coefficient. Normality of data was confirmed using the Shapiro-Wilk test. Correlation was tested by calculating Pearson's R for both means and standard errors.

4.6.2 Study level outcome for collagen production

Normalized mean differences θ_i were calculated as $\theta_i = \frac{\theta i^e - \theta i^c}{\theta i^c}$, where θi^c was collagen production in control and θi^e in experimental conditions.

4.7 Heterogeneity analysis

Heterogeneity was calculated using H² and I² statistic. The H² statistic is independent of the number of datasets and is calculated as $H^2 = \frac{Q}{df}$, where Q is the measure of total variation and df is the degree of freedom. The I² statistic is used to describe variance on a relative scale and is a transformation of the H² statistic as $I^2 = (\frac{Q-df}{Q}) \times 100\%$. The contribution of individual studies to the data pool heterogeneity was assessed using single study and cumulative study exclusion analysis. In cumulative study exclusion analysis, the homogeneity threshold T_H was calculated, which estimates the percentage of studies that need to be removed to attain homogeneity. Publication bias was assessed with funnel plot, generated by plotting the inverse standard error, $1/se(\theta_i)$ against the individual study effect size, θ_i .

4.8 Meta-analysis

Random effects models are generally preferred in basic research (Mikolajewicz and Komarova, 2019) because of high amounts of experimental and biological variability. However, given the clear distribution of study level effects around the fixed effects estimate for both outcomes, a fixed effects model was used to plot subsequent forest plots.

The study-level outcomes were used to generate a forest plot to achieve a summary outcome, $\hat{\theta}$ as $\hat{\theta} = \frac{\sum_{i}^{N} (\theta \times w_i)}{\sum_{i}^{N} w_i}$, where N is the number of datasets, and the weight assigned to each study (w_i) was

calculated as $w_i = \frac{1}{(se(\theta_i))^2}$. The 95% confidence interval was calculated as $\pm CI = \pm z_{1-\frac{\alpha}{2}} \times se(\hat{\theta})$. Outliers were identified using the ROUT method.

4.9 Subgroup analysis and meta-regression

Subgroup analysis and meta-regression (Mikolajewicz and Komarova, 2019) were carried out to identify factors related to cell donor, donor-site, experimental conditions, and stretching parameters that account for heterogeneity and influence the overall outcomes. Subgroup analysis was used for categorical variables. A fixed-effects meta-regression model was used when sufficient data for continuous variables was present was employed: $\theta_i = \beta_o + \beta_{inter} X_i + \varepsilon_i$, where β_o is the intercept, β_{inter} is the coefficient of the slope that describes the relationship between the covariate and the outcome, and ε_I is the within-study error. \mathbb{R}^2 was calculated to evaluate the model's goodness of fit.

Author contributions

Conceptualization: LvH, CD, YdJ, PvZ MG, SK Data Curation: LvH, CD, YM, YdJ, JK, SK Formal Analysis: LvH, CD, YM, YdJ, JK, MG, SK Funding Acquisition: PvZ, MG, SK Investigation: LvH, CD, SK Methodology: LvH, CD, SK Project Administration: LvH, SK Resources: LvH, CD, JK, SK Software: LvH, CD, JK, SK Supervision: PvZ, MG, SK Validation: LvH, CD, IK, SK Visualization: LvH, CD, SK Writing - Original Draft Preparation: LvH, CD, YM, IK, YdJ, PvZ, MG, and SK Writing—Review and Editing: LvH, CD, YM, IK, YdJ, JK, PvZ, MG, and SK.

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Conflict of interest

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Supplementary material

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